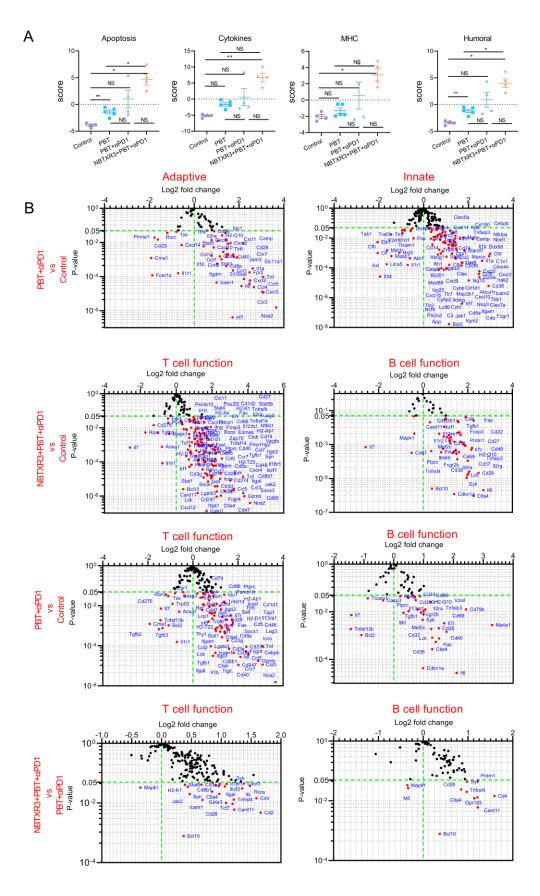
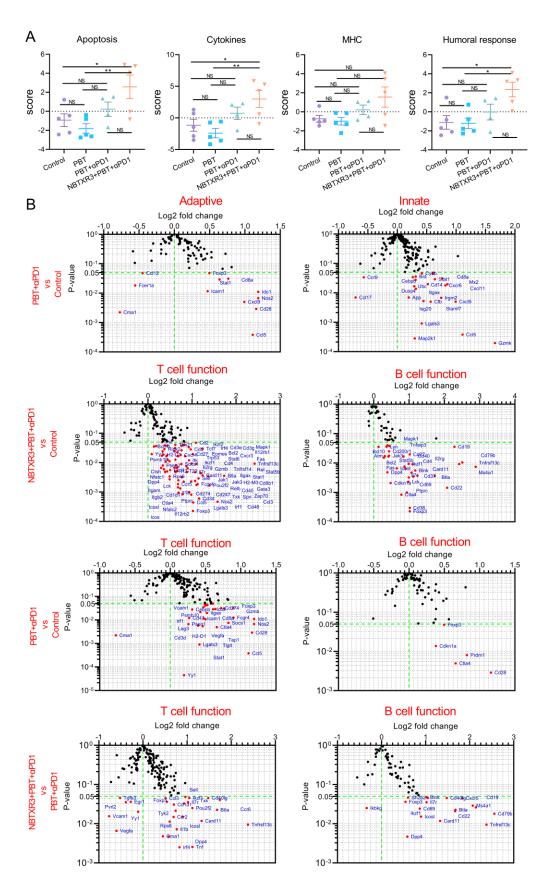


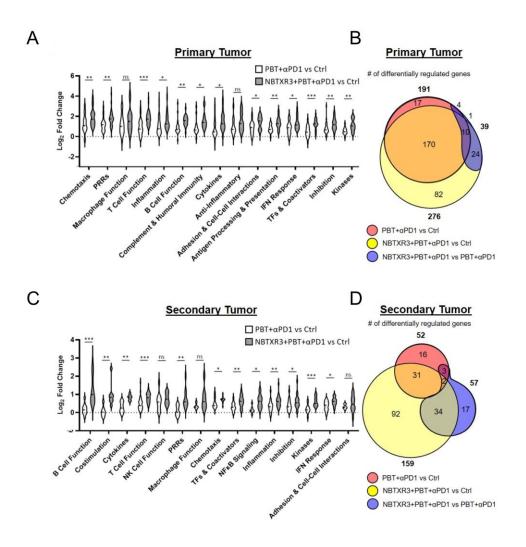
Supplemental Figure 1. Average growth of primary and secondary tumors (n=5) treated with various combinations of NBTXR3, PBT, and  $\alpha$ PD1. Female 129/SvEv syngeneic mice aged 8-12 weeks were inoculated with 344SQR cells on the right and left legs to establish primary tumors on day 0 and secondary tumors on day 4, respectively. On day 7, primary tumors received intratumoral injections of NBTXR3, followed by two 12Gy proton beam radiation fractions on days 8 and 9. Mice were administered 200  $\mu$ g of  $\alpha$ PD1 via intraperitoneal injections on days 7, 10, and 14. Data are presented as mean  $\pm$  standard error of the mean (SEM). A p-value < 0.05 was considered statistically significant. \*\*\*\*P < 0.0001, NS not significant.



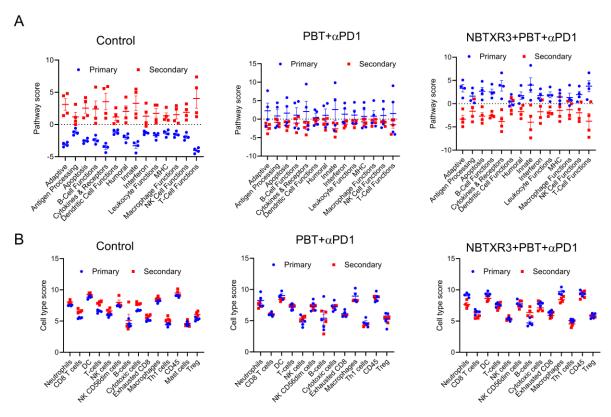
Supplemental Figure 2. Triple therapy of NBTXR3, PBT, and  $\alpha$ PD1 modulates the expression of immune-related genes in favor of antitumor immune response in the irradiated (Primary) tumors. (A) Activity scores of different immune pathways. (B) Changes in gene expression in adaptive, innate pathways, T cell function, and B cell function. Primary tumors were harvested from mice (n=4-5) treated with different combinations of NBTXR3, PBT, and  $\alpha$ PD1 10 days post irradiation. Total RNAs were extracted from the tumors, followed by an analysis of immune-related genes with a nCounter PanCancer Immune Profiling Panel. The gene expression data were then analyzed with the PanCancer Immune Profiling Advanced Analysis Module. Data are expressed as means  $\pm$  standard error of the mean (SEM). P< 0.05 was considered statistically significant. \*P< 0.05, \*\*P< 0.01, \*\*\*P< 0.001, NS not significant.



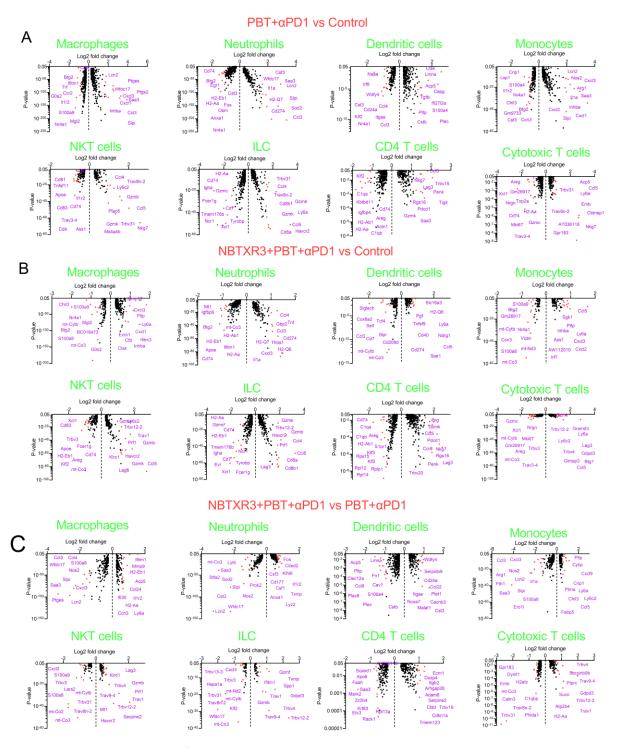
Supplemental Figure 3. Triple therapy of NBTXR3, PBT, and  $\alpha$ PD1 modulates the expression of immune-related genes in favor of antitumor immune response in the unirradiated (Secondary) tumors. (A) Activity scores of different immune pathways. (B) Changes in gene expression in adaptive, innate pathways, T cell function, and B cell function. Secondary tumors were harvested from mice (n=4-5) treated with different combinations of NBTXR3, PBT, and  $\alpha$ PD1 10 days post irradiation. Total RNAs were extracted from the tumors, followed by an analysis of immune-related genes with a nCounter PanCancer Immune Profiling Panel. The gene expression data were then analyzed with the PanCancer Immune Profiling Advanced Analysis Module. Data are expressed as means  $\pm$  standard error of the mean (SEM). P<0.05 was considered statistically significant. \*P<0.05, \*\*P<0.01, NS not significant.



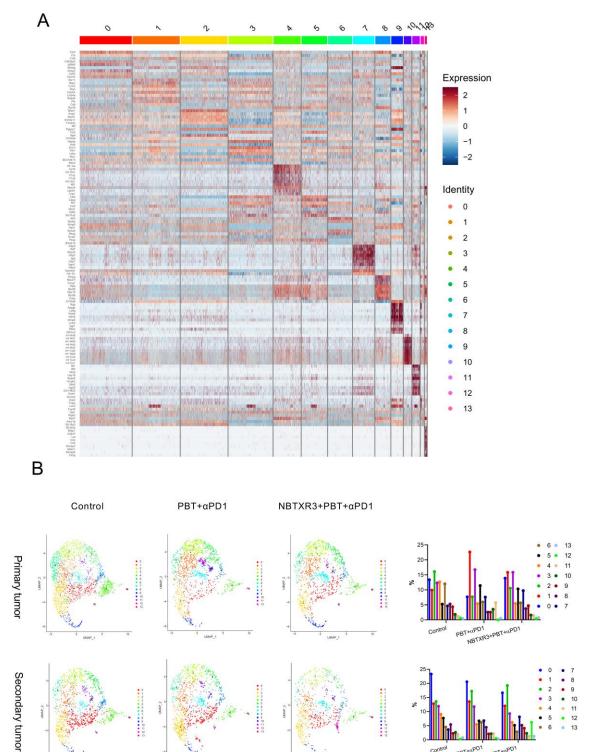
Supplemental Figure 4. Functional distribution of immuno-related genes differentially regulated between experimental groups. Gene transcripts identified as significantly upregulated in either the PBT+ $\alpha$ PD1 or NBTXR3+PBT+ $\alpha$ PD1 group relative to the control from the NanoString data (Figure 2, Figure 3, and Supp. Figure 2 and 3) were manually assigned to functional categories based on their known functions and associations. (A) Top 15 differentially expressed pathways in the primary tumors. (B) Genes differentially upregulated in the primary tumors in PBT+ $\alpha$ PD1 relative to Control mice (red), NBTXR3+PBT+ $\alpha$ PD1 relative to Control mice (green), and NBTXR3+PBT+ $\alpha$ PD1 relative to PBT+ $\alpha$ PD1 mice (blue). The area of each diagram is proportional to the number of genes contained therein, as is the area of overlap (i.e., number of genes shared) between them. (C) Top 15 differentially expressed pathways in the secondary tumors. (D) Genes differentially upregulated in the secondary tumors.



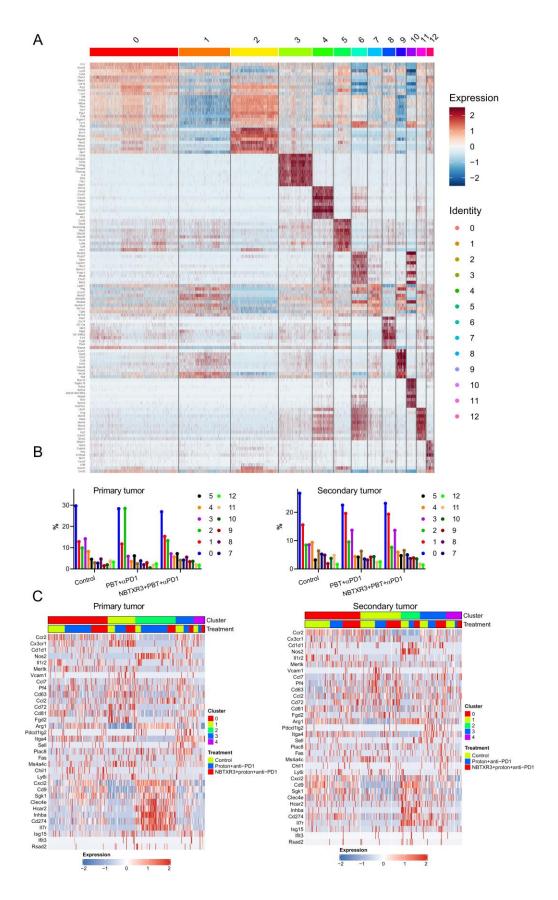
**Supplemental Figure 5. (A)** Comparison of immune pathway activities, and **(B)** relative cell abundance between the primary and secondary tumors.



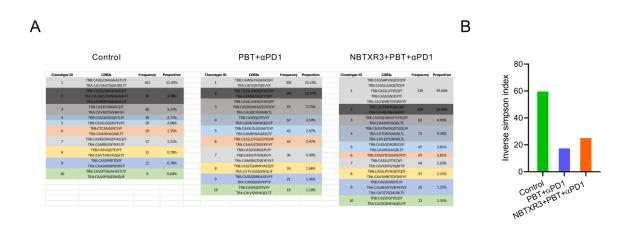
**Supplemental Figure 6.** Significantly changed gene expression in macrophages, neutrophils, DCs, monocytes, NKT cells, ILC, CD4<sup>+</sup> T cells, and cytotoxic T cells in the primary tumors when treated with **(A)** Control vs PBT+ $\alpha$ PD1, **(B)** Control vs NBTXR3+PBT+ $\alpha$ PD1, and **(C)** PBT+ $\alpha$ PD1 vs NBTXR3+PBT+ $\alpha$ PD1.



PBT+aPD1 NBTXR3+PBT+aPD1 **Supplemental Figure 7. (A)** Heatmap showing row-scaled expression of differently expressed genes per cluster for all neutrophils. **(B)** UMAP visualization and percentages of subclusters of neutrophils.

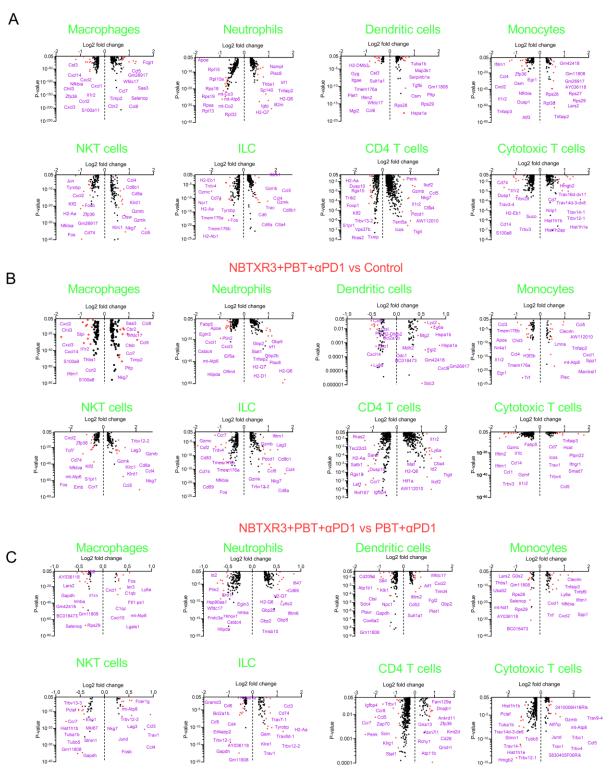


**Supplemental Figure 8. (A)** Heatmap showing row-scaled expression of differently expressed genes per cluster for all macrophages. **(B)** Percentages of subclusters of macrophages. **(C)** Gene set enrichment analysis performed on the top 5 macrophages in both primary and secondary tumors.

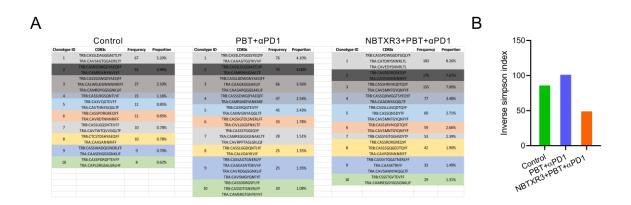


Supplemental Figure 9. (A) Top 10 TCR $\alpha$  and TCR $\beta$  pairs. (B) inverse Simpson index in T cell populations. Mice (n=5) inoculated with 344SQR tumors were treated with dual therapy of PBT+ $\alpha$ PD1 or triple therapy of NBTXR3+PBT+ $\alpha$ PD1 shown in **Figure 1a**, immune cells extracted from the irradiated tumors 17 days post radiation were analyzed via scRNAseq.

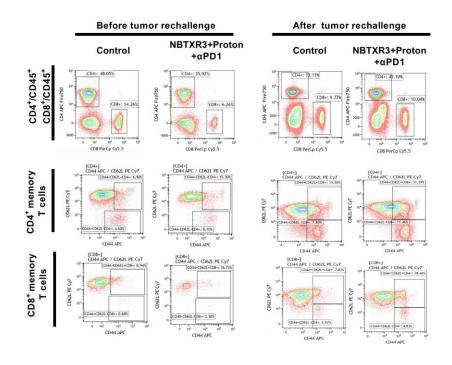
## PBT+αPD1 vs Control



**Supplemental Figure 10.** Significantly changed gene expression in macrophages, neutrophils, DCs, monocytes, NKT cells, ILC, CD4<sup>+</sup> T cells, and cytotoxic T cells in the secondary tumors when treated with **(A)** Control vs PBT+ $\alpha$ PD1, **(B)** Control vs NBTXR3+PBT+ $\alpha$ PD1, and **(C)** PBT+ $\alpha$ PD1 vs NBTXR3+PBT+ $\alpha$ PD1.



Supplemental Figure 11. (A) Top 10 TCR $\alpha$  and TCR $\beta$  pairs. (B) Inverse Simpson index in T cell populations. Mice (n=5) inoculated with 344SQR tumors were treated with dual therapy of PBT+ $\alpha$ PD1 or triple therapy of NBTXR3+PBT+ $\alpha$ PD1 shown in Figure 1A, immune cells extracted from the unirradiated tumors 9 days post radiation were analyzed via scRNAseq.



Supplemental Figure 12. Representative FACS graphs of memory T cells before and after tumor rechallenge.

## **Supplemental Tables**

## Supplemental Table 1: Genes Differentially Upregulated in Primary (Irradiated) Tumor

|        | PBT+αPD1 vs Ctrl |  |                                   |  |  |
|--------|------------------|--|-----------------------------------|--|--|
| Gene   | Log2 Fold Change | Full Name                                  | Notable Aliases                   | Function   |  |
|        |                  | Adhes                                      | sion & Cell-Cell Intera           | actions  |  |
| Cd37   | 1.26             | Cluster of differentiation 37              | Tetraspanin-26                    | Cell surface glycoprotein known to complex with integrins and other transmembrane 4 superfamily proteins; may play a role in T cell- B cell interactions |  |
| Cd47   | 0.412            | Cluster of differentiation 47              | Integrin-associated protein (IAP) | Partners with membrane integrins to serve as an inhibitor of phagocytosis  |  |
| Cyfip2 | 1.26             | Cytoplasmic FMR1-<br>interacting protein 2 |                                   | Involved in T cell adhesion and p53/TP53-<br>dependent induction of apoptosis  |  |
| lcam1  | 0.997            | Intracellular adhesion molecule 1          | CD54                              | Cell surface glycoprotein that serves as strong adhesive ligand for LFA-1; important for leukocyte mobility and costimulation                            |  |

| ltga1  | 0.886 | Integrin subunit      | CD49a; very late     | Alpha 1 subunit for common integrin                   |
|--------|-------|-----------------------|----------------------|---|
|        |       | alpha 1               | activation protein 1 | receptors; pairs with the $\beta 1$ subunit to form a |
|        |       |                       | (VLA-1)              | cell-surface receptor for collagen and laminin;       |
|        |       |                       |                      | involved in cell-cell adhesion and may play a         |
|        |       |                       |                      | role in inflammation and fibrosis                     |
| Itgal  | 1.2   | Integrin alpha L      |                      | Pairs with ITGB2 to form lymphocyte function-         |
|        |       |                       |                      | associated antigen-1 (LFA-1), a common                |
|        |       |                       |                      | leukocyte adhesion molecule and                       |
|        |       |                       |                      | costimulatory receptor                                |
| Itgam  | 1.45  | Integrin alpha M      | CD11b                | Pairs with CD18 to form Mac-1 aka                     |
|        |       |                       |                      | complement receptor 3; mediates leukocyte             |
|        |       |                       |                      | activation, adhesion, chemotaxis, migration,          |
|        |       |                       |                      | phagocytosis, and cell-mediated cytotoxicity;         |
|        |       |                       |                      | serves as a macrophage marker                         |
| Itgb2  | 1.25  | Integrin subunit beta |                      | Pairs with ITGAL to form a receptor for ICAM1,        |
|        |       | 2                     |                      | with ITGAM or ITGAX for iC3b and fibronectin          |
| Jam3   | 1.9   | Junctional adhesion   |                      | Immunoglobulin that mediates tight junctions          |
|        |       | molecule C            |                      | between endothelial cells; mediates                   |
|        |       |                       |                      | transepithelial migration of PMNs                     |
| Lgals3 | 0.794 | Galectin 3            |                      | Galactose-specific lectin that binds IgE;             |
|        |       |                       |                      | involved in acute inflammatory responses,             |
|        |       |                       |                      | including neutrophil activation and adhesion,         |
|        |       |                       |                      | chemoattraction of monocytes macrophages,             |

|               |       |                       |                   | opsonization of apoptotic neutrophils, and    |
|---------------|-------|-----------------------|-------------------|---|
|               |       |                       |                   | activation of mast cells                      |
| Map2k1        | 0.393 | Dual specificity      | MAPK/ERK kinase 1 | Essential component of the MAP kinase signal  |
|               |       | mitogen-activated     | (MEK1)            | transduction pathway; participates in         |
|               |       | protein kinase kinase |                   | numerous biological functions, including cell |
|               |       | 1                     |                   | growth, adhesion, survival, differentiation,  |
|               |       |                       |                   | transcription, metabolism, and cytoskeletal   |
|               |       |                       |                   | remodeling                                    |
| <u>S100a8</u> | 2.22  | S100 calcium-binding  | Calgranulin A     | Calcium- and zinc-binding protein involved in |
|               |       | protein A8            |                   | pro-inflammatory, antimicrobial, oxidant-     |
|               |       |                       |                   | scavenging and apoptosis-inducing activities; |
|               |       |                       |                   | can induce neutrophil chemotaxis, adhesion,   |
|               |       |                       |                   | phagocytosis, and degranulation;              |
|               |       |                       |                   | predominantly found as calprotectin           |
|               |       |                       |                   | (S100A8/A9) which has a wide plethora of      |
|               |       |                       |                   | intra- and extracellular functions, including |
|               |       |                       |                   | adhesion, apoptosis, autophagy, cytoskeletal  |
|               |       |                       |                   | remodeling, cytokine production, chemotaxis,  |
|               |       |                       |                   | migration, inflammation, arachidonic acid     |
|               |       |                       |                   | metabolism, oxidant-scavenging, and PRR       |
|               |       |                       |                   | signaling                                     |
| Sell          | 1.08  | L-selectin            |                   | Mediates cell adhesion by binding to          |
|               |       |                       |                   | glycoproteins on neighboring cells            |

| Thy1         | 0.719 | Thy-1 T cell antigen  |                      | Cell surface glycoprotein involved in cell adhesion and communication in immune and |
|--------------|-------|-----------------------|----------------------|---|
|              |       |                       |                      | nerve cells   |
|              |       | Antige                | n Processing & Prese | entation  |
| <u>Cd1d1</u> | 0.845 | Cluster of            |                      | Murine non-classical class I MHC; primarily   |
| Calai        | 0.043 | differentiation 1 D1  |                      | presents lipid and glycolipid Ags   |
| <u>Cd74</u>  | 0.643 | Cluster of            | MHC class II         | Stabilizes peptide-free class II αβ   |
|              |       | differentiation 74    | gamma chain          | heterodimers during MHC-Ag complex  |
|              |       |                       |                      | formation   |
| <u>Ctss</u>  | 1.43  | Cathepsin S           |                      | Lysosomal protease that participates in   |
|              |       |                       |                      | processing of Ag by MHC class II  |
| Cyfip2       | 1.26  | Cytoplasmic FMR1-     |                      | Involved in T cell adhesion and p53/TP53-   |
|              |       | interacting protein 2 |                      | dependent induction of apoptosis  |
| H2-Ab1       | 0.902 | Histocompatibility 2, |                      | MHC class II molecule; presents Ags to CD4 <sup>+</sup> T                           |
|              |       | class II antigen A,   |                      | cells   |
|              |       | beta 1                |                      |   |
| <u>H2-D1</u> | 0.437 | Histocompatibility 2, |                      | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T                            |
|              |       | D region locus 1      |                      | cells   |
| H2-Q10       | 1.24  | Histocompatibility 2, |                      | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T                            |
|              |       | Q region locus 10     |                      | cells   |

| <u>H2-T23</u>  | 0.993 | Histocompatibility 2,  Q region locus 10 |                    | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T |
|----------------|-------|--|--------------------|--|
|                |       |  |                    |  |
| Psmb10         | 0.461 | Proteasome subunit                       |                    | 20S core β subunit of the proteasome                     |
|                |       | beta type 10                             |                    | involved in Ag processing to generate class I            |
|                |       |  |                    | binding peptides   |
| Tap1           | 0.749 | Transporter antigen                      | Really interesting | ATP-binding cassette transporter that pumps              |
|                |       | peptide 1                                | new gene 4         | degraded cytosolic peptides from the cytosol             |
|                |       |  | (RING4)            | to the ER for packaging into MHC class I                 |
|                |       |  |                    | molecules  |
|                |       |  | Anti-Inflammatory  |  |
|                |       | l I                                      |                    |  |
| <u>Cd200r1</u> | 0.709 | Cluster of                               |                    | Inhibitory receptor for the CD200/OX2 cell               |
|                |       | differentiation 200                      |                    | surface glycoprotein; limits inflammation by             |
|                |       | receptor 1                               |                    | inhibiting the expression of proinflammatory             |
|                |       |  |                    | molecules including TNF $\alpha$ , IFNs, and iNOS        |
| <u>Cd274</u>   | 2.37  | Cluster of                               | Programmed cell    | Ubiquitously expressed ligand for co-                    |
|                |       | differentiation 274                      | death receptor     | inhibitory receptor PD-1; upregulated by                 |
|                |       |  | ligand 1 (PD-L1)   | tumors as an immune evasion strategy                     |
| Ctla4          | 1.29  | Cytotoxic T                              | CD152              | Inhibitory receptor that blocks CD28                     |
|                |       | lymphocyte antigen 4                     |                    | costimulation by competitively binding its               |
|                |       |  |                    | ligands CD80 and CD86                                    |
| Lag3           | 1.73  | Lymphocyte                               | CD223              | Inhibitory receptor on activated T cells; binds          |
|                |       | activating gene 3                        |                    | to ligands, such as FGL1; constitutively                 |

| <u>Mefv</u> | 1.79  | Mediterranean fever                            | Marenostrin; pyrin | expressed on a subset of regulatory Tregs and contributes to their suppressive function; acts as a negative regulator of plasmacytoid DC activation  Involved in the regulation of innate immunity and the inflammatory response in response to |
|-------------|-------|--|--------------------|---|
|             |       |  |                    | IFNy; both stimulates and restrains the inflammasome; also acts as a mediator of pyroptosis   |
| Tnfaip3     | 0.804 | Tumor necrosis factor, alpha-induced protein 3 |                    | Ubiquitin-editing enzyme that complexes with  ITCH to degrade inflammatory signaling  components in the TNF, IL1, and TLR  pathways; targets TRAF2, TRAF6, and IKK  |
|             |       |  | Apoptosis          |   |
| Casp8       | 0.314 | Caspase 8                                      |                    | Cysteine-aspartic acid protease; cleaves and activates effector caspases CASP3, -4, -6, -7, -9, and -10; initiates pyroptosis through cleavage of gasdermin-D; mediates noncanonical cleavage of IL-1β in DCs and macrophages                   |
| Cyfip2      | 1.26  | Cytoplasmic FMR1-<br>interacting protein 2     |                    | Involved in T cell adhesion and p53/TP53-<br>dependent induction of apoptosis   |

| Eas           | 1.52 | Fragment anestes:    |                   | Cell surface death receptor; interaction with    |
|---------------|------|----------------------|-------------------|--|
| <u>Fas</u>    | 1.52 | Fragment apoptosis   |                   | Cen surface death receptor; interaction with     |
|               |      | stimulating          |                   | FAS-ligand triggers an apoptotic signaling       |
|               |      |                      |                   | cascade; also activates NFкВ, ERK1, and          |
|               |      |                      |                   | МАРК8  |
|               |      |                      |                   |  |
| <u>S100a8</u> | 2.22 | S100 calcium-binding | Calgranulin A     | Calcium- and zinc-binding protein involved in    |
|               |      | protein A8           |                   | pro-inflammatory, antimicrobial, oxidant-        |
|               |      |                      |                   | scavenging and apoptosis-inducing activities;    |
|               |      |                      |                   | can induce neutrophil chemotaxis, adhesion,      |
|               |      |                      |                   | phagocytosis, and degranulation;                 |
|               |      |                      |                   | predominantly found as calprotectin              |
|               |      |                      |                   | (S100A8/A9) which has a wide plethora of         |
|               |      |                      |                   | intra- and extracellular functions, including    |
|               |      |                      |                   | adhesion, apoptosis, autophagy, cytoskeletal     |
|               |      |                      |                   | remodeling, cytokine production, chemotaxis,     |
|               |      |                      |                   | migration, inflammation, arachidonic acid        |
|               |      |                      |                   | metabolism, oxidant-scavenging, and PRR          |
|               |      |                      |                   | signaling  |
| Tnfsf14       | 1.39 | TNF superfamily      | CD258; LIGHT;     | Cytokine that binds to TNFRSF3/LTBR and          |
|               |      | member 14            | Herpesvirus entry | TNFRSF14/HVEM; delivers costimulation to T       |
|               |      |                      | mediator ligand   | cells; triggers apoptosis of various tumor cells |
|               |      |                      | (HVEML)           |  |
|               |      |                      | (114 - 141-)      |  |
|               |      |                      | Autophagy         |  |
|               |      |                      |                   |  |

| Irgm2          | 0.453 | Immunity-related     | Interferon-         | Function not fully known, but most likely     |
|----------------|-------|----------------------|---------------------|---|
|                |       | GTPase family M      | inducible protein 1 | regulates autophagy and pro-inflammatory      |
|                |       | member 2             | (IFI1)              | cytokine production                           |
| <u>\$100a8</u> | 2.22  | S100 calcium-binding | Calgranulin A       | Calcium- and zinc-binding protein involved in |
|                |       | protein A8           |                     | pro-inflammatory, antimicrobial, oxidant-     |
|                |       |                      |                     | scavenging and apoptosis-inducing activities; |
|                |       |                      |                     | can induce neutrophil chemotaxis, adhesion,   |
|                |       |                      |                     | phagocytosis, and degranulation;              |
|                |       |                      |                     | predominantly found as calprotectin           |
|                |       |                      |                     | (S100A8/A9) which has a wide plethora of      |
|                |       |                      |                     | intra- and extracellular functions, including |
|                |       |                      |                     | adhesion, apoptosis, autophagy, cytoskeletal  |
|                |       |                      |                     | remodeling, cytokine production, chemotaxis,  |
|                |       |                      |                     | migration, inflammation, arachidonic acid     |
|                |       |                      |                     | metabolism, oxidant-scavenging, and PRR       |
|                |       |                      |                     | signaling                                     |
| <u>Ubc</u>     | 0.263 | Polyubiquitin C      |                     | Serves various roles, including innate        |
|                |       |                      |                     | immunity, DNA repair, and stimulation of      |
|                |       |                      |                     | autophagy and the proteasomal response        |
|                |       | R                    | Cell-associated Gen | Δς.   |
|                |       | Ü                    | cen associated dell |   |
| <u>Cd37</u>    | 1.26  | Cluster of           | Tetraspanin-26      | Cell surface glycoprotein known to complex    |
|                |       | differentiation 37   |                     | with integrins and other transmembrane 4      |

|             |       |                     |                      | superfamily proteins; may play a role in T cell-  |
|-------------|-------|---------------------|----------------------|---|
|             |       |                     |                      | B cell interactions                               |
| <u>Cd48</u> | 1.5   | Cluster of          | B-lymphocyte         | B cell-specific cellular differentiation Ag; when |
|             |       | differentiation 48  | activation marker    | bound to CD2, promotes T cell activation, and     |
|             |       |                     | (BLAST-1); signaling | the formation of lipid rafts and caveolae for     |
|             |       |                     | lymphocytic          | macrophages                                       |
|             |       |                     | activation molecule  |   |
|             |       |                     | 2 (SLAMF2)           |   |
| <u>Cd69</u> | 0.994 | Cluster of          | C-type lectin        | Signal transmitting receptor in lymphocytes,      |
|             |       | differentiation 69  | domain family 2,     | NK cells, and platelets; induced upon T cell      |
|             |       |                     | member C             | activation; involved in lymphocyte                |
|             |       |                     |                      | proliferation                                     |
| Cd79b       | 2.43  | Cluster of          | B29                  | One of the two flanking proteins that initiate    |
|             |       | differentiation 79b |                      | signaling downstream of the BCR                   |
| Icosl       | 0.484 | Inducible T cell    | CD275                | Ligand for T cell-specific co-receptor ICOS; also |
|             |       | costimulator ligand |                      | induces B cell proliferation and plasma cell      |
|             |       |                     |                      | differentiation                                   |
| Fcgr1       | 1.16  | Fc fragment of IgG  | CD64                 | High affinity receptor for the Fc region of γ-    |
|             |       | receptor la         |                      | lgs; functions in both innate and adaptive        |
|             |       |                     |                      | immune responses                                  |
| Fcgr2b      | 0.915 | Fc fragment of IgG  | CD32                 | Low affinity receptor for the Fc region of        |
|             |       | receptor IIb        |                      | complexed or aggregated γ-lgs; involved in a      |
|             |       |                     |                      | variety of effector and regulatory functions      |

|        |       |                       |            | such as phagocytosis of immune complexes          |
|--------|-------|-----------------------|------------|---|
|        |       |                       |            | and modulation of Ab production by B cells;       |
|        |       |                       |            | essential for the maintenance of humoral          |
|        |       |                       |            | tolerance; acts as a late checkpoint at the       |
|        |       |                       |            | level of class-switched memory B cells,           |
|        |       |                       |            | plasmablasts or plasma cells; regulates           |
|        |       |                       |            | plasma-cell homeostasis and survival              |
| Mef2c  | 0.998 | Myocyte enhancer      |            | Transcriptional activator that binds specifically |
|        |       | factor 2c             |            | to the MEF2 element present in the regulatory     |
|        |       |                       |            | regions of many muscle-specific genes;            |
|        |       |                       |            | controls cardiac morphogenesis and                |
|        |       |                       |            | myogenesis, and is also involved in vascular      |
|        |       |                       |            | development; required for B cell survival and     |
|        |       |                       |            | proliferation in response to BCR stimulation,     |
|        |       |                       |            | efficient IgG1 Ab responses to T cell-            |
|        |       |                       |            | dependent Ags, and for normal induction of        |
|        |       |                       |            | GC B cells  |
| Ms4a1  | 3.3   | Membrane spanning     | CD20; Bp35 | B cell-specific membrane protein that             |
|        |       | 4-domains A1          |            | functions as a store-operated Ca(2+) channel      |
|        |       |                       |            | component, promoting Ca(2+) influx after BCR      |
|        |       |                       |            | activation  |
|        |       |                       |            |   |
| Pik3cd | 0.584 | Phosphatidylinositol- |            | A subunit of PI3K; acts downstream of TLR4,       |
|        |       | 4,5-bisphosphate 3-   |            | TCR, BCR, and CD40; contributes to T helper       |
|        |       |                       |            | cell expansion, mast cell development, and        |
|        |       |                       |            |   |

|             |       | kinase catalytic      |                     | neutrophil chemotaxis, extravasation, and                |
|-------------|-------|-----------------------|---------------------|--|
|             |       | subunit delta isoform |                     | respiratory burst  |
| Spn         | 0.72  | Sialophorin           | Leukosialin; CD43   | Cell surface sialoglycoprotein expressed by T            |
|             |       |                       |                     | cells, B cells, monocytes, and granulocytes;             |
|             |       |                       |                     | promotes lymph node localization in T cells;             |
|             |       |                       |                     | shunts T cells away from the T <sub>H</sub> 2 phenotype  |
|             |       |                       |                     | and towards T <sub>H</sub> 1; promotes the expression of |
|             |       |                       |                     | IFNγ in CD4 <sup>+</sup> T cells                         |
| Syk         | 0.969 | Spleen-associated     |                     | Critical kinase that transmits signals from the          |
|             |       | tyrosine kinase       |                     | TCR and BCR  |
|             |       |                       | Cell Cycle          |  |
| Cdkn1a      | 0.986 | Cyclin dependent      | p21; CDK-           | Binds to and inhibits cyclin-dependent kinase            |
|             |       | kinase inhibitor 1A   | interaction protein | activity, preventing phosphorylation of critical         |
|             |       |                       | 1 (CIP1)            | cyclin-dependent kinase substrates and                   |
|             |       |                       |                     | blocking cell cycle progression                          |
|             |       |                       | Chemotaxis          |  |
| Ccl2        | 1.4   | C-C motif chemokine   |                     | Chemoattractant ligand for CCR2 and -4;                  |
|             |       | ligand 2              |                     | attracts monocytes and basophils                         |
| <u>Ccl3</u> | 3.09  | C-C motif chemokine   | Macrophage          | Chemoattractant ligand for CCR1, -4, and -5              |
|             |       | ligand 3              | inflammatory        |  |
|             |       |                       | protein 1α (MIP1α)  |  |

| Ccl4     | 2.66  | C-C motif chemokine | Macrophage         | Chemoattractant for NK cells and monocytes;    |
|----------|-------|---------------------|--------------------|--|
|          |       | ligand 4            | inflammatory       | binds to CCR5 receptors                        |
|          |       |                     | protein 1β (MIP1β) |  |
|          |       |                     |                    |  |
| Ccl5     | 2.47  | C-C motif chemokine | Regulated upon     | Chemoattractant ligand for CCR1, -3, -4, and - |
|          |       | ligand 5            | activation,        | 5; attracts blood monocytes, memory T helper   |
|          |       |                     | normally T-        | cells and eosinophils; causes the release of   |
|          |       |                     | expressed, and     | histamine from basophils and activates         |
|          |       |                     | presumably         | eosinophils                                    |
|          |       |                     | secreted (RANTES)) |  |
|          |       |                     |                    |  |
| Ccl7     | 1.22  | C-C motif chemokine | Monocte            | General chemokine that recruits leukocytes to  |
|          |       | ligand 7            | chemotactic        | infected tissues; mainly observed in monocyte  |
|          |       |                     | protein 3 (MCP3)   | mobilization                                   |
| Ccl8     | 1.14  | C-C motif chemokine | Monocyte           | General chemokine that recruits leukocytes to  |
| <u> </u> | 1.1.  | ligand 8            | chemoattractant    | infected tissues                               |
|          |       | ligaliu o           |                    | illiected tissues                              |
|          |       |                     | protein 2 (MCP2)   |  |
| Ccl12    | 0.773 | C-C motif chemokine | Monocyte           | Chemoattractant specific for eosinophils,      |
|          |       | ligand 12           | chemotactic        | monocytes, and lymphocytes; found primarily    |
|          |       |                     | protein 5 (MCP5)   | in the lymph nodes and thymus, but can be      |
|          |       |                     |                    | strongly expressed by macrophages              |
|          |       |                     |                    |  |
| Ccr1     | 1.73  | C-C motif chemokine | MIP1α receptor     | Receptor for CCL3, -5, -7, and -23             |
|          |       | receptor 1          |                    |  |
|          |       |                     |                    |  |

| Ccr7   | 1.74  | C-C chemokine receptor type 7      | CD197  | Chemokine receptor that activates B and T cells and promotes their homing to secondary lymphoid organs; also stimulates DC expression of MHC class I and II |
|--------|-------|------------------------------------|--|---|
| Ccrl2  | 1.44  | C-C chemokine receptor-like 2      |  | Stabilizes TLR4 surface expression on macrophages   |
| Cxcl1  | 1.5   | C-X-C motif chemokine ligand 1     | GRO1 oncogene  | Chemoattractant ligand for CXCR2; plays a role in inflammation and as a chemoattractant for neutrophils   |
| Cxcl2  | 2.49  | C-X-C motif chemokine ligand 2     | Macrophage inflammatory protein 2-alpha (MIP2α); GRO2 oncogene | Chemokine produced by activated monocytes and neutrophils and expressed at sites of inflammation  |
| Cxcl3  | 3.04  | C-X-C motif chemokine ligand 3     | GRO3 oncogene  | Ligand for CXCR2; attracts neutrophils  |
| Cxcl9  | 0.774 | C-X-C motif<br>chemokine ligand 9  | Humig  | Chemoattractant ligand for CXCR3; attracts activated T cells  |
| Cxcl10 | 0.874 | C-X-C motif<br>chemokine ligand 10 | IFNγ-induced<br>protein 10 (IP-10)                             | Macrophage, DC, T cell, and NK cell chemattractant secreted by several cell types in response to IFNγ; binds to CXCR3                                       |

| Cxcl11 | 0.954 | C-X-C motif chemokine ligand 11    |                           | Dominant ligand for CXCR3; attracts activated  T cells; strongly induced by IFNγ   |
|--------|-------|------------------------------------|---------------------------|--|
| Cxcl16 | 0.961 | C-X-C motif<br>chemokine ligand 16 |                           | Chemoattractant for T cells and NKT cells produced by DCs in response to IFN $\gamma$ and TNF $\alpha$   |
| Cxcr2  | 1.62  | C-X-C motif chemokine receptor     | CD182; IL-8<br>receptor B | Receptor for IL-8 and CXCL3; powerful chemoattractant for neutrophils  |
| Cxcr4  | 0.947 | C-X-C motif chemokine receptor     | CD184; fusin              | Alpha-chemokine receptor specific for SDF1  aka CXCL12   |
| Cxcr6  | 1.69  | C-X-C motif chemokine receptor     | CD186                     | Receptor for the C-X-C chemokine CXCL16; expressed in several T lymphocyte subsets and bone marrow stromal cells   |
| Isg15  | 2.02  | Interferon-stimulated gene 15      |                           | Ubiquitin-like protein that binds intracellular target proteins upon activation by IFN $\alpha$ or $\beta$ ; can also be secreted to induce NK cell proliferation, act as a chemoattractant for neutrophils, and induce IFN $\gamma$ upon binding to ITGAL/ITGB2 |
| Itgam  | 1.45  | Integrin alpha M                   | CD11b                     | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration,   |

|                |       |                                 |                    | phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker  |
|----------------|-------|---------------------------------|--------------------|--|
| <u>\$100a8</u> | 2.22  | S100 calcium-binding protein A8 | Calgranulin A      | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |
|                |       | Comple                          | ement & Humoral Im | nmunity  |
| C1s1           | 0.85  | Complement component 1s         |                    | Serine protease that enzymatically cleaves C4 and C2   |
| C3             | 0.656 | Complement component 3          |                    | Cleaved by C3 convertase to form C3a and C3b, an anaphalotoxin and an opsonizing agent, respectively   |
| C4b            | 0.624 | Complement<br>component 4B      |                    | Mediates interactions between Ab-bound Ags and other complement components   |

| <u>Cfd</u> | 2.97  | Complement factor D  | Adipsin            | Chymotrypsin-family peptidase that cleaves     |  |  |
|------------|-------|----------------------|--------------------|--|--|--|
| <u>Cra</u> | 2.97  | Complement factor D  | Adipsin            | factor B when the latter is complexed with     |  |  |
|            |       |                      |                    | factor C3b, activating C3 convertase           |  |  |
| Fcgr1      | 1.16  | Fc fragment of IgG   | CD64               | High affinity receptor for the Fc region of γ- |  |  |
|            |       | receptor la          |                    | Igs; functions in both innate and adaptive     |  |  |
|            |       |                      |                    | immune responses                               |  |  |
| Fcgr2b     | 0.915 | Fc fragment of IgG   | CD32               | Low affinity receptor for the Fc region of     |  |  |
|            |       | receptor IIb         |                    | complexed or aggregated γ-lgs; involved in a   |  |  |
|            |       |                      |                    | variety of effector and regulatory functions   |  |  |
|            |       |                      |                    | such as phagocytosis of immune complexes       |  |  |
|            |       |                      |                    | and modulation of Ab production by B cells;    |  |  |
|            |       |                      |                    | essential for the maintenance of humoral       |  |  |
|            |       |                      |                    | tolerance; acts as a late checkpoint at the    |  |  |
|            |       |                      |                    | level of class-switched memory B cells,        |  |  |
|            |       |                      |                    | plasmablasts or plasma cells; regulates        |  |  |
|            |       |                      |                    | plasma-cell homeostasis and survival           |  |  |
| Fcgr4      | 2.36  | Fragment             | Fc receptor-like 3 | Putative mouse ortholog to human FcγRIIIA      |  |  |
|            |       | crystallizable gamma | (Fcrl3); CD16-2    |  |  |  |
|            |       | receptor 4           |                    |  |  |  |
|            |       |                      | Costimulation      |  |  |  |
|            |       |                      |                    |  |  |  |

| <u>Cd28</u>  | 1.51  | Cluster of          |       | Essential T cell co-receptor that enhances T        |
|--------------|-------|---------------------|-------|---|
|              |       | differentiation 28  |       | cell activation, proliferation, cytokine            |
|              |       |                     |       | production, and survival; binds to CD80 and         |
|              |       |                     |       | CD86  |
| <u>Cd40</u>  | 1.77  | Cluster of          |       | APC-expressed costimulatory protein that            |
|              |       | differentiation 40  |       | binds to CD40L on CD4 <sup>+</sup> T cells, causing |
|              |       |                     |       | activation of both                                  |
| <u>Cd86</u>  | 0.886 | Cluster of          | B7-2  | One of the two ligands for the CD28                 |
|              |       | differentiation 86  |       | costimulatory receptor and the CTLA4                |
|              |       |                     |       | inhibitory receptor, the other being CD80           |
| <u>lcam1</u> | 0.997 | Intracellular       | CD54  | Cell surface glycoprotein that serves as strong     |
|              |       | adhesion molecule 1 |       | adhesive ligand for LFA-1; important for            |
|              |       |                     |       | leukocyte mobility and costimulation                |
| <u>lcos</u>  | 1.58  | Inducible T cell    | CD278 | Enhances all basic T cell responses to foreign      |
|              |       | costimulator        |       | Ag; essential both for efficient interaction        |
|              |       |                     |       | between T and B cells and for normal Ab             |
|              |       |                     |       | responses to T cell-dependent Ags                   |
| Icosl        | 0.484 | Inducible T cell    | CD275 | Ligand for T cell-specific co-receptor ICOS; also   |
|              |       | costimulator ligand |       | induces B cell proliferation and plasma cell        |
|              |       |                     |       | differentiation                                     |

| Ptprc       | 0.739 | Protein tyrosine    | CD45; leukocyte   | Delivers costimulation during T cell activation   |
|-------------|-------|---------------------|-------------------|---|
| . 46. 5     |       |                     |                   |   |
|             |       | phosphatase         | common antigen    | upon binding to its ligand DPP4;                  |
|             |       | receptor type C     | (LCA)             | dephosphorylates Lyn and suppresses JAK           |
|             |       |                     |                   | kinases   |
|             |       |                     |                   |   |
| Tnfsf14     | 1.39  | TNF superfamily     | CD258; LIGHT;     | Cytokine that binds to TNFRSF3/LTBR and           |
|             |       | member 14           | Herpesvirus entry | TNFRSF14/HVEM; delivers costimulation to T        |
|             |       |                     | mediator ligand   | cells; triggers apoptosis of various tumor cells  |
|             |       |                     | (HVEML)           |   |
|             |       |                     |                   |   |
|             |       |                     | Cytokines         |   |
|             |       |                     |                   |   |
| <u>ll1a</u> | 2.61  | Interleukin 1 alpha | Hematopoietin-1   | Cytokine produced by monocytes and                |
|             |       |                     |                   | macrophages in response to cell injury;           |
|             |       |                     |                   | stimulates thymocyte proliferation by             |
|             |       |                     |                   | inducing IL-2 release; also stimulates B cell     |
|             |       |                     |                   | maturation and proliferation, and fibroblast      |
|             |       |                     |                   | growth factor activity                            |
|             |       |                     |                   | <b>3</b> • • • • • • • • • • • • • • • • • • •    |
| <u>II1b</u> | 1.3   | Interleukin 1 beta  | Catabolin         | One of the two primary inflammatory               |
|             |       |                     |                   | cytokines produced by the inflammasome (the       |
|             |       |                     |                   | other one being IL-18); induces neutrophil        |
|             |       |                     |                   |   |
|             |       |                     |                   | influx and activation, T cell activation and      |
|             |       |                     |                   | cytokine production, B cell activation and Ab     |
|             |       |                     |                   | production, fibroblast proliferation, and         |
|             |       |                     |                   | collagen production; synergizes with IL-12 to     |
|             |       |                     |                   | induce IFNγ synthesis from T <sub>H</sub> 1 cells |
|             |       |                     |                   |   |

| ll2ra | 1.06  | Interleukin 2 receptor subunit alpha   | CD25                        | Alpha chain of the IL-2 receptor   |
|-------|-------|--|-----------------------------|--|
| II6   | 2.11  | Interleukin 6                          |                             | Pro-inflammatory cytokine that signals through the JAK and STAT pathways   |
| II7r  | 1.69  | Interleukin 7 receptor                 | CD127                       | Receptor for IL-7  |
| Mif   | 0.328 | Macrophage migration inhibitory factor | L-dopachrome<br>tautomerase | Pro-inflammatory cytokine that promotes  macrophage function through suppression of anti-inflammatory effects of glucocorticoids   |
| Tgfb1 | 0.498 | Transforming growth factor beta 1      |                             | Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to neurodegeneration; can induce EMT and cell migration in various cell types; frequently acts as an immunosuppressive cytokine in the TME |
| Tnf   | 3.07  | Tumor necrosis<br>factor               | Cachectin                   | Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines   |

| Tnfsf14       | 1.39 | TNF superfamily      | CD258; LIGHT;      | Cytokine that binds to TNFRSF3/LTBR and          |
|---------------|------|----------------------|--------------------|--|
|               |      | member 14            | Herpesvirus entry  | TNFRSF14/HVEM; delivers costimulation to T       |
|               |      |                      | mediator ligand    | cells; triggers apoptosis of various tumor cells |
|               |      |                      | (HVEML)            |  |
|               |      |                      |                    |  |
|               |      |                      | Cytotoxicity       |  |
| <u>Gzmb</u>   | 2.9  | Granzyme B           | Fragmentin 2       | Abundant protease in the cytosolic granules of   |
|               |      |                      |                    | cytotoxic T and NK cells that activates          |
|               |      |                      |                    | caspase-mediated cell death when delivered       |
|               |      |                      |                    | into the target cell through the immunological   |
|               |      |                      |                    | synapse  |
| <u>Gzmk</u>   | 2.49 | Granzyme K           | Tryptase II        | Granule-secreted, pro-apoptotic serine           |
| <u>UZITIK</u> | 2.49 | Granzyme K           | пурсазеп           |  |
|               |      |                      |                    | protease found in the cytoplasmic granules of    |
|               |      |                      |                    | CTLs   |
| Fcgr4         | 2.36 | Fragment             | Fc receptor-like 3 | Putative mouse ortholog to human FcyRIIIA        |
|               |      | crystallizable gamma | (Fcrl3); CD16-2    |  |
|               |      | receptor 4           |                    |  |
|               |      |                      |                    |  |
| Itgam         | 1.45 | Integrin alpha M     | CD11b              | Pairs with CD18 to form Mac-1 aka                |
|               |      |                      |                    | complement receptor 3; mediates leukocyte        |
|               |      |                      |                    | activation, adhesion, chemotaxis, migration,     |
|               |      |                      |                    | phagocytosis, and cell-mediated cytotoxicity;    |
|               |      |                      |                    | serves as a macrophage marker                    |
|               |      |                      |                    |  |

| <u>\$100a8</u> | 2.22  | S100 calcium-binding | Calgranulin A           | Calcium- and zinc-binding protein involved in    |
|----------------|-------|----------------------|-------------------------|--|
|                |       | protein A8           |                         | pro-inflammatory, antimicrobial, oxidant-        |
|                |       |                      |                         | scavenging and apoptosis-inducing activities;    |
|                |       |                      |                         | can induce neutrophil chemotaxis, adhesion,      |
|                |       |                      |                         | phagocytosis, and degranulation;                 |
|                |       |                      |                         | predominantly found as calprotectin              |
|                |       |                      |                         | (S100A8/A9) which has a wide plethora of         |
|                |       |                      |                         | intra- and extracellular functions, including    |
|                |       |                      |                         | adhesion, apoptosis, autophagy, cytoskeletal     |
|                |       |                      |                         | remodeling, cytokine production, chemotaxis,     |
|                |       |                      |                         | migration, inflammation, arachidonic acid        |
|                |       |                      |                         | metabolism, oxidant-scavenging, and PRR          |
|                |       |                      |                         | signaling  |
|                |       |                      | Dendritic Cell Function |  |
|                |       | ,                    | renantic cen ranctic    | JII  |
| Casp8          | 0.314 | Caspase 8            |                         | Cysteine-aspartic acid protease; cleaves and     |
|                |       |                      |                         | activates effector caspases CASP3, -4, -6, -7, - |
|                |       |                      |                         | 9, and -10; initiates pyroptosis through         |
|                |       |                      |                         | cleavage of gasdermin-D; mediates                |
|                |       |                      |                         | noncanonical cleavage of IL-1β in DCs and        |
|                |       |                      |                         | macrophages                                      |
| Tigit          | 0.861 | T cell               |                         | Binds with high affinity to the poliovirus       |
|                |       | immunoreceptor       |                         | receptor, causing increased secretion of IL-10,  |
|                |       | with Ig and ITIM     |                         | decreased secretion of IL-12B, and               |
|                |       | domains              |                         |  |
|                |       |                      |                         |  |

|            |                      |  |                                | suppressing T cell activation by promoting the generation of mature immunoregulatory DCs   |  |  |  |
|------------|----------------------|--|--------------------------------|--|--|--|--|
|            | Growth/Proliferation |  |                                |  |  |  |  |
| Lcn2       | 1.69                 | Lipocalin 2  | Neutrophil gelatinase-         | Neutrophil-secreted factor that sequesters iron-containing siderophores; also functions  |  |  |  |
|            |                      |  | associated lipocalin<br>(NGAL) | as a growth factor   |  |  |  |
| Map2k1     | 0.393                | Dual specificity mitogen-activated protein kinase kinase | MAPK/ERK kinase 1 (MEK1)       | Essential component of the MAP kinase signal transduction pathway; participates in numerous biological functions, including cell growth, adhesion, survival, differentiation, transcription, metabolism, and cytoskeletal remodeling   |  |  |  |
| Tgfb1      | 0.498                | Transforming growth factor beta 1                        |                                | Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to neurodegeneration; can induce EMT and cell migration in various cell types; frequently acts as an immunosuppressive cytokine in the TME |  |  |  |
| <u>Yy1</u> | 0.196                | Yin yang 1   |                                | Ubiquitous factor that serves as a transcriptional "switch", either promoting or   |  |  |  |

|             |       |                                    |                        | repressing the transcription of numerous genes through the selective recruitment of either histone deacetylases or acetyltransferases; plays a fundamental role in diverse processes, such as differentiation, replication, and cellular proliferation |
|-------------|-------|------------------------------------|------------------------|--|
|             |       |                                    | Inflammation           |  |
| Bst2        | 1.13  | Bone marrow stromal cell antigen 2 | Tethrin; CD317         | IFN-induced antiviral factor that blocks budding of enveloped viruses by directly tethering nascent virions to the membranes of infected cells   |
| Casp8       | 0.314 | Caspase 8                          |                        | Cysteine-aspartic acid protease; cleaves and activates effector caspases CASP3, -4, -6, -7, -9, and -10; initiates pyroptosis through cleavage of gasdermin-D; mediates noncanonical cleavage of IL-1β in DCs and macrophages                          |
| <u>Cd38</u> | 1.41  | Cluster of differentiation 38      | ADP-ribosyl cyclase  1 | Synthesizes the second messengers cyclic  ADP-ribose and NADPH; appears to play a  critical role in inflammation, although its exact  immunological function(s) remain(s) poorly  defined  |

| Cebpb       | 0.724 | CCAAT/enhancer-<br>binding protein beta |           | Critical macrophage TF that promotes expression of several acute-phase and inflammatory cytokine genes, including II6   |
|-------------|-------|---|-----------|---|
| <u>Ctsh</u> | 0.906 | Cathepsin H                             |           | Lysosomal protease; increased in macrophages in response to IFNy  |
| II1b        | 1.3   | Interleukin 1 beta                      | Catabolin | One of the two primary inflammatory cytokines produced by the inflammasome (the other one being IL-18); induces neutrophil influx and activation, T cell activation and cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNy synthesis from T <sub>H</sub> 1 cells |
| II6         | 2.11  | Interleukin 6                           |           | Pro-inflammatory cytokine that signals through the JAK and STAT pathways  |
| <u>Irf1</u> | 1.2   | Interferon regulatory factor 1          |           | Transcriptional regulator that promotes inflammatory innate and adaptive immune responses   |
| <u>Irf7</u> | 1.71  | Interferon regulatory factor 7          |           | Key transcriptional regulator of type I IFN-dependent immune responses; promotes $transcription\ of\ IFN\alpha\ and\ -\beta$  |

| Into         | 0.044 | Interferen                     |                     | TE that regulates of linear a committee set in  |
|--------------|-------|--------------------------------|---------------------|---|
| <u>Irf8</u>  | 0.841 | Interferon regulatory factor 8 |                     | TF that regulates of lineage commitment in myeloid cell maturation; promotes monocyte |
|              |       | Tactor 8                       |                     | myelola celi maturation; promotes monocyte  |
|              |       |                                |                     | and plasmacytoid DC development   |
| Irgm2        | 0.453 | Immunity-related               | Interferon-         | Function not fully known, but most likely   |
|              |       | GTPase family M                | inducible protein 1 | regulates autophagy and pro-inflammatory  |
|              |       | member 2                       | (IFI1)              | cytokine production   |
| <u>Isg15</u> | 2.02  | Interferon-stimulated          |                     | Ubiquitin-like protein that binds intracellular                                       |
|              |       | gene 15                        |                     | target proteins upon activation by IFN $\alpha$ or $\beta$ ;                          |
|              |       |                                |                     | can also be secreted to induce NK cell  |
|              |       |                                |                     | proliferation, act as a chemoattractant for   |
|              |       |                                |                     | neutrophils, and induce IFNγ upon binding to  |
|              |       |                                |                     | ITGAL/ITGB2   |
| Isg20        | 0.904 | Interferon-stimulated          |                     | IFN-induced antiviral exoribonuclease that  |
|              |       | gene 20                        |                     | acts on ssRNA with minor activity towards   |
|              |       |                                |                     | ssDNA   |
| Jak1         | 0.337 | Janus kinase 1                 |                     | Essential tyrosine kinase involved signal   |
|              |       |                                |                     | transduction in type I and II cytokines and IFNs                                      |
| Mif          | 0.328 | Macrophage                     | L-dopachrome        | Pro-inflammatory cytokine that promotes   |
|              |       | migration inhibitory           | tautomerase         | macrophage function through suppression of  |
|              |       | factor                         |                     | anti-inflammatory effects of glucocorticoids  |
| Mefv         | 1.79  | Mediterranean fever            | Marenostrin; pyrin  | Involved in the regulation of innate immunity   |
|              |       |                                |                     | and the inflammatory response in response to  |
|              |       |                                |                     | IFNγ; both stimulates and restrains the   |
|              |       | 1                              | l                   |   |

|               |      |                        |                   | inflammasome; also acts as a mediator of      |
|---------------|------|------------------------|-------------------|---|
|               |      |                        |                   | pyroptosis                                    |
| Nlrp3         | 1.47 | NACHT domain-,         | Cryopyrin         | PRR with a wide diversity of recognized       |
|               |      | leucine-rich repeat-,  |                   | targets that activates the NLRP3              |
|               |      | and PYD-containing     |                   | inflammasome consisting of NLRP3, PYCARD,     |
|               |      | protein 3              |                   | and caspase-1/-8                              |
| Nos2          | 3.66 | Inducible nitric oxide |                   | Produces reactive oxygen species and          |
|               |      | synthase (iNOS)        |                   | contributes to inflammatory cytokine          |
|               |      |                        |                   | production                                    |
| <u>S100a8</u> | 2.22 | S100 calcium-binding   | Calgranulin A     | Calcium- and zinc-binding protein involved in |
|               |      | protein A8             |                   | pro-inflammatory, antimicrobial, oxidant-     |
|               |      |                        |                   | scavenging and apoptosis-inducing activities; |
|               |      |                        |                   | can induce neutrophil chemotaxis, adhesion,   |
|               |      |                        |                   | phagocytosis, and degranulation;              |
|               |      |                        |                   | predominantly found as calprotectin           |
|               |      |                        |                   | (S100A8/A9) which has a wide plethora of      |
|               |      |                        |                   | intra- and extracellular functions, including |
|               |      |                        |                   | adhesion, apoptosis, autophagy, cytoskeletal  |
|               |      |                        |                   | remodeling, cytokine production, chemotaxis,  |
|               |      |                        |                   | migration, inflammation, arachidonic acid     |
|               |      |                        |                   | metabolism, oxidant-scavenging, and PRR       |
|               |      |                        |                   | signaling                                     |
| Spn           | 0.72 | Sialophorin            | Leukosialin; CD43 | Cell surface sialoglycoprotein expressed by T |
|               |      |                        |                   | cells, B cells, monocytes, and granulocytes;  |

| Tbk1          | 0.767 | TANK-binding kinase                                  |                                   | promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNγ in CD4 <sup>+</sup> T cells  Coordinates the activation of IRF3 and NFκB and induction of type I IFNs |
|---------------|-------|--|-----------------------------------|--|
| Tnf           | 3.07  | Tumor necrosis<br>factor                             | Cachectin                         | Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines   |
| Traf6         | 0.697 | Tumor necrosis  factor receptor- associated factor 6 |                                   | Adaptor protein that acts in the CD40 signaling cascade; promotes inflammation, IL- $ 6 \text{, and TNF} \alpha $  |
|               |       |  | Inhibition                        |  |
| <u>Bcl2l1</u> | 0.665 | B cell lymphoma 2                                    | Protein<br>phosphatase 1          | Potent inhibitor of caspase-mediated cell death  |
| <u>Cd47</u>   | 0.412 | Cluster of differentiation 47                        | Integrin-associated protein (IAP) | Partners with membrane integrins to serve as an inhibitor of phagocytosis  |
| Cd200r1       | 0.709 | Cluster of differentiation 200 receptor 1            |                                   | Inhibitory receptor for the CD200/OX2 cell surface glycoprotein; limits inflammation by inhibiting the expression of proinflammatory molecules including TNF $\alpha$ , IFNs, and iNOS   |

| <u>Cd274</u> | 2.37  | Cluster of           | Programmed cell     | Ubiquitously expressed ligand for co-            |
|--------------|-------|----------------------|---------------------|--|
|              |       | differentiation 274  | death receptor      | inhibitory receptor PD-1; upregulated by         |
|              |       |                      | ligand 1 (PD-L1)    | tumors as an immune evasion strategy             |
| Cdkn1a       | 0.986 | Cyclin dependent     | p21; CDK-           | Binds to and inhibits cyclin-dependent kinase    |
|              |       | kinase inhibitor 1A  | interaction protein | activity, preventing phosphorylation of critical |
|              |       |                      | 1 (CIP1)            | cyclin-dependent kinase substrates and           |
|              |       |                      |                     | blocking cell cycle progression                  |
| <u>Ctla4</u> | 1.29  | Cytotoxic T          | CD152               | Inhibitory receptor that blocks CD28             |
|              |       | lymphocyte antigen 4 |                     | costimulation by competitively binding its       |
|              |       |                      |                     | ligands CD80 and CD86                            |
| Cyld         | 0.381 | Cylindromatosis      |                     | Inhibits NFkB activation by deubiquitinating     |
|              |       | lysine 63            |                     | upstream signaling factors; inhibits Wnt         |
|              |       | deubiquitinase       |                     | signaling; restricts polyubiquitination of RIPK1 |
|              |       |                      |                     | and -2, thereby limiting necroptosis             |
| ldo1         | 1.39  | Indoleamine 2,3-     |                     | Initiates catabolism of tryptophan; limits       |
|              |       | dioxygenase 1        |                     | immunopathology by inhibiting T cell division    |
| <u>Lag3</u>  | 1.73  | Lymphocyte           | CD223               | Inhibitory receptor on activated T cells; binds  |
|              |       | activating gene 3    |                     | to ligands, such as FGL1; constitutively         |
|              |       |                      |                     | expressed on a subset of regulatory Tregs and    |
|              |       |                      |                     | contributes to their suppressive function; acts  |
|              |       |                      |                     | as a negative regulator of plasmacytoid DC       |
|              |       |                      |                     | activation                                       |
|              |       |                      |                     |  |

| Nfkbia       | 1.12  | Nuclear factor kappa  B inhibitor alpha             |       | Inhibits activity of REL dimers by masking of their nuclear localization signals   |
|--------------|-------|---|-------|--|
| NIrc5        | 0.633 | NLR family CARD domain containing 5                 | NOD27 | Inhibits NFKB and type I IFN signaling  pathways; may also regulate the type II IFN  signaling pathway   |
| Socs1        | 1.27  | Suppressor of cytokine signaling 1                  |       | Inhibits JAK proteins; negative regulator of IL-   |
| Tank         | 0.743 | TRAF family member-<br>associated NFkB<br>activator |       | Inhibitory protein that sequesters TRAFs in the cytoplasm, constitutively binds TBK1, and serves as a negative regulator of NFkB   |
| Tgfb1        | 0.498 | Transforming growth factor beta 1                   |       | Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to neurodegeneration; can induce EMT and cell migration in various cell types; frequently acts as an immunosuppressive cytokine in the TME |
| <u>Tigit</u> | 0.861 | T cell immunoreceptor with Ig and ITIM domains      |       | Binds with high affinity to the poliovirus receptor, causing increased secretion of IL-10, decreased secretion of IL-12B, and suppressing T cell activation by promoting the generation of mature immunoregulatory DCs   |

|              | Interferon Response |   |  |   |  |
|--------------|---------------------|---|--|---|--|
| Bst2         | 1.13                | Bone marrow stromal cell antigen 2          | Tethrin; CD317   | IFN-induced antiviral factor that blocks budding of enveloped viruses by directly   |  |
|              |                     | _   |  | tethering nascent virions to the membranes of infected cells  |  |
| <u>Ctsh</u>  | 0.906               | Cathepsin H                                 |  | Lysosomal protease; increased in macrophages in response to IFNγ  |  |
| <u>Ifih1</u> | 1.05                | Interferon induced with helicase C domain 1 | Helicard; melanoma differentiation- associated protein | PRR for cytoplasmic dsRNA; upon target recognition, associates with MAVS to activate TNK1 and IKBKE, which phosphorylate IRF3 and -7, which, in turn, activate transcription of |  |
|              |                     |   | 5 (MDA5)   | IFNα and -β   |  |
| <u>Irf1</u>  | 1.2                 | Interferon regulatory factor 1              |  | Transcriptional regulator that promotes inflammatory innate and adaptive immune responses   |  |
| <u>Irf7</u>  | 1.71                | Interferon regulatory factor 7              |  | Key transcriptional regulator of type I IFN-dependent immune responses; promotes $transcription\ of\ IFN\alpha\ and\ -\beta$  |  |
| <u>Irf8</u>  | 0.841               | Interferon regulatory factor 8              |  | TF that regulates of lineage commitment in myeloid cell maturation; promotes monocyte and plasmacytoid DC development   |  |

| Isg15 | 2.02  | Interferon-stimulated |                    | Ubiquitin-like protein that binds intracellular              |
|-------|-------|-----------------------|--------------------|--|
|       |       | gene 15               |                    | target proteins upon activation by IFN $\alpha$ or $\beta$ ; |
|       |       |                       |                    | can also be secreted to induce NK cell                       |
|       |       |                       |                    | proliferation, act as a chemoattractant for                  |
|       |       |                       |                    | neutrophils, and induce IFNγ upon binding to                 |
|       |       |                       |                    | ITGAL/ITGB2  |
| Isg20 | 0.904 | Interferon-stimulated |                    | IFN-induced antiviral exoribonuclease that                   |
|       |       | gene 20               |                    | acts on ssRNA with minor activity towards                    |
|       |       |                       |                    | ssDNA  |
| Jak1  | 0.337 | Janus kinase 1        |                    | Essential tyrosine kinase involved signal                    |
|       |       |                       |                    | transduction in type I and II cytokines and IFNs             |
| Mefv  | 1.79  | Mediterranean fever   | Marenostrin; pyrin | Involved in the regulation of innate immunity                |
|       |       |                       |                    | and the inflammatory response in response to                 |
|       |       |                       |                    | IFNγ; both stimulates and restrains the                      |
|       |       |                       |                    | inflammasome; also acts as a mediator of                     |
|       |       |                       |                    | pyroptosis   |
| Tbk1  | 0.767 | TANK-binding kinase   |                    | Coordinates the activation of IRF3 and NFkB                  |
|       |       | 1                     |                    | and induction of type I IFNs                                 |
|       |       |                       | Ion Transport      |  |
| Арр   | 0.458 | Amyloid-beta          |                    | Cell surface receptor and transmembrane                      |
|       |       | precursor protein     |                    | precursor protein that is cleaved by secretases              |
|       |       |                       |                    | to form a number of peptides; involved in cell               |

|              |       |  |                  | mobility, copper homeostasis, and oxidative stress   |
|--------------|-------|--|------------------|--|
| Ms4a1        | 3.3   | Membrane spanning 4-domains A1                               | CD20; Bp35       | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation |
| Slc11a1      | 1.46  | Natural resistance-<br>associated<br>macrophage protein<br>1 |                  | Macrophage-specific metal ion transporter; uptakes divalent metal cations to neutralize  ROSs  |
|              |       |  | IRAKs & TRAFs    |  |
| <u>Irak2</u> | 0.943 | Interleukin-1 receptor-associated kinase 2                   |                  | Adaptor protein involved in TLR and IL-1 signaling   |
| <u>Traf6</u> | 0.697 | Tumor necrosis factor receptor- associated factor 6          |                  | Adaptor protein that acts in the CD40 signaling cascade; promotes inflammation, IL- $ 6\text{, and TNF} \alpha $                           |
|              |       |  | JAK-STAT Pathway |  |
| <u>Jak1</u>  | 0.337 | Janus kinase 1   |                  | Essential tyrosine kinase involved signal transduction in type I and II cytokines and IFNs   |

| Stat1      | 1.02  | Signal transducer and |                    | Transcriptional activator that mediates          |
|------------|-------|-----------------------|--------------------|--|
| Stati      | 1.02  |                       |                    | ·  |
|            |       | activator of          |                    | cellular responses to IFNs, cytokines, and       |
|            |       | transcription 1       |                    | other growth factors                             |
| Stat3      | 0.418 | Signal transducer and |                    | Transcriptional activator of genes involved in   |
|            |       | activator of          |                    | cell growth and apoptosis; activated by JAKs     |
|            |       | transcription 3       |                    |  |
|            |       |                       | Kinases            |  |
|            |       |                       |                    |  |
| <u>Lck</u> | 1.29  | Lymphocyte cell       |                    | Src family tyrosine kinase that acts as one of   |
|            |       | kinase                |                    | the main signaling intermediaries downstream     |
|            |       |                       |                    | of the TCR; constitutively associated with the   |
|            |       |                       |                    | cytoplasmic portion of CD4                       |
| Jak1       | 0.337 | Janus kinase 1        |                    | Essential tyrosine kinase involved signal        |
|            |       |                       |                    | transduction in type I and II cytokines and IFNs |
| Map2k1     | 0.393 | Dual specificity      | MAPK/ERK kinase 1  | Essential component of the MAP kinase signal     |
|            |       | mitogen-activated     | (MEK1)             | transduction pathway; participates in            |
|            |       | protein kinase kinase |                    | numerous biological functions, including cell    |
|            |       | 1                     |                    | growth, adhesion, survival, differentiation,     |
|            |       |                       |                    | transcription, metabolism, and cytoskeletal      |
|            |       |                       |                    | remodeling                                       |
| Syk        | 0.969 | Spleen-associated     |                    | Critical kinase that transmits signals from the  |
|            |       | tyrosine kinase       |                    | TCR and BCR                                      |
|            |       |                       | Lysosomal Activity |  |
|            |       |                       | Lysosomar Activity |  |
|            |       |                       |                    |  |

| Camp        | 2.19  | Cathelicidin          | Polypeptide stored in the lysosomes of           |
|-------------|-------|-----------------------|--|
|             |       | antimicrobial peptide | macrophages and PMNs that digests                |
|             |       |                       | phagocytosed cells                               |
| <u>Ctsh</u> | 0.906 | Cathepsin H           | Lysosomal protease; increased in                 |
|             |       |                       | macrophages in response to IFNγ                  |
| <u>Ctss</u> | 1.43  | Cathepsin S           | Lysosomal protease that participates in          |
|             |       |                       | processing of Ag by MHC class II                 |
|             |       | <u></u>               | Macrophage Function                              |
|             |       |                       |  |
| Camp        | 2.19  | Cathelicidin          | Polypeptide stored in the lysosomes of           |
|             |       | antimicrobial peptide | macrophages and PMNs that digests                |
|             |       |                       | phagocytosed cells                               |
| Casp8       | 0.314 | Caspase 8             | Cysteine-aspartic acid protease; cleaves and     |
|             |       |                       | activates effector caspases CASP3, -4, -6, -7, - |
|             |       |                       | 9, and -10; initiates pyroptosis through         |
|             |       |                       | cleavage of gasdermin-D; mediates                |
|             |       |                       | noncanonical cleavage of IL-1β in DCs and        |
|             |       |                       | macrophages                                      |
| <u>Cd14</u> | 1.11  | Cluster of            | PRR that recognizes LPS; mostly found on         |
|             |       | differentiation 14    | macrophages                                      |
| Cebpb       | 0.724 | CCAAT/enhancer-       | Critical macrophage TF that promotes             |
|             |       | binding protein beta  | expression of several acute-phase and            |
|             |       |                       | inflammatory cytokine genes, including II6       |

| Clec5a      | 1.18  | C-Type lectin domain  | Myeloid DAP12-       | Critical macrophage receptor for dengue virus |
|-------------|-------|-----------------------|----------------------|---|
|             |       | family 5, member a    | associating lectin-1 | serotypes 1-4; positive regulator of          |
|             |       |                       |                      | osteoclastogenesis                            |
| <u>Ctsh</u> | 0.906 | Cathepsin H           |                      | Lysosomal protease; increased in              |
|             |       |                       |                      | macrophages in response to IFNγ               |
| Itgam       | 1.45  | Integrin alpha M      | CD11b                | Pairs with CD18 to form Mac-1 aka             |
|             |       |                       |                      | complement receptor 3; mediates leukocyte     |
|             |       |                       |                      | activation, adhesion, chemotaxis, migration,  |
|             |       |                       |                      | phagocytosis, and cell-mediated cytotoxicity; |
|             |       |                       |                      | serves as a macrophage marker                 |
| Marco       | 2.92  | Macrophage            |                      | A PRR that recognizes LDL                     |
|             |       | receptor with         |                      |   |
|             |       | collagenous structure |                      |   |
| Mif         | 0.328 | Macrophage            | L-dopachrome         | Pro-inflammatory cytokine that promotes       |
|             |       | migration inhibitory  | tautomerase          | macrophage function through suppression of    |
|             |       | factor                |                      | anti-inflammatory effects of glucocorticoids  |
| Slamf7      | 1.82  | Signaling lymphocytic |                      | A super-activator of macrophages and a        |
|             |       | activation molecule   |                      | strong promoter of phagocytosis; binds to     |
|             |       | family member 7       |                      | CD74  |
| Slc11a1     | 1.46  | Natural resistance-   |                      | Macrophage-specific metal ion transporter;    |
|             |       | associated            |                      | uptakes divalent metal cations to neutralize  |
|             |       |                       |                      | ROSs  |
|             |       |                       |                      |   |

|             |       | I                     | I                        |  |
|-------------|-------|-----------------------|--------------------------|--|
|             |       | macrophage protein    |                          |  |
|             |       | 1                     |                          |  |
|             |       |                       |                          |  |
| <u>Tnf</u>  | 3.07  | Tumor necrosis        | Cachectin                | Inflammatory cytokine mainly produced by       |
|             |       | factor                |                          | macrophages; binds to TNFRSF1A/TNFR1 and       |
|             |       |                       |                          | TNFRSF1B/TNFBR; capable of inducing cell       |
|             |       |                       |                          | death in certain tumor cell lines              |
|             |       |                       | Metabolism               |  |
|             |       |                       |                          |  |
| Abca1       | 1.35  | ATP-binding cassette  |                          | Membrane-associated cholesterol efflux         |
|             |       | transporter A1        |                          | pump   |
|             |       |                       |                          |  |
| Abcg1       | 1.4   | ATP-binding cassette  |                          | Membrane-associated cholesterol efflux         |
|             |       | transporter G1        |                          | pump   |
| <u>Cd36</u> | 2.9   | Cluster of            | Fatty acid               | Class B scavenger receptor that mediates fatty |
|             |       |                       |                          |  |
|             |       | differentiation 36    | translocase (FAT)        | acid uptake                                    |
| Map2k1      | 0.393 | Dual specificity      | MAPK/ERK kinase 1        | Essential component of the MAP kinase signal   |
|             |       | mitogen-activated     | (MEK1)                   | transduction pathway; participates in          |
|             |       | protein kinase kinase |                          | numerous biological functions, including cell  |
|             |       | 1                     |                          | growth, adhesion, survival, differentiation,   |
|             |       |                       |                          | transcription, metabolism, and cytoskeletal    |
|             |       |                       |                          | remodeling                                     |
|             |       |                       | Migration/Motility       |  |
|             |       |                       | wings actionly widelifty |  |
|             |       |                       |                          |  |

| Арр            | 0.458 | Amyloid-beta         |               | Cell surface receptor and transmembrane         |
|----------------|-------|----------------------|---------------|---|
|                |       | precursor protein    |               | precursor protein that is cleaved by secretases |
|                |       |                      |               | to form a number of peptides; involved in cell  |
|                |       |                      |               | mobility, copper homeostasis, and oxidative     |
|                |       |                      |               | stress  |
| lcam1          | 0.997 | Intracellular        | CD54          | Cell surface glycoprotein that serves as strong |
|                |       | adhesion molecule 1  |               | adhesive ligand for LFA-1; important for        |
|                |       |                      |               | leukocyte mobility and costimulation            |
| Itgam          | 1.45  | Integrin alpha M     | CD11b         | Pairs with CD18 to form Mac-1 aka               |
|                |       |                      |               | complement receptor 3; mediates leukocyte       |
|                |       |                      |               | activation, adhesion, chemotaxis, migration,    |
|                |       |                      |               | phagocytosis, and cell-mediated cytotoxicity;   |
|                |       |                      |               | serves as a macrophage marker                   |
| Jam3           | 1.9   | Junctional adhesion  |               | Immunoglobulin that mediates tight junctions    |
|                |       | molecule C           |               | between endothelial cells; mediates             |
|                |       |                      |               | transepithelial migration of PMNs               |
| <u>\$100a8</u> | 2.22  | S100 calcium-binding | Calgranulin A | Calcium- and zinc-binding protein involved in   |
|                |       | protein A8           |               | pro-inflammatory, antimicrobial, oxidant-       |
|                |       |                      |               | scavenging and apoptosis-inducing activities;   |
|                |       |                      |               | can induce neutrophil chemotaxis, adhesion,     |
|                |       |                      |               | phagocytosis, and degranulation;                |
|                |       |                      |               | predominantly found as calprotectin             |
|                |       |                      |               | (S100A8/A9) which has a wide plethora of        |
|                |       |                      |               | intra- and extracellular functions, including   |

| Tgfb1       | 0.498 | Transforming growth factor beta 1  |  | adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling  Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to neurodegeneration; can induce EMT and cell migration in various cell types; frequently acts as an immunosuppressive cytokine in the TME |
|-------------|-------|------------------------------------|--|---|
|             |       |                                    | Neutrophil Function                                |   |
| Camp        | 2.19  | Cathelicidin antimicrobial peptide |  | Polypeptide stored in the lysosomes of macrophages and PMNs that digests phagocytosed cells   |
| <u>Fpr2</u> | 2.47  | Formyl peptide<br>receptor 2       | Lipoxin A4 receptor                                | Low affinity receptor for N-formyl-methionyl peptides; activates neutrophils  |
| Lcn2        | 1.69  | Lipocalin 2                        | Neutrophil gelatinase- associated lipocalin (NGAL) | Neutrophil-secreted factor that sequesters iron-containing siderophores; also functions as a growth factor  |

| Ncf4           | 0.833 | Neutrophil cytosolic  | SH3 and PX        | Cytosolic regulatory component of the         |
|----------------|-------|-----------------------|-------------------|---|
|                |       | factor 4              | domain-containing | superoxide-producing phagocyte NADPH-         |
|                |       |                       | protein 4         | oxidase, a multicomponent enzyme system       |
|                |       |                       | (SH3PXD4)         | important for host defense                    |
| Pik3cd         | 0.584 | Phosphatidylinositol- |                   | A subunit of PI3K; acts downstream of TLR4,   |
|                |       | 4,5-bisphosphate 3-   |                   | TCR, BCR, and CD40; contributes to T helper   |
|                |       | kinase catalytic      |                   | cell expansion, mast cell development, and    |
|                |       | subunit delta isoform |                   | neutrophil chemotaxis, extravasation, and     |
|                |       |                       |                   | respiratory burst                             |
| <u>\$100a8</u> | 2.22  | S100 calcium-binding  | Calgranulin A     | Calcium- and zinc-binding protein involved in |
|                |       | protein A8            |                   | pro-inflammatory, antimicrobial, oxidant-     |
|                |       |                       |                   | scavenging and apoptosis-inducing activities; |
|                |       |                       |                   | can induce neutrophil chemotaxis, adhesion,   |
|                |       |                       |                   | phagocytosis, and degranulation;              |
|                |       |                       |                   | predominantly found as calprotectin           |
|                |       |                       |                   | (S100A8/A9) which has a wide plethora of      |
|                |       |                       |                   | intra- and extracellular functions, including |
|                |       |                       |                   | adhesion, apoptosis, autophagy, cytoskeletal  |
|                |       |                       |                   | remodeling, cytokine production, chemotaxis,  |
|                |       |                       |                   | migration, inflammation, arachidonic acid     |
|                |       |                       |                   | metabolism, oxidant-scavenging, and PRR       |
|                |       |                       |                   | signaling                                     |
|                |       |                       | NFkB Signaling    |   |
|                |       |                       |                   |   |

| Ikbkg       | 0.223 | Inhibitor of nuclear  | NFĸB essential   | Regulatory subunit of the IKK core complex     |
|-------------|-------|-----------------------|------------------|--|
|             |       | factor kappa B kinase | modifier (NEMO)  | that phosphorylates inhibitors of NFKB thus    |
|             |       | subunit gamma         |                  | leading to the dissociation of the             |
|             |       |                       |                  | inhibitor/NFĸB complex and ultimately the      |
|             |       |                       |                  | degradation of the inhibitor                   |
|             |       |                       |                  | degradation of the immotor                     |
| Rel         | 0.74  | Avian                 | c-Rel            | One of the NFκB family TFs; important for B    |
|             |       | reticuloendotheliosis |                  | cell and Treg development                      |
|             |       | viral oncogene        |                  |  |
|             |       | homolog               |                  |  |
|             |       |                       |                  |  |
| Relb        | 0.443 | Avian                 |                  | One of the NFkB family TFs; controls lymphoid  |
|             |       | reticuloendotheliosis |                  | development, DC biology, and noncanonical      |
|             |       | viral oncogene        |                  | NFkB signaling                                 |
|             |       | homolog B             |                  |  |
|             |       |                       |                  |  |
|             |       |                       | NK Cell Function |  |
| <u>Cd69</u> | 0.994 | Cluster of            | C-type lectin    | Signal transmitting receptor in lymphocytes,   |
|             |       | differentiation 69    | domain family 2, | NK cells, and platelets; induced upon T cell   |
|             |       |                       | member C         | activation; involved in lymphocyte             |
|             |       |                       |                  | proliferation                                  |
|             |       |                       |                  |  |
| <u>Gzmb</u> | 2.9   | Granzyme B            | Fragmentin 2     | Abundant protease in the cytosolic granules of |
|             |       |                       |                  | cytotoxic T and NK cells that activates        |
|             |       |                       |                  | caspase-mediated cell death when delivered     |
|             |       |                       |                  | into the target cell through the immunological |
|             |       |                       |                  | synapse  |
|             |       |                       |                  |  |

| <u>Gzmk</u>   | 2.49 | Granzyme K                              | Tryptase II                            | Granule-secreted, pro-apoptotic serine protease found in the cytoplasmic granules of CTLs  |
|---------------|------|---|--|--|
|               |      | Patte                                   | ern Recognition Rece                   | eptors   |
| Ccrl2         | 1.44 | C-C chemokine<br>receptor-like 2        |  | Stabilizes TLR4 surface expression on macrophages  |
| <u>Cd14</u>   | 1.11 | Cluster of differentiation 14           |  | PRR that recognizes LPS; mostly found on macrophages   |
| <u>Cd180</u>  | 1.28 | Cluster of differentiation 180          |  | Heterodimeric binding partner of Ly86 that participates in LPS binding in APCs   |
| <u>Clec4n</u> | 1.32 | C-type lectin domain family 4, member N | Dectin-2                               | PRR specific for Mycobacterial mannose-<br>capped lipoarabinomannan  |
| <u>Clec5a</u> | 1.18 | C-Type lectin domain family 5, member a | Myeloid DAP12-<br>associating lectin-1 | Critical macrophage receptor for dengue virus serotypes 1-4; positive regulator of osteoclastogenesis  |
| Clec7a        | 1.83 | C-Type lectin domain family 7, member a | Dectin-1                               | PRR specific for $\beta$ -1,3- and $\beta$ -1,6-linked glucans from fungi and plants; necessary for the TLR2-mediated inflammatory response and for TLR2-mediated activation of NF- $\kappa$ B |

| <u>Ddx58</u> | 0.528 | DExD/H-box helicase   | Retinoic acid-      | Cytoplasmic PRR that recognizes dsRNA; can        |
|--------------|-------|-----------------------|---------------------|---|
|              |       | 58                    | inducible gene I    | promote T cell-independent B cell activation;     |
|              |       |                       | (RIG-I)             | uses MAVS as an adaptor                           |
|              |       |                       |                     |   |
| Fpr2         | 2.47  | Formyl peptide        | Lipoxin A4 receptor | Low affinity receptor for N-formyl-methionyl      |
|              |       | receptor 2            |                     | peptides; activates neutrophils                   |
| Ifih1        | 1.05  | Interferon induced    | Helicard;           | PRR for cytoplasmic dsRNA; upon target            |
|              |       | with helicase C       | melanoma            | recognition, associates with MAVS to activate     |
|              |       | domain 1              | differentiation-    | TNK1 and IKBKE, which phosphorylate IRF3          |
|              |       |                       | associated protein  | and -7, which, in turn, activate transcription of |
|              |       |                       | 5 (MDA5)            | IFNα and -β                                       |
| 1.05         | 0.570 |                       |                     |   |
| <u>Ly96</u>  | 0.578 | Lymphocyte antigen    | Myeloid             | Heterodimeric binding partner of TLR4 that        |
|              |       | 96                    | differentiation     | participates in LPS binding                       |
|              |       |                       | factor 2 (MD-2)     |   |
| Marco        | 2.92  | Macrophage            |                     | A PRR that recognizes LDL                         |
|              |       | receptor with         |                     |   |
|              |       | collagenous structure |                     |   |
|              |       |                       |                     |   |
| <u>Myd88</u> | 0.64  | Myeloid               |                     | Key adaptor in the TLR signaling pathways;        |
|              |       | differentiation       |                     | interacts with all TLRs except TLR3; activates    |
|              |       | primary response 88   |                     | NFkB and IRFs                                     |
| Nlrp3        | 1.47  | NACHT domain-,        | Cryopyrin           | PRR with a wide diversity of recognized           |
|              |       | leucine-rich repeat-, |                     | targets that activates the NLRP3                  |
|              |       | 1                     |                     |   |

|                |       | and PYD-containing   |               | inflammasome consisting of NLRP3, PYCARD,     |
|----------------|-------|----------------------|---------------|---|
|                |       | protein 3            |               | and caspase-1/-8                              |
| Nod1           | 0.947 | Nucleotide binding   |               | Intracellular PRR that recognizes             |
|                |       | oligomerization      |               | peptidoglycan-derived muropeptides and        |
|                |       | domain containing 1  |               | Shigella effector proteins                    |
| <u>\$100a8</u> | 2.22  | S100 calcium-binding | Calgranulin A | Calcium- and zinc-binding protein involved in |
|                |       | protein A8           |               | pro-inflammatory, antimicrobial, oxidant-     |
|                |       |                      |               | scavenging and apoptosis-inducing activities; |
|                |       |                      |               | can induce neutrophil chemotaxis, adhesion,   |
|                |       |                      |               | phagocytosis, and degranulation;              |
|                |       |                      |               | predominantly found as calprotectin           |
|                |       |                      |               | (S100A8/A9) which has a wide plethora of      |
|                |       |                      |               | intra- and extracellular functions, including |
|                |       |                      |               | adhesion, apoptosis, autophagy, cytoskeletal  |
|                |       |                      |               | remodeling, cytokine production, chemotaxis,  |
|                |       |                      |               | migration, inflammation, arachidonic acid     |
|                |       |                      |               | metabolism, oxidant-scavenging, and PRR       |
|                |       |                      |               | signaling                                     |
| Ticam2         | 1.18  | TIR domain-          |               | Sorting adapter in various innate immune      |
|                |       | containing adaptor   |               | signaling cascades; bridges TLR2 and MyD88    |
|                |       | molecule 2           |               |   |
| Tlr2           | 0.853 | Toll-like receptor 2 | CD282         | Surface PRR that binds to various lipid-      |
|                |       |                      |               | containing PAMPs                              |

| <u>Tlr4</u> | 0.571 | Toll-like receptor 4               | CD284        | Surface PRR that recognizes LPS; pairs with LY96 and CD14; acts via MYD88, TIRAP, and TRAF6, leading to NFκB activation, cytokine secretion, and the inflammatory response; in complex with TLR6, promotes sterile inflammation in monocytes/macrophages in response to oxLDL or amyloid-β 42 |
|-------------|-------|------------------------------------|--------------|---|
| Tlr6        | 1.42  | Toll like receptor 6               | CD286        | Surface PRR that recognizes diacylated and triacylated lipopeptides   |
| Tlr7        | 1.34  | Toll-like receptor 7               | CD287        | Endosomic PRR that recognizes ssRNA   |
| Tlr8        | 0.964 | Toll-like receptor 8               | CD288        | Endosomic PRR that recognizes ssRNA   |
| Tlr9        | 1.41  | Toll like receptor 9               | CD289        | Endosomic PRR that recognizes unmethylated  CpG dinucleotides   |
|             |       |                                    | Phagocytosis |   |
| Camp        | 2.19  | Cathelicidin antimicrobial peptide |              | Polypeptide stored in the lysosomes of macrophages and PMNs that digests phagocytosed cells   |
| Itgam       | 1.45  | Integrin alpha M                   | CD11b        | Pairs with CD18 to form Mac-1 aka  complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker   |

| Ncf4           | 0.833 | Neutrophil cytosolic  | SH3 and PX         | Cytosolic regulatory component of the           |
|----------------|-------|-----------------------|--------------------|---|
|                |       | factor 4              | domain-containing  | superoxide-producing phagocyte NADPH-           |
|                |       |                       | protein 4          | oxidase, a multicomponent enzyme system         |
|                |       |                       | (SH3PXD4)          | important for host defense                      |
| <u>\$100a8</u> | 2.22  | S100 calcium-binding  | Calgranulin A      | Calcium- and zinc-binding protein involved in   |
|                |       | protein A8            |                    | pro-inflammatory, antimicrobial, oxidant-       |
|                |       |                       |                    | scavenging and apoptosis-inducing activities;   |
|                |       |                       |                    | can induce neutrophil chemotaxis, adhesion,     |
|                |       |                       |                    | phagocytosis, and degranulation;                |
|                |       |                       |                    | predominantly found as calprotectin             |
|                |       |                       |                    | (S100A8/A9) which has a wide plethora of        |
|                |       |                       |                    | intra- and extracellular functions, including   |
|                |       |                       |                    | adhesion, apoptosis, autophagy, cytoskeletal    |
|                |       |                       |                    | remodeling, cytokine production, chemotaxis,    |
|                |       |                       |                    | migration, inflammation, arachidonic acid       |
|                |       |                       |                    | metabolism, oxidant-scavenging, and PRR         |
|                |       |                       |                    | signaling                                       |
| Slamf7         | 1.82  | Signaling lymphocytic |                    | A super-activator of macrophages and a          |
|                |       | activation molecule   |                    | strong promoter of phagocytosis; binds to       |
|                |       | family member 7       |                    | CD74  |
|                |       | ROS                   | Generation & Prote | ction   |
| Δ :-:          | 0.450 | A                     |                    | Call austana nacentan and the                   |
| App            | 0.458 | Amyloid-beta          |                    | Cell surface receptor and transmembrane         |
|                |       | precursor protein     |                    | precursor protein that is cleaved by secretases |
|                |       |                       |                    | to form a number of peptides; involved in cell  |

|             |       |  |  | mobility, copper homeostasis, and oxidative stress   |
|-------------|-------|--|--|--|
| Cybb        | 1.32  | Cytochrome b-245<br>heavy chain        | Nox2   | Part of the NADPH oxidase process; generates superoxides   |
| Ncf4        | 0.833 | Neutrophil cytosolic factor 4          | SH3 and PX domain-containing protein 4 (SH3PXD4) | Cytosolic regulatory component of the superoxide-producing phagocyte NADPH-oxidase, a multicomponent enzyme system important for host defense  |
| Nos2        | 3.66  | Inducible nitric oxide synthase (iNOS) |  | Produces reactive oxygen species and contributes to inflammatory cytokine production   |
|             |       |  | T Cell Function                                  |  |
| Cd3d        | 1.81  | Cluster of differentiation 3 delta     |  | Component of the TCR-CD3 complex; upon phosphorylation by Lck, serves as a docking station for downstream TCR signaling adaptors   |
| <u>Cd3e</u> | 1.6   | Cluster of differentiation 3 epsilon   |  | Component of the TCR-CD3 complex; initiates the TCR-CD3 complex assembly by forming the two heterodimers CD3δ/CD3ε and CD3γ/CD3ε; also participates in internalization and cell surface down-regulation of TCR-CD3 |

|              |      |                                    |      | complexes via endocytosis sequences present in CD3ε cytosolic region  |
|--------------|------|------------------------------------|------|---|
| Cd3g         | 1.54 | Cluster of differentiation 3       |      | Component of the TCR-CD3 complex; plays an essential role in the dynamic regulation of TCR expression at the cell surface   |
| <u>Cd5</u>   | 1.37 | Cluster of differentiation 5       | LEU1 | Type-I transmembrane glycoprotein found on the surface of T and B cells; may act as a receptor in regulating T cell proliferation   |
| <u>Cd8a</u>  | 1.43 | Cluster of differentiation 8 alpha | LEU2 | Alpha chain of the CD8 coreceptor, which binds to MHC class I; in T cells, functions primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells |
| <u>Cd8b1</u> | 1.2  | Cluster of differentiation 8 beta  |      | Beta chain of the CD8 coreceptor, which binds to MHC class I  |
| <u>Cd28</u>  | 1.51 | Cluster of differentiation 28      |      | Essential T cell co-receptor that enhances T  cell activation, proliferation, cytokine  production, and survival; binds to CD80 and  CD86   |

| <u>Cd37</u>  | 1.26  | Cluster of            | Tetraspanin-26   | Cell surface glycoprotein known to complex        |
|--------------|-------|-----------------------|------------------|---|
|              |       | differentiation 37    |                  | with integrins and other transmembrane 4          |
|              |       |                       |                  | superfamily proteins; may play a role in T cell-  |
|              |       |                       |                  | B cell interactions                               |
| 0.150        | 0.004 |                       |                  |   |
| <u>Cd69</u>  | 0.994 | Cluster of            | C-type lectin    | Signal transmitting receptor in lymphocytes,      |
|              |       | differentiation 69    | domain family 2, | NK cells, and platelets; induced upon T cell      |
|              |       |                       | member C         | activation; involved in lymphocyte                |
|              |       |                       |                  | proliferation                                     |
| <u>Cd247</u> | 2.02  | Cluster of            | T cell surface   | Central intracellular signaling chain of the TCR, |
|              |       | differentiation 247   | glycoprotein CD3 | to which downstream signaling adaptors dock       |
|              |       |                       | zeta chain       |   |
|              |       |                       |                  |   |
| Cyfip2       | 1.26  | Cytoplasmic FMR1-     |                  | Involved in T cell adhesion and p53/TP53-         |
|              |       | interacting protein 2 |                  | dependent induction of apoptosis                  |
| <u>Gzmb</u>  | 2.9   | Granzyme B            | Fragmentin 2     | Abundant protease in the cytosolic granules of    |
|              |       |                       |                  | cytotoxic T and NK cells that activates           |
|              |       |                       |                  | caspase-mediated cell death when delivered        |
|              |       |                       |                  | into the target cell through the immunological    |
|              |       |                       |                  | synapse   |
| <u>Gzmk</u>  | 2.49  | Granzyme K            | Tryptase II      | Granule-secreted, pro-apoptotic serine            |
|              |       |                       |                  | protease found in the cytoplasmic granules of     |
|              |       |                       |                  | CTLs  |
|              |       |                       |                  |   |

| <u>lcos</u> | 1.58  | Inducible T cell      | CD278             | Enhances all basic T cell responses to foreign  |
|-------------|-------|-----------------------|-------------------|---|
|             |       | costimulator          |                   | Ag; essential both for efficient interaction    |
|             |       |                       |                   | between T and B cells and for normal Ab         |
|             |       |                       |                   | responses to T cell-dependent Ags               |
| <u>Lck</u>  | 1.29  | Lymphocyte cell       |                   | Src family tyrosine kinase that acts as one of  |
|             |       | kinase                |                   | the main signaling intermediaries downstream    |
|             |       |                       |                   | of the TCR; constitutively associated with the  |
|             |       |                       |                   | cytoplasmic portion of CD4                      |
| Lcp1        | 0.867 | Lymphocyte cytosolic  | Plastin-2         | Actin-binding protein that promotes T cell      |
|             |       | protein 1             |                   | activation in response to costimulation         |
|             |       |                       |                   | through TCR/CD3 and CD2 or CD28; assists        |
|             |       |                       |                   | with IL2RA transport to the cell surface        |
| Pik3cd      | 0.584 | Phosphatidylinositol- |                   | A subunit of PI3K; acts downstream of TLR4,     |
|             |       | 4,5-bisphosphate 3-   |                   | TCR, BCR, and CD40; contributes to T helper     |
|             |       | kinase catalytic      |                   | cell expansion, mast cell development, and      |
|             |       | subunit delta isoform |                   | neutrophil chemotaxis, extravasation, and       |
|             |       |                       |                   | respiratory burst                               |
| Ptprc       | 0.739 | Protein tyrosine      | CD45; leukocyte   | Delivers costimulation during T cell activation |
|             |       | phosphatase           | common antigen    | upon binding to its ligand DPP4;                |
|             |       | receptor type C       | (LCA)             | dephosphorylates Lyn and suppresses JAK         |
|             |       |                       |                   | kinases   |
| Spn         | 0.72  | Sialophorin           | Leukosialin; CD43 | Cell surface sialoglycoprotein expressed by T   |
|             |       |                       |                   | cells, B cells, monocytes, and granulocytes;    |

| Syk         | 0.969 | Spleen-associated tyrosine kinase | promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNγ in CD4 <sup>+</sup> T cells  Critical kinase that transmits signals from the TCR and BCR |
|-------------|-------|-----------------------------------|---|
|             |       | Transcri                          | iption Factors & Coactivators   |
|             |       |                                   |   |
| Cebpb       | 0.724 | CCAAT/enhancer-                   | Critical macrophage TF that promotes  |
|             |       | binding protein beta              | expression of several acute-phase and   |
|             |       |                                   | inflammatory cytokine genes, including II6  |
| <u>Irf1</u> | 1.2   | Interferon regulatory             | Transcriptional regulator that promotes   |
|             |       | factor 1                          | inflammatory innate and adaptive immune   |
|             |       |                                   | responses   |
| <u>Irf7</u> | 1.71  | Interferon regulatory             | Key transcriptional regulator of type I IFN-  |
|             |       | factor 7                          | dependent immune responses; promotes  |
|             |       |                                   | transcription of IFN $\alpha$ and - $\beta$   |
| <u>Irf8</u> | 0.841 | Interferon regulatory             | TF that regulates of lineage commitment in  |
|             |       | factor 8                          | myeloid cell maturation; promotes monocyte  |
|             |       |                                   | and plasmacytoid DC development   |
| Mef2c       | 0.998 | Myocyte enhancer                  | Transcriptional activator that binds specifically   |
|             |       | factor 2c                         | to the MEF2 element present in the regulatory   |
|             |       |                                   | regions of many muscle-specific genes;  |

|       |       |  |                            | controls cardiac morphogenesis and myogenesis, and is also involved in vascular development; required for B cell survival and proliferation in response to BCR stimulation, efficient IgG1 Ab responses to T cell-dependent Ags, and for normal induction of GC B cells   |
|-------|-------|--|----------------------------|---|
| Stat1 | 1.02  | Signal transducer and activator of transcription 1 |                            | Transcriptional activator that mediates cellular responses to IFNs, cytokines, and other growth factors   |
| Stat3 | 0.418 | Signal transducer and activator of transcription 3 |                            | Transcriptional activator of genes involved in cell growth and apoptosis; activated by JAKs   |
| Yy1   | 0.196 | Yin yang 1   |                            | Ubiquitous factor that serves as a transcriptional "switch", either promoting or repressing the transcription of numerous genes through the selective recruitment of either histone deacetylases or acetyltransferases; plays a fundamental role in diverse processes, such as differentiation, replication, and cellular proliferation |
| Zbp1  | 1.27  | Z-DNA binding protein 1                            | Tumor stroma and activated | Key innate sensor that recognizes and binds Z-RNA structures, which are produced by a number of viruses, and induces type-I IFN   |

|             |       |  | macrophage<br>protein DLM-1 | production; key activator of cellular $necroptosis; promotes \ \text{IL-1}\alpha \ induction \ in \ an}$ $NLRP3-inflammasome-independent \ manner$                          |
|-------------|-------|--|-----------------------------|---|
|             |       |  | Ubiquitin Regulation        | n   |
| <u>Cyld</u> | 0.381 | Cylindromatosis  lysine 63  deubiquitinase       |                             | Inhibits NFkB activation by deubiquitinating upstream signaling factors; inhibits Wnt signaling; restricts polyubiquitination of RIPK1 and -2, thereby limiting necroptosis |
| Tnfaip3     | 0.804 | Tumor necrosis  factor, alpha-induced  protein 3 |                             | Ubiquitin-editing enzyme that complexes with  ITCH to degrade inflammatory signaling  components in the TNF, IL1, and TLR  pathways; targets TRAF2, TRAF6, and IKK          |
| Ubc         | 0.263 | Polyubiquitin C                                  |                             | Serves various roles, including innate immunity, DNA repair, and stimulation of autophagy and the proteasomal response  |

|      | NBTXR3+PBT+αPD1 vs Ctrl |           |                     |          |
|------|-------------------------|-----------|---------------------|----------|
| Gene | Log2<br>fold<br>change  | Full Name | Notable Aliases     | Function |
|      |                         |           | Acute Phase Respons | se       |

| <u>App</u>  | 0.801 | Amyloid-beta       |                        | Cell surface receptor and transmembrane          |
|-------------|-------|--------------------|------------------------|--|
|             |       | precursor protein  |                        | precursor protein that is cleaved by secretases  |
|             |       |                    |                        | to form a number of peptides; involved in cell   |
|             |       |                    |                        | mobility, copper homeostasis, and oxidative      |
|             |       |                    |                        | stress   |
| Psen2       | 0.848 | Presenilin-2       |                        | Putative catalytic subunit of the gamma-         |
|             |       |                    |                        | secretase complex, which cleaves integral        |
|             |       |                    |                        | membrane proteins such as APP; also involved     |
|             |       |                    |                        | in Ca(2+) homeostasis between the ER and the     |
|             |       |                    |                        | mitochondria                                     |
| Serping1    | 0.72  | Serpin family G    | C1-inhibitor           | Inflammation-induced acute phase protein         |
|             |       | member 1           |                        | that inhibits C1r and C1s proteases in the C1    |
|             |       |                    |                        | complex  |
|             |       | Adhes              | sion & Cell-Cell Inter | actions  |
| Cd2         | 1.68  | Cluster of         | Leukocyte              | Interacts with LFA-3 and CD48 to mediate         |
|             |       | differentiation 2  | functional antigen     | adhesion between T cells and other cell types    |
|             |       |                    | 2 (LFA-2)              |  |
| <u>Cd37</u> | 2.07  | Cluster of         | Tetraspanin-26         | Cell surface glycoprotein known to complex       |
|             |       | differentiation 37 |                        | with integrins and other transmembrane 4         |
|             |       |                    |                        | superfamily proteins; may play a role in T cell- |
|             |       |                    |                        | B cell interactions                              |

| <u>Cd47</u>  | 0.743 | Cluster of            | Integrin-associated  | Partners with membrane integrins to serve as     |
|--------------|-------|-----------------------|----------------------|--|
|              |       | differentiation 47    | protein (IAP)        | an inhibitor of phagocytosis                     |
| <u>Cd97</u>  | 0.748 | Cluster of            | BL-Ac[F2]            | GPCR that promotes granulocyte adhesion and      |
|              |       | differentiation 97    |                      | migration; activates T cells via binding to      |
|              |       |                       |                      | CD55; stimulates angiogenesis through            |
|              |       |                       |                      | binding integrin counterreceptors on             |
|              |       |                       |                      | endothelial cells                                |
| Cyfip2       | 1.74  | Cytoplasmic FMR1-     |                      | Involved in T cell adhesion and p53/TP53-        |
|              |       | interacting protein 2 |                      | dependent induction of apoptosis                 |
| <u>lcam1</u> | 1.69  | Intracellular         | CD54                 | Cell surface glycoprotein that serves as strong  |
|              |       | adhesion molecule 1   |                      | adhesive ligand for LFA-1; important for         |
|              |       |                       |                      | leukocyte mobility and costimulation             |
| Ifitm1       | 0.626 | Interferon-induced    | CD225                | IFN-induced antiviral protein implicated in cell |
|              |       | transmembrane         |                      | adhesion and control of cell growth and          |
|              |       | protein 1             |                      | migration  |
| ltga1        | 0.653 | Integrin subunit      | CD49a; very late     | Alpha 1 subunit for common integrin              |
|              |       | alpha 1               | activation protein 1 | receptors; pairs with the β1 subunit to form a   |
|              |       |                       | (VLA-1)              | cell-surface receptor for collagen and laminin;  |
|              |       |                       |                      | involved in cell-cell adhesion and may play a    |
|              |       |                       |                      | role in inflammation and fibrosis                |
| <u>Itgal</u> | 2.16  | Integrin alpha L      |                      | Pairs with ITGB2 to form lymphocyte function-    |
|              |       |                       |                      | associated antigen-1 (LFA-1), a common           |

|              |      |                                 |       | leukocyte adhesion molecule and costimulatory receptor  |
|--------------|------|---------------------------------|-------|---|
| <u>Itgam</u> | 2.19 | Integrin alpha M                | CD11b | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker  |
| <u>Itgax</u> | 1.47 | Integrin alpha X                | CD11c | Adhesion molecule; signature marker of Ag-<br>presenting DCs  |
| <u>ltgb2</u> | 1.87 | Integrin subunit beta           |       | Pairs with ITGAL to form a receptor for ICAM1, with ITGAM or ITGAX for iC3b and fibronectin   |
| Jam3         | 1.54 | Junctional adhesion  molecule C |       | Immunoglobulin that mediates tight junctions between endothelial cells; mediates transepithelial migration of PMNs; promotes chemotaxis of vascular endothelial cells and stimulates angiogenesis   |
| Lgals3       | 1.19 | Galectin 3                      |       | Galactose-specific lectin that binds IgE; involved in acute inflammatory responses, including neutrophil activation and adhesion, chemoattraction of monocytes macrophages, opsonization of apoptotic neutrophils, and activation of mast cells |

| Map2k1         | 0.266 | Dual specificity      | MAPK/ERK kinase 1 | Essential component of the MAP kinase signal     |
|----------------|-------|-----------------------|-------------------|--|
|                |       | mitogen-activated     | (MEK1)            | transduction pathway; participates in            |
|                |       | protein kinase kinase |                   | numerous biological functions, including cell    |
|                |       | 1                     |                   | growth, adhesion, survival, differentiation,     |
|                |       |                       |                   | transcription, metabolism, and cytoskeletal      |
|                |       |                       |                   | remodeling                                       |
| <u>\$100a8</u> | 2.3   | S100 calcium-binding  | Calgranulin A     | Calcium- and zinc-binding protein involved in    |
|                |       | protein A8            |                   | pro-inflammatory, antimicrobial, oxidant-        |
|                |       |                       |                   | scavenging and apoptosis-inducing activities;    |
|                |       |                       |                   | can induce neutrophil chemotaxis, adhesion,      |
|                |       |                       |                   | phagocytosis, and degranulation;                 |
|                |       |                       |                   | predominantly found as calprotectin              |
|                |       |                       |                   | (S100A8/A9) which has a wide plethora of         |
|                |       |                       |                   | intra- and extracellular functions, including    |
|                |       |                       |                   | adhesion, apoptosis, autophagy, cytoskeletal     |
|                |       |                       |                   | remodeling, cytokine production, chemotaxis,     |
|                |       |                       |                   | migration, inflammation, arachidonic acid        |
|                |       |                       |                   | metabolism, oxidant-scavenging, and PRR          |
|                |       |                       |                   | signaling  |
| Sell           | 1.78  | L-selectin            |                   | Mediates cell adhesion by binding to             |
|                |       |                       |                   | glycoproteins on neighboring cells               |
| Thbs1          | 0.589 |                       |                   | Adhesive glycoprotein that mediates cell-to-     |
|                |       |                       |                   | cell and cell-to-matrix interactions; ligand for |
|                |       | Thrombospondin 1      |                   | CD36   |

| Thy1           | 1.25  | Thy-1 T cell antigen            |               | Cell surface glycoprotein involved in cell adhesion and communication in immune and nerve cells   |
|----------------|-------|---------------------------------|---------------|---|
|                |       |                                 | Angiogenesis  |   |
| <u>Cd97</u>    | 0.748 | Cluster of differentiation 97   | BL-Ac[F2]     | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells  |
| <u>Itgam</u>   | 2.19  | Integrin alpha M                | CD11b         | Pairs with CD18 to forms Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker   |
| Jam3           | 1.54  | Junctional adhesion molecule C  |               | Immunoglobulin that mediates tight junctions between endothelial cells; mediates transepithelial migration of PMNs  |
| <u>\$100a8</u> | 2.3   | S100 calcium-binding protein A8 | Calgranulin A | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin |

|              |       |                      | (S100A8/A9) which has a wide plethora of         |
|--------------|-------|----------------------|--|
|              |       |                      | intra- and extracellular functions, including    |
|              |       |                      | adhesion, apoptosis, autophagy, cytoskeletal     |
|              |       |                      | remodeling, cytokine production, chemotaxis,     |
|              |       |                      | migration, inflammation, arachidonic acid        |
|              |       |                      | metabolism, oxidant-scavenging, and PRR          |
|              |       |                      | signaling  |
| Tgfb1        | 0.803 | Transforming growth  | Multifunctional protein that regulates the       |
|              |       | factor beta 1        | growth and differentiation of various cell       |
|              |       |                      | types and is involved in various processes,      |
|              |       |                      | such as normal development, immune               |
|              |       |                      | function, microglia function and responses to    |
|              |       |                      | neurodegeneration; can induce EMT and cell       |
|              |       |                      | migration in various cell types; frequently acts |
|              |       |                      | as an immunosuppressive cytokine in the TME      |
| <u>Vegfa</u> | 0.592 | Vascular endothelial | Glycosylated mitogen that promotes vascular      |
|              |       | growth factor A      | permeability, vasculogenesis, angiogenesis,      |
|              |       |                      | and cell migration                               |
|              |       |                      |  |
|              |       | Antigei              | n Processing & Presentation                      |
| <u>Cd1d1</u> | 1.45  | Cluster of           | Murine non-classical class I MHC; primarily      |
|              |       | differentiation 1 D1 | presents lipid and glycolipid Ags                |
| <u>Cd1d2</u> | 1.64  | Cluster of           | Pairs with CD1d1 to form the murine non-         |
|              |       | differentiation 1 D2 | classical class I MHC, CD1d; primarily presents  |
|              |       |                      |  |

|               |       |  |                          | lipid and glycolipid Ags; essential for NKT cell development; presents shorter acyl chain Ags than CD1d1 |
|---------------|-------|--|--------------------------|--|
| <u>Cd74</u>   | 1.22  | Cluster of differentiation 74                          | MHC class II gamma chain | Stabilizes peptide-free class II αβ heterodimers during MHC-Ag complex formation                         |
| Clec4a2       | 1.75  | C-type lectin domain family 4 member A2                |                          | PRR that, upon binding mannose or fucose, is endocytosed and processed in the Ag presentation pathway    |
| Ctss          | 2.53  | Cathepsin S  |                          | Lysosomal protease that participates in processing of Ag by MHC class II                                 |
| Cyfip2        | 1.74  | Cytoplasmic FMR1-<br>interacting protein 2             |                          | Involved in T cell adhesion and p53/TP53-<br>dependent induction of apoptosis                            |
| H2-Aa         | 1.09  | Histocompatibility 2,<br>class II antigen A,<br>alpha  |                          | MHC class II molecule; presents Ags to CD4 <sup>+</sup> T cells  |
| <u>H2-Ab1</u> | 1.34  | Histocompatibility 2,<br>class II antigen A,<br>beta 1 |                          | MHC class II molecule; presents Ags to CD4 <sup>+</sup> T cells  |
| <u>H2-D1</u>  | 0.841 | Histocompatibility 2,  D region locus 1                |                          | MHC class I molecule; presents Ags to CD8+ T cells   |

| <u>H2-K1</u>  | 0.421 | Histocompatibility 2, |                    | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T |
|---------------|-------|-----------------------|--------------------|--|
|               |       | K1, K region          |                    | cells  |
| H2-Q10        | 2.19  | Histocompatibility 2, |                    | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T |
|               |       | Q region locus 10     |                    | cells  |
| <u>H2-T23</u> | 1.23  | Histocompatibility 2, |                    | MHC class I molecule; presents Ags to CD8+ T             |
|               |       | Q region locus 10     |                    | cells  |
| Psmb10        | 0.571 | Proteasome subunit    |                    | 20S core $\beta$ subunit of the proteasome involved      |
|               |       | beta type 10          |                    | in Ag processing to generate class I binding             |
|               |       |                       |                    | peptides   |
| Tap1          | 1.1   | Transporter antigen   | Really interesting | ATP-binding cassette transporter that pumps              |
|               |       | peptide 1             | new gene 4         | degraded cytosolic peptides from the cytosol             |
|               |       |                       | (RING4)            | to the ER for packaging into MHC class I                 |
|               |       |                       |                    | molecules  |
|               |       |                       | Anti-Inflammatory  |  |
| Cd200r1       | 1.18  | Cluster of            |                    | Inhibitory receptor for the CD200/OX2 cell               |
|               |       | differentiation 200   |                    | surface glycoprotein; limits inflammation by             |
|               |       | receptor 1            |                    | inhibiting the expression of proinflammatory             |
|               |       |                       |                    | molecules including TNFα, IFNs, and iNOS                 |
| <u>Cd274</u>  | 3.02  | Cluster of            | Programmed cell    | Ubiquitously expressed ligand for co-inhibitory          |
|               |       | differentiation 274   | death receptor     | receptor PD-1; upregulated by tumors as an               |
|               |       |                       | ligand 1 (PD-L1)   | immune evasion strategy                                  |

| Ctla4    | 2.26  | Cytotoxic T  lymphocyte antigen   | CD152        | Inhibitory receptor that blocks CD28 costimulation by competitively binding its  |
|----------|-------|-----------------------------------|--------------|--|
|          |       | 4                                 |              | ligands CD80 and CD86  |
| Pdcd1    | 0.768 | Programmed cell death 1           | CD279; PD-1  | Checkpoint receptor; inhibits T and NK cell activation   |
| Pdcd1lg2 | 1.45  | Programmed cell death 1 ligand 2  |              |  |
| Lag3     | 2.17  | Lymphocyte activating gene 3      | CD223        | Inhibitory receptor on activated T cells; binds to ligands, such as FGL1; constitutively expressed on a subset of regulatory Tregs and contributes to their suppressive function; acts as a negative regulator of plasmacytoid DC activation |
| Nfkb1    | 0.349 | Nuclear factor kappa  B subunit 1 | p105/p50     | One of the NFkB family TFs; inhibits inflammation  |
| Serping1 | 0.72  | Serpin family G<br>member 1       | C1-inhibitor | Inflammation-induced acute phase protein that inhibits C1r and C1s proteases in the C1 complex   |
| Tgfb1    | 0.803 | Transforming growth factor beta 1 |              | Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to                           |

|              |       |                   |                | neurodegeneration; can induce EMT and cell       |
|--------------|-------|-------------------|----------------|--|
|              |       |                   |                | migration in various cell types; frequently acts |
|              |       |                   |                | as an immunosuppressive cytokine in the TME      |
| <u>Tigit</u> | 1.28  | T cell            |                | Binds with high affinity to the poliovirus       |
|              |       | immunoreceptor    |                | receptor, causing increased secretion of IL-10,  |
|              |       | with Ig and ITIM  |                | decreased secretion of IL-12B, and               |
|              |       | domains           |                | suppressing T cell activation by promoting the   |
|              |       |                   |                | generation of mature immunoregulatory DCs        |
| Tnfaip3      | 1.34  | Tumor necrosis    |                | Ubiquitin-editing enzyme that complexes with     |
|              |       | factor, alpha-    |                | ITCH to degrade inflammatory signaling           |
|              |       | induced protein 3 |                | components in the TNF, IL1, and TLR              |
|              |       |                   |                | pathways; targets TRAF2, TRAF6, and IKK          |
| Tollip       | 0.413 | Toll interacting  |                | Inhibitory adaptor protein; recruits IRAK1 to    |
|              |       | protein           |                | the IL-1 receptor complex and inhibitively       |
|              |       |                   |                | phosphorylates it                                |
|              |       |                   | Apoptosis      |  |
| Casp1        | 0.882 | Caspase 1         | Interleukin 1β | Cysteine-aspartic acid protease that mediates    |
|              |       |                   | convertase     | cleavage-based activation of IL-1β and IL-18;    |
|              |       |                   |                | serves as the central enzymatic core of the      |
|              |       |                   |                | inflammasome; also induces apoptosis             |
| Casp8        | 0.536 | Caspase 8         |                | Cysteine-aspartic acid protease; cleaves and     |
|              |       |                   |                | activates effector caspases CASP3, -4, -6, -7, - |
|              |       |                   |                | 9, and -10; initiates pyroptosis through         |

|             |       |  |   | cleavage of gasdermin-D; mediates noncanonical cleavage of IL-1β in DCs and macrophages  |
|-------------|-------|--|---|--|
| Cyfip2      | 1.74  | Cytoplasmic FMR1-<br>interacting protein 2 |   | Involved in T cell adhesion and p53/TP53-<br>dependent induction of apoptosis  |
| <u>Fas</u>  | 1.4   | Fragment apoptosis stimulating             |   | Cell surface death receptor; interaction with  FAS-ligand triggers an apoptotic signaling  cascade; also activates NFkB, ERK1, and  MAPK8  |
| <u>Fasl</u> | 2.01  | Fas ligand                                 | CD178; CD95L; apoptosis antigen ligand (APTL)  Tumor necrosis factor ligand superfamily member 6 (TNFSF6) | Ligand for the Fas death receptor; involved in CTL and NK cell-mediated apoptosis  |
| Prkcd       | 0.892 | Protein kinase C<br>delta                  |   | Ca(2+)-independent, phospholipid- and DAG-dependent serine/threonine-protein kinase that promotes apoptosis in response to DNA damage but inhibits it during cytokine receptor-initiated cell death; required for oxygen radical production by NADPH oxidase |

| <u>S100a8</u>  | 2.3   | S100 calcium-binding | Calgranulin A       | Calcium- and zinc-binding protein involved in    |
|----------------|-------|----------------------|---------------------|--|
|                |       | protein A8           |                     | pro-inflammatory, antimicrobial, oxidant-        |
|                |       |                      |                     | scavenging and apoptosis-inducing activities;    |
|                |       |                      |                     | can induce neutrophil chemotaxis, adhesion,      |
|                |       |                      |                     | phagocytosis, and degranulation;                 |
|                |       |                      |                     | predominantly found as calprotectin              |
|                |       |                      |                     | (S100A8/A9) which has a wide plethora of         |
|                |       |                      |                     | intra- and extracellular functions, including    |
|                |       |                      |                     | adhesion, apoptosis, autophagy, cytoskeletal     |
|                |       |                      |                     | remodeling, cytokine production, chemotaxis,     |
|                |       |                      |                     | migration, inflammation, arachidonic acid        |
|                |       |                      |                     | metabolism, oxidant-scavenging, and PRR          |
|                |       |                      |                     | signaling  |
| - C C .        | 0.40  |                      | 00000 11000         |  |
| Tnfsf14        | 2.19  | TNF superfamily      | CD258; LIGHT;       | Cytokine that binds to TNFRSF3/LTBR and          |
|                |       | member 14            | Herpesvirus entry   | TNFRSF14/HVEM; delivers costimulation to T       |
|                |       |                      | mediator ligand     | cells; triggers apoptosis of various tumor cells |
|                |       |                      | (HVEML)             |  |
|                |       |                      | Autophagy           |  |
|                |       |                      | Autophugy           |  |
| Irgm2          | 0.832 | Immunity-related     | Interferon-         | Function not fully known, but most likely        |
|                |       | GTPase family M      | inducible protein 1 | regulates autophagy and pro-inflammatory         |
|                |       | member 2             | (IFI1)              | cytokine production                              |
| <u>\$100a8</u> | 2.3   | S100 calcium-binding | Calgranulin A       | Calcium- and zinc-binding protein involved in    |
|                |       | protein A8           |                     | pro-inflammatory, antimicrobial, oxidant-        |
|                |       |                      |                     | scavenging and apoptosis-inducing activities;    |

|            |       |                 |                     | can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including |
|------------|-------|-----------------|---------------------|---|
|            |       |                 |                     | adhesion, apoptosis, autophagy, cytoskeletal  |
|            |       |                 |                     | remodeling, cytokine production, chemotaxis,  |
|            |       |                 |                     | migration, inflammation, arachidonic acid   |
|            |       |                 |                     | metabolism, oxidant-scavenging, and PRR   |
|            |       |                 |                     | signaling   |
| <u>Ubc</u> | 0.653 | Polyubiquitin C |                     | Serves various roles, including innate  |
|            |       |                 |                     | immunity, DNA repair, and stimulation of  |
|            |       |                 |                     | autophagy and the proteasomal response  |
|            |       |                 | B Cell Function     |   |
| Blnk       | 1.54  | B cell linker   | Src homology 1      | Functions as a central linker protein   |
|            |       |                 | domain-containing   | downstream of the B cell receptor, bridging   |
|            |       |                 | leukocyte protein   | SYK kinase to a multitude of signaling  |
|            |       |                 | of 65 kDa (SLP-65); | pathways and regulating biological outcomes   |
|            |       |                 | Ly57                | of B cell function and development; plays a   |
|            |       |                 |                     | role in the activation of ERK/EPHB2, MAP  |
|            |       |                 |                     | kinase p38 and JNK; modulates AP1 activation;   |
|            |       |                 |                     | important for the activation of NFkB and NFAT   |

| <u>Btk</u>  | 1.2  | Bruton's tyrosine  |                      | Crucial kinase in B cell receptor signal          |
|-------------|------|--------------------|----------------------|---|
|             |      | kinase             |                      | transmission and B cell activation                |
| <u>Cd22</u> | 1.67 | Cluster of         | Sialic acid-binding  | Mediates interactions between B cells; binds      |
|             |      | differentiation 22 | Ig-like lectin 2     | CD45  |
|             |      |                    | (SIGLEC2)            |   |
|             |      |                    | B lymphocyte cell    |   |
|             |      |                    | adhesion molecule    |   |
|             |      |                    | (BL-CAM)             |   |
| <u>Cd37</u> | 2.07 | Cluster of         | Tetraspanin-26       | Cell surface glycoprotein known to complex        |
|             |      | differentiation 37 |                      | with integrins and other transmembrane 4          |
|             |      |                    |                      | superfamily proteins; may play a role in T cell-  |
|             |      |                    |                      | B cell interactions                               |
| <u>Cd48</u> | 2.08 | Cluster of         | B-lymphocyte         | B cell-specific cellular differentiation Ag; when |
|             |      | differentiation 48 | activation marker    | bound to CD2, promotes T cell activation, and     |
|             |      |                    | (BLAST-1); signaling | the formation of lipid rafts and caveolae for     |
|             |      |                    | lymphocytic          | macrophages                                       |
|             |      |                    | activation molecule  |   |
|             |      |                    | 2 (SLAMF2)           |   |
| <u>Cd69</u> | 1.46 | Cluster of         | C-type lectin        | Signal transmitting receptor in lymphocytes,      |
|             |      | differentiation 69 | domain family 2,     | NK cells, and platelets; induced upon T cell      |
|             |      |                    | member C             | activation; involved in lymphocyte                |
|             |      |                    |                      | proliferation                                     |

| <u>Cd79b</u>  | 1.91  | Cluster of differentiation 79b     | B29                                      | One of the two flanking proteins that initiate signaling downstream of the BCR  |
|---------------|-------|------------------------------------|--|---|
| Cxcl13        | 1.58  | C-X-C motif<br>chemokine ligand 13 | BLC, BCA-1                               | B cell chemokine induced by type I interferons; participates in GC formation  |
| Fcgr1         | 2.08  | Fc fragment of IgG receptor Ia     | CD64                                     | High affinity receptor for the Fc region of $\gamma$ -lgs; functions in both innate and adaptive immune responses   |
| Fcgr2b        | 1.41  | Fc fragment of IgG receptor IIb    | CD32                                     | Low affinity receptor for the Fc region of complexed or aggregated y-lgs; involved in a variety of effector and regulatory functions such as phagocytosis of immune complexes and modulation of Ab production by B cells; essential for the maintenance of humoral tolerance; acts as a late checkpoint at the level of class-switched memory B cells, plasmablasts or plasma cells; regulates plasma-cell homeostasis and survival |
| <u>Gpr183</u> | 0.932 | G protein-coupled receptor 183     | EBV-induced G protein-coupled receptor 2 | Lymphocyte GPCR that acts as a chemotactic receptor for B cells, T cells, splenic DCs, monocytes/macrophages, and astrocytes  |
| <u>lkzf1</u>  | 1.19  | IKAROS family zinc finger 1        |  | Transcriptional regulator of hematopoietic cell differentiation; plays a role in T and B cell development   |

| <u>Lyn</u>   | 0.921 | Lck/Yes-related  |            | Src family tyrosine kinase that potentiates  |
|--------------|-------|--|------------|--|
|              |       | novel kinase   |            | signaling from the B cell receptor and CD40  |
| Mef2c        | 1.24  | Myocyte enhancer   |            | Transcriptional activator that binds specifically  |
|              |       | factor 2c  |            | to the MEF2 element present in the regulatory  |
|              |       |  |            | regions of many muscle-specific genes;   |
|              |       |  |            | controls cardiac morphogenesis and   |
|              |       |  |            | myogenesis, and is also involved in vascular   |
|              |       |  |            | development; required for B cell survival and  |
|              |       |  |            | proliferation in response to BCR stimulation,  |
|              |       |  |            | efficient IgG1 Ab responses to T cell-   |
|              |       |  |            | dependent Ags, and for normal induction of   |
|              |       |  |            | GC B cells   |
|              |       |  |            | 30.5 30.0  |
| <u>Ms4a1</u> | 2.26  | Membrane spanning  | CD20; Bp35 | B cell-specific membrane protein that  |
| <u>Ms4a1</u> | 2.26  | Membrane spanning 4-domains A1   | CD20; Bp35 |  |
| <u>Ms4a1</u> | 2.26  |  | CD20; Bp35 | B cell-specific membrane protein that  |
| <u>Ms4a1</u> | 2.26  |  | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel   |
| Ms4a1 Pik3cd | 2.26  |  | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR  |
|              |       | 4-domains A1   | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation   |
|              |       | 4-domains A1  Phosphatidylinositol-                                      | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation  A subunit of PI3K; acts downstream of TLR4,  |
|              |       | 4-domains A1  Phosphatidylinositol- 4,5-bisphosphate 3-                  | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation  A subunit of PI3K; acts downstream of TLR4, TCR, BCR, and CD40; contributes to T helper  |
|              |       | 4-domains A1  Phosphatidylinositol- 4,5-bisphosphate 3- kinase catalytic | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation  A subunit of PI3K; acts downstream of TLR4, TCR, BCR, and CD40; contributes to T helper cell expansion, mast cell development, and |

| Pik3cg       | 1.21 | Phosphatidylinositol-  |                    | A subunit of PI3K; modulates leukocyte                   |
|--------------|------|------------------------|--------------------|--|
|              |      | 4,5-bisphosphate 3-    |                    | chemotaxis to inflammatory sites and in                  |
|              |      | kinase catalytic       |                    | response to chemoattractant agents                       |
|              |      |                        |                    | response to enemoattractant agents                       |
|              |      | subunit gamma          |                    |  |
|              |      | isoform                |                    |  |
| Pou2f2       | 1.81 | POU domain class 2,    |                    | TF that regulates Ab and IL-6 expression in B            |
|              |      | transcription factor 2 |                    | cells  |
| Prdm1        | 1.77 | Positive regulatory    | B lymphocyte-      | TF that plays a role in the development,                 |
|              |      | domain I-binding       | induced            | retention, and long-term establishment of T              |
|              |      | factor                 | maturation protein | cell, NK cell, and NK-T cells in non-lymphoid            |
|              |      |                        | (BLIMP1)           | organs; drives the maturation of B cell into Ig          |
|              |      |                        |                    | secreting cells  |
| <u>Spn</u>   | 1.63 | Sialophorin            | Leukosialin; CD43  | Cell surface sialoglycoprotein expressed by T            |
|              |      |                        |                    | cells, B cells, monocytes, and granulocytes;             |
|              |      |                        |                    | promotes lymph node localization in T cells;             |
|              |      |                        |                    | shunts T cells away from the T <sub>H</sub> 2 phenotype  |
|              |      |                        |                    | and towards T <sub>H</sub> 1; promotes the expression of |
|              |      |                        |                    | IFNγ in CD4 <sup>+</sup> T cells                         |
| <u>Syk</u>   | 1.95 | Spleen-associated      |                    | Critical kinase that transmits signals from the          |
|              |      | tyrosine kinase        |                    | TCR and BCR  |
| <u>Zap70</u> | 1.72 | Zeta chain of T cell   |                    | Tyrosine kinase that plays an essential role in          |
|              |      | receptor associated    |                    | regulation of the adaptive immune response;              |
|              |      | protein kinase 70      |                    | regulates motility, adhesion, proliferation,             |

|               |      |   |  | differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  |
|---------------|------|---|--|--|
|               |      |   | Cell Cycle   |  |
|               |      |   | •  |  |
| Ccnd3  Cdkn1a | 1.39 | Cyclin D3  Cyclin dependent kinase inhibitor 1A | p21; CDK-<br>interaction protein<br>1 (CIP1)               | Regulatory component of the cyclin D3-CDK4  complex that inhibitively phosphorylates  members of the retinoblastoma protein  family; regulates the cell-cycle during G1/S  transition  Binds to and inhibits cyclin-dependent kinase activity, preventing phosphorylation of critical cyclin-dependent kinase substrates and blocking cell cycle progression |
|               |      |   | Chemotaxis   |  |
|               |      |   | CHEIHOLAXIS  |  |
| Ccl2          | 1.9  | C-C motif chemokine                             |  | Chemoattractant ligand for CCR2 and -4; attracts monocytes and basophils   |
| Ccl3          | 3.74 | C-C motif chemokine ligand 3                    | Macrophage inflammatory protein $1\alpha$ (MIP $1\alpha$ ) | Chemoattractant ligand for CCR1, -4, and -5  |

| <u>Ccl4</u>  | 3.44  | C-C motif chemokine | Macrophage         | Chemoattractant for NK cells and monocytes;    |
|--------------|-------|---------------------|--------------------|--|
|              |       | ligand 4            | inflammatory       | binds to CCR5 receptors                        |
|              |       |                     | protein 1β (MIP1β) |  |
|              |       |                     |                    |  |
| <u>Ccl5</u>  | 3.21  | C-C motif chemokine | Regulated upon     | Chemoattractant ligand for CCR1, -3, -4, and - |
|              |       | ligand 5            | activation,        | 5; attracts blood monocytes, memory T helper   |
|              |       |                     | normally T-        | cells and eosinophils; causes the release of   |
|              |       |                     | expressed, and     | histamine from basophils and activates         |
|              |       |                     | presumably         | eosinophils                                    |
|              |       |                     | secreted (RANTES)) |  |
|              |       |                     |                    |  |
| <u>Ccl7</u>  | 1.57  | C-C motif chemokine | Monocyte           | General chemokine that recruits leukocytes to  |
|              |       | ligand 7            | chemotactic        | infected tissues; mainly observed in monocyte  |
|              |       |                     | protein 3 (MCP3)   | mobilization                                   |
| Ccl8         | 1.56  | C-C motif chemokine | Monocyte           | General chemokine that recruits leukocytes to  |
|              |       | ligand 8            | chemoattractant    | infected tissues                               |
|              |       | liganu o            |                    | infected tissues                               |
|              |       |                     | protein 2 (MCP2)   |  |
| <u>Ccl11</u> | 0.972 | C-C motif chemokine | Eotaxin            | Chemoattractant for eosinophils                |
|              |       | ligand 11           |                    |  |
|              |       |                     |                    |  |
| <u>Ccl12</u> | 0.874 | C-C motif chemokine | Monocyte           | Chemoattractant specific for eosinophils,      |
|              |       | ligand 12           | chemotactic        | monocytes, and lymphocytes; found primarily    |
|              |       |                     | protein 5 (MCP5)   | in the lymph nodes and thymus, but can be      |
|              |       |                     |                    | strongly expressed by macrophages              |
|              |       |                     |                    |  |

| Ccl24 |       | C-C motif chemokine               | Eotaxin-2      | Chemoattractant for resting T cells and   |
|-------|-------|-----------------------------------|----------------|---|
|       | 1.25  | ligand 24                         |                | eosinophils   |
| Ccr1  | 2.27  | C-C motif chemokine receptor 1    | MIP1α receptor | Receptor for CCL3, -5, -7, and -23  |
| Ccr3  | 1.42  | C-C motif chemokine<br>receptor 3 | CD193          | Receptor for a variety of chemokines, including CCL11, CCL26, CCL7, CCL13, CCL5 (RANTES), and CCL15; signals through Ca(2+) flux                            |
| Ccr5  | 1.37  | C-C motif chemokine<br>receptor 5 | CD195          | Receptor for a number of inflammatory CC-chemokines, including CCL3/MIP1 $\alpha$ , CCL4/MIP1 $\beta$ , and RANTES; signals via Ca(2+) flux                 |
| Ccr7  | 2.19  | C-C chemokine<br>receptor type 7  | CD197          | Chemokine receptor that activates B and T cells and promotes their homing to secondary lymphoid organs; also stimulates DC expression of MHC class I and II |
| Ccr9  | 0.792 | C-C motif chemokine<br>receptor 9 |                | Receptor for CCL25; increases intracellular  Ca(2+) levels upon ligand binding  |
| Ccrl2 | 2.19  | C-C chemokine<br>receptor-like 2  |                | Stabilizes TLR4 surface expression on macrophages   |

| <u>Cxcl1</u> | 1.94  | C-X-C motif                           | GRO1 oncogene      | Chemoattractant ligand for CXCR2; plays a role           |
|--------------|-------|---------------------------------------|--------------------|--|
|              |       | chemokine ligand 1                    |                    | in inflammation and as a chemoattractant for neutrophils |
|              |       |                                       |                    | ·  |
| Cxcl2        | 3.12  | C-X-C motif                           | Macrophage         | Chemokine produced by activated monocytes                |
|              |       | chemokine ligand 2                    | inflammatory       | and neutrophils and expressed at sites of                |
|              |       |                                       | protein 2-alpha    | inflammation   |
|              |       |                                       | (MIP2α); GRO2      |  |
|              |       |                                       | oncogene           |  |
| Cxcl3        | 3.93  | C-X-C motif                           | GRO3 oncogene      | Ligand for CXCR2; attracts neutrophils                   |
|              |       | chemokine ligand 3                    |                    |  |
| Cxcl9        | 0.947 | C-X-C motif                           | Humig              | Chemoattractant ligand for CXCR3; attracts               |
|              |       | chemokine ligand 9                    |                    | activated T cells  |
|              |       | , , , , , , , , , , , , , , , , , , , |                    |  |
| Cxcl10       | 1.32  | C-X-C motif                           | IFNγ-induced       | Macrophage, DC, T cell, and NK cell                      |
|              |       | chemokine ligand 10                   | protein 10 (IP-10) | chemoattractant secreted by several cell types           |
|              |       |                                       |                    | in response to IFNγ; binds to CXCR3                      |
| Cxcl11       | 1.24  | C-X-C motif                           |                    | Dominant ligand for CXCR3; attracts activated            |
|              |       | chemokine ligand 11                   |                    | T cells; strongly induced by IFNγ                        |
| Cxcl12       | 1.29  | C-X-C motif                           | Stromal cell-      | Ubiquitously expressed chemokine that acts a             |
|              |       | chemokine ligand 12                   | derived factor 1   | strong chemoattractant for lymphocytes                   |
|              |       |                                       | (SDF1)             |  |
|              |       |                                       |                    |  |
| Cxcl13       | 1.58  | C-X-C motif                           | BLC, BCA-1         | B cell chemokine induced by type I                       |
|              |       | chemokine ligand 13                   |                    | interferons; participates in GC formation                |
|              |       |                                       |                    |  |

| Cxcl16        | 1.68  | C-X-C motif         |                 | Chemoattractant for T cells and NKT cells                    |
|---------------|-------|---------------------|-----------------|--|
|               |       | chemokine ligand 16 |                 | produced by DCs in response to IFNγ and                      |
|               |       |                     |                 | ΤΝΓα   |
|               |       |                     |                 |  |
| Cxcr2         | 2.57  | C-X-C motif         | CD182; IL-8     | Receptor for IL-8 and CXCL3; powerful                        |
|               |       | chemokine receptor  | receptor B      | chemoattractant for neutrophils                              |
|               |       | 2                   |                 |  |
| Cxcr3         | 2.23  | C-X-C motif         | CD183           | Induces integrin activation, cytoskeletal                    |
|               |       | chemokine receptor  |                 | remodeling, and chemotaxis; expressed by T                   |
|               |       | 3                   |                 | cells and NK cells; prominently expressed in                 |
|               |       |                     |                 | effector and memory T cells                                  |
| Cxcr4         | 1.74  | C-X-C motif         | CD184; fusin    | Alpha-chemokine receptor specific for SDF1                   |
|               |       | chemokine receptor  |                 | aka CXCL12   |
|               |       | 4                   |                 |  |
|               |       |                     |                 |  |
| <u>Cxcr6</u>  | 2.3   | C-X-C motif         | CD186           | Receptor for the C-X-C chemokine CXCL16;                     |
|               |       | chemokine receptor  |                 | expressed in several T lymphocyte subsets and                |
|               |       | 6                   |                 | bone marrow stromal cells                                    |
| <u>Gpr183</u> | 0.932 | G protein-coupled   | EBV-induced G   | Lymphocyte GPCR that acts as a chemotactic                   |
|               |       | receptor 183        | protein-coupled | receptor for B cells, T cells, splenic DCs,                  |
|               |       |                     | receptor 2      | monocytes/macrophages, and astrocytes                        |
| <u>Isg15</u>  | 2.41  | Interferon-         |                 | Ubiquitin-like protein that binds intracellular              |
|               |       | stimulated gene 15  |                 | target proteins upon activation by IFN $\alpha$ or $\beta$ ; |
|               |       |                     |                 | can also be secreted to induce NK cell                       |
|               |       |                     |                 | proliferation, act as a chemoattractant for                  |

|                |      |  |               | neutrophils, and induce IFNγ upon binding to  ITGAL/ITGB2  |
|----------------|------|--|---------------|--|
| <u>Itgam</u>   | 2.19 | Integrin alpha M   | CD11b         | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker   |
| Pik3cd         | 1.19 | Phosphatidylinositol-<br>4,5-bisphosphate 3-<br>kinase catalytic<br>subunit delta<br>isoform |               | A subunit of PI3K; acts downstream of TLR4,  TCR, BCR, and CD40; contributes to T helper  cell expansion, mast cell development, and  neutrophil chemotaxis, extravasation, and  respiratory burst   |
| <u>\$100a8</u> | 2.3  | S100 calcium-binding protein A8  | Calgranulin A | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid |

| Xcl1        | 0.805 | X-C motif chemokine ligand 1          | Lymphotaxin        | metabolism, oxidant-scavenging, and PRR signaling  Chemoattractant for lymphocytes but not monocytes or neutrophils; mediates medullary accumulation of thymic DCs and contributes to Treg development |
|-------------|-------|---------------------------------------|--------------------|--|
|             |       | Comple                                | ement & Humoral In | nmunity  |
| <u>C1qa</u> | 0.818 | Complement C1q subcomponent subunit A |                    | A chain of the C1q complex, which acts as the Ag-Ab-binding subunit of the C1 complex  |
| <u>C1qb</u> | 1.17  | Complement C1q subcomponent subunit B |                    | B chain of the C1q complex, which acts as the Ag-Ab-binding subunit of the C1 complex  |
| <u>C1ra</u> | 0.793 | Complement C1r-A subcomponent         |                    | Proteolytic subunit of the C1 complex that enzymatically cleaves C1s   |
| <u>C1s1</u> | 1.07  | Complement component 1s               | C1 esterase        | Serine protease that enzymatically cleaves C4 and C2   |
| <u>C2</u>   | 0.757 | Complement component 2                |                    | Cleaved by activated factor C1 into two fragments: C2b and C2a; C2a combines with C4b to generate C3 or C5 convertase  |

| <u>C3</u>    | 1.17  | Complement         |                    | Cleaved by C3 convertase to form C3a and       |
|--------------|-------|--------------------|--------------------|--|
|              |       | component 3        |                    | C3b, an anaphalotoxin and an opsonizing        |
|              |       |                    |                    | agent, respectively                            |
| <u>C3ar1</u> | 1.5   | Complement         |                    | GPCR that binds to C3a, activating chemotaxis, |
|              |       | component 3a       |                    | granule enzyme release, superoxide anion       |
|              |       | receptor 1         |                    | production, and bacterial opsonization         |
| C4b          | 0.684 | Complement         |                    | Mediates interactions between Ab-bound Ags     |
|              |       | component 4B       |                    | and other complement components                |
| <u>C6</u>    | 1.02  | Complement         |                    | Part of the membrane attack complex            |
|              |       | component 6        |                    |  |
| <u>Cd55</u>  | 2     | Cluster of         | Complement         | Cell surface glycoprotein that interacts with  |
|              |       | differentiation 55 | decay-accelerating | surface-bound C4b and inhibits its conversion  |
|              |       |                    | factor             | of C2 to C2b                                   |
| <u>Cfb</u>   | 1.37  | Complement factor  |                    | Alternate complement pathway component;        |
|              |       | В                  |                    | when cleaved, produces a serine protease that  |
|              |       |                    |                    | binds to C3b to form C3 convertase             |
| Cfd          | 3.27  | Complement factor  | Adipsin            | Chymotrypsin-family peptidase that cleaves     |
|              |       | D                  |                    | factor B when the latter is complexed with     |
|              |       |                    |                    | factor C3b, activating C3 convertase           |
| Cfp          | 1.5   | Complement factor  |                    | Alternate complement pathway component;        |
|              |       | properdin          |                    | when cleaved, produces a serine protease that  |
|              |       |                    |                    | binds to C3b to form C3 convertase             |
|              |       |                    |                    |  |

| Fcgr1       | 2.08 | Fc fragment of IgG   | CD64               | High affinity receptor for the Fc region of γ-lgs; |
|-------------|------|----------------------|--------------------|--|
|             |      | receptor la          |                    | functions in both innate and adaptive immune       |
|             |      |                      |                    | responses  |
| Fcgr2b      | 1.41 | Fc fragment of IgG   | CD32               | Low affinity receptor for the Fc region of         |
|             |      | receptor IIb         |                    | complexed or aggregated γ-lgs; involved in a       |
|             |      | ,                    |                    | variety of effector and regulatory functions       |
|             |      |                      |                    | such as phagocytosis of immune complexes           |
|             |      |                      |                    | and modulation of Ab production by B cells;        |
|             |      |                      |                    | essential for the maintenance of humoral           |
|             |      |                      |                    | tolerance; acts as a late checkpoint at the level  |
|             |      |                      |                    | of class-switched memory B cells,                  |
|             |      |                      |                    | plasmablasts or plasma cells; regulates            |
|             |      |                      |                    | plasma-cell homeostasis and survival               |
| Fcgr4       | 3.21 | Fragment             | Fc receptor-like 3 | Putative mouse ortholog to human FcγRIIIA          |
|             |      | crystallizable gamma | (Fcrl3); CD16-2    |  |
|             |      | receptor 4           |                    |  |
|             |      |                      |                    |  |
|             |      |                      | Costimulation      |  |
| <u>Cd28</u> | 2.48 | Cluster of           |                    | Essential T cell co-receptor that enhances T       |
|             |      | differentiation 28   |                    | cell activation, proliferation, cytokine           |
|             |      |                      |                    | production, and survival; binds to CD80 and        |
|             |      |                      |                    | CD86   |
|             |      |                      |                    |  |

| <u>Cd40</u>  | 1.9  | Cluster of          |                 | APC-expressed costimulatory protein that            |
|--------------|------|---------------------|-----------------|---|
|              |      | differentiation 40  |                 | binds to CD40L on CD4 <sup>+</sup> T cells, causing |
|              |      |                     |                 | activation of both                                  |
| <u>Cd80</u>  | 1.04 | Cluster of          | B7-1            | One of the two ligands for the CD28                 |
|              |      | differentiation 80  |                 | costimulatory receptor and the CTLA4                |
|              |      |                     |                 | inhibitory receptor, the other being CD86           |
| <u>Cd86</u>  | 0.98 | Cluster of          | B7-2            | One of the two ligands for the CD28                 |
|              |      | differentiation 86  |                 | costimulatory receptor and the CTLA4                |
|              |      |                     |                 | inhibitory receptor, the other being CD80           |
| <u>lcam1</u> | 1.69 | Intracellular       | CD54            | Cell surface glycoprotein that serves as strong     |
|              |      | adhesion molecule 1 |                 | adhesive ligand for LFA-1; important for            |
|              |      |                     |                 | leukocyte mobility and costimulation                |
| Icos         | 2.45 | Inducible T cell    | CD278           | Enhances all basic T cell responses to foreign      |
|              |      | costimulator        |                 | Ag; essential both for efficient interaction        |
|              |      |                     |                 | between T and B cells and for normal Ab             |
|              |      |                     |                 | responses to T cell-dependent Ags                   |
| <u>Ptprc</u> | 1.5  | Protein tyrosine    | CD45; leukocyte | Delivers costimulation during T cell activation     |
|              |      | phosphatase         | common antigen  | upon binding to its ligand DPP4;                    |
|              |      | receptor type C     | (LCA)           | dephosphorylates Lyn and suppresses JAK             |
|              |      |                     |                 | kinases   |

| Tnfsf14   | 2.19  | TNF superfamily      | CD258; LIGHT;      | Cytokine that binds to TNFRSF3/LTBR and          |
|-----------|-------|----------------------|--------------------|--|
|           |       | member 14            | Herpesvirus entry  | TNFRSF14/HVEM; delivers costimulation to T       |
|           |       |                      | mediator ligand    | cells; triggers apoptosis of various tumor cells |
|           |       |                      | (HVEML)            |  |
| Tnfrsf4   | 1.61  | Tumor necrosis       | OX40; CD134        | Receptor for TNFSF4/OX40L/GP34;                  |
| <u></u>   | 1.01  | factor receptor      | ON 10, 02 10 1     | costimulatory molecule implicated in long-       |
|           |       | superfamily member   |                    | term T cell immunity; activates NFkB through     |
|           |       |                      |                    |  |
|           |       | 4                    |                    | its interaction with adaptor proteins TRAF2      |
|           |       |                      |                    | and TRAF5; suppresses apoptosis through          |
|           |       |                      |                    | upregulation of BCL2                             |
| Tnfrsf11a | 1.16  | Tumor necrosis       | Receptor activator | Recruits TRAFs and activates NFkB and JNK        |
|           |       | factor receptor      | of NFkB (RANK)     | activation upon binding to RANKL on the          |
|           |       | superfamily member   |                    | surface of T cells                               |
|           |       | 11A                  |                    |  |
| Tnfrsf14  | 0.731 | Tumor necrosis       |                    |  |
|           |       | factor receptor      |                    |  |
|           |       | superfamily member   |                    |  |
|           |       | 14                   |                    |  |
|           |       |                      |                    |  |
|           |       |                      | Cytokines          |  |
| Csf1      | 0.698 | Macrophage colony-   |                    | Cytokine that promote activation and survival    |
|           |       | stimulating factor 1 |                    | of monocytes                                     |

| Csf1r         | 1.11  | Macrophage colony-   | CD115           | Receptor for CSF1; promotes release of          |
|---------------|-------|----------------------|-----------------|---|
|               |       | stimulating factor 1 |                 | inflammatory cytokines in response to IL-34     |
|               |       | receptor             |                 | and CSF1  |
| <u>Ifnar1</u> | 0.731 | Interferon-          |                 | Component of the receptor for type I IFNs,      |
|               |       | alpha/beta receptor  |                 | binding of which activates the JAK-STAT         |
|               |       | alpha chain          |                 | pathway   |
| Ifng          | 1.93  | Interferon gamma     |                 | T and NK cell-secreted inflammatory cytokine    |
|               |       |                      |                 | that stimulates cytolysis, activates            |
|               |       |                      |                 | macrophages, and stimulates MHC class II        |
|               |       |                      |                 | expression                                      |
| Ifngr1        | 0.707 | Interferon gamma     | CD54            | One of the two components of the IFNy           |
|               |       | receptor 1           |                 | receptor; stimulates activation of the          |
|               |       |                      |                 | JAK/STAT signaling pathway                      |
| <u>II1a</u>   | 3.29  | Interleukin 1 alpha  | Hematopoietin-1 | Cytokine produced by monocytes and              |
|               |       |                      |                 | macrophages in response to cell injury;         |
|               |       |                      |                 | stimulates thymocyte proliferation by inducing  |
|               |       |                      |                 | IL-2 release; also stimulates B cell maturation |
|               |       |                      |                 | and proliferation, and fibroblast growth factor |
|               |       |                      |                 | activity  |
| <u>II1b</u>   | 2.04  | Interleukin 1 beta   | Catabolin       | One of the two primary inflammatory             |
|               |       |                      |                 | cytokines produced by the inflammasome (the     |
|               |       |                      |                 | other one being IL-18); induces neutrophil      |
|               |       |                      |                 | influx and activation, T cell activation and    |

| <u>Il1rap</u> | 1.27 | Interleukin 1 receptor accessory protein |   | cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNγ synthesis from T <sub>H</sub> 1 cells  Co-receptor for several ligands, including IL-1R1 in the IL-1 pathway, IL-RL1 in the IL-33 pathway, IL-1RL2 in the IL-36 pathway; signaling involves Tollip, MyD88, IRAK1, and |
|---------------|------|--|---|---|
|               |      |  |   | IRAK2   |
| II2ra         | 2    | Interleukin 2 receptor subunit alpha     | CD25  | Alpha chain of the IL-2 receptor  |
| <u>ll2rg</u>  | 1.53 | Interleukin 2 receptor subunit gamma     | Common gamma<br>chain; CD132                | Common subunit for the receptors for a variety of interleukins, including IL-2, -4, -7, and -21   |
| <u>II6</u>    | 2.7  | Interleukin 6                            |   | Pro-inflammatory cytokine that signals through the JAK and STAT pathways  |
| <u>117r</u>   | 2.16 | Interleukin 7 receptor                   | CD127                                       | Receptor for IL-7   |
| <u>II10</u>   | 1.01 | Interleukin 10                           | Cytokine synthesis inhibitory factor (CSIF) | Major immunoregulatory cytokine that inhibits production of pro-inflammatory cytokines, including GM-CSF, G-CSF, IL-1α, IL-1β, IL-6, IL-8, and TNFα; also interferes with Ag  |

| <u>ll12rb1</u> | 1.07  | Interleukin 12<br>receptor subunit<br>beta 1 | CD212     | presentation by reducing expression of MHC  class II and costimulatory molecules, thereby  inhibiting their ability to induce T cell  activation  Cytokine receptor component that associates  with IL12RB2 to IL23R   |
|----------------|-------|--|-----------|--|
| <u>ll13ra1</u> | 0.717 | Interleukin 13 receptor subunit alpha 1      | CD213a1   | Pairs with IL4RA to form to form the IL-13 receptor; may mediate the signaling processes that lead to the activation of JAK1, STAT3, and STAT6 induced by IL-4 and -13   |
| <u>ll18r1</u>  | 1.36  | Interleukin 18 receptor 1                    | CD218a    | Receptor for IL-18   |
| Tgfb1          | 0.803 | Transforming growth factor beta 1            |           | Multifunctional protein that regulates the growth and differentiation of various cell types and is involved in various processes, such as normal development, immune function, microglia function and responses to neurodegeneration; can induce EMT and cell migration in various cell types; frequently acts as an immunosuppressive cytokine in the TME |
| Tnf            | 3.73  | Tumor necrosis<br>factor                     | Cachectin | Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and  |

|                |       |                    |                    | TNFRSF1B/TNFBR; capable of inducing cell         |
|----------------|-------|--------------------|--------------------|--|
|                |       |                    |                    | death in certain tumor cell lines                |
| Tnfsf14        | 2.19  | TNF superfamily    | CD258; LIGHT;      | Cytokine that binds to TNFRSF3/LTBR and          |
|                |       | member 14          | Herpesvirus entry  | TNFRSF14/HVEM; delivers costimulation to T       |
|                |       |                    | mediator ligand    | cells; triggers apoptosis of various tumor cells |
|                |       |                    | (HVEML)            |  |
| <u>Tnfrsf4</u> | 1.61  | Tumor necrosis     | OX40; CD134        | Receptor for TNFSF4/OX40L/GP34;                  |
|                |       | factor receptor    |                    | costimulatory molecule implicated in long-       |
|                |       | superfamily member |                    | term T cell immunity; activates NFkB through     |
|                |       | 4                  |                    | its interaction with adaptor proteins TRAF2      |
|                |       |                    |                    | and TRAF5; suppresses apoptosis through          |
|                |       |                    |                    | upregulation of BCL2                             |
| Tnfrsf11a      | 1.16  | Tumor necrosis     | Receptor activator | Recruits TRAFs and activates NFkB and JNK        |
|                |       | factor receptor    | of NFκB (RANK)     | activation upon binding to RANKL on the          |
|                |       | superfamily member |                    | surface of T cells                               |
|                |       | 11A                |                    |  |
| Tnfrsf14       | 0.731 | Tumor necrosis     |                    |  |
|                |       | factor receptor    |                    |  |
|                |       | superfamily member |                    |  |
|                |       | 14                 |                    |  |
|                |       |                    | Cytotoxicity       |  |
|                |       |                    |                    |  |
| <u>Gzmb</u>    | 3.8   | Granzyme B         | Fragmentin 2       | Abundant protease in the cytosolic granules of   |
|                |       |                    |                    | cytotoxic T and NK cells that activates caspase- |

|                |      |  |                                       | mediated cell death when delivered into the target cell through the immunological synapse  |
|----------------|------|--|---------------------------------------|--|
| <u>Gzmk</u>    | 3.36 | Granzyme K                               | Tryptase II                           | Granule-secreted, pro-apoptotic serine protease found in the cytoplasmic granules of CTLs  |
| Fcgr4          | 3.21 | Fragment crystallizable gamma receptor 4 | Fc receptor-like 3<br>(Fcrl3); CD16-2 | Putative mouse ortholog to human FcγRIIIA  |
| <u>Itgam</u>   | 2.19 | Integrin alpha M                         | CD11b                                 | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker   |
| <u>\$100a8</u> | 2.3  | S100 calcium-binding protein A8          | Calgranulin A                         | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid |

|               |                         |  |  | metabolism, oxidant-scavenging, and PRR signaling   |  |  |  |
|---------------|-------------------------|--|--|---|--|--|--|
|               | Dendritic Cell Function |  |  |   |  |  |  |
| <u>Casp8</u>  | 0.536                   | Caspase 8                                      |  | Cysteine-aspartic acid protease; cleaves and activates effector caspases CASP3, -4, -6, -7, -9, and -10; initiates pyroptosis through cleavage of gasdermin-D; mediates noncanonical cleavage of IL-1β in DCs and macrophages |  |  |  |
| Cxcl16        | 1.68                    | C-X-C motif<br>chemokine ligand 16             |  | Chemoattractant for T cells and NKT cells produced by DCs in response to IFN $\gamma$ and TNF $\alpha$  |  |  |  |
| <u>Gpr183</u> | 0.932                   | G protein-coupled receptor 183                 | EBV-induced G protein-coupled receptor 2 | Lymphocyte GPCR that acts as a chemotactic receptor for B cells, T cells, splenic DCs, monocytes/macrophages, and astrocytes  |  |  |  |
| Tigit         | 1.28                    | T cell immunoreceptor with Ig and ITIM domains |  | Binds with high affinity to the poliovirus receptor, causing increased secretion of IL-10, decreased secretion of IL-12B, and suppressing T cell activation by promoting the generation of mature immunoregulatory DCs        |  |  |  |
| <u>Itgax</u>  | 1.47                    | Integrin alpha X                               | CD11c                                    | Adhesion molecule; signature marker of Agpresenting DCs   |  |  |  |

|              | Growth/Proliferation |                       |                      |  |  |  |
|--------------|----------------------|-----------------------|----------------------|--|--|--|
| Lcn2         | 1.97                 | Lipocalin 2           | Neutrophil           | Neutrophil-secreted factor that sequesters       |  |  |
|              |                      |                       | gelatinase-          | iron-containing siderophores; also functions as  |  |  |
|              |                      |                       | associated lipocalin | a growth factor                                  |  |  |
|              |                      |                       | (NGAL)               |  |  |  |
|              |                      |                       |                      |  |  |  |
| Map2k1       | 0.266                | Dual specificity      | MAPK/ERK kinase 1    | Essential component of the MAP kinase signal     |  |  |
|              |                      | mitogen-activated     | (MEK1)               | transduction pathway; participates in            |  |  |
|              |                      | protein kinase kinase |                      | numerous biological functions, including cell    |  |  |
|              |                      | 1                     |                      | growth, adhesion, survival, differentiation,     |  |  |
|              |                      |                       |                      | transcription, metabolism, and cytoskeletal      |  |  |
|              |                      |                       |                      | remodeling                                       |  |  |
|              |                      |                       |                      |  |  |  |
| Tgfb1        | 0.803                | Transforming growth   |                      | Multifunctional protein that regulates the       |  |  |
|              |                      | factor beta 1         |                      | growth and differentiation of various cell       |  |  |
|              |                      |                       |                      | types and is involved in various processes,      |  |  |
|              |                      |                       |                      | such as normal development, immune               |  |  |
|              |                      |                       |                      | function, microglia function and responses to    |  |  |
|              |                      |                       |                      | neurodegeneration; can induce EMT and cell       |  |  |
|              |                      |                       |                      | migration in various cell types; frequently acts |  |  |
|              |                      |                       |                      | as an immunosuppressive cytokine in the TME      |  |  |
| <u>Vegfa</u> | 0.592                | Vascular endothelial  |                      | Glycosylated mitogen that promotes vascular      |  |  |
|              |                      | growth factor A       |                      | permeability, vasculogenesis, angiogenesis,      |  |  |
|              |                      |                       |                      | and cell migration                               |  |  |
|              | Hematopoiesis        |                       |                      |  |  |  |
|              |                      |                       |                      |  |  |  |

|               |      | 1                    |                |  |
|---------------|------|----------------------|----------------|--|
| <u>Hck</u>    | 1.76 | Hematopoietic cell   |                | Src family tyrosine kinase that mediates         |
|               |      | kinase               |                | secretory lysosome mobilization,                 |
|               |      |                      |                | degranulation, and activation of NADPH           |
|               |      |                      |                | oxidase  |
|               |      |                      |                |  |
| <u>lkzf1</u>  | 1.19 | IKAROS family zinc   |                | Transcriptional regulator of hematopoietic cell  |
|               |      | finger 1             |                | differentiation; plays a role in T and B cell    |
|               |      |                      |                | development                                      |
|               |      |                      |                |  |
| <u>Jam3</u>   | 1.54 | Junctional adhesion  |                | Immunoglobulin that mediates tight junctions     |
|               |      | molecule C           |                | between endothelial cells; mediates              |
|               |      |                      |                | transepithelial migration of PMNs; promotes      |
|               |      |                      |                | chemotaxis of vascular endothelial cells and     |
|               |      |                      |                | stimulates angiogenesis                          |
|               |      |                      |                |  |
| <u>Stat5b</u> | 0.31 | Signal transducer    |                | Carries out a dual function: signal transduction |
|               |      | and activator of     |                | and activation of transcription; positively      |
|               |      | transcription 5b     |                | regulates hematopoietic/erythroid                |
|               |      |                      |                | differentiation.                                 |
|               |      |                      |                |  |
|               |      |                      | Inflammation   |  |
| Bst2          | 1.31 | Bone marrow          | Tethrin; CD317 | IFN-induced antiviral factor that blocks         |
|               |      | stromal cell antigen |                | budding of enveloped viruses by directly         |
|               |      | 2                    |                | tethering nascent virions to the membranes of    |
|               |      |                      |                | infected cells                                   |
|               |      |                      |                | 11112  |
|               |      | 1                    |                |  |

| <u>Casp1</u> | 0.882 | Caspase 1            | Interleukin 1β      | Cysteine-aspartic acid protease that mediates     |
|--------------|-------|----------------------|---------------------|---|
|              |       |                      | convertase          | cleavage-based activation of IL-1β and IL-18;     |
|              |       |                      |                     | serves as the central enzymatic core of the       |
|              |       |                      |                     | inflammasome; also induces apoptosis              |
| Casp8        | 0.536 | Caspase 8            |                     | Cysteine-aspartic acid protease; cleaves and      |
|              |       |                      |                     | activates effector caspases CASP3, -4, -6, -7, -  |
|              |       |                      |                     | 9, and -10; initiates pyroptosis through          |
|              |       |                      |                     | cleavage of gasdermin-D; mediates                 |
|              |       |                      |                     | noncanonical cleavage of IL-1β in DCs and         |
|              |       |                      |                     | macrophages                                       |
| <u>Cd38</u>  | 2.02  | Cluster of           | ADP-ribosyl cyclase | Synthesizes the second messengers cyclic          |
|              |       | differentiation 38   | 1                   | ADP-ribose and NADPH; appears to play a           |
|              |       |                      |                     | critical role in inflammation, although its exact |
|              |       |                      |                     | immunological function(s) remain(s) poorly        |
|              |       |                      |                     | defined   |
| <u>Cebpb</u> | 1.28  | CCAAT/enhancer-      |                     | Critical macrophage TF that promotes              |
|              |       | binding protein beta |                     | expression of several acute-phase and             |
|              |       |                      |                     | inflammatory cytokine genes, including II6        |
| <u>Csf1r</u> | 1.11  | Macrophage colony-   | CD115               | Receptor for CSF1; promotes release of            |
|              |       | stimulating factor 1 |                     | inflammatory cytokines in response to IL-34       |
|              |       | receptor             |                     | and CSF1  |
| <u>Ctsh</u>  | 1.25  | Cathepsin H          |                     | Lysosomal protease; increased in                  |
|              |       |                      |                     | macrophages in response to IFNγ                   |
|              |       |                      |                     |   |

| <u>Ifnar1</u>  | 0.731 | Interferon-<br>alpha/beta receptor     |           | Component of the receptor for type I IFNs, binding of which activates the JAK-STAT   |
|----------------|-------|--|-----------|--|
|                |       | alpha chain                            |           | pathway  |
| Ifng           | 1.93  | Interferon gamma                       |           | T and NK cell-secreted inflammatory cytokine that stimulates cytolysis, activates macrophages, and stimulates MHC class II expression  |
| Ifngr1         | 0.707 | Interferon gamma receptor 1            | CD54      | One of the two components of the IFNy receptor; stimulates activation of the JAK/STAT signaling pathway  |
| <u>II1b</u>    | 2.04  | Interleukin 1 beta                     | Catabolin | One of the two primary inflammatory cytokines produced by the inflammasome (the other one being IL-18); induces neutrophil influx and activation, T cell activation and cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNy synthesis from TH1 cells |
| <u>II6</u>     | 2.7   | Interleukin 6                          |           | Pro-inflammatory cytokine that signals through the JAK and STAT pathways   |
| <u>ll12rb1</u> | 1.07  | Interleukin 12 receptor subunit beta 1 | CD212     | Cytokine receptor component that associates with IL12RB2 to IL23R  |

| <u>ll18r1</u> | 1.36  | Interleukin 18<br>receptor 1              | CD218a                                 | Receptor for IL-18  |
|---------------|-------|---|--|---|
| <u>Irf1</u>   | 1.71  | Interferon regulatory factor 1            |  | Transcriptional regulator that promotes inflammatory innate and adaptive immune responses   |
| <u>Irf4</u>   | 1.2   | Interferon regulatory factor 4            |  | Transcriptional activator that complexes with  BATF and binds ISREs within the promoters of  multiple genes involved in inflammation  |
| <u>Irf7</u>   | 2.22  | Interferon regulatory factor 7            |  | Key transcriptional regulator of type I IFN-dependent immune responses; promotes $transcription\ of\ IFN\alpha\ and\ -\beta$  |
| <u>Irf8</u>   | 1.12  | Interferon regulatory factor 8            |  | TF that regulates of lineage commitment in myeloid cell maturation; promotes monocyte and plasmacytoid DC development   |
| Irgm2         | 0.832 | Immunity-related GTPase family M member 2 | Interferon- inducible protein 1 (IFI1) | Function not fully known, but most likely regulates autophagy and pro-inflammatory cytokine production  |
| <u>Isg15</u>  | 2.41  | Interferon-<br>stimulated gene 15         |  | Ubiquitin-like protein that binds intracellular target proteins upon activation by IFN $\alpha$ or $\beta$ ; can also be secreted to induce NK cell proliferation, act as a chemoattractant for |

|       |       |                        |                    | neutrophils, and induce IFNγ upon binding to     |
|-------|-------|------------------------|--------------------|--|
|       | 0.055 |                        |                    |  |
| Isg20 | 0.855 | Interferon-            |                    | IFN-induced antiviral exoribonuclease that       |
|       |       | stimulated gene 20     |                    | acts on ssRNA with minor activity towards        |
|       |       |                        |                    | ssDNA  |
| Jak1  | 0.709 | Janus kinase 1         |                    | Essential tyrosine kinase involved signal        |
|       |       |                        |                    | transduction in type I and II cytokines and IFNs |
| Jak2  | 0.797 | Janus kinase 2         |                    | Tyrosine kinase that participates in IFN and     |
|       |       |                        |                    | IL6ST signaling cascades                         |
| Mefv  | 2.79  | Mediterranean fever    | Marenostrin; pyrin | Involved in the regulation of innate immunity    |
|       |       |                        |                    | and the inflammatory response in response to     |
|       |       |                        |                    | IFNγ; both stimulates and restrains the          |
|       |       |                        |                    | inflammasome; also acts as a mediator of         |
|       |       |                        |                    | pyroptosis                                       |
| Nlrp3 | 1.75  | NACHT domain-,         | Cryopyrin          | PRR with a wide diversity of recognized          |
|       |       | leucine-rich repeat-,  |                    | targets that activates the NLRP3                 |
|       |       | and PYD-containing     |                    | inflammasome consisting of NLRP3, PYCARD,        |
|       |       | protein 3              |                    | and caspase-1/-8                                 |
| Nos2  | 4.12  | Inducible nitric oxide |                    | Produces reactive oxygen species and             |
|       |       | synthase (iNOS)        |                    | contributes to inflammatory cytokine             |
|       |       |                        |                    | production                                       |
|       |       |                        |                    |  |

| <u>Pik3cd</u> | 1.19 | Phosphatidylinositol- |               | A subunit of PI3K; acts downstream of TLR4,   |
|---------------|------|-----------------------|---------------|---|
|               |      | 4,5-bisphosphate 3-   |               | TCR, BCR, and CD40; contributes to T helper   |
|               |      | kinase catalytic      |               | cell expansion, mast cell development, and    |
|               |      | subunit delta         |               | neutrophil chemotaxis, extravasation, and     |
|               |      | isoform               |               | respiratory burst                             |
|               |      |                       |               |   |
| Pik3cg        | 1.21 | Phosphatidylinositol- |               | A subunit of PI3K; modulates leukocyte        |
|               |      | 4,5-bisphosphate 3-   |               | chemotaxis to inflammatory sites and in       |
|               |      | kinase catalytic      |               | response to chemoattractant agents            |
|               |      | subunit gamma         |               |   |
|               |      | isoform               |               |   |
|               |      |                       |               |   |
| <u>S100a8</u> | 2.3  | S100 calcium-binding  | Calgranulin A | Calcium- and zinc-binding protein involved in |
|               |      | protein A8            |               | pro-inflammatory, antimicrobial, oxidant-     |
|               |      |                       |               | scavenging and apoptosis-inducing activities; |
|               |      |                       |               | can induce neutrophil chemotaxis, adhesion,   |
|               |      |                       |               | phagocytosis, and degranulation;              |
|               |      |                       |               | predominantly found as calprotectin           |
|               |      |                       |               | (S100A8/A9) which has a wide plethora of      |
|               |      |                       |               | intra- and extracellular functions, including |
|               |      |                       |               | adhesion, apoptosis, autophagy, cytoskeletal  |
|               |      |                       |               | remodeling, cytokine production, chemotaxis,  |
|               |      |                       |               | migration, inflammation, arachidonic acid     |
|               |      |                       |               | metabolism, oxidant-scavenging, and PRR       |
|               |      |                       |               | signaling                                     |
|               |      |                       |               |   |

| Son   1.63   Sialophorin   Leukosialin; CD43   Cell surface sialoglycoprotein expressed by T   cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>1</sub> 2 phenotype and towards T <sub>1</sub> 1; promotes the expression of IFNγ in CD4* T cells    Tok1  |               |       |                     |                   |   |
|---|---------------|-------|---------------------|-------------------|---|
| Tok1 1.05 TANK-binding kinase 1 Coordinates the activation of IRF3 and NFκB and induction of type I IFNs  Tof 3.73 Tumor necrosis Cachectin Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines  Traf6 0.755 Tumor necrosis factor receptor-associated factor 6 Inhibition  Inhibition  Bci211 0.729 B cell lymphoma 2 Protein phosphatase 1 Potent inhibitor of caspase-mediated cell death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  | <u>Spn</u>    | 1.63  | Sialophorin         | Leukosialin; CD43 | cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype |
| 1 and induction of type I IFNs  To and induction of type I IFNs  Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines  Trafe 0.755 Tumor necrosis factor receptor-associated factor 6 TNFα  Inhibition  Bci2i1 0.729 B cell lymphoma 2 Protein Potent inhibitor of caspase-mediated cell death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  |               |       |                     |                   |   |
| Traf6 3.73 Tumor necrosis Cachectin Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines  Traf6 0.755 Tumor necrosis factor receptorassociated factor 6 Inhibition  Inhibition  Bci2l1 0.729 B cell lymphoma 2 Protein phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates   | <u>Tbk1</u>   | 1.05  | TANK-binding kinase |                   | Coordinates the activation of IRF3 and NFkB   |
| factor macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines  Traf6 0.755 Tumor necrosis factor receptor-associated factor 6 Inhibition  Inhibition  Bcl2l1 0.729 B cell lymphoma 2 Protein Potent inhibitor of caspase-mediated cell phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  |               |       | 1                   |                   | and induction of type I IFNs  |
| TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines  Traf6 0.755 Tumor necrosis Adaptor protein that acts in the CD40 signaling cascade; promotes inflammation, IL-6, and TNFα  Inhibition  Bcl2l1 0.729 B cell lymphoma 2 Protein Potent inhibitor of caspase-mediated cell like 1 phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates   | <u>Tnf</u>    | 3.73  | Tumor necrosis      | Cachectin         | Inflammatory cytokine mainly produced by  |
| Description   Cend3   Cyclin D3   Cascade; promotes inflammation   Cend3   Complex that a complex that inhibitively phosphorylates   Cend3   Complex that inhibitively phosphorylates   Cend3   Cen |               |       | factor              |                   | macrophages; binds to TNFRSF1A/TNFR1 and  |
| Traf6       0.755       Tumor necrosis       Adaptor protein that acts in the CD40 signaling cascade; promotes inflammation, IL-6, and TNFα         Inhibition         Bcl2l1       0.729       B cell lymphoma 2 like 1       Potent inhibitor of caspase-mediated cell death         Ccnd3       0.254       Cyclin D3       Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  |               |       |                     |                   | TNFRSF1B/TNFBR; capable of inducing cell  |
| factor receptor- associated factor 6  Inhibition  Bcl2l1 0.729 B cell lymphoma 2 Protein Potent inhibitor of caspase-mediated cell like 1 phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  |               |       |                     |                   | death in certain tumor cell lines   |
| Bcl2l1   0.729   B cell lymphoma 2   Protein   Potent inhibitor of caspase-mediated cell   like 1   phosphatase 1   death     Ccnd3   0.254   Cyclin D3   Regulatory component of the cyclin D3-CDK4   complex that inhibitively phosphorylates   | <u>Traf6</u>  | 0.755 | Tumor necrosis      |                   | Adaptor protein that acts in the CD40 signaling   |
| Inhibition    Bcl2l1  |               |       | factor receptor-    |                   | cascade; promotes inflammation, IL-6, and   |
| Bcl2l1 0.729 B cell lymphoma 2 Protein Potent inhibitor of caspase-mediated cell like 1 phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates  |               |       | associated factor 6 |                   | ΤΝΓα  |
| like 1 phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates   |               |       |                     | Inhibition        |   |
| like 1 phosphatase 1 death  Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates   |               |       |                     |                   |   |
| Ccnd3 0.254 Cyclin D3 Regulatory component of the cyclin D3-CDK4 complex that inhibitively phosphorylates   | <u>Bcl2l1</u> | 0.729 | B cell lymphoma 2   | Protein           | Potent inhibitor of caspase-mediated cell   |
| complex that inhibitively phosphorylates  |               |       | like 1              | phosphatase 1     | death   |
|   | Ccnd3         | 0.254 | Cyclin D3           |                   | Regulatory component of the cyclin D3-CDK4  |
| members of the retinoblastoma protein   |               |       |                     |                   | complex that inhibitively phosphorylates  |
|   |               |       |                     |                   | members of the retinoblastoma protein   |

|              |       |                     |                     | family; regulates the cell-cycle during G1/S     |
|--------------|-------|---------------------|---------------------|--|
|              |       |                     |                     | transition                                       |
| <u>Cd47</u>  | 0.743 | Cluster of          | Integrin-associated | Partners with membrane integrins to serve as     |
|              |       | differentiation 47  | protein (IAP)       | an inhibitor of phagocytosis                     |
| Cd200r1      | 1.18  | Cluster of          |                     | Inhibitory receptor for the CD200/OX2 cell       |
|              |       | differentiation 200 |                     | surface glycoprotein; limits inflammation by     |
|              |       | receptor 1          |                     | inhibiting the expression of proinflammatory     |
|              |       |                     |                     | molecules including TNFα, IFNs, and iNOS         |
| <u>Cd274</u> | 2.49  | Cluster of          | Programmed cell     | Ubiquitously expressed ligand for co-inhibitory  |
|              |       | differentiation 274 | death receptor      | receptor PD-1; upregulated by tumors as an       |
|              |       |                     | ligand 1 (PD-L1)    | immune evasion strategy                          |
| Cdkn1a       | 1.39  | Cyclin dependent    | p21; CDK-           | Binds to and inhibits cyclin-dependent kinase    |
|              |       | kinase inhibitor 1A | interaction protein | activity, preventing phosphorylation of critical |
|              |       |                     | 1 (CIP1)            | cyclin-dependent kinase substrates and           |
|              |       |                     |                     | blocking cell cycle progression                  |
| <u>Ctla4</u> | 2.26  | Cytotoxic T         | CD152               | Inhibitory receptor that blocks CD28             |
|              |       | lymphocyte antigen  |                     | costimulation by competitively binding its       |
|              |       | 4                   |                     | ligands CD80 and CD86                            |
| Cyld         | 0.595 | Cylindromatosis     |                     | Inhibits NFkB activation by deubiquitinating     |
|              |       | lysine 63           |                     | upstream signaling factors; inhibits Wnt         |
|              |       | deubiquitinase      |                     | signaling; restricts polyubiquitination of RIPK1 |
|              |       |                     |                     | and -2, thereby limiting necroptosis             |
|              |       |                     |                     |  |

| 1-1-4        | 4.22 | Indalas : 22        |        | Indicates catalogue Co. 1 10 to  |
|--------------|------|---------------------|--------|--|
| <u>ldo1</u>  | 1.39 | Indoleamine 2,3-    |        | Initiates catabolism of tryptophan; limits                                   |
|              |      | dioxygenase 1       |        | immunopathology by inhibiting T cell division                                |
|              |      |                     |        |  |
| <u>ll1r2</u> | 1.07 | Interleukin 1       | CD121b | Non-signaling receptor for IL-1 $\alpha$ , - $\beta$ , and RN;               |
|              |      | receptor type II    |        | serves as a decoy receptor by competitive                                    |
|              |      |                     |        | binding to IL-1 $\beta$ and preventing its binding to                        |
|              |      |                     |        | IL1R1  |
|              |      |                     |        |  |
| <u>Irak3</u> | 1.89 | Interleukin-1       |        | Adaptor protein that negatively regulates TLR                                |
|              |      | receptor-associated |        | signaling; predominantly expressed in  |
|              |      | kinase 3            |        | monocytes and macrophages  |
|              |      |                     |        |  |
| Foxp3        | 1.56 | Forkhead box P3     | DIETER | Master TF for Tregs; represses expression of                                 |
|              |      |                     |        | II2 and Ifng; activates expression of Tnfrsf18,                              |
|              |      |                     |        | Il2ra, and Ctla4   |
|              |      |                     |        | ·  |
| Fcgr2b       | 1.41 | Fc fragment of IgG  | CD32   | Low affinity receptor for the Fc region of                                   |
|              |      | receptor IIb        |        | complexed or aggregated γ-lgs; involved in a                                 |
|              |      |                     |        | variety of effector and regulatory functions                                 |
|              |      |                     |        | such as phagocytosis of immune complexes                                     |
|              |      |                     |        | and modulation of Ab production by B cells;                                  |
|              |      |                     |        | essential for the maintenance of humoral                                     |
|              |      |                     |        | tolerance; acts as a late checkpoint at the level                            |
|              |      |                     |        | of class-switched memory B cells,  |
|              |      |                     |        |  |
|              |      |                     |        | plasmablasts or plasma cells; regulates                                      |
|              |      |                     |        |  |
|              |      |                     |        | plasmablasts or plasma cells; regulates plasma-cell homeostasis and survival |

| Klrg1         | 1.08  | Killer cell lectin-like | C-type lectin       | NK and T cell inhibitory receptor; binds to non- |
|---------------|-------|-------------------------|---------------------|--|
|               |       | receptor, subfamily     | domain family 15,   | MHC ligands                                      |
|               |       | G, member 1             | member A            |  |
|               |       |                         | (CLEC15A)           |  |
|               |       |                         | Mast cell function- |  |
|               |       |                         | associated antigen  |  |
|               |       |                         | (MAFA)              |  |
|               |       |                         | (IVIALA)            |  |
| Lag3          | 2.17  | Lymphocyte              | CD223               | Inhibitory receptor on activated T cells; binds  |
|               |       | activating gene 3       |                     | to ligands, such as FGL1; constitutively         |
|               |       |                         |                     | expressed on a subset of regulatory Tregs and    |
|               |       |                         |                     | contributes to their suppressive function; acts  |
|               |       |                         |                     | as a negative regulator of plasmacytoid DC       |
|               |       |                         |                     | activation                                       |
| <u>Nfkbia</u> | 1.65  | Nuclear factor kappa    |                     | Inhibits activity of REL dimers by masking of    |
|               |       | B inhibitor alpha       |                     | their nuclear localization signals               |
|               |       |                         |                     | -  |
| Nlrc5         | 0.956 | NLR family CARD         | NOD27               | Inhibits NFкВ and type I IFN signaling           |
|               |       | domain containing 5     |                     | pathways; may also regulate the type II IFN      |
|               |       |                         |                     | signaling pathway                                |
| Pdcd1         | 0.768 | Programmed cell         | CD279; PD-1         | Checkpoint receptor; inhibits T and NK cell      |
|               |       | death 1                 |                     | activation                                       |
|               |       |                         |                     |  |
| Pdcd1lg2      | 1.45  | Programmed cell         |                     |  |
|               |       | death 1 ligand 2        |                     |  |
|               |       |                         |                     |  |

| Serping1     | 0.72  | Serpin family G      | C1-inhibitor | Inflammation-induced acute phase protein          |
|--------------|-------|----------------------|--------------|---|
|              |       | member 1             |              | that inhibits C1r and C1s proteases in the C1     |
|              |       |                      |              | complex   |
|              |       |                      |              |   |
| Socs1        | 1.93  | Suppressor of        |              | Inhibits JAK proteins; negative regulator of IL-6 |
|              |       | cytokine signaling 1 |              |   |
| Socs3        | 0.874 | Suppressor of        |              | Inhibits IL6ST and JAK2; negative regulator of    |
|              |       | cytokine signaling 3 |              | IL-6  |
| <u>Sh2b2</u> | 1.64  | Src homology 2B      |              | Adapter protein for several members of the        |
|              |       | adaptor protein 2    |              | tyrosine kinase receptor family; involved in      |
|              |       |                      |              | multiple signaling pathways; may be involved      |
|              |       |                      |              | in coupling from immunoreceptor to Ras            |
|              |       |                      |              | signaling; acts as a negative regulator of        |
|              |       |                      |              | cytokine signaling in collaboration with CBL;     |
|              |       |                      |              | may induce cytoskeletal reorganization via        |
|              |       |                      |              | interaction with Vav3                             |
| <u>Tank</u>  | 0.975 | TRAF family          |              | Inhibitory protein that sequesters TRAFs in the   |
|              |       | member-associated    |              | cytoplasm, constitutively binds TBK1, and         |
|              |       | NFkB activator       |              | serves as a negative regulator of NFkB            |
| Tgfb1        | 0.803 | Transforming growth  |              | Multifunctional protein that regulates the        |
|              |       | factor beta 1        |              | growth and differentiation of various cell        |
|              |       |                      |              | types and is involved in various processes,       |
|              |       |                      |              | such as normal development, immune                |
|              |       |                      |              | function, microglia function and responses to     |

|              |       |                      |                     | neurodegeneration; can induce EMT and cell       |
|--------------|-------|----------------------|---------------------|--|
|              |       |                      |                     | migration in various cell types; frequently acts |
|              |       |                      |                     | as an immunosuppressive cytokine in the TME      |
| <u>Tigit</u> | 1.28  | T cell               |                     | Binds with high affinity to the poliovirus       |
|              |       | immunoreceptor       |                     | receptor, causing increased secretion of IL-10,  |
|              |       | with Ig and ITIM     |                     | decreased secretion of IL-12B, and               |
|              |       | domains              |                     | suppressing T cell activation by promoting the   |
|              |       |                      |                     | generation of mature immunoregulatory DCs        |
| Tollip       | 0.413 | Toll interacting     |                     | Inhibitory adaptor protein; recruits IRAK1 to    |
|              |       | protein              |                     | the IL-1 receptor complex and inhibitively       |
|              |       |                      |                     | phosphorylates it                                |
|              |       |                      | Interferon Response | e  |
|              |       |                      |                     |  |
| Bst2         | 1.31  | Bone marrow          | Tethrin; CD317      | IFN-induced antiviral factor that blocks         |
|              |       | stromal cell antigen |                     | budding of enveloped viruses by directly         |
|              |       | 2                    |                     | tethering nascent virions to the membranes of    |
|              |       |                      |                     | infected cells                                   |
| <u>Ctsh</u>  | 1.25  | Cathepsin H          |                     | Lysosomal protease; increased in                 |
|              |       |                      |                     | macrophages in response to IFNγ                  |
| Ifih1        | 1.43  | Interferon induced   | Helicard;           | PRR for cytoplasmic dsRNA; upon target           |
|              |       | with helicase C      | melanoma            | recognition, associates with MAVS to activate    |
|              |       | domain 1             | differentiation-    | TNK1 and IKBKE, which phosphorylate IRF3         |
|              |       |                      |                     |  |

|               |       |                       | associated protein | and -7, which, in turn, activate transcription of |
|---------------|-------|-----------------------|--------------------|---|
|               |       |                       | 5 (MDA5)           | IFNα and -β                                       |
|               |       |                       |                    |   |
| <u>Ifitm1</u> | 0.626 | Interferon-induced    | CD225              | IFN-induced antiviral protein implicated in cell  |
|               |       | transmembrane         |                    | adhesion and control of cell growth and           |
|               |       | protein 1             |                    | migration   |
| <u>Ifnar1</u> | 0.731 | Interferon-           |                    | Component of the receptor for type I IFNs,        |
|               |       | alpha/beta receptor   |                    | binding of which activates the JAK-STAT           |
|               |       | alpha chain           |                    | pathway   |
| <u>Ifng</u>   | 1.93  | Interferon gamma      |                    | T and NK cell-secreted inflammatory cytokine      |
|               |       |                       |                    | that stimulates cytolysis, activates              |
|               |       |                       |                    | macrophages, and stimulates MHC class II          |
|               |       |                       |                    | expression  |
| Ifngr1        | 0.707 | Interferon gamma      | CD54               | One of the two components of the IFNy             |
|               |       | receptor 1            |                    | receptor; stimulates activation of the            |
|               |       |                       |                    | JAK/STAT signaling pathway                        |
| <u>Irf1</u>   | 1.71  | Interferon regulatory |                    | Transcriptional regulator that promotes           |
|               |       | factor 1              |                    | inflammatory innate and adaptive immune           |
|               |       |                       |                    | responses   |
| Irf4          | 1.2   | Interferon regulatory |                    | Transcriptional activator that complexes with     |
|               |       | factor 4              |                    | BATF and binds ISREs within the promoters of      |
|               |       |                       |                    | multiple genes involved in inflammation           |

| <u>Irf7</u>  | 2.22  | Interferon regulatory factor 7 |                     | Key transcriptional regulator of type I IFN-<br>dependent immune responses; promotes |
|--------------|-------|--------------------------------|---------------------|--|
|              |       |                                |                     | transcription of IFN $\alpha$ and - $\beta$  |
| <u>Irf8</u>  | 1.12  | Interferon regulatory          |                     | TF that regulates of lineage commitment in   |
|              |       | factor 8                       |                     | myeloid cell maturation; promotes monocyte   |
|              |       |                                |                     | and plasmacytoid DC development  |
| Irgm2        | 0.832 | Immunity-related               | Interferon-         | Function not fully known, but most likely  |
|              |       | GTPase family M                | inducible protein 1 | regulates autophagy and pro-inflammatory   |
|              |       | member 2                       | (IFI1)              | cytokine production  |
| <u>Isg15</u> | 2.41  | Interferon-                    |                     | Ubiquitin-like protein that binds intracellular                                      |
|              |       | stimulated gene 15             |                     | target proteins upon activation by IFN $\alpha$ or $\beta$ ;                         |
|              |       |                                |                     | can also be secreted to induce NK cell   |
|              |       |                                |                     | proliferation, act as a chemoattractant for  |
|              |       |                                |                     | neutrophils, and induce IFNγ upon binding to   |
|              |       |                                |                     | ITGAL/ITGB2  |
| Isg20        | 0.855 | Interferon-                    |                     | IFN-induced antiviral exoribonuclease that   |
|              |       | stimulated gene 20             |                     | acts on ssRNA with minor activity towards  |
|              |       |                                |                     | ssDNA  |
| Jak1         | 0.709 | Janus kinase 1                 |                     | Essential tyrosine kinase involved signal  |
|              |       |                                |                     | transduction in type I and II cytokines and IFNs                                     |
| Mefv         | 2.79  | Mediterranean fever            | Marenostrin; pyrin  | Involved in the regulation of innate immunity  |
|              |       |                                |                     | and the inflammatory response in response to   |
|              |       |                                |                     | IFNγ; both stimulates and restrains the  |
|              |       |                                |                     |  |

| Mx2 Tbk1       | 1.03  | Myxovirus resistance protein 2  TANK-binding kinase             |               | inflammasome; also acts as a mediator of pyroptosis  IFN-induced dynamin-like GTPase with potent antiviral activity against HIV-1  Coordinates the activation of IRF3 and NFκB and induction of type I IFNs                          |
|----------------|-------|---|---------------|--|
|                |       |   | Ion Transport |  |
| Арр            | 0.801 | Amyloid-beta precursor protein                                  |               | Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; involved in cell mobility, copper homeostasis, and oxidative stress  |
| Ms4a1  Slc11a1 | 2.79  | Membrane spanning  4-domains A1  Natural resistance- associated | CD20; Bp35    | B cell-specific membrane protein that  functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation  Macrophage-specific metal ion transporter; uptakes divalent metal cations to neutralize |
|                |       | associated macrophage protein 1                                 | IRAKs & TRAFs | ROSs   |

| <u>Irak1</u> | 0.24  | Interleukin-1       | Adaptor protein involved in TLR and IL-1         |
|--------------|-------|---------------------|--|
|              |       | receptor-associated | signaling; recruited to TLRs by MyD88 and        |
|              |       | kinase 1            | phosphorylated by IRAK4; promotes the            |
|              |       |                     | degradation of TIRAP                             |
|              |       |                     |  |
| <u>Irak2</u> | 1.49  | Interleukin-1       | Adaptor protein involved in TLR and IL-1         |
|              |       | receptor-associated | signaling  |
|              |       | kinase 2            |  |
|              |       |                     |  |
| <u>Irak3</u> | 1.89  | Interleukin-1       | Adaptor protein that negatively regulates TLR    |
|              |       | receptor-associated | signaling; predominantly expressed in            |
|              |       | kinase 3            | monocytes and macrophages                        |
|              |       |                     |  |
| <u>Traf6</u> | 0.755 | Tumor necrosis      | Adaptor protein that acts in the CD40 signaling  |
|              |       | factor receptor-    | cascade; promotes inflammation, IL-6, and        |
|              |       | associated factor 6 | ΤΝΓα   |
|              |       |                     |  |
|              |       |                     | JAK-STAT Pathway                                 |
| Jak1         | 0.709 | Janus kinase 1      | Essential tyrosine kinase involved signal        |
|              |       |                     | transduction in type I and II cytokines and IFNs |
|              |       |                     |  |
| Jak2         | 0.797 | Janus kinase 2      | Tyrosine kinase that participates in IFN and     |
|              |       |                     | IL6ST signaling cascades                         |
|              |       |                     |  |
| Stat1        | 1.53  | Signal transducer   | Transcriptional activator that mediates cellular |
|              |       | and activator of    | responses to IFNs, cytokines, and other          |
|              |       | transcription 1     | growth factors                                   |
|              |       |                     |  |

| Stat3         | 0.649 | Signal transducer  | Transcriptional activator of genes involved in                        |
|---------------|-------|--------------------|---|
|               |       | and activator of   | cell growth and apoptosis; activated by JAKs                          |
|               |       | transcription 3    |   |
|               |       |                    |   |
| Stat4         | 1.25  | Signal transducer  | Essential TF for T <sub>H</sub> 1 CD4 <sup>+</sup> T cell development |
|               |       | and activator of   | and IFNγ production; also promotes                                    |
|               |       | transcription 4    | expression of MyD88   |
| <u>Stat5b</u> | 0.31  | Signal transducer  | Carries out a dual function: signal transduction                      |
|               |       | and activator of   | and activation of transcription; positively                           |
|               |       | transcription 5b   | regulates hematopoietic/erythroid                                     |
|               |       |                    | differentiation.  |
|               |       |                    |   |
| <u>Stat6</u>  | 0.571 | Signal transducer  | Essential TF for T <sub>H</sub> 2 CD4+ T cell and                     |
|               |       | and activator of   | macrophage function and M2 macrophage                                 |
|               |       | transcription 6    | polarization  |
|               |       |                    | 10  |
|               |       |                    | Kinases   |
| <u>Btk</u>    | 1.2   | Bruton's tyrosine  | Crucial kinase in B cell receptor signal                              |
|               |       | kinase             | transmission and B cell activation                                    |
| Hck           | 1.76  | Hematopoietic cell | Src family tyrosine kinase that mediates                              |
|               |       | kinase             | secretory lysosome mobilization,                                      |
|               |       |                    | degranulation, and activation of NADPH                                |
|               |       |                    | oxidase   |
|               |       |                    |   |

| <u>Itk</u> | 2.06  | Interleukin-2-        | LYK | Key actor in the TCR signaling cascade;          |
|------------|-------|-----------------------|-----|--|
|            |       | inducible T cell      |     | phosphorylates PLCγ1, LAT, and LCP2              |
|            |       | kinase                |     |  |
|            |       | Killase               |     |  |
| <u>Lck</u> | 1.98  | Lymphocyte cell       |     | Src family tyrosine kinase that acts as one of   |
|            |       | kinase                |     | the main signaling intermediaries downstream     |
|            |       |                       |     | of the TCR; constitutively associated with the   |
|            |       |                       |     | cytoplasmic portion of CD4                       |
| <u>Lyn</u> | 0.921 | Tyrosine-protein      |     | Src family tyrosine kinase that potentiates      |
|            |       | kinase Lyn            |     | signaling from the B cell receptor and CD40      |
| Jak1       | 0.709 | Janus kinase 1        |     | Essential tyrosine kinase involved signal        |
|            |       |                       |     | transduction in type I and II cytokines and IFNs |
| Jak2       | 0.797 | Janus kinase 2        |     | Tyrosine kinase that participates in IFN and     |
|            |       |                       |     | IL6ST signaling cascades                         |
| Pik3cd     | 1.19  | Phosphatidylinositol- |     | A subunit of PI3K; acts downstream of TLR4,      |
|            |       | 4,5-bisphosphate 3-   |     | TCR, BCR, and CD40; contributes to T helper      |
|            |       | kinase catalytic      |     | cell expansion, mast cell development, and       |
|            |       | subunit delta         |     | neutrophil chemotaxis, extravasation, and        |
|            |       | isoform               |     | respiratory burst                                |
| Pik3cg     | 1.21  | Phosphatidylinositol- |     | A subunit of PI3K; modulates leukocyte           |
|            |       | 4,5-bisphosphate 3-   |     | chemotaxis to inflammatory sites and in          |
|            |       | kinase catalytic      |     | response to chemoattractant agents               |
|            |       |                       |     |  |

|              |       | subunit gamma        |   |
|--------------|-------|----------------------|---|
|              |       | isoform              |   |
|              |       |                      |   |
| Prkcd        | 0.892 | Protein kinase C     | Ca(2+)-independent, phospholipid- and DAG-      |
|              |       | delta                | dependent serine/threonine-protein kinase       |
|              |       |                      | that promotes apoptosis in response to DNA      |
|              |       |                      | damage but inhibits it during cytokine          |
|              |       |                      | receptor-initiated cell death; required for     |
|              |       |                      | oxygen radical production by NADPH oxidase      |
| Syk          | 1.95  | Spleen-associated    | Critical kinase that transmits signals from the |
|              |       | tyrosine kinase      | TCR and BCR                                     |
| Tbk1         | 1.05  | TANK-binding kinase  | Coordinates the activation of IRF3 and NFкВ     |
|              |       | 1                    | and induction of type I IFNs                    |
| <u>Zap70</u> | 1.72  | Zeta chain of T cell | Tyrosine kinase that plays an essential role in |
|              |       | receptor associated  | regulation of the adaptive immune response;     |
|              |       | protein kinase 70    | regulates motility, adhesion, proliferation,    |
|              |       |                      | differentiation, and cytokine expression in T   |
|              |       |                      | cells; contributes to the development and       |
|              |       |                      | activation of primary B cells; phosphorylates   |
|              |       |                      | at least two essential adapter proteins: LAT    |
|              |       |                      | and LCP2, leading in turn to a large number of  |
|              |       |                      | signaling molecules being recruited             |
|              |       |                      | Lysosomal Activity                              |
|              |       |                      |   |

| Ctsh         | 1.25  | Cathepsin H          |                      | Lysosomal protease; increased in               |
|--------------|-------|----------------------|----------------------|--|
| Ctsii        | 1.23  | Cathepsiiiii         |                      |  |
|              |       |                      |                      | macrophages in response to IFNγ                |
|              |       |                      |                      |  |
| <u>Ctss</u>  | 2.53  | Cathepsin S          |                      | Lysosomal protease that participates in        |
|              |       |                      |                      | processing of Ag by MHC class II               |
|              |       |                      |                      |  |
| <u>Hck</u>   | 1.76  | Hematopoietic cell   |                      | Src family tyrosine kinase that mediates       |
|              |       | kinase               |                      | secretory lysosome mobilization, degranlation, |
|              |       |                      |                      | and activation of NADPH oxidase                |
|              |       |                      |                      |  |
|              |       |                      | Macrophage Functio   | n  |
|              |       |                      |                      |  |
| <u>Cd14</u>  | 1.86  | Cluster of           |                      | PRR that recognizes LPS; mostly found on       |
|              |       | differentiation 14   |                      | macrophages                                    |
|              |       |                      |                      | , ,  |
| Csf1         | 0.698 | Macrophage colony-   |                      | Cytokine that promote activation and survival  |
|              |       | stimulating factor 1 |                      | of monocytes                                   |
|              |       |                      |                      | 3  |
| Csf1r        | 1.11  | Macrophage colony-   | CD115                | Receptor for CSF1; promotes release of         |
|              |       | stimulating factor 1 |                      | inflammatory cytokines in response to IL-34    |
|              |       |                      |                      |  |
|              |       | receptor             |                      | and CSF1                                       |
| <u>Cebpb</u> | 1.28  | CCAAT/enhancer-      |                      | Critical macrophage TF that promotes           |
| cenhn        | 1.20  |                      |                      |  |
|              |       | binding protein beta |                      | expression of several acute-phase and          |
|              |       |                      |                      | inflammatory cytokine genes, including II6     |
|              |       |                      |                      |  |
| Clec5a       | 1.73  | C-Type lectin domain | Myeloid DAP12-       | Critical macrophage receptor for dengue virus  |
|              |       | family 5, member a   | associating lectin-1 | serotypes 1-4; positive regulator of           |
|              |       |                      |                      | osteoclastogenesis                             |
|              |       |                      |                      | osteodiustogenesis                             |
|              |       |                      |                      |  |

| <u>Ctsh</u>   | 1.25  | Cathepsin H                             |                 | Lysosomal protease; increased in macrophages in response to IFNγ |
|---------------|-------|---|-----------------|--|
| Cxcl24        |       | C-X-C motif                             |                 | Macrophage-produced cytokine; function                           |
|               |       | chemokine ligand 24                     |                 | unknown  |
| <u>Gpr183</u> | 0.932 | G protein-coupled                       | EBV-induced G   | Lymphocyte GPCR that acts as a chemotactic                       |
|               |       | receptor 183                            | protein-coupled | receptor for B cells, T cells, splenic DCs,                      |
|               |       |   | receptor 2      | monocytes/macrophages, and astrocytes                            |
| <u>Irak3</u>  | 1.89  | Interleukin-1                           |                 | Adaptor protein that negatively regulates TLR                    |
|               |       | receptor-associated                     |                 | signaling; predominantly expressed in                            |
|               |       | kinase 3                                |                 | monocytes and macrophages  |
| <u>Itgam</u>  | 2.19  | Integrin alpha M                        | CD11b           | Pairs with CD18 to form Mac-1 aka                                |
|               |       |   |                 | complement receptor 3; mediates leukocyte                        |
|               |       |   |                 | activation, adhesion, chemotaxis, migration,                     |
|               |       |   |                 | phagocytosis, and cell-mediated cytotoxicity;                    |
|               |       |   |                 | serves as a macrophage marker                                    |
| Marco         | 4.23  | Macrophage                              |                 | A PRR that recognizes LDL  |
|               |       | receptor with                           |                 |  |
|               |       | collagenous                             |                 |  |
|               |       | structure                               |                 |  |
| Slamf7        | 2.32  | Signaling                               |                 | A super-activator of macrophages and a                           |
|               |       | lymphocytic                             |                 | strong promoter of phagocytosis; binds to                        |
|               |       | , |                 | CD74   |

| <u>Slc11a1</u> | 2.2    | activation molecule family member 7  Natural resistance- associated macrophage protein  1 |   | Macrophage-specific metal ion transporter; uptakes divalent metal cations to neutralize ROSs   |
|----------------|--------|---|---|--|
| <u>Tnf</u>     | 3.73   | Tumor necrosis<br>factor  | Cachectin  MAP Kinase Signalin                        | Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines |
|                |        |   | WAT KINGSE SIGNAMI                                    | 5  |
| Mapk1          | -0.398 | Mitogen-activated protein kinase 1  | Extracellular signal-<br>regulated kinase 2<br>(ERK2) | Serine/threonine kinase that acts as an essential component of the MAP kinase signal transduction pathway  |
| Mapk3          | -0.347 | Mitogen-activated protein kinase 3  | Extracellular signal-<br>regulated kinase 1<br>(ERK1) | Serine/threonine kinase that acts as an essential component of the MAP kinase signal transduction pathway  |
| Mapk14         | 0.289  | Mitogen-activated protein kinase 14   |   | One of the four p38 MAPKs; key kinase in the cascades of cellular responses evoked by extracellular stimuli such as proinflammatory cytokines                |

| Map2k1   | 0.266 | Dual specificity        | MAPK/ERK kinase 1  | Essential component of the MAP kinase signal     |
|----------|-------|-------------------------|--------------------|--|
|          |       | mitogen-activated       | (MEK1)             | transduction pathway; participates in            |
|          |       | protein kinase kinase   |                    | numerous biological functions, including cell    |
|          |       | 1                       |                    | growth, adhesion, survival, differentiation,     |
|          |       |                         |                    | transcription, metabolism, and cytoskeletal      |
|          |       |                         |                    | remodeling                                       |
| Map4k2   | 0.666 | Mitogen-activated       |                    | Essential component of the MAP kinase signal     |
|          |       | protein kinase kinase   |                    | transduction pathway downstream of TRAF6;        |
|          |       | kinase kinase 2         |                    | upstream activator of the SAP/JNK signaling      |
|          |       |                         |                    | pathway;   |
| Mapkapk2 | 0.409 | MAP kinase-             |                    | Serine/threonine-protein kinase involved in      |
|          |       | activated protein       |                    | cytokine production, endocytosis,                |
|          |       | kinase 2                |                    | reorganization of the cytoskeleton, cell         |
|          |       |                         |                    | migration, cell cycle control, chromatin         |
|          |       |                         |                    | remodeling, DNA damage response, and             |
|          |       |                         |                    | transcriptional regulation                       |
|          |       |                         | Mast Cell Function |  |
|          |       |                         |                    |  |
| Klrg1    | 1.08  | Killer cell lectin-like | C-type lectin      | NK and T cell inhibitory receptor; binds to non- |
|          |       | receptor, subfamily     | domain family 15,  | MHC ligands                                      |
|          |       | G, member 1             | member A           |  |
|          |       |                         | (CLEC15A)          |  |
|          |       |                         |                    |  |

|        |      |                       | Mast cell function- |   |
|--------|------|-----------------------|---------------------|---|
|        |      |                       | associated antigen  |   |
|        |      |                       | (MAFA)              |   |
| Lgals3 | 1.19 | Galectin 3            |                     | Galactose-specific lectin that binds IgE;     |
|        |      |                       |                     | involved in acute inflammatory responses,     |
|        |      |                       |                     | including neutrophil activation and adhesion, |
|        |      |                       |                     | chemoattraction of monocytes macrophages,     |
|        |      |                       |                     | opsonization of apoptotic neutrophils, and    |
|        |      |                       |                     | activation of mast cells                      |
| Pik3cd | 1.19 | Phosphatidylinositol- |                     | A subunit of PI3K; acts downstream of TLR4,   |
|        |      | 4,5-bisphosphate 3-   |                     | TCR, BCR, and CD40; contributes to T helper   |
|        |      | kinase catalytic      |                     | cell expansion, mast cell development, and    |
|        |      | subunit delta         |                     | neutrophil chemotaxis, extravasation, and     |
|        |      | isoform               |                     | respiratory burst                             |
| Pik3cg | 1.21 | Phosphatidylinositol- |                     | A subunit of PI3K; modulates leukocyte        |
|        |      | 4,5-bisphosphate 3-   |                     | chemotaxis to inflammatory sites and in       |
|        |      | kinase catalytic      |                     | response to chemoattractant agents            |
|        |      | subunit gamma         |                     |   |
|        |      | isoform               |                     |   |
|        |      |                       | Metabolism          |   |
| Abca1  | 2.26 | ATP-binding cassette  |                     | Membrane-associated cholesterol efflux pump   |
|        |      | transporter A1        |                     |   |

| Abcg1          | 1.98  | ATP-binding cassette transporter G1                        |                                 | Membrane-associated cholesterol efflux pump   |
|----------------|-------|--|---------------------------------|---|
| <u>Cd36</u>    | 3.27  | Cluster of differentiation 36                              | Fatty acid<br>translocase (FAT) | Class B scavenger receptor that mediates fatty acid uptake  |
| Map2k1         | 0.266 | Dual specificity mitogen-activated protein kinase kinase 1 | MAPK/ERK kinase 1 (MEK1)        | Essential component of the MAP kinase signal transduction pathway; participates in numerous biological functions, including cell growth, adhesion, survival, differentiation, transcription, metabolism, and cytoskeletal remodeling  |
| <u>\$100a8</u> | 2.3   | S100 calcium-binding protein A8                            | Calgranulin A                   | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin  (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |

|              | Migration/Motility |                                   |           |  |  |  |
|--------------|--------------------|-----------------------------------|-----------|--|--|--|
| Арр          | 0.801              | Amyloid-beta precursor protein    |           | Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; involved in cell mobility, copper homeostasis, and oxidative stress            |  |  |
| <u>Cd97</u>  | 0.748              | Cluster of differentiation 97     | BL-Ac[F2] | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells                 |  |  |
| lcam1        | 1.69               | Intracellular adhesion molecule 1 | CD54      | Cell surface glycoprotein that serves as strong adhesive ligand for LFA-1; important for leukocyte mobility and costimulation  |  |  |
| <u>Itgam</u> | 2.19               | Integrin alpha M                  | CD11b     | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker |  |  |
| Jam3         | 1.54               | Junctional adhesion molecule C    |           | Immunoglobulin that mediates tight junctions between endothelial cells; mediates transepithelial migration of PMNs; promotes   |  |  |

|               |       |                      |               | chemotaxis of vascular endothelial cells and     |
|---------------|-------|----------------------|---------------|--|
|               |       |                      |               |  |
|               |       |                      |               | stimulates angiogenesis                          |
| <u>S100a8</u> | 2.3   | S100 calcium-binding | Calgranulin A | Calcium- and zinc-binding protein involved in    |
| <u> </u>      |       |                      | ou.g. uu      |  |
|               |       | protein A8           |               | pro-inflammatory, antimicrobial, oxidant-        |
|               |       |                      |               | scavenging and apoptosis-inducing activities;    |
|               |       |                      |               | can induce neutrophil chemotaxis, adhesion,      |
|               |       |                      |               | phagocytosis, and degranulation;                 |
|               |       |                      |               | predominantly found as calprotectin              |
|               |       |                      |               | (S100A8/A9) which has a wide plethora of         |
|               |       |                      |               | intra- and extracellular functions, including    |
|               |       |                      |               | adhesion, apoptosis, autophagy, cytoskeletal     |
|               |       |                      |               | remodeling, cytokine production, chemotaxis,     |
|               |       |                      |               | migration, inflammation, arachidonic acid        |
|               |       |                      |               | metabolism, oxidant-scavenging, and PRR          |
|               |       |                      |               | signaling  |
| Tgfb1         | 0.803 | Transforming growth  |               | Multifunctional protein that regulates the       |
|               |       | factor beta 1        |               | growth and differentiation of various cell       |
|               |       |                      |               | types and is involved in various processes,      |
|               |       |                      |               | such as normal development, immune               |
|               |       |                      |               | function, microglia function and responses to    |
|               |       |                      |               | neurodegeneration; can induce EMT and cell       |
|               |       |                      |               | migration in various cell types; frequently acts |
|               |       |                      |               | as an immunosuppressive cytokine in the TME      |
|               |       |                      |               |  |

| <u>Vegfa</u>  | 0.592 | Vascular endothelial  |                      | Glysosylated mitogen that promotes vascular     |
|---------------|-------|-----------------------|----------------------|---|
|               |       | growth factor A       |                      | permeability, vasculogenesis, angiogenesis,     |
|               |       |                       |                      | and cell migration                              |
|               |       |                       |                      |   |
|               |       |                       | Neutrophil Function  | 1   |
| Fpr2          | 3.4   | Formyl peptide        | Lipoxin A4 receptor  | Low affinity receptor for N-formyl-methionyl    |
|               |       | receptor 2            |                      | peptides; activates neutrophils                 |
| Lcn2          | 1.97  | Lipocalin 2           | Neutrophil           | Neutrophil-secreted factor that sequesters      |
|               |       |                       | gelatinase-          | iron-containing siderophores; also functions as |
|               |       |                       | associated lipocalin | a growth factor                                 |
|               |       |                       | (NGAL)               |   |
|               |       |                       |                      |   |
| Ncf4          | 1.34  | Neutrophil cytosolic  | SH3 and PX           | Cytosolic regulatory component of the           |
|               |       | factor 4              | domain-containing    | superoxide-producing phagocyte NADPH-           |
|               |       |                       | protein 4            | oxidase, a multicomponent enzyme system         |
|               |       |                       | (SH3PXD4)            | important for host defense                      |
|               |       |                       |                      |   |
| <u>Pik3cd</u> | 1.19  | Phosphatidylinositol- |                      | A subunit of PI3K; acts downstream of TLR4,     |
|               |       | 4,5-bisphosphate 3-   |                      | TCR, BCR, and CD40; contributes to T helper     |
|               |       | kinase catalytic      |                      | cell expansion, mast cell development, and      |
|               |       | subunit delta         |                      | neutrophil chemotaxis, extravasation, and       |
|               |       | isoform               |                      | respiratory burst                               |
|               |       |                       |                      |   |
| <u>S100a8</u> | 2.3   | S100 calcium-binding  | Calgranulin A        | Calcium- and zinc-binding protein involved in   |
|               |       | protein A8            |                      | pro-inflammatory, antimicrobial, oxidant-       |
|               |       |                       |                      | scavenging and apoptosis-inducing activities;   |
|               |       |                       |                      | can induce neutrophil chemotaxis, adhesion,     |
|               |       |                       |                      |   |

|        |       |   |   | phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR   |
|--------|-------|---|---|---|
|        |       |   |   | signaling   |
|        |       |   | NFkB Signaling                            |   |
| Bcl10  | 0.393 | B cell lymphoma/leukemia 10                   |   | Activates NFκB via ubiquitination of ΙΚΚγ   |
| Card11 | 1.15  | Caspase recruitment  domain family  member 11 | Bcl10-interacting MAGUK protein 3 (BIMP3) | Adapter protein that plays a key role in adaptive immune response by transducing the activation of NFkB downstream of TCR and BCR engagement; upon activation in response to TCR or BCR triggering, homooligomerizes to form a nucleating helical template that recruits BCL10 via CARD-CARD interaction, thereby promoting polymerization of BCL10 and subsequent recruitment of MALT1, leading to IKK phosphorylation and |

|              |       |  |          | degradation and release of NFkB proteins for nuclear translocation  |
|--------------|-------|--|----------|---|
| Cxcr3        | 2.23  | C-X-C motif chemokine receptor                             | CD183    | Induces integrin activation, cytoskeletal remodeling, and chemotaxis; expressed by T cells and NK cells; prominently expressed in effector and memory T cells |
| <u>Ikbkb</u> | 0.362 | Inhibitor of nuclear factor kappa B kinase subunit beta    |          | Part of the IKK complex that inhibits IκBα and permits NFκB nuclear localization  |
| <u>Ikbke</u> | 1.36  | Inhibitor of nuclear factor kappa B kinase subunit epsilon |          | Noncanonical IKB kinase; plays an essential role in regulating inflammatory through the activation of the type I IFN, NFKB, and STAT signaling                |
| Nfkb1        | 0.349 | Nuclear factor kappa  B subunit 1                          | p105/p50 | One of the NFκB family TFs; inhibits inflammation   |
| Rel          | 1.47  | Avian reticuloendotheliosis viral oncogene homolog         | c-Rel    | One of the NFkB family TFs; important for B  cell and Treg development  |
| Relb         | 0.86  | Avian reticuloendotheliosis                                |          | One of the NFkB family TFs; controls lymphoid development, DC biology, and noncanonical NFkB signaling  |

| cytotoxic T and NK cells that activates caspase- mediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      | viral oncogene     |                  |  |
|--|-------------|------|--------------------|------------------|--|
| Cd8a 2.02 Cluster of differentiation 8 alpha LEU2 Alpha chain of the CD8 coreceptor, which binds to MHC class I; in T cells, functions primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8a homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69 1.46 Cluster of domain family 2, member C Signal transmitting receptor in lymphocytes, NK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse |             |      | homolog B          |                  |  |
| differentiation 8 alpha binds to MHC class I; in T cells, functions primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69 1.46 Cluster of differentiation 69 domain family 2, member C  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspase- mediated cell death when delivered into the target cell through the immunological synapse  |             |      |                    | NK Cell Function |  |
| differentiation 8 alpha binds to MHC class I; in T cells, functions primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69 1.46 Cluster of differentiation 69 domain family 2, member C  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspase- mediated cell death when delivered into the target cell through the immunological synapse  |             |      |                    |                  |  |
| alpha primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69 1.46 Cluster of domain family 2, MK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  | <u>Cd8a</u> | 2.02 | Cluster of         | LEU2             | Alpha chain of the CD8 coreceptor, which         |
| molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69  1.46 Cluster of differentiation 69 domain family 2, member C  Macells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb  3.8 Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk  3.36 Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      | differentiation 8  |                  | binds to MHC class I; in T cells, functions      |
| presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69  1.46  Cluster of domain family 2, MK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb  3.8  Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk  3.36  Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      | alpha              |                  | primarily as a coreceptor for MHC class I        |
| surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Cd69  1.46  Cluster of C-type lectin domain family 2, MK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb  3.8  Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk  3.36  Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | molecule:peptide complex; in NK cells, the       |
| allowing conjugation and lysis of multiple target cells  Cd69  1.46  Cluster of C-type lectin Signal transmitting receptor in lymphocytes, MK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation  Gzmb  3.8  Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk  3.36  Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | presence of CD8α homodimers at the cell          |
| Cd69  1.46 Cluster of differentiation 69 domain family 2, member C  G2mb  3.8 Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  G2mk  3.36 Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | surface provides a survival mechanism            |
| Cd69  1.46  Cluster of differentiation 69  domain family 2, member C  Gzmb  3.8  Granzyme B  Fragmentin 2  Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk  3.36  Granzyme K  Tryptase II  Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | allowing conjugation and lysis of multiple       |
| differentiation 69 domain family 2, MK cells, and platelets; induced upon T cell member C activation; involved in lymphocyte proliferation  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | target cells                                     |
| member C activation; involved in lymphocyte proliferation  Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine   | <u>Cd69</u> | 1.46 | Cluster of         | C-type lectin    | Signal transmitting receptor in lymphocytes,     |
| Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      | differentiation 69 | domain family 2, | NK cells, and platelets; induced upon T cell     |
| Gzmb 3.8 Granzyme B Fragmentin 2 Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspasemediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      |                    | member C         | activation; involved in lymphocyte               |
| cytotoxic T and NK cells that activates caspase- mediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | proliferation                                    |
| mediated cell death when delivered into the target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine   | <u>Gzmb</u> | 3.8  | Granzyme B         | Fragmentin 2     | Abundant protease in the cytosolic granules of   |
| target cell through the immunological synapse  Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine   |             |      |                    |                  | cytotoxic T and NK cells that activates caspase- |
| Gzmk 3.36 Granzyme K Tryptase II Granule-secreted, pro-apoptotic serine  |             |      |                    |                  | mediated cell death when delivered into the      |
|  |             |      |                    |                  | target cell through the immunological synapse    |
| protease found in the cytoplasmic granules of  | <u>Gzmk</u> | 3.36 | Granzyme K         | Tryptase II      | Granule-secreted, pro-apoptotic serine           |
|  |             |      |                    |                  | protease found in the cytoplasmic granules of    |
| CTLs   |             |      |                    |                  | CTLs   |

| Klrg1  | 1.08 | Killer cell lectin-like | C-type lectin       | NK and T cell inhibitory receptor; binds to non- |
|--------|------|-------------------------|---------------------|--|
|        |      | receptor, subfamily     | domain family 15,   | MHC ligands                                      |
|        |      | G, member 1             | member A            |  |
|        |      |                         | (CLEC15A)           |  |
|        |      |                         | Mast cell function- |  |
|        |      |                         | associated antigen  |  |
|        |      |                         | (MAFA)              |  |
| Pik3cd | 1.19 | Phosphatidylinositol-   |                     | A subunit of PI3K; acts downstream of TLR4,      |
|        |      | 4,5-bisphosphate 3-     |                     | TCR, BCR, and CD40; contributes to T helper      |
|        |      | kinase catalytic        |                     | cell expansion, mast cell development, and       |
|        |      | subunit delta           |                     | neutrophil chemotaxis, extravasation, and        |
|        |      | isoform                 |                     | respiratory burst                                |
| Pik3cg | 1.21 | Phosphatidylinositol-   |                     | A subunit of PI3K; modulates leukocyte           |
|        |      | 4,5-bisphosphate 3-     |                     | chemotaxis to inflammatory sites and in          |
|        |      | kinase catalytic        |                     | response to chemoattractant agents               |
|        |      | subunit gamma           |                     |  |
|        |      | isoform                 |                     |  |
| Prdm1  | 1.77 | Positive regulatory     | B lymphocyte-       | TF that plays a role in the development,         |
|        |      | domain I-binding        | induced             | retention, and long-term establishment of T      |
|        |      | factor                  | maturation protein  | cell, NK cell, and NK-T cells in non-lymphoid    |
|        |      |                         | (BLIMP1)            | organs; drives the maturation of B cell into Ig  |
|        |      |                         |                     | secreting cells                                  |
|        |      |                         |                     |  |

| <u>Tcf7</u>   | 1.32 | Transcription factor 7                     |  | HMG box TF predominantly expressed by T cells that drives their development, although also involved in NK cell development; activates transcription through a Wnt/β-catenin signaling pathway  |
|---------------|------|--|--|--|
|               |      | Patte                                      | ern Recognition Rece                   | eptors   |
| <u>Cd14</u>   | 1.86 | Cluster of differentiation 14              |  | PRR that recognizes LPS; mostly found on macrophages   |
| <u>Cd180</u>  | 2    | Cluster of differentiation 180             |  | Heterodimeric binding partner of Ly86 that participates in LPS binding in APCs   |
| Clec4a2       | 1.75 | C-type lectin domain<br>family 4 member A2 |  | PRR that, upon binding mannose or fucose, is endocytosed and processed in the Ag presentation pathway  |
| <u>Clec4n</u> | 1.82 | C-type lectin domain family 4, member N    | Dectin-2                               | PRR specific for Mycobacterial mannose-<br>capped lipoarabinomannan  |
| Clec5a        | 1.73 | C-Type lectin domain family 5, member a    | Myeloid DAP12-<br>associating lectin-1 | Critical macrophage receptor for dengue virus serotypes 1-4; positive regulator of osteoclastogenesis  |
| Clec7a        | 2.64 | C-Type lectin domain family 7, member a    | Dectin-1                               | PRR specific for $\beta$ -1,3- and $\beta$ -1,6-linked glucans from fungi and plants; necessary for the TLR2-mediated inflammatory response and for TLR2-mediated activation of NF- $\kappa$ B |

| Ddx58        | 0.921 | DExD/H-box helicase | Retinoic acid-      | Cytoplasmic PRR that recognizes dsRNA; can        |
|--------------|-------|---------------------|---------------------|---|
|              |       | 58                  | inducible gene I    | promote T cell-independent B cell activation;     |
|              |       |                     | (RIG-I)             | uses MAVS as an adaptor                           |
| Fpr2         | 3.4   | Formyl peptide      | Lipoxin A4 receptor | Low affinity receptor for N-formyl-methionyl      |
|              |       | receptor 2          |                     | peptides; activates neutrophils                   |
| <u>Ifih1</u> | 1.43  | Interferon induced  | Helicard;           | PRR for cytoplasmic dsRNA; upon target            |
|              |       | with helicase C     | melanoma            | recognition, associates with MAVS to activate     |
|              |       | domain 1            | differentiation-    | TNK1 and IKBKE, which phosphorylate IRF3          |
|              |       |                     | associated protein  | and -7, which, in turn, activate transcription of |
|              |       |                     | 5 (MDA5)            | IFNα and -β                                       |
| <u>Ly86</u>  | 0.948 | Lymphocyte antigen  | Myeloid             | Heterodimeric binding partner of CD180 that       |
|              |       | 86                  | differentiation     | participates in LPS binding in APCs               |
|              |       |                     | factor 1 (MD-1)     |   |
| <u>Ly96</u>  | 0.606 | Lymphocyte antigen  | Myeloid             | Heterodimeric binding partner of TLR4 that        |
|              |       | 96                  | differentiation     | participates in LPS binding                       |
|              |       |                     | factor 2 (MD-2)     |   |
| Marco        | 4.23  | Macrophage          |                     | A PRR that recognizes LDL                         |
|              |       | receptor with       |                     |   |
|              |       | collagenous         |                     |   |
|              |       | structure           |                     |   |

| Myd88         | 0.978 | Myeloid               |               | Key adaptor in the TLR signaling pathways;     |
|---------------|-------|-----------------------|---------------|--|
|               |       | differentiation       |               | interacts with all TLRs except TLR3; activates |
|               |       |                       |               |  |
|               |       | primary response 88   |               | NFkB and IRFs                                  |
| Nlrp3         | 1.75  | NACHT domain-,        | Cryopyrin     | PRR with a wide diversity of recognized        |
|               |       | leucine-rich repeat-, |               | targets that activates the NLRP3               |
|               |       | and PYD-containing    |               | inflammasome consisting of NLRP3, PYCARD,      |
|               |       | protein 3             |               | and caspase-1/-8                               |
| Nod1          | 1.02  | Nucleotide binding    |               | Intracellular PRR that recognizes              |
|               |       | oligomerization       |               | peptidoglycan-derived muropeptides and         |
|               |       | domain containing 1   |               | Shigella effector proteins                     |
| Nod2          | 1.7   | Nucleotide-binding    |               | PRR specific for muramyl dipeptide (MDP);      |
|               |       | oligomerization       |               | upon binding to its ligand, recruits RIPK2 and |
|               |       | domain containing 2   |               | triggers MAPK and NFкB signaling               |
| <u>S100a8</u> | 2.3   | S100 calcium-binding  | Calgranulin A | Calcium- and zinc-binding protein involved in  |
|               |       | protein A8            |               | pro-inflammatory, antimicrobial, oxidant-      |
|               |       |                       |               | scavenging and apoptosis-inducing activities;  |
|               |       |                       |               | can induce neutrophil chemotaxis, adhesion,    |
|               |       |                       |               | phagocytosis, and degranulation;               |
|               |       |                       |               | predominantly found as calprotectin            |
|               |       |                       |               | (S100A8/A9) which has a wide plethora of       |
|               |       |                       |               | intra- and extracellular functions, including  |
|               |       |                       |               | adhesion, apoptosis, autophagy, cytoskeletal   |
|               |       |                       |               | remodeling, cytokine production, chemotaxis,   |
|               |       |                       |               | migration, inflammation, arachidonic acid      |

|             |      |   |              | metabolism, oxidant-scavenging, and PRR signaling   |
|-------------|------|---|--------------|---|
| Ticam2      | 1.79 | TIR domain-<br>containing adaptor<br>molecule 2 |              | Sorting adapter in various innate immune signaling cascades; bridges TLR2 and MyD88   |
| Tlr2        | 1.49 | Toll-like receptor 2                            | CD282        | Surface PRR that binds to various lipid-<br>containing PAMPs  |
| Tlr4        | 1.08 | Toll-like receptor 4                            | CD284        | Surface PRR that recognizes LPS; pairs with LY96 and CD14; acts via MYD88, TIRAP, and TRAF6, leading to NFκB activation, cytokine secretion, and the inflammatory response; in complex with TLR6, promotes sterile inflammation in monocytes/macrophages in response to oxLDL or amyloid-β 42 |
| <u>Tlr6</u> | 2.37 | Toll like receptor 6                            | CD286        | Surface PRR that recognizes diacylated and triacylated lipopeptides   |
| Tlr7        | 2.19 | Toll-like receptor 7                            | CD287        | Endosomic PRR that recognizes ssRNA   |
| Tlr8        | 1.65 | Toll-like receptor 8                            | CD288        | Endosomic PRR that recognizes ssRNA   |
| <u>Tir9</u> | 1.65 | Toll like receptor 9                            | CD289        | Endosomic PRR that recognizes unmethylated  CpG dinucleotides   |
|             |      |   | Phagocytosis |   |

| Fcgr2b        | 1.41 | Fragment             | CD32              | Low affinity receptor for the Fc region of        |
|---------------|------|----------------------|-------------------|---|
|               |      | crystallizable of    |                   | complexed or aggregated γ-lgs; involved in a      |
|               |      | immunoglobulin       |                   | variety of effector and regulatory functions      |
|               |      | gamma receptor IIb   |                   | such as phagocytosis of immune complexes          |
|               |      |                      |                   | and modulation of Ab production by B cells;       |
|               |      |                      |                   | essential for the maintenance of humoral          |
|               |      |                      |                   | tolerance; acts as a late checkpoint at the level |
|               |      |                      |                   | of class-switched memory B cells,                 |
|               |      |                      |                   | plasmablasts or plasma cells; regulates           |
|               |      |                      |                   | plasma-cell homeostasis and survival              |
| <u>Itgam</u>  | 2.19 | Integrin alpha M     | CD11b             | Pairs with CD18 to form Mac-1 aka                 |
|               |      |                      |                   | complement receptor 3; mediates leukocyte         |
|               |      |                      |                   | activation, adhesion, chemotaxis, migration,      |
|               |      |                      |                   | phagocytosis, and cell-mediated cytotoxicity;     |
|               |      |                      |                   | serves as a macrophage marker                     |
| Ncf4          | 1.34 | Neutrophil cytosolic | SH3 and PX        | Cytosolic regulatory component of the             |
|               |      | factor 4             | domain-containing | superoxide-producing phagocyte NADPH-             |
|               |      |                      | protein 4         | oxidase, a multicomponent enzyme system           |
|               |      |                      | (SH3PXD4)         | important for host defense                        |
| <u>S100a8</u> | 2.3  | S100 calcium-binding | Calgranulin A     | Calcium- and zinc-binding protein involved in     |
|               |      | protein A8           |                   | pro-inflammatory, antimicrobial, oxidant-         |
|               |      |                      |                   | scavenging and apoptosis-inducing activities;     |
|               |      |                      |                   | can induce neutrophil chemotaxis, adhesion,       |
|               |      |                      |                   | phagocytosis, and degranulation;                  |

|             |      |   |  | predominantly found as calprotectin  (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |
|-------------|------|---|--|---|
| Slamf7      | 2.32 | Signaling lymphocytic activation molecule family member 7 |  | A super-activator of macrophages and a strong promoter of phagocytosis; binds to CD74   |
|             |      | ROS   | Generation & Prote                               | ction   |
| <u>Cybb</u> | 2.37 | Cytochrome b-245 heavy chain                              | Nox2   | Part of the NADPH oxidase process; generates superoxides  |
| Ncf4        | 1.34 | Neutrophil cytosolic factor 4                             | SH3 and PX domain-containing protein 4 (SH3PXD4) | Cytosolic regulatory component of the superoxide-producing phagocyte NADPH-oxidase, a multicomponent enzyme system important for host defense   |
| Nos2        | 4.12 | Inducible nitric oxide synthase (iNOS)                    |  | Produces reactive oxygen species and contributes to inflammatory cytokine production  |

| Prkcd | 0.892 | Protein kinase C<br>delta  |  | Ca(2+)-independent, phospholipid- and DAG- dependent serine/threonine-protein kinase that promotes apoptosis in response to DNA damage but inhibits it during cytokine receptor-initiated cell death; required for oxygen radical production by NADPH oxidase  |
|-------|-------|--|--|--|
|       |       |  | Stress Response                        |  |
| Aft2  |       | Activator of Fe+ transcription 2  Amyloid-beta precursor protein |  | Iron-regulated transcriptional activator; activates genes involved in intracellular iron use; required for iron homeostasis and oxidative stress resistance  Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; involved in cell mobility, copper homeostasis, and oxidative stress |
|       |       |  | T Cell Function                        |  |
| Cd2   | 1.68  | Cluster of differentiation 2                                     | Leukocyte functional antigen 2 (LFA-2) | Interacts with LFA-3 and CD48 to mediate adhesion between T cells and other cell types   |

| delta  Cd3e  Cd3e  Cluster of differentiation 3 epsilon  CD3y/CD3e; also participates in internalizat and cell surface down-regulation of TCR-CL complexes via endocytosis sequences presenting to the TCR-CD3 complex; plays essential role in the dynamic regulation of Tale and cell surface  Cd4  1.68  Cluster of differentiation 4  Cd4  Cd4  Cd4  Cd4  Cd4  Cd5  Component of the TCR-CD3 complex; plays essential role in the dynamic regulation of Tale and cell surface  Cd4  Cd4  Cd4  Cd4  Cd4  Cd4  Cluster of differentiation 4  Cd4  Cd4  Cd4  Cd4  Cd5  Cluster of differentiation 4  Cd5  Cd7  Camponent of the TCR-CD3 complex; plays essential role in the dynamic regulation of Tale and the cell surface | <u>Cd3d</u> | 2.3  | Cluster of        |      | Component of the TCR-CD3 complex; upon          |
|---|-------------|------|-------------------|------|---|
| Cd3e  2.27 Cluster of differentiation 3 the TCR-CD3 complex assembly by forming two heterodimers CD36/CD3ɛ and CD3y/CD3ɛ; also participates in internalizat and cell surface down-regulation of TCR-CI complexes via endocytosis sequences press in CD3ɛ cytosolic region  Cd3g  2.01 Cluster of differentiation 3 essential role in the dynamic regulation of T expression at the cell surface  Cd4  1.68 Cluster of differentiation 4 Signature helper T cell marker; binds to MI class II and provides necessary costimulation for T cell activation   |             |      | differentiation 3 |      | phosphorylation by Lck, serves as a docking     |
| differentiation 3 epsilon  the TCR-CD3 complex assembly by forming two heterodimers CD3δ/CD3ε and CD3γ/CD3ε; also participates in internalizat and cell surface down-regulation of TCR-Ci complexes via endocytosis sequences prese in CD3ε cytosolic region  Cd3g  2.01  Cluster of differentiation 3 gamma  expression at the cell surface  Cd4  1.68  Cluster of differentiation 4  Class II and provides necessary costimulation for T cell activation  |             |      | delta             |      | station for downstream TCR signaling adaptors   |
| epsilon  two heterodimers CD36/CD3ɛ and  CD3y/CD3ɛ; also participates in internalizat  and cell surface down-regulation of TCR-Cl  complexes via endocytosis sequences prese  in CD3ɛ cytosolic region  Cd3g  2.01  Cluster of  differentiation 3  gamma  essential role in the dynamic regulation of T  expression at the cell surface  Cd4  1.68  Cluster of  differentiation 4  Class II and provides necessary costimulation  for T cell activation   | Cd3e        | 2.27 | Cluster of        |      | Component of the TCR-CD3 complex; initiates     |
| CD3y/CD3e; also participates in internalizat and cell surface down-regulation of TCR-CI complexes via endocytosis sequences press in CD3e cytosolic region  Cd3g  2.01 Cluster of   |             |      | differentiation 3 |      | the TCR-CD3 complex assembly by forming the     |
| and cell surface down-regulation of TCR-CIC complexes via endocytosis sequences prese in CD3ɛ cytosolic region  Cd3g  2.01  Cluster of  differentiation 3  gamma  cssential role in the dynamic regulation of T expression at the cell surface  Cd4  1.68  Cluster of  differentiation 4  Class II and provides necessary costimulation for T cell activation   |             |      | epsilon           |      | two heterodimers CD3δ/CD3ε and                  |
| complexes via endocytosis sequences prese in CD3ε cytosolic region  Cd3g  2.01 Cluster of   |             |      |                   |      | CD3γ/CD3ε; also participates in internalization |
| in CD3ɛ cytosolic region  Cd3g  2.01  Cluster of  differentiation 3  gamma  Cd4  1.68  Cluster of  differentiation 4  Cd4  Component of the TCR-CD3 complex; plays essential role in the dynamic regulation of T expression at the cell surface  Signature helper T cell marker; binds to Mi class II and provides necessary costimulation for T cell activation  |             |      |                   |      | and cell surface down-regulation of TCR-CD3     |
| Cd3g  2.01 Cluster of Component of the TCR-CD3 complex; plays essential role in the dynamic regulation of T gamma  expression at the cell surface  Cd4  1.68 Cluster of Signature helper T cell marker; binds to Mi class II and provides necessary costimulation for T cell activation   |             |      |                   |      | complexes via endocytosis sequences present     |
| differentiation 3 gamma expression at the cell surface  Cd4 1.68 Cluster of differentiation 4 Class II and provides necessary costimulation for T cell activation   |             |      |                   |      | in CD3ε cytosolic region                        |
| gamma expression at the cell surface  Cd4 1.68 Cluster of Signature helper T cell marker; binds to MI class II and provides necessary costimulation for T cell activation   | Cd3g        | 2.01 | Cluster of        |      | Component of the TCR-CD3 complex; plays an      |
| Cd4  1.68  Cluster of  differentiation 4  Class II and provides necessary costimulation  for T cell activation  |             |      | differentiation 3 |      | essential role in the dynamic regulation of TCR |
| differentiation 4 class II and provides necessary costimulation for T cell activation   |             |      | gamma             |      | expression at the cell surface                  |
| for T cell activation   | <u>Cd4</u>  | 1.68 | Cluster of        |      | Signature helper T cell marker; binds to MHC    |
|   |             |      | differentiation 4 |      | class II and provides necessary costimulation   |
| Cd5   2.56   Cluster of   LEU1   Type-I transmembrane glycoprotein found  |             |      |                   |      | for T cell activation                           |
|   | <u>Cd5</u>  | 2.56 | Cluster of        | LEU1 | Type-I transmembrane glycoprotein found on      |
| differentiation 5 the surface of T and B cells; may act as a  |             |      | differentiation 5 |      | the surface of T and B cells; may act as a      |
| receptor in regulating T cell proliferation   |             |      |                   |      | receptor in regulating T cell proliferation     |
| Cd8a 2.02 Cluster of LEU2 Alpha chain of the CD8 coreceptor, which  | <u>Cd8a</u> | 2.02 | Cluster of        | LEU2 | Alpha chain of the CD8 coreceptor, which        |
| differentiation 8 binds to MHC class I; in T cells, functions   |             |      | differentiation 8 |      | binds to MHC class I; in T cells, functions     |
| alpha primarily as a coreceptor for MHC class I   |             |      | alpha             |      | primarily as a coreceptor for MHC class I       |
| molecule:peptide complex; in NK cells, th   |             |      |                   |      | molecule:peptide complex; in NK cells, the      |

| <u>Cd8b1</u> | 2.14  | Cluster of                    |   | presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells  Beta chain of the CD8 coreceptor, which binds |
|--------------|-------|-------------------------------|---|--|
|              |       | differentiation 8<br>beta 1   |   | to MHC class I   |
| <u>Cd28</u>  | 2.48  | Cluster of differentiation 28 |   | Essential T cell co-receptor that enhances T cell activation, proliferation, cytokine production, and survival; binds to CD80 and CD86   |
| <u>Cd69</u>  | 1.46  | Cluster of differentiation 69 | C-type lectin<br>domain family 2,<br>member C | Signal transmitting receptor in lymphocytes,  NK cells, and platelets; induced upon T cell  activation; involved in lymphocyte  proliferation  |
| <u>Cd37</u>  | 2.07  | Cluster of differentiation 37 | Tetraspanin-26                                | Cell surface glycoprotein known to complex with integrins and other transmembrane 4 superfamily proteins; may play a role in T cell- B cell interactions                             |
| <u>Cd97</u>  | 0.748 | Cluster of differentiation 97 | BL-Ac[F2]                                     | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through  |

|               |       |                       |                  | binding integrin counterreceptors on                      |
|---------------|-------|-----------------------|------------------|---|
|               |       |                       |                  | endothelial cells   |
| <u>Cd247</u>  | 2.49  | Cluster of            | T cell surface   | Central intracellular signaling chain of the TCR,         |
|               |       | differentiation 247   | glycoprotein CD3 | to which downstream signaling adaptors dock               |
|               |       |                       | zeta chain       |   |
| Cxcr3         | 2.23  | C-X-C motif           | CD183            | Induces integrin activation, cytoskeletal                 |
|               |       | chemokine receptor    |                  | remodeling, and chemotaxis; expressed by T                |
|               |       | 3                     |                  | cells and NK cells; prominently expressed in              |
|               |       |                       |                  | effector and memory T cells                               |
| Cyfip2        | 1.74  | Cytoplasmic FMR1-     |                  | Involved in T cell adhesion and p53/TP53-                 |
|               |       | interacting protein 2 |                  | dependent induction of apoptosis                          |
| Gata3         | 1.2   | GATA binding          |                  | Transcriptional activator that binds to the               |
|               |       | protein 3             |                  | enhancer of the TCR $\alpha$ and $\delta$ genes; required |
|               |       |                       |                  | for T <sub>H</sub> 2 differentiation following immune and |
|               |       |                       |                  | inflammatory responses                                    |
| <u>Gpr183</u> | 0.932 | G protein-coupled     | EBV-induced G    | Lymphocyte GPCR that acts as a chemotactic                |
|               |       | receptor 183          | protein-coupled  | receptor for B cells, T cells, splenic DCs,               |
|               |       |                       | receptor 2       | monocytes/macrophages, and astrocytes                     |
| Gzmb          | 3.8   | Granzyme B            | Fragmentin 2     | Abundant protease in the cytosolic granules of            |
|               |       |                       |                  | cytotoxic T and NK cells that activates caspase-          |
|               |       |                       |                  | mediated cell death when delivered into the               |
|               |       |                       |                  | target cell through the immunological synapse             |

| <u>Gzmk</u>    | 3.36 | Granzyme K         | Tryptase II         | Granule-secreted, pro-apoptotic serine           |
|----------------|------|--------------------|---------------------|--|
|                |      |                    |                     | protease found in the cytoplasmic granules of    |
|                |      |                    |                     | CTLs   |
|                |      |                    |                     |  |
| <u>Eomes</u>   | 1.04 | Eomesodermin       | T-box brain protein | Transcriptional activator critical for           |
|                |      |                    | 2 (TBR2)            | development; involved in CD8 <sup>+</sup> T cell |
|                |      |                    |                     | differentiation                                  |
| <u>lcos</u>    | 2.45 | Inducible T cell   | CD278               | Enhances all basic T cell responses to foreign   |
|                |      | costimulator       |                     | Ag; essential both for efficient interaction     |
|                |      |                    |                     | between T and B cells and for normal Ab          |
|                |      |                    |                     | responses to T cell-dependent Ags                |
| <u>lkzf1</u>   | 1.19 | IKAROS family zinc |                     | Transcriptional regulator of hematopoietic cell  |
|                |      | finger 1           |                     | differentiation; plays a role in T and B cell    |
|                |      |                    |                     | development                                      |
| <u>ll12rb1</u> | 1.07 | Interleukin 12     | CD212               | Cytokine receptor component that associates      |
|                |      | receptor subunit   |                     | with IL12RB2 to IL23R                            |
|                |      | beta 1             |                     |  |
| <u>Itk</u>     | 2.06 | Interleukin-2-     | LYK                 | Key actor in the TCR signaling cascade;          |
|                |      | inducible T cell   |                     | phosphorylates PLCy1, LAT, and LCP2              |
|                |      | kinase             |                     |  |
|                |      |                    |                     |  |
| <u>Lck</u>     | 1.98 | Lymphocyte cell    |                     | Src family tyrosine kinase that acts as one of   |
|                |      | kinase             |                     | the main signaling intermediaries downstream     |
|                |      |                    |                     |  |

|              |       |  |  | of the TCR; constitutively associated with the cytoplasmic portion of CD4  |
|--------------|-------|--|--|--|
| Lcp1         | 1.62  | Lymphocyte cytosolic protein 1   | Plastin-2  | Actin-binding protein that promotes T cell activation in response to costimulation through TCR/CD3 and CD2 or CD28; assists with IL2RA transport to the cell surface                           |
| Nfatc2       | 0.971 | Nuclear factor of activated T cells,   |  | Cytosolic component of the NFAT TF complex; mediates induction of IL-2, IL-3, IL-4, TNF $\alpha$ ,   |
|              |       | cytoplasmic 2  |  | and GM-CSF   |
| Pik3cd       | 1.19  | Phosphatidylinositol-<br>4,5-bisphosphate 3-<br>kinase catalytic<br>subunit delta<br>isoform |  | A subunit of PI3K; acts downstream of TLR4, TCR, BCR, and CD40; contributes to T helper cell expansion, mast cell development, and neutrophil chemotaxis, extravasation, and respiratory burst |
| Pik3cg       | 1.21  | Phosphatidylinositol-<br>4,5-bisphosphate 3-<br>kinase catalytic<br>subunit gamma<br>isoform |  | A subunit of PI3K; modulates leukocyte chemotaxis to inflammatory sites and in response to chemoattractant agents  |
| <u>Prdm1</u> | 1.77  | Positive regulatory  domain I-binding  factor  | B lymphocyte-<br>induced<br>maturation protein<br>(BLIMP1) | TF that plays a role in the development, retention, and long-term establishment of T cell, NK cell, and NK-T cells in non-lymphoid   |

|              |      |  |  | organs; drives the maturation of B cell into Ig secreting cells   |
|--------------|------|--|--|---|
| <u>Ptprc</u> | 1.5  | Protein tyrosine  phosphatase  receptor type C   | CD45; leukocyte<br>common antigen<br>(LCA) | Delivers costimulation during T cell activation  upon binding to its ligand DPP4;  dephosphorylates Lyn and suppresses JAK  kinases   |
| <u>Rora</u>  | 1.69 | Retinoic acid receptor-related orphan receptor A |  | Nuclear receptor that binds hormone response elements upstream of several genes to enhance the expression of those genes; plays an essential role in the development of type 2 ILCs   |
| <u>Spn</u>   | 1.63 | Sialophorin                                      | Leukosialin; CD43                          | Cell surface sialoglycoprotein expressed by T cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNγ in CD4 <sup>+</sup> T cells |
| <u>Syk</u>   | 1.95 | Spleen-associated tyrosine kinase                |  | Critical kinase that transmits signals from the TCR and BCR   |
| Tbx21        | 1.5  | T-box transcription factor 21                    |  | Initiates $T_H1$ lineage development from naïve $T_H$ precursor cells both by activating $T_H1$ genetic programs and by repressing the opposing $T_H2$ and $T_H17$ genetic programs   |

| transcription through a Wnt/β-catenin signaling pathway  2ap70  1.72  Zeta chain of T cell receptor associated protein kinase 70  regulation of the adaptive immune response; regulates motility, adhesion, proliferation, differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT   | Tcf7         | 1.32 | Transcription factor |                      | HMG box TF predominantly expressed by T          |
|---|--------------|------|----------------------|----------------------|--|
| transcription through a Wnt/β-catenin signaling pathway  2ap70  1.72 Zeta chain of T cell receptor associated protein kinase 70  protein kinase 70  differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb  1.28 CCAAT/enhancer-binding protein beta  Critical macrophage TF that promotes expression of several acute-phase and  |              |      | 7                    |                      | cells that drives their development, although    |
| Transcription Factors & Coactivators  2 Zeta chain of T cell receptor associated protein kinase 70  Tyrosine kinase that plays an essential role in regulation of the adaptive immune response; regulates motility, adhesion, proliferation, differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer-binding protein beta expression of several acute-phase and    |              |      |                      |                      | also involved in NK cell development; activates  |
| Transcription Factors & Coactivators  1.72 Zeta chain of T cell receptor associated protein kinase 70  Tyrosine kinase that plays an essential role in regulation of the adaptive immune response; regulates motility, adhesion, proliferation, differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer-binding protein beta expression of several acute-phase and |              |      |                      |                      | transcription through a Wnt/β-catenin            |
| receptor associated protein kinase 70 regulation of the adaptive immune response; regulates motility, adhesion, proliferation, differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer-binding protein beta expression of several acute-phase and  |              |      |                      |                      | signaling pathway                                |
| protein kinase 70  regulates motility, adhesion, proliferation, differentiation, and cytokine expression in T  cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb  1.28 CCAAT/enhancer- binding protein beta  Critical macrophage TF that promotes expression of several acute-phase and  | <u>Zap70</u> | 1.72 | Zeta chain of T cell |                      | Tyrosine kinase that plays an essential role in  |
| differentiation, and cytokine expression in T cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta Critical macrophage TF that promotes expression of several acute-phase and   |              |      | receptor associated  |                      | regulation of the adaptive immune response;      |
| cells; contributes to the development and activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta Critical macrophage TF that promotes expression of several acute-phase and   |              |      | protein kinase 70    |                      | regulates motility, adhesion, proliferation,     |
| activation of primary B cells; phosphorylates at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta Critical macrophage TF that promotes expression of several acute-phase and   |              |      |                      |                      | differentiation, and cytokine expression in T    |
| at least two essential adapter proteins: LAT and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta expression of several acute-phase and  |              |      |                      |                      | cells; contributes to the development and        |
| Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer-binding protein beta and LCP2, leading in turn to a large number of signaling molecules being recruited  Transcription Factors & Coactivators  Critical macrophage TF that promotes expression of several acute-phase and   |              |      |                      |                      | activation of primary B cells; phosphorylates    |
| Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta expression of several acute-phase and   |              |      |                      |                      | at least two essential adapter proteins: LAT     |
| Transcription Factors & Coactivators  Cebpb 1.28 CCAAT/enhancer- binding protein beta Critical macrophage TF that promotes expression of several acute-phase and  |              |      |                      |                      | and LCP2, leading in turn to a large number of   |
| Cebpb       1.28       CCAAT/enhancer-       Critical macrophage TF that promotes         binding protein beta       expression of several acute-phase and  |              |      |                      |                      | signaling molecules being recruited              |
| binding protein beta expression of several acute-phase and  |              |      | Transcr              | iption Factors & Coa | ctivators  |
|   | <u>Cebpb</u> | 1.28 | CCAAT/enhancer-      |                      | Critical macrophage TF that promotes             |
| inflammatory cytokine genes, including II6  |              |      | binding protein beta |                      | expression of several acute-phase and            |
|   |              |      |                      |                      | inflammatory cytokine genes, including II6       |
| Eomes 1.04 Eomesodermin T-box brain protein Transcriptional activator critical for  | Eomes        | 1.04 | Eomesodermin         | T-box brain protein  | Transcriptional activator critical for           |
| 2 (TBR2) development; involved in CD8 <sup>+</sup> T cell   |              |      |                      | 2 (TBR2)             | development; involved in CD8 <sup>+</sup> T cell |
| differentiation   |              |      |                      |                      | differentiation                                  |

| Foxp3        | 1.56 | Forkhead box P3       | DIETER | Master TF for Tregs; represses expression of              |
|--------------|------|-----------------------|--------|---|
|              |      |                       |        | II2 and Ifng; activates expression of Tnfrsf18,           |
|              |      |                       |        |   |
|              |      |                       |        | Il2ra, and Ctla4  |
| Gata3        | 1.2  | GATA binding          |        | Transcriptional activator that binds to the               |
|              |      | protein 3             |        | enhancer of the TCR $\alpha$ and $\delta$ genes; required |
|              |      |                       |        | for T <sub>H</sub> 2 differentiation following immune and |
|              |      |                       |        | inflammatory responses                                    |
| Irf1         | 1.71 | Interferon regulatory |        | Transcriptional regulator that promotes                   |
|              |      | factor 1              |        | inflammatory innate and adaptive immune                   |
|              |      | ractor 1              |        |   |
|              |      |                       |        | responses   |
| <u>Irf4</u>  | 1.2  | Interferon regulatory |        | Transcriptional activator that complexes with             |
|              |      | factor 4              |        | BATF and binds ISREs within the promoters of              |
|              |      |                       |        | multiple genes involved in inflammation                   |
| <u>Irf7</u>  | 2.22 | Interferon regulatory |        | Key transcriptional regulator of type I IFN-              |
|              |      | factor 7              |        | dependent immune responses; promotes                      |
|              |      |                       |        | transcription of IFNα and -β                              |
| <u>Irf8</u>  | 1.12 | Interferon regulatory |        | TF that regulates of lineage commitment in                |
|              |      | factor 8              |        | myeloid cell maturation; promotes monocyte                |
|              |      |                       |        | and plasmacytoid DC development                           |
| <u>Ikzf1</u> | 1.19 | IKAROS family zinc    |        | Transcriptional regulator of hematopoietic cell           |
|              |      | finger 1              |        | differentiation; plays a role in T and B cell             |
|              |      | iiikei 1              |        |   |
|              |      |                       |        | development   |
|              |      |                       |        |   |

| Mef2c  | 1.24  | Myocyte enhancer       |                    | Transcriptional activator that binds specifically |
|--------|-------|------------------------|--------------------|---|
|        |       | factor 2c              |                    | to the MEF2 element present in the regulatory     |
|        |       |                        |                    | regions of many muscle-specific genes;            |
|        |       |                        |                    | controls cardiac morphogenesis and                |
|        |       |                        |                    | myogenesis, and is also involved in vascular      |
|        |       |                        |                    | development; required for B cell survival and     |
|        |       |                        |                    | proliferation in response to BCR stimulation,     |
|        |       |                        |                    | efficient IgG1 Ab responses to T cell-            |
|        |       |                        |                    | dependent Ags, and for normal induction of        |
|        |       |                        |                    | GC B cells  |
| Nfatc2 | 0.971 | Nuclear factor of      |                    | Cytosolic component of the NFAT TF complex;       |
|        |       | activated T cells,     |                    | mediates induction of IL-2, IL-3, IL-4, TNFα,     |
|        |       | cytoplasmic 2          |                    | and GM-CSF  |
| Nfkb1  | 0.349 | Nuclear factor kappa   | p105/p50           | One of the NFkB family TFs; inhibits              |
|        |       | B subunit 1            |                    | inflammation                                      |
| Pou2f2 | 1.81  | POU domain class 2,    |                    | TF that regulates Ab and IL-6 expression in B     |
|        |       | transcription factor 2 |                    | cells   |
|        |       |                        |                    |   |
| Prdm1  | 1.77  | Positive regulatory    | B lymphocyte-      | TF that plays a role in the development,          |
|        |       | domain I-binding       | induced            | retention, and long-term establishment of T       |
|        |       | factor                 | maturation protein | cell, NK cell, and NK-T cells in non-lymphoid     |
|        |       |                        | (BLIMP1)           | organs; drives the maturation of B cell into Ig   |
|        |       |                        |                    | secreting cells                                   |
|        |       |                        |                    |   |

| Stat1 | 1.53  | Signal transducer    | Transcriptional activator that mediates cellular                      |
|-------|-------|----------------------|---|
|       |       | and activator of     | responses to IFNs, cytokines, and other                               |
|       |       | transcription 1      | growth factors  |
|       |       |                      |   |
| Stat3 | 0.649 | Signal transducer    | Transcriptional activator of genes involved in                        |
|       |       | and activator of     | cell growth and apoptosis; activated by JAKs                          |
|       |       | transcription 3      |   |
|       |       |                      |   |
| Stat4 | 1.25  | Signal transducer    | Essential TF for T <sub>H</sub> 1 CD4 <sup>+</sup> T cell development |
|       |       | and activator of     | and IFNγ production; also promotes                                    |
|       |       | transcription 4      | expression of MyD88   |
|       |       |                      |   |
| Stat6 | 0.571 | Signal transducer    | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and         |
|       |       | and activator of     | macrophage function and M2 macrophage                                 |
|       |       | transcription 6      | polarization  |
|       |       |                      |   |
| Tbx21 | 1.5   | T-box transcription  | Initiates T <sub>H</sub> 1 lineage development from naïve             |
|       |       | factor 21            | $T_H$ precursor cells both by activating $T_H 1$                      |
|       |       |                      | genetic programs and by repressing the                                |
|       |       |                      | opposing T <sub>H</sub> 2 and T <sub>H</sub> 17 genetic programs      |
|       |       |                      |   |
| Tcf7  | 1.32  | Transcription factor | HMG box TF predominantly expressed by T                               |
|       |       | 7                    | cells that drives their development, although                         |
|       |       |                      | also involved in NK cell development; activates                       |
|       |       |                      | transcription through a Wnt/β-catenin                                 |
|       |       |                      | signaling pathway   |
|       |       |                      | Signaling patriway  |
|       |       |                      |   |

| <b>-</b> 1 - |       |                   |                      |   |
|--------------|-------|-------------------|----------------------|---|
| Zbp1         | 1.86  | Z-DNA binding     | Tumor stroma and     | Key innate sensor that recognizes and binds Z-      |
|              |       | protein 1         | activated            | RNA structures, which are produced by a             |
|              |       |                   | macrophage           | number of viruses, and induces type-I IFN           |
|              |       |                   | protein DLM-1        | production; key activator of cellular               |
|              |       |                   |                      | necroptosis; promotes IL-1 $\alpha$ induction in an |
|              |       |                   |                      | NLRP3-inflammasome-independent manner               |
|              |       |                   | Ubiquitin Regulation | 1   |
|              |       |                   |                      |   |
| Bcl10        | 0.393 | B cell            |                      | Activates NFκB via ubiquitination of ΙΚΚγ           |
|              |       | lymphoma/leukemia |                      |   |
|              |       | 10                |                      |   |
|              |       |                   |                      |   |
| Cyld         | 0.595 | Cylindromatosis   |                      | Inhibits NFkB activation by deubiquitinating        |
|              |       | lysine 63         |                      | upstream signaling factors; inhibits Wnt            |
|              |       | deubiquitinase    |                      | signaling; restricts polyubiquitination of RIPK1    |
|              |       |                   |                      | and -2, thereby limiting necroptosis                |
| Tnfaip3      | 1.34  | Tumor necrosis    |                      | Ubiquitin-editing enzyme that complexes with        |
| Tillaip3     | 1.54  |                   |                      |   |
|              |       | factor, alpha-    |                      | ITCH to degrade inflammatory signaling              |
|              |       | induced protein 3 |                      | components in the TNF, IL1, and TLR                 |
|              |       |                   |                      | pathways; targets TRAF2, TRAF6, and IKK             |
| Ubc          | 0.653 | Polyubiquitin C   |                      | Serves various roles, including innate              |
|              |       |                   |                      | immunity, DNA repair, and stimulation of            |
|              |       |                   |                      | autophagy and the proteasomal response              |
|              |       |                   |                      |   |

|        | NBTXR3+PBT+αPD1 vs PBT+αPD1 |                       |                        |   |  |  |
|--------|-----------------------------|-----------------------|------------------------|---|--|--|
| Gene   | Log2 fold                   | Full Name             | Notable Aliases        | Function  |  |  |
|        | change                      |                       |                        |   |  |  |
|        |                             |                       |                        |   |  |  |
|        |                             | Adhes                 | ion & Cell-Cell Intera | actions   |  |  |
| Cd2    | 1.66                        | Cluster of            | T cell surface         | Interacts with LFA-3 and CD48 to mediate                      |  |  |
|        |                             | differentiation 2     | antigen CD2            | adhesion between T cells and other cell types                 |  |  |
| lcam1  | 0.688                       | Intracellular         | CD54                   | Cell surface glycoprotein that serves as strong               |  |  |
|        |                             | adhesion molecule 1   |                        | adhesive ligand for LFA-1; important for                      |  |  |
|        |                             |                       |                        | leukocyte mobility and costimulation                          |  |  |
| Itgal  | 0.959                       | Integrin alpha L      |                        | Pairs with ITGB2 to form LFA-1, a common                      |  |  |
|        |                             |                       |                        | leukocyte adhesion molecule and                               |  |  |
|        |                             |                       |                        | costimulatory receptor  |  |  |
|        |                             | Antiger               | n Processing & Prese   | entation  |  |  |
| H2-K1  | 0.439                       | Histocompatibility 2, |                        | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T      |  |  |
| 11Z-KI | 0.433                       | K1, K region          |                        | cells   |  |  |
|        |                             | KI, KTEGIOH           |                        | cens  |  |  |
|        | Anti-Inflammatory           |                       |                        |   |  |  |
| Ctla4  | 0.969                       | Cytotoxic T           | CD152                  | Inhibitory receptor that blocks CD28                          |  |  |
|        |                             | lymphocyte antigen    |                        | costimulation by competitively binding its                    |  |  |
|        |                             | 4                     |                        | ligands CD80 and CD86   |  |  |
| Stat6  | 0.392                       | Signal transducer     |                        | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and |  |  |
|        |                             | and activator of      |                        | macrophage function and M2 macrophage                         |  |  |

|           |       | transcription 6     |                     | polarization   |  |  |
|-----------|-------|---------------------|---------------------|--|--|--|
| Autophagy |       |                     |                     |  |  |  |
| Ubc       | 0.39  | Polyubiquitin C     |                     | Serves various roles, including innate                   |  |  |
|           |       |                     |                     | immunity, DNA repair, and stimulation of                 |  |  |
|           |       |                     |                     | autophagy and the proteasomal response                   |  |  |
|           |       | В                   | Cell-associated Gen | es   |  |  |
| Gpr183    | 1.14  | G protein-coupled   | EBV-induced G       | Lymphocyte GPCR that acts as a chemotactic               |  |  |
|           |       | receptor 183        | protein-coupled     | receptor for B cells, T cells, splenic DCs,              |  |  |
|           |       |                     | receptor 2          | monocytes/macrophages, and astrocytes                    |  |  |
| Prdm1     | 1.22  | Positive regulatory | B lymphocyte-       | TF that plays a role in the development,                 |  |  |
|           |       | domain I-binding    | induced             | retention, and long-term establishment of T              |  |  |
|           |       | factor              | maturation protein  | cell, NK cell, and NK-T cells in non-lymphoid            |  |  |
|           |       |                     | (BLIMP1)            | organs; drives the maturation of B cell into Ig          |  |  |
|           |       |                     |                     | secreting cells  |  |  |
| Spn       | 0.91  | Sialophorin         | Leukosialin; CD43   | Cell surface sialoglycoprotein expressed by T            |  |  |
|           |       |                     |                     | cells, B cells, monocytes, and granulocytes;             |  |  |
|           |       |                     |                     | promotes lymph node localization in T cells;             |  |  |
|           |       |                     |                     | shunts T cells away from the T <sub>H</sub> 2 phenotype  |  |  |
|           |       |                     |                     | and towards T <sub>H</sub> 1; promotes the expression of |  |  |
|           |       |                     |                     | IFNγ in CD4 <sup>+</sup> T cells                         |  |  |
| Syk       | 0.978 | Spleen-associated   |                     | Critical kinase that transmits signals from the          |  |  |
|           |       | tyrosine kinase     |                     | TCR and BCR  |  |  |

|             | Chemotaxis |   |  |   |  |  |
|-------------|------------|---|--|---|--|--|
| Ccr9        | 0.918      | C-C motif chemokine receptor 9                  |  | Receptor for CCL25; increases intracellular  Ca(2+) levels upon ligand binding  |  |  |
| Cxcr3       | 1.38       | C-X-C motif chemokine receptor                  | CD183                                    | Induces integrin activation, cytoskeletal remodeling, and chemotaxis; expressed by T cells and NK cells; prominently expressed in effector and memory T cells |  |  |
| Gpr183      | 1.14       | G protein-coupled receptor 183                  | EBV-induced G protein-coupled receptor 2 | Lymphocyte GPCR that acts as a chemotactic receptor for B cells, T cells, splenic DCs, monocytes/macrophages, and astrocytes                                  |  |  |
|             |            | Comple  | ement & Humoral Im                       | nmunity   |  |  |
| <u>C2</u>   | 0.885      | Complement component 2                          |  | Serine protease that binds to C4b to form the C4bC2 complex   |  |  |
| <u>Cd55</u> | 1.31       | Cluster of differentiation 55                   | Complement  decay-accelerating  factor   | Cell surface glycoprotein that interacts with surface-bound C4b and inhibits its conversion of C2 to C2b  |  |  |
| <u>Cfp</u>  | 0.922      | Complement factor properdin                     |  | Alternate complement pathway component; when cleaved, produces a serine protease that binds to C3b to form C3 convertase                                      |  |  |
| Fcer1a      | 1.18       | Fragment<br>crystallizable of<br>immunoglobulin |  | High affinity receptor for IgE; responsible for initiating the allergic response  |  |  |

|               |       | epsilon receptor 1a                                   |                                   |  |  |  |  |
|---------------|-------|---|-----------------------------------|--|--|--|--|
| Costimulation |       |   |                                   |  |  |  |  |
| <u>Cd28</u>   | 0.848 | Cluster of differentiation 28                         |                                   | Essential T cell co-receptor that enhances T cell activation, proliferation, cytokine production, and survival; binds to CD80 and CD86   |  |  |  |
| <u>lcam1</u>  | 0.688 | Intracellular adhesion molecule 1                     | CD54                              | Cell surface glycoprotein that serves as strong adhesive ligand for LFA-1; important for leukocyte mobility and costimulation  |  |  |  |
| <u>Itgal</u>  | 0.959 | Integrin alpha L                                      |                                   | Pairs with ITGB2 to form lymphocyte functionassociated antigen-1 (LFA-1), a common leukocyte adhesion molecule and costimulatory receptor  |  |  |  |
| Tnfrsf11a     | 1.18  | Tumor necrosis factor receptor superfamily member 11A | Receptor activator of NFkB (RANK) | Recruits TRAFs and activates NFkB and JNK activation upon binding to RANKL on the surface of T cells   |  |  |  |
| Tnfrsf4       | 1.21  | Tumor necrosis factor receptor superfamily member 4   | OX40; CD134                       | Receptor for TNFSF4/OX40L/GP34;  costimulatory molecule implicated in long- term T cell immunity; activates NFkB through its interaction with adaptor proteins TRAF2 and TRAF5; suppresses apoptosis through |  |  |  |

|               |               |                       |                   | upregulation of BCL2                           |  |  |  |
|---------------|---------------|-----------------------|-------------------|--|--|--|--|
| Cytokines     |               |                       |                   |  |  |  |  |
| <u>ll1rl1</u> | 1.22          | Interleukin 1         |                   | Receptor for IL-33; recruits MyD88, IRAK1,     |  |  |  |
|               |               | receptor-like 1       |                   | IRAK4, and TRAF6; activates ERK1, ERK2, and    |  |  |  |
|               |               |                       |                   | MAPK14   |  |  |  |
| <u>ll18r1</u> | 1.09          | Interleukin 18        | CD218a            | Receptor for IL-18                             |  |  |  |
|               |               | receptor 1            |                   |  |  |  |  |
|               | Hematopoiesis |                       |                   |  |  |  |  |
| <u>lkzf2</u>  | 1.06          | IKAROS family zinc    |                   | Hematopoietic cell-specific TF involved in     |  |  |  |
|               |               | finger protein 2      |                   | early hematopoietic development                |  |  |  |
|               | Inflammation  |                       |                   |  |  |  |  |
| <u>Irf7</u>   | 0.513         | Interferon regulatory |                   | Key transcriptional regulator of type I IFN-   |  |  |  |
|               |               | factor 7              |                   | dependent immune responses; promotes           |  |  |  |
|               |               |                       |                   | transcription of IFN $\alpha$ and - $\beta$    |  |  |  |
| Jak2          | 0.473         | Janus kinase 2        |                   | Tyrosine kinase that participates in IFN and   |  |  |  |
|               |               |                       |                   | IL6ST signaling cascades                       |  |  |  |
| Nod2          | 0.978         | Nucleotide-binding    |                   | PRR specific for muramyl dipeptide (MDP);      |  |  |  |
|               |               | oligomerization       |                   | upon binding to its ligand, recruits RIPK2 and |  |  |  |
|               |               | domain containing 2   |                   | triggers MAPK and NFкВ signaling               |  |  |  |
| <u>Spn</u>    | 0.91          | Sialophorin           | Leukosialin; CD43 | Cell surface sialoglycoprotein expressed by T  |  |  |  |
|               |               |                       |                   | cells, B cells, monocytes, and granulocytes;   |  |  |  |

|            |       |  | Inhibition          | promotes lymph node localization in T cells; shunts T cells away from the $T_{H}2$ phenotype and towards $T_{H}1$ ; promotes the expression of IFN $\gamma$ in CD4 $^{+}$ T cells |
|------------|-------|--|---------------------|---|
| Ctla4      | 0.969 | Cytotoxic T  lymphocyte antigen  4           | CD152               | Inhibitory receptor that blocks CD28  costimulation by competitively binding its  ligands CD80 and CD86   |
|            |       |  | Interferon Response |   |
| Irf7       | 0.513 | Interferon regulatory factor 7               |                     | Key transcriptional regulator of type I IFN-dependent immune responses; promotes $transcription\ of\ IFN\alpha\ and\ -\beta$  |
| Jak2       | 0.473 | Janus kinase 2                               |                     | Tyrosine kinase that participates in IFN and  IL6ST signaling cascades  |
|            |       |  | Kinases             |   |
| <u>Itk</u> | 1.35  | Interleukin-2-<br>inducible T cell<br>kinase | LYK                 | Key actor in the TCR signaling cascade; phosphorylates PLCγ1, LAT, and LCP2   |
| Jak2       | 0.473 | Janus kinase 2                               |                     | Tyrosine kinase that participates in IFN and  IL6ST signaling cascades  |
| <u>Syk</u> | 0.978 | Spleen-associated                            |                     | Critical kinase that transmits signals from the   |

|        |                | tyrosine kinase                               |  | TCR and BCR  |  |  |  |
|--------|----------------|---|--|--|--|--|--|
|        | NFκB Signaling |   |  |  |  |  |  |
| Bcl10  | 0.371          | B cell lymphoma/leukemia                      |  | Activates NFκB via ubiquitination of ΙΚΚγ  |  |  |  |
| Card11 | 1.22           | Caspase recruitment  domain family  member 11 | Bcl10-interacting  MAGUK protein 3  (BIMP3)                | Adapter protein that plays a key role in adaptive immune response by transducing the activation of NFkB downstream of TCR and  |  |  |  |
|        |                |   |  | BCR engagement; upon activation in response to TCR or BCR triggering, homooligomerizes to form a nucleating helical template that recruits BCL10 via CARD-CARD interaction, thereby promoting polymerization of BCL10 and subsequent recruitment of MALT1, leading to IKK phosphorylation and degradation and release of NFkB proteins for |  |  |  |
|        |                |   |  | nuclear translocation  |  |  |  |
|        |                |   | NK Cell Function   |  |  |  |  |
| Prdm1  | 1.22           | Positive regulatory  domain I-binding  factor | B lymphocyte-<br>induced<br>maturation protein<br>(BLIMP1) | TF that plays a role in the development, retention, and long-term establishment of T cell, NK cell, and NK-T cells in non-lymphoid organs; drives the maturation of B cell into Ig   |  |  |  |
|        |                |   |  | secreting cells  |  |  |  |

| Tcf7         | 1.19  | Transcription factor 7 |                 | HMG box TF predominantly expressed by T cells that drives their development, although also involved in NK cell development; activates transcription through a Wnt/β- |
|--------------|-------|------------------------|-----------------|--|
|              |       |                        |                 | catenin signaling pathway  |
|              |       |                        | T Cell Function |  |
| Cd2          | 1.66  | Cluster of             | T cell surface  | Interacts with LFA-3 and CD48 to mediate   |
|              |       | differentiation 2      | antigen CD2     | adhesion between T cells and other cell types  |
| <u>Cd3e</u>  | 0.669 | Cluster of             |                 | Component of the TCR-CD3 complex; initiates  |
|              |       | differentiation 3      |                 | the TCR-CD3 complex assembly by forming  |
|              |       | epsilon                |                 | the two heterodimers CD3 $\delta$ /CD3 $\epsilon$ and  |
|              |       |                        |                 | CD3γ/CD3ε; also participates in internalization  |
|              |       |                        |                 | and cell surface down-regulation of TCR-CD3  |
|              |       |                        |                 | complexes via endocytosis sequences present  |
|              |       |                        |                 | in CD3ε cytosolic region   |
| <u>Cd4</u>   | 1.59  | Cluster of             |                 | Signature helper T cell marker; binds to MHC   |
|              |       | differentiation 4      |                 | class II and provides necessary costimulation  |
|              |       |                        |                 | for T cell activation  |
| <u>Cd8b1</u> | 0.856 | Cluster of             |                 | Beta chain of the CD8 coreceptor, which binds  |
|              |       | differentiation 8 beta |                 | to MHC class I   |
|              |       | 1                      |                 |  |
| <u>Cd28</u>  | 0.848 | Cluster of             |                 | Essential T cell co-receptor that enhances T   |
|              |       |                        |                 | cell activation, proliferation, cytokine   |

|               |      | differentiation 28                            |  | production, and survival; binds to CD80 and CD86  |
|---------------|------|---|--|---|
| Gata3         | 1.08 | GATA binding protein 3                        |  | Transcriptional activator that binds to the enhancer of the TCR $\alpha$ and $\delta$ genes; required for TH2 differentiation following immune and inflammatory responses   |
| <u>Gpr183</u> | 1.14 | G protein-coupled receptor 183                | EBV-induced G protein-coupled receptor 2                   | Lymphocyte GPCR that acts as a chemotactic receptor for B cells, T cells, splenic DCs, monocytes/macrophages, and astrocytes  |
| <u>Itk</u>    | 1.35 | Interleukin-2-<br>inducible T cell<br>kinase  | LYK  | Key actor in the TCR signaling cascade; phosphorylates PLCγ1, LAT, and LCP2   |
| Prdm1         | 1.22 | Positive regulatory  domain I-binding  factor | B lymphocyte-<br>induced<br>maturation protein<br>(BLIMP1) | TF that plays a role in the development, retention, and long-term establishment of T cell, NK cell, and NK-T cells in non-lymphoid organs; drives the maturation of B cell into Ig secreting cells  |
| Spn           | 0.91 | Sialophorin                                   | Leukosialin; CD43  | Cell surface sialoglycoprotein expressed by T cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNγ in CD4 <sup>+</sup> T cells |

| <u>Syk</u>   | 0.978 | Spleen-associated     |                     | Critical kinase that transmits signals from the           |
|--------------|-------|-----------------------|---------------------|---|
|              |       | tyrosine kinase       |                     | TCR and BCR   |
| Tcf7         | 1.19  | Transcription factor  |                     | HMG box TF predominantly expressed by T                   |
|              |       | 7                     |                     | cells that drives their development, although             |
|              |       |                       |                     | also involved in NK cell development;                     |
|              |       |                       |                     | activates transcription through a Wnt/ $\beta$ -          |
|              |       |                       |                     | catenin signaling pathway                                 |
|              |       | Transcri              | ption Factors & Coa | ctivators   |
|              |       |                       |                     |   |
| <u>Gata3</u> | 1.08  | GATA binding protein  |                     | Transcriptional activator that binds to the               |
|              |       | 3                     |                     | enhancer of the TCR $\alpha$ and $\delta$ genes; required |
|              |       |                       |                     | for T <sub>H</sub> 2 differentiation following immune and |
|              |       |                       |                     | inflammatory responses                                    |
| Ikzf2        | 1.06  | IKAROS family zinc    |                     | Hematopoietic cell-specific TF involved in                |
|              |       | finger protein 2      |                     | early hematopoietic development                           |
| <u>Irf7</u>  | 0.513 | Interferon regulatory |                     | Key transcriptional regulator of type I IFN-              |
|              |       | factor 7              |                     | dependent immune responses; promotes                      |
|              |       |                       |                     | transcription of IFN $\alpha$ and - $\beta$               |
| Prdm1        | 1.22  | Positive regulatory   | B lymphocyte-       | TF that plays a role in the development,                  |
|              |       | domain I-binding      | induced             | retention, and long-term establishment of T               |
|              |       | factor                | maturation protein  | cell, NK cell, and NK-T cells in non-lymphoid             |
|              |       |                       | (BLIMP1)            | organs; drives the maturation of B cell into Ig           |
|              |       |                       |                     | secreting cells   |
|              |       |                       |                     |   |

| Stat6      | 0.392 | Signal transducer    | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and |
|------------|-------|----------------------|---|
|            |       | and activator of     | macrophage function and M2 macrophage                         |
|            |       | transcription 6      | polarization  |
| Tcf7       | 1.19  | Transcription factor | HMG box TF predominantly expressed by T                       |
|            |       | 7                    | cells that drives their development, although                 |
|            |       |                      | also involved in NK cell development;                         |
|            |       |                      | activates transcription through a Wnt/β-                      |
|            |       |                      | catenin signaling pathway                                     |
|            |       |                      | Jbiquitin Regulation  |
|            |       |                      |   |
| <u>Ubc</u> | 0.39  | Polyubiquitin C      | Serves various roles, including innate                        |
|            |       |                      | immunity, DNA repair, and stimulation of                      |
|            |       |                      | autophagy and the proteasomal response                        |

## Supplemental Table 2: Genes Differentially Upregulated in Secondary (Unirradiated) Tumor

|      |                  |                                | PBT+αPD1 vs Ctrl    |   |
|------|------------------|--------------------------------|---------------------|---|
| Gene | Log2 fold change | Full Name                      | Notable Aliases     | Function  |
|      |                  | A                              | cute Phase Response |   |
| Арр  | 0.196            | Amyloid-beta precursor protein |                     | Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; |

| <u>Psen1</u> | 0.258 | Presenilin-1                            |                                   | involved in cell mobility, copper homeostasis, and oxidative stress  Catalytic subunit of the gamma-secretase complex, which cleaves integral membrane proteins such as APP; also involved in Ca(2+) |
|--------------|-------|---|-----------------------------------|--|
|              |       | Adhasi                                  | on & Cell-Cell Interact           | homeostasis  |
|              |       | Auriesi                                 | on & cen-cen interact             | uons   |
| <u>Cd47</u>  | 0.269 | Cluster of differentiation 47           | Integrin-associated protein (IAP) | Partners with membrane integrins to serve as an inhibitor of phagocytosis  |
|              |       |   |                                   |  |
| <u>Itgax</u> | 0.521 | Integrin alpha X                        | CD11c                             | Adhesion molecule; signature marker of Agpresenting DCs  |
| <u>lcam1</u> | 0.473 | Intracellular adhesion molecule 1       | CD54                              | Cell surface glycoprotein that serves as strong adhesive ligand for LFA-1; important for leukocyte mobility and costimulation  |
| Vcam1        | 0.49  | Vascular cell adhesion molecule 1       | CD106                             | Endothelial-cell adhesion molecule that binds to ITGA4/ITGB1 on leukocytes and mediates both adhesion and signal transduction  |
|              |       | Antigen                                 | Processing & Present              | tation   |
| H2-D1        | 0.461 | Histocompatibility 2,  D region locus 1 |                                   | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T cells   |

| Psmb10      | 0.427 | Proteasome subunit   |                      | 20S core β subunit of the proteasome          |
|-------------|-------|----------------------|----------------------|---|
|             |       | beta type 10         |                      | involved in Ag processing to generate class I |
|             |       |                      |                      | binding peptides                              |
|             |       |                      |                      |   |
| <u>Tap1</u> | 0.5   | Transporter antigen  | Really interesting   | ATP-binding cassette transporter that         |
|             |       | peptide 1            | new gene 4 (RING4)   | pumps degraded cytosolic peptides from        |
|             |       |                      |                      | the cytosol to the ER for packaging into      |
|             |       |                      |                      | MHC class I molecules                         |
|             |       |                      | A                    |   |
|             |       |                      | Autophagy            |   |
| Irgm2       | 0.747 | Immunity-related     | Interferon-inducible | Function not fully known, but most likely     |
|             |       | GTPase family M      | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory      |
|             |       | member 2             |                      | cytokine production                           |
|             |       |                      |                      |   |
| Ubc         | 0.289 | Polyubiquitin C      |                      | Serves various roles, including innate        |
|             |       |                      |                      | immunity, DNA repair, and stimulation of      |
|             |       |                      |                      | autophagy and the proteasomal response        |
|             |       | Comple               | ment & Humoral Imm   | nunity  |
|             |       | ·                    |                      | ·   |
| Cfb         | 0.567 |                      |                      | Alternate complement pathway                  |
|             |       |                      |                      | component; when cleaved, produces a           |
|             |       |                      |                      | serine protease that binds to C3b to form     |
|             |       | Complement factor B  |                      | C3 convertase                                 |
| Fac::A      | 0.000 | Fun account.         | Fa wasantan lilia 2  | Dutating manage outled to the bound of 5000   |
| Fcgr4       | 0.898 | Fragment             | Fc receptor-like 3   | Putative mouse ortholog to human FcγRIIIA     |
|             |       | crystallizable gamma | (Fcrl3); CD16-2      |   |
|             |       | receptor 4           |                      |   |
|             |       |                      |                      |   |

| Icam1 | 0.473 | Intracellular       | CD54                 | Cell surface glycoprotein that serves as     |
|-------|-------|---------------------|----------------------|--|
|       |       | adhesion molecule 1 |                      | strong adhesive ligand for LFA-1; important  |
|       |       |                     |                      | for leukocyte mobility and costimulation     |
|       |       |                     |                      |  |
|       |       |                     | Costimulation        |  |
| Cd28  | 1.16  | Cluster of          |                      | Essential T cell co-receptor that enhances T |
|       |       | differentiation 28  |                      | cell activation, proliferation, cytokine     |
|       |       |                     |                      | production, and survival; binds to CD80 and  |
|       |       |                     |                      | CD86   |
| Icos  | 0.608 | Inducible T cell    | CD278                | Enhances all basic T cell responses to       |
|       |       | costimulator        |                      | foreign Ag; essential both for efficient     |
|       |       |                     |                      | interaction between T and B cells and for    |
|       |       |                     |                      | normal Ab responses to T cell-dependent      |
|       |       |                     |                      | Ags  |
|       |       |                     | Chemotaxis           |  |
|       |       |                     |                      |  |
| Ccl5  | 1.11  | C-C motif chemokine | Regulated upon       | Chemoattractant ligand for CCR1, -3, -4, and |
|       |       | ligand 5            | activation, normally | -5; attracts blood monocytes, memory T       |
|       |       |                     | T-expressed, and     | helper cells and eosinophils; causes the     |
|       |       |                     | presumably           | release of histamine from basophils and      |
|       |       |                     | secreted (RANTES)    | activates eosinophils                        |
|       | 0.523 | C-C motif chemokine | MIP1α receptor       | Receptor for CCL3, -5, -7, and -23           |
| Ccr1  |       | receptor 1          |                      |  |

| CIO    | 0.992 | C-X-C motif           | Humig                | Chemoattractant ligand for CXCR3; attracts    |
|--------|-------|-----------------------|----------------------|---|
| Cxcl9  |       | chemokine ligand 9    |                      | activated T cells                             |
|        | 0.852 | C-X-C motif           |                      | Dominant ligand for CXCR3; attracts           |
| Cxcl11 |       | chemokine ligand 11   |                      | activated T cells; strongly induced by IFNγ   |
|        | 0.817 | C-X-C motif           | CD186                | Receptor for the C-X-C chemokine CXCL16;      |
|        |       | chemokine receptor    |                      | expressed in several T lymphocyte subsets     |
| Cxcr6  |       | 6                     |                      | and bone marrow stromal cells                 |
|        |       | G                     | irowth/Proliferation |   |
|        |       |                       | ,                    |   |
| Map2k1 | 0.297 | Dual specificity      | MAPK/ERK kinase 1    | Essential component of the MAP kinase         |
|        |       | mitogen-activated     | (MEK1)               | signal transduction pathway; participates in  |
|        |       | protein kinase kinase |                      | numerous biological functions, including cell |
|        |       | 1                     |                      | growth, adhesion, survival, differentiation,  |
|        |       |                       |                      | transcription, metabolism, and cytoskeletal   |
|        |       |                       |                      | remodeling                                    |
| Stat1  | 0.641 | Signal transducer and |                      | Transcriptional activator that mediates       |
|        |       | activator of          |                      | cellular responses to IFNs, cytokines, and    |
|        |       | transcription 1       |                      | other growth factors                          |
|        | 0.397 | Vascular endothelial  |                      | Glysosylated mitogen that promotes            |
|        |       | growth factor A       |                      | vascular permeability, vasculogenesis,        |
| Vegfa  |       |                       |                      | angiogenesis, and cell migration              |
| Yy1    | 0.197 | Yin yang 1            |                      | Ubiquitous factor that serves as a            |
|        |       |                       |                      | transcriptional "switch", either promoting    |

|        |       | I                     |                      |  |
|--------|-------|-----------------------|----------------------|--|
|        |       |                       |                      | or repressing the transcription of numerous    |
|        |       |                       |                      | genes through the selective recruitment of     |
|        |       |                       |                      | either histone deacetylases or                 |
|        |       |                       |                      | acetyltransferases; plays a fundamental role   |
|        |       |                       |                      | in diverse processes, such as differentiation, |
|        |       |                       |                      | replication, and cellular proliferation        |
|        |       |                       | Inflammation         |  |
|        |       |                       |                      |  |
| Cebpb  | 0.312 | CCAAT/enhancer-       |                      | Critical macrophage TF that promotes           |
|        |       | binding protein beta  |                      | expression of several acute-phase and          |
|        |       |                       |                      | inflammatory cytokine genes, including II6     |
| Irf1   | 0.471 | Interferon regulatory |                      | Transcriptional regulator that promotes        |
|        |       | factor 1              |                      | inflammatory innate and adaptive immune        |
|        |       |                       |                      | responses                                      |
|        |       |                       |                      |  |
| Irgm2  | 0.747 | Immunity-related      | Interferon-inducible | Function not fully known, but most likely      |
|        |       | GTPase family M       | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory       |
|        |       | member 2              |                      | cytokine production                            |
| lsg20  | 0.446 | Interferon-stimulated |                      | IFN-induced antiviral exoribonuclease that     |
|        |       | gene 20               |                      | acts on ssRNA with minor activity towards      |
|        |       |                       |                      | ssDNA  |
| Lgals3 | 0.415 | Galectin 3            |                      | Galactose-specific lectin that binds IgE;      |
| LEGISS | 0.413 | Galecting             |                      | -  |
|        |       |                       |                      | involved in acute inflammatory responses,      |
|        |       |                       |                      | including neutrophil activation and            |
|        |       |                       |                      | adhesion, chemoattraction of                   |
|        |       |                       |                      |  |

|        |       |                        |                       | monocytes/macrophages, opsonization of      |
|--------|-------|------------------------|-----------------------|---|
|        |       |                        |                       | apoptotic neutrophils, and activation of    |
|        |       |                        |                       | mast cells                                  |
| Nos2   | 1.19  | Inducible nitric oxide |                       | Produces reactive oxygen species and        |
|        |       | synthase (iNOS)        |                       | contributes to inflammatory cytokine        |
|        |       |                        |                       | production                                  |
|        |       |                        | Inhibition            |   |
| Cd47   | 0.269 | Cluster of             | Integrin-associated   | Partners with membrane integrins to serve   |
|        |       | differentiation 47     | protein (IAP)         | as an inhibitor of phagocytosis             |
| Cd274  | 0.723 | Cluster of             | Programmed cell       | Ubiquitously expressed ligand for co-       |
|        |       | differentiation 274    | death receptor        | inhibitory receptor PD-1; upregulated by    |
|        |       |                        | ligand 1 (PD-L1)      | tumors as an immune evasion strategy        |
| Cdkn1a | 0.381 | Cyclin dependent       | p21; CDK-             | Binds to and inhibits cyclin-dependent      |
|        |       | kinase inhibitor 1A    | interaction protein 1 | kinase activity, preventing phosphorylation |
|        |       |                        | (CIP1)                | of critical cyclin-dependent kinase         |
|        |       |                        |                       | substrates and blocking cell cycle          |
|        |       |                        |                       | progression                                 |
| Ctla4  | 0.66  | Cytotoxic T            | CD152                 | Inhibitory receptor that blocks CD28        |
|        |       | lymphocyte antigen 4   |                       | costimulation by competitively binding its  |
|        |       |                        |                       | ligands CD80 and CD86                       |
| Dusp4  | 0.318 | Dual specificity       |                       | Inactivates ERK1, ERK2, and JNK             |
|        |       | phosphatase 4          |                       |   |
|        |       |                        |                       |   |

| Foxp3 | 0.496               | Forkhead box P3                                | DIETER | Master TF for Tregs; represses expression of II2 and Ifng; activates expression of Tnfrsf18, II2ra, and Ctla4  |  |
|-------|---------------------|--|--------|--|--|
| ldo1  | 1.19                | Indoleamine 2,3-<br>dioxygenase 1              |        | Initiates catabolism of tryptophan; limits immunopathology by inhibiting T cell division   |  |
| Lag3  | 0.527               | Lymphocyte activating gene 3                   | CD223  | Inhibitory receptor on activated T cells; binds to ligands, such as FGL1; constitutively expressed on a subset of regulatory Tregs and contributes to their suppressive function; acts as a negative regulator of plasmacytoid DC activation |  |
| Socs1 | 0.826               | Suppressor of cytokine signaling 1             |        | Inhibits JAK proteins; negative regulator of IL-6  |  |
| Tigit | 0.5                 | T cell immunoreceptor with Ig and ITIM domains |        | Binds with high affinity to the poliovirus receptor, causing increased secretion of IL- 10, decreased secretion of IL-12B, and suppressing T cell activation by promoting the generation of mature immunoregulatory DCs                      |  |
|       | Interferon Response |  |        |  |  |

| Irf1  | 0.471 | Interferon regulatory |                      | Transcriptional regulator that promotes    |
|-------|-------|-----------------------|----------------------|--|
|       |       | factor 1              |                      | inflammatory innate and adaptive immune    |
|       |       |                       |                      | responses                                  |
|       |       |                       |                      |  |
| Irgm2 | 0.747 | Immunity-related      | Interferon-inducible | Function not fully known, but most likely  |
|       |       | GTPase family M       | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory   |
|       |       | member 2              |                      | cytokine production                        |
| lsg20 | 0.446 | Interferon-stimulated |                      | IFN-induced antiviral exoribonuclease that |
|       |       | gene 20               |                      | acts on ssRNA with minor activity towards  |
|       |       |                       |                      | ssDNA                                      |
| Mx2   | 0.602 | Myxovirus resistance  |                      | IFN-induced dynamin-like GTPase with       |
|       |       | protein 2             |                      | potent antiviral activity against HIV-1    |
| Stat1 | 0.641 | Signal transducer and |                      | Transcriptional activator that mediates    |
|       |       | activator of          |                      | cellular responses to IFNs, cytokines, and |
|       |       | transcription 1       |                      | other growth factors                       |
|       |       | N                     | Nacrophage Function  |  |
| Cd14  | 0.264 | Cluster of            |                      | PRR that recognizes LPS; mostly found on   |
|       |       | differentiation 14    |                      | macrophages                                |
| Cebpb | 0.312 | CCAAT/enhancer-       |                      | Critical macrophage TF that promotes       |
|       |       | binding protein beta  |                      | expression of several acute-phase and      |
|       |       |                       |                      | inflammatory cytokine genes, including Il6 |

| Slamf7 | 0.615 | Signaling lymphocytic |                      | A super-activator of macrophages and a       |
|--------|-------|-----------------------|----------------------|--|
|        |       | activation molecule   |                      | strong promoter of phagocytosis; binds to    |
|        |       | family member 7       |                      | CD74   |
|        |       |                       |                      |  |
|        |       |                       | Migration/Motility   |  |
|        |       |                       |                      |  |
| App    | 0.196 | Amyloid-beta          |                      | Cell surface receptor and transmembrane      |
|        |       | precursor protein     |                      | precursor protein that is cleaved by         |
|        |       |                       |                      | secretases to form a number of peptides;     |
|        |       |                       |                      | involved in cell mobility, copper            |
|        |       |                       |                      | homeostasis, and oxidative stress            |
|        |       |                       |                      |  |
| Ccl5   | 1.11  | C-C motif chemokine   | Regulated upon       | Chemoattractant ligand for CCR1, -3, -4, and |
|        |       | ligand 5              | activation, normally | -5; attracts blood monocytes, memory T       |
|        |       |                       | T-expressed, and     | helper cells and eosinophils; causes the     |
|        |       |                       | presumably           | release of histamine from basophils and      |
|        |       |                       | secreted (RANTES))   | activates eosinophils                        |
|        |       |                       | (,                   | 333.333                                      |
| Cxcl9  | 0.992 | C-X-C motif           | Humig                | Chemoattractant ligand for CXCR3; attracts   |
|        |       | chemokine ligand 9    |                      | activated T cells                            |
|        |       |                       |                      |  |
| Cxcl11 | 0.852 | C-X-C motif           |                      | Dominant ligand for CXCR3; attracts          |
|        |       | chemokine ligand 11   |                      | activated T cells; strongly induced by IFNγ  |
|        |       |                       |                      |  |
| Cxcr6  | 0.817 | C-X-C motif           | CD186                | Receptor for the C-X-C chemokine CXCL16;     |
|        |       | chemokine receptor    |                      | expressed in several T lymphocyte subsets    |
|        |       | 6                     |                      | and bone marrow stromal cells                |
|        |       |                       |                      | 22.2   |
|        |       |                       |                      |  |

| Lgals3 | 0.415 | Galectin 3           |                  | Galactose-specific lectin that binds IgE;       |
|--------|-------|----------------------|------------------|---|
|        |       |                      |                  | involved in acute inflammatory responses,       |
|        |       |                      |                  | including neutrophil activation and             |
|        |       |                      |                  | adhesion, chemoattraction of                    |
|        |       |                      |                  | monocytes/macrophages, opsonization of          |
|        |       |                      |                  | apoptotic neutrophils, and activation of        |
|        |       |                      |                  | mast cells                                      |
|        | 0.397 | Vascular endothelial |                  | Glysosylated mitogen that promotes              |
|        |       | growth factor A      |                  | vascular permeability, vasculogenesis,          |
| Vegfa  |       |                      |                  | angiogenesis, and cell migration                |
|        |       |                      | NK Cell Function |   |
|        |       |                      |                  |   |
| Cd8a   | 0.766 | Cluster of           | LEU2             | Alpha chain of the CD8 coreceptor, which        |
|        |       | differentiation 8    |                  | binds to MHC class I; in T cells, functions     |
|        |       | alpha                |                  | primarily as a coreceptor for MHC class I       |
|        |       |                      |                  | molecule:peptide complex; in NK cells, the      |
|        |       |                      |                  | presence of CD8 $\alpha$ homodimers at the cell |
|        |       |                      |                  | surface provides a survival mechanism           |
|        |       |                      |                  | allowing conjugation and lysis of multiple      |
|        |       |                      |                  | target cells                                    |
| Gzmb   | 0.895 | Granzyme B           | Fragmentin 2     | Abundant protease in the cytosolic granules     |
|        |       |                      |                  | of cytotoxic T and NK cells that activates      |
|        |       |                      |                  | caspase-mediated cell death when                |
|        |       |                      |                  |   |

|       |       |                        |                      | delivered into the target cell through the immunological synapse |
|-------|-------|------------------------|----------------------|--|
| Gzmk  | 1.66  | Granzyme K             | Tryptase II          | Granule-secreted, pro-apoptotic serine                           |
|       |       |                        |                      | protease found in the cytoplasmic granules                       |
|       |       |                        |                      | of CTLs  |
| Prdm1 | 0.82  | Positive regulatory    | B lymphocyte-        | TF that plays a role in the development,                         |
|       |       | domain I-binding       | induced maturation   | retention, and long-term establishment of T                      |
|       |       | factor                 | protein (BLIMP1)     | cell, NK cell, and NK-T cells in non-lymphoid                    |
|       |       |                        |                      | organs; drives the maturation of B cell into                     |
|       |       |                        |                      | lg secreting cells   |
|       |       | ROS                    | Generation & Protect | ion  |
| Cybb  | 0.418 | Cytochrome b-245       | Nox2                 | Part of the NADPH oxidase process;                               |
|       |       | heavy chain            |                      | generates superoxides  |
| Nos2  | 1.19  | Inducible nitric oxide |                      | Produces reactive oxygen species and                             |
|       |       | synthase (iNOS)        |                      | contributes to inflammatory cytokine                             |
|       |       |                        |                      | production   |
|       |       |                        | T Cell Function      |  |
| Cd3d  | 0.606 | Cluster of             |                      | Component of the TCR-CD3 complex; upon                           |
|       |       | differentiation 3      |                      | phosphorylation by Lck, serves as a docking                      |
|       |       | delta                  |                      | station for downstream TCR signaling                             |
|       |       |                        |                      | adaptors   |
|       |       |                        |                      |  |

| Cd8a | 0.766 | Cluster of differentiation 8 alpha | LEU2         | Alpha chain of the CD8 coreceptor, which binds to MHC class I; in T cells, functions primarily as a coreceptor for MHC class I molecule:peptide complex; in NK cells, the presence of CD8α homodimers at the cell surface provides a survival mechanism allowing conjugation and lysis of multiple target cells |
|------|-------|------------------------------------|--------------|---|
| Cd28 | 1.16  | Cluster of differentiation 28      |              | cell activation, proliferation, cytokine production, and survival; binds to CD80 and CD86   |
| Gzmb | 0.895 | Granzyme B                         | Fragmentin 2 | Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspase-mediated cell death when delivered into the target cell through the immunological synapse  |
| Gzmk | 1.66  | Granzyme K                         | Tryptase II  | Granule-secreted, pro-apoptotic serine protease found in the cytoplasmic granules of CTLs   |
| Icos | 0.608 | Inducible T cell costimulator      | CD278        | Enhances all basic T cell responses to foreign Ag; essential both for efficient interaction between T and B cells and for   |

| Prdm1 | 0.82  | Positive regulatory domain I-binding factor | B lymphocyte-<br>induced maturation<br>protein (BLIMP1) | normal Ab responses to T cell-dependent  Ags  TF that plays a role in the development, retention, and long-term establishment of T  cell, NK cell, and NK-T cells in non-lymphoid organs; drives the maturation of B cell into  Ig secreting cells |
|-------|-------|---|---|--|
|       |       | Transcrip                                   | otion Factors & Coacti                                  | vators   |
|       |       |   |   |  |
| Cebpb | 0.312 | CCAAT/enhancer-                             |   | Critical macrophage TF that promotes   |
|       |       | binding protein beta                        |   | expression of several acute-phase and  |
|       |       |   |   | inflammatory cytokine genes, including II6   |
| Irf1  | 0.471 | Interferon regulatory                       |   | Transcriptional regulator that promotes  |
|       |       | factor 1                                    |   | inflammatory innate and adaptive immune  |
|       |       |   |   | responses  |
| Prdm1 | 0.82  | Positive regulatory                         | B lymphocyte-   | TF that plays a role in the development,   |
|       |       | domain I-binding                            | induced maturation                                      | retention, and long-term establishment of T  |
|       |       | factor                                      | protein (BLIMP1)  | cell, NK cell, and NK-T cells in non-lymphoid  |
|       |       |   |   | organs; drives the maturation of B cell into   |
|       |       |   |   | Ig secreting cells   |
| Stat1 | 0.641 | Signal transducer and                       |   | Transcriptional activator that mediates  |
|       |       | activator of                                |   | cellular responses to IFNs, cytokines, and   |
|       |       | transcription 1                             |   | other growth factors   |

| Yy1 | 0.197 | Yin yang 1 | Ubiquitous factor that serves as a             |
|-----|-------|------------|--|
|     |       |            | transcriptional "switch", either promoting     |
|     |       |            | or repressing the transcription of numerous    |
|     |       |            | genes through the selective recruitment of     |
|     |       |            | either histone deacetylases or                 |
|     |       |            | acetyltransferases; plays a fundamental role   |
|     |       |            | in diverse processes, such as differentiation, |
|     |       |            | replication, and cellular proliferation        |
|     |       |            |  |

|       |                  | NBT                            | XR3+PBT+αPD1 vs Ct  | ri  |
|-------|------------------|--------------------------------|---------------------|---|
| Gene  | Log2 fold change | Full Name                      | Notable Aliases     | Function  |
|       |                  | Ac                             | cute Phase Response |   |
| Арр   | 0.157            | Amyloid-beta precursor protein |                     | Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; involved in cell mobility, copper homeostasis, and oxidative stress |
| Psen2 | 0.337            | Presenilin-2                   |                     | Putative catalytic subunit of the gamma-<br>secretase complex, which cleaves integral<br>membrane proteins such as APP; also  |

|        |       |                       |                        | Ţ  |
|--------|-------|-----------------------|------------------------|--|
|        |       |                       |                        | involved in Ca(2+) homeostasis between the     |
|        |       |                       |                        | ER and the mitochondria                        |
|        |       | Adhesio               | on & Cell-Cell Interac | tions  |
|        |       |                       |                        |  |
| Cd37   | 1.12  | Cluster of            | Tetraspanin-26         | Cell surface glycoprotein known to complex     |
|        |       | differentiation 37    |                        | with integrins and other transmembrane 4       |
|        |       |                       |                        | superfamily proteins; may play a role in T     |
|        |       |                       |                        | cell-B cell interactions                       |
| Cd97   | 0.29  | Cluster of            | BL-Ac[F2]              | GPCR that promotes granulocyte adhesion        |
|        |       | differentiation 97    |                        | and migration; activates T cells via binding   |
|        |       |                       |                        | to CD55; stimulates angiogenesis through       |
|        |       |                       |                        | binding integrin counterreceptors on           |
|        |       |                       |                        | endothelial cells                              |
| Cyfip2 | 0.877 | Cytoplasmic FMR1-     |                        | Involved in T cell adhesion and p53/TP53-      |
|        |       | interacting protein 2 |                        | dependent induction of apoptosis               |
| Dpp4   | 0.388 | Dipeptidyl-peptidase  | CD26                   | Cell surface glycoprotein receptor involved    |
|        |       | 4                     |                        | in the costimulatory signal essential for TCR- |
|        |       |                       |                        | mediated T cell activation; serine             |
|        |       |                       |                        | exopeptidase that cleaves various              |
|        |       |                       |                        | substrates, thereby inactivating them;         |
|        |       |                       |                        | involved in the migration and invasion of      |
|        |       |                       |                        | endothelial cells into the ECM; also           |
|        |       |                       |                        | regulates lymphocyte-epithelial cell           |
|        |       |                       |                        | adhesion                                       |
|        |       |                       |                        |  |

| Itgam  | 0.681 | Integrin alpha M      | CD11b             | Pairs with CD18 to form Mac-1 aka             |
|--------|-------|-----------------------|-------------------|---|
|        |       |                       |                   | complement receptor 3; mediates leukocyte     |
|        |       |                       |                   | activation, adhesion, chemotaxis, migration,  |
|        |       |                       |                   | phagocytosis, and cell-mediated               |
|        |       |                       |                   | cytotoxicity; serves as a macrophage marker   |
| Itgax  | 0.617 | Integrin alpha X      | CD11c             | Adhesion molecule; signature marker of Ag-    |
|        |       |                       |                   | presenting DCs                                |
| Itgb2  | 0.735 | Integrin subunit beta |                   | Pairs with ITGAL to form a receptor for       |
|        |       | 2                     |                   | ICAM1, with ITGAM or ITGAX for iC3b and       |
|        |       |                       |                   | fibronectin                                   |
| Lgals3 | 0.299 | Galectin 3            |                   | Galactose-specific lectin that binds IgE;     |
|        |       |                       |                   | involved in acute inflammatory responses,     |
|        |       |                       |                   | including neutrophil activation and           |
|        |       |                       |                   | adhesion, chemoattraction of monocytes        |
|        |       |                       |                   | macrophages, opsonization of apoptotic        |
|        |       |                       |                   | neutrophils, and activation of mast cells     |
| Map2k1 | 0.181 | Dual specificity      | MAPK/ERK kinase 1 | Essential component of the MAP kinase         |
|        |       | mitogen-activated     | (MEK1)            | signal transduction pathway; participates in  |
|        |       | protein kinase kinase |                   | numerous biological functions, including cell |
|        |       | 1                     |                   | growth, adhesion, survival, differentiation,  |
|        |       |                       |                   | transcription, metabolism, and cytoskeletal   |
|        |       |                       |                   | remodeling                                    |
|        |       |                       |                   |   |

| Sell   | 1.15  | L-selectin                                 | Mediates cell adhesion by binding to glycoproteins on neighboring cells   |
|--------|-------|--|---|
|        |       | Antigen                                    | Processing & Presentation   |
| Cd1d1  | 0.628 | Cluster of differentiation 1 D1            | Murine non-classical class I MHC; primarily presents lipid and glycolipid Ags                                   |
| Ctss   | 0.736 | Cathepsin S                                | Lysosomal protease that participates in processing of Ag by MHC class II  |
| Cyfip2 | 0.877 | Cytoplasmic FMR1-<br>interacting protein 2 | Involved in T cell adhesion and p53/TP53-<br>dependent induction of apoptosis                                   |
| H2-D1  | 0.353 | Histocompatibility 2,  D region locus 1    | MHC class I molecule; presents Ags to CD8 <sup>+</sup> T cells  |
| H2-M3  | 0.603 | Histocompatibility 2,  M region locus 3    | MHC class Ib molecule; presents Ags to  CD8 <sup>+</sup> T cells, with a preference for N-  formylated peptides |
| H2-T23 | 0.67  | Histocompatibility 2, Q region locus 10    | MHC class I molecule; presents Ags to CD8+  T cells   |
| Psmb10 | 0.446 | Proteasome subunit<br>beta type 10         | 20S core β subunit of the proteasome involved in Ag processing to generate class I binding peptides             |
|        |       |  | Anti-Inflammatory   |

| Bcl2      | 0.418 | B cell lymphoma 2                              |   | Outer mitochondrial membrane protein that inhibits apoptosis and autophagy; may attenuate inflammation by impairing inflammasome formation   |  |  |  |
|-----------|-------|--|---|--|--|--|--|
| Cd200r1   | 0.436 | Cluster of differentiation 200 receptor 1      |   | Inhibitory receptor for the CD200/OX2 cell surface glycoprotein; limits inflammation by inhibiting the expression of proinflammatory molecules including TNF $\alpha$ , IFNs, and iNOS |  |  |  |
| Ctla4     | 0.786 | Cytotoxic T lymphocyte antigen 4               | CD152                                       | Inhibitory receptor that blocks CD28 costimulation by competitively binding its ligands CD80 and CD86  |  |  |  |
| Tnfaip3   | 0.387 | Tumor necrosis factor, alpha-induced protein 3 |   | Ubiquitin-editing enzyme that complexes with ITCH to degrade inflammatory signaling components in the TNF, IL1, and TLR pathways; targets TRAF2, TRAF6, and IKK                        |  |  |  |
| Apoptosis |       |  |   |  |  |  |  |
| Bid       | 0.285 | BH3 interacting domain death agonist           | Desmocollin type 4, apoptotic death agonist | Induces caspases and apoptosis; counters the protective effect of BCL2, allowing release of cytochrome C   |  |  |  |

| Casp1                   | 0.417 | Caspase 1             | Interleukin 1β       | Cysteine-aspartic acid protease that               |  |  |  |
|-------------------------|-------|-----------------------|----------------------|--|--|--|--|
|                         |       |                       | convertase           | mediates cleavage-based activation of IL-1 $\beta$ |  |  |  |
|                         |       |                       |                      | and IL-18; serves as the central enzymatic         |  |  |  |
|                         |       |                       |                      | core of the inflammasome; also induces             |  |  |  |
|                         |       |                       |                      | apoptosis  |  |  |  |
| Cyfip2                  | 0.877 | Cytoplasmic FMR1-     |                      | Involved in T cell adhesion and p53/TP53-          |  |  |  |
|                         |       | interacting protein 2 |                      | dependent induction of apoptosis                   |  |  |  |
| Fas                     | 0.901 | Fragment apoptosis    |                      | Cell surface death receptor; interaction with      |  |  |  |
|                         |       | stimulating           |                      | FAS-ligand triggers an apoptotic signaling         |  |  |  |
|                         |       |                       |                      | cascade; also activates NFkB, ERK1, and            |  |  |  |
|                         |       |                       |                      | MAPK8  |  |  |  |
| Trp53                   | 0.403 | Transformation-       |                      | TF that induces cell-cycle arrest and              |  |  |  |
|                         |       | related protein 53    |                      | apoptosis through stimulation of Fas               |  |  |  |
|                         |       |                       |                      | expression   |  |  |  |
| B Cell-associated Genes |       |                       |                      |  |  |  |  |
|                         |       |                       |                      |  |  |  |  |
| Blnk                    | 1.18  | B cell linker         | Src homology 1       | Functions as a central linker protein              |  |  |  |
|                         |       |                       | domain-containing    | downstream of the B cell receptor, bridging        |  |  |  |
|                         |       |                       | leukocyte protein of | SYK kinase to a multitude of signaling             |  |  |  |
|                         |       |                       | 65 kDa (SLP-65);     | pathways and regulating biological                 |  |  |  |
|                         |       |                       | Ly57                 | outcomes of B cell function and                    |  |  |  |
|                         |       |                       |                      | development; plays a role in the activation        |  |  |  |
|                         |       |                       |                      | of ERK/EPHB2, MAP kinase p38 and JNK;              |  |  |  |

|      |       |                               |   | modulates AP1 activation; important for the activation of NFκB and NFAT   |
|------|-------|-------------------------------|---|---|
| Btk  | 0.615 | Bruton's tyrosine<br>kinase   |   | Crucial kinase in B cell receptor signal transmission and B cell activation   |
| Btla | 1.73  | B and T lymphocyte attenuator | CD272   | Inhibitory cell surface protein that inhibits T cell function by binding to B7H4 and TNFRSF14   |
| Cd19 | 2.29  | Cluster of differentiation 19 | B-lymphocyte<br>surface antigen B4  | BCR coreceptor; activates signaling pathways that lead to the activation of PI3K and Ca(2+) flux  |
| Cd22 | 2.08  | Cluster of differentiation 22 | Sialic acid-binding  Ig-like lectin 2  (SIGLEC2)  B lymphocyte cell adhesion molecule  (BL-CAM) | Mediates interactions between B cells; binds CD45   |
| Cd37 | 1.12  | Cluster of differentiation 37 | Tetraspanin-26  | Cell surface glycoprotein known to complex with integrins and other transmembrane 4 superfamily proteins; may play a role in T cell-B cell interactions |
| Cd48 | 0.639 | Cluster of differentiation 48 | B-lymphocyte activation marker  | B cell-specific cellular differentiation Ag;<br>when bound to CD2, promotes T cell  |

| Cd69  | 0.997 | Cluster of                           | (BLAST-1); signaling lymphocytic activation molecule 2 (SLAMF2) C-type lectin | activation, and the formation of lipid rafts and caveolae for macrophages  Signal transmitting receptor in lymphocytes, |
|-------|-------|--------------------------------------|---|---|
| Cuos  | 0.557 | differentiation 69                   | domain family 2,  | NK cells, and platelets; induced upon T cell activation; involved in lymphocyte proliferation                           |
| Cd79b | 2.51  | Cluster of differentiation 79b       | B29   | One of the two flanking proteins that initiate signaling downstream of the BCR  |
| Fcgr1 | 0.456 | Fc fragment of IgG receptor Ia       | CD64  | High affinity receptor for the Fc region of γ- Igs; functions in both innate and adaptive immune responses              |
| IcosI | 0.655 | Inducible T cell costimulator ligand | CD275   | Ligand for T cell-specific co-receptor ICOS; also induces B cell proliferation and plasma cell differentiation          |
| lkzf1 | 0.994 | IKAROS family zinc finger 1          |   | Transcriptional regulator of hematopoietic cell differentiation; plays a role in T and B cell development               |
| Lyn   | 0.359 | Lck/Yes-related novel kinase         |   | Src family tyrosine kinase that potentiates signaling from the B cell receptor and CD40                                 |

| Ms4a1  | 2.89  | Membrane spanning      | CD20; Bp35        | B cell-specific membrane protein that                             |
|--------|-------|------------------------|-------------------|---|
|        |       | 4-domains A1           |                   | functions as a store-operated Ca(2+)                              |
|        |       |                        |                   | channel component, promoting Ca(2+)                               |
|        |       |                        |                   | influx after BCR activation                                       |
|        |       |                        |                   |   |
| Pik3cd | 0.741 | Phosphatidylinositol-  |                   | A subunit of PI3K; acts downstream of TLR4,                       |
|        |       | 4,5-bisphosphate 3-    |                   | TCR, BCR, and CD40; contributes to T helper                       |
|        |       | kinase catalytic       |                   | cell expansion, mast cell development, and                        |
|        |       | subunit delta isoform  |                   | neutrophil chemotaxis, extravasation, and                         |
|        |       |                        |                   | respiratory burst   |
|        |       |                        |                   |   |
| Pik3cg | 0.387 | Phosphatidylinositol-  |                   | A subunit of PI3K; modulates leukocyte                            |
|        |       | 4,5-bisphosphate 3-    |                   | chemotaxis to inflammatory sites and in                           |
|        |       | kinase catalytic       |                   | response to chemoattractant agents                                |
|        |       | subunit gamma          |                   |   |
|        |       | isoform                |                   |   |
|        |       |                        |                   |   |
| Pou2f2 | 1.4   | POU domain class 2,    |                   | TF that regulates Ab and IL-6 expression in B                     |
|        |       | transcription factor 2 |                   | cells   |
| Spn    | 0.812 | Sialophorin            | Leukosialin; CD43 | Cell surface sialoglycoprotein expressed by                       |
|        |       |                        |                   | T cells, B cells, monocytes, and                                  |
|        |       |                        |                   | granulocytes; promotes lymph node                                 |
|        |       |                        |                   | localization in T cells; shunts T cells away                      |
|        |       |                        |                   |   |
|        |       |                        |                   | from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; |
|        |       |                        |                   | promotes the expression of IFNγ in CD4 <sup>+</sup> T             |
|        |       |                        |                   | cells   |
|        |       |                        |                   |   |

| Tnfrsf13c | 2.42  | TNF receptor          |                     | B cell-activating factor; enhances B cell      |
|-----------|-------|-----------------------|---------------------|--|
|           |       | superfamily member    |                     | survival in vitro; regulator of the peripheral |
|           |       | 13C                   |                     | B cell population                              |
| Zap70     | 0.772 | Zeta chain of T cell  |                     | Tyrosine kinase that plays an essential role   |
|           |       | receptor associated   |                     | in regulation of the adaptive immune           |
|           |       | protein kinase 70     |                     | response; regulates motility, adhesion,        |
|           |       |                       |                     | proliferation, differentiation, and cytokine   |
|           |       |                       |                     | expression in T cells; contributes to the      |
|           |       |                       |                     | development and activation of primary B        |
|           |       |                       |                     | cells; phosphorylates at least two essential   |
|           |       |                       |                     | adapter proteins: LAT and LCP2, leading in     |
|           |       |                       |                     | turn to a large number of signaling            |
|           |       |                       |                     | molecules being recruited                      |
|           |       |                       | Cell Cycle          |  |
| Atm       | 0.29  | Ataxia telangiectasia |                     | Serine/threonine protein kinase that           |
|           |       | mutated               |                     | activates checkpoint signaling upon DSBs,      |
|           |       |                       |                     | apoptosis, and genotoxic stresses; acts as a   |
|           |       |                       |                     | master controller for cell cycle checkpoint    |
|           |       |                       |                     | signaling pathways required for the DNA        |
|           |       |                       |                     | damage response and genomic stability          |
| Cdkn1a    | 0.478 | Cyclin dependent      | p21; CDK-           | Binds to and inhibits cyclin-dependent         |
|           |       | kinase inhibitor 1A   | interaction protein | kinase activity, preventing phosphorylation    |
|           |       |                       | 1 (CIP1)            | of critical cyclin-dependent kinase            |

|       |            |                                   |  | substrates and blocking cell cycle progression   |  |  |  |
|-------|------------|-----------------------------------|--|--|--|--|--|
|       | Chemotaxis |                                   |  |  |  |  |  |
| Ccl3  | 0.891      | C-C motif chemokine<br>ligand 3   | Macrophage inflammatory protein $1\alpha$ (MIP1 $\alpha$ )                         | Chemoattractant ligand for CCR1, -4, and -5  |  |  |  |
| Ccl4  | 0.609      | C-C motif chemokine ligand 4      | Macrophage inflammatory protein 1β (MIP1β)   | Chemoattractant for NK cells and monocytes; binds to CCR5 receptors  |  |  |  |
| Ccl5  | 1.03       | C-C motif chemokine               | Regulated upon activation, normally T-expressed, and presumably secreted (RANTES)) | Chemoattractant ligand for CCR1, -3, -4, and -5; attracts blood monocytes, memory T helper cells and eosinophils; causes the release of histamine from basophils and activates eosinophils |  |  |  |
| Ccr1  | 0.523      | C-C motif chemokine receptor 1    | MIP1α receptor   | Receptor for CCL3, -5, -7, and -23   |  |  |  |
| Ccr5  | 0.68       | C-C motif chemokine<br>receptor 5 | CD195  | Receptor for a number of inflammatory CC-chemokines, including CCL3/MIP1 $\alpha$ , CCL4/MIP1 $\beta$ , and RANTES; signals via Ca(2+) flux  |  |  |  |
| Ccrl2 | 0.338      | C-C chemokine<br>receptor-like 2  |  | Stabilizes TLR4 surface expression on macrophages  |  |  |  |

| Cv12   | 0.000 | C V C +:t           | N.A                | Chamadina madus di busatis d                 |
|--------|-------|---------------------|--------------------|--|
| Cxcl2  | 0.609 | C-X-C motif         | Macrophage         | Chemokine produced by activated              |
|        |       | chemokine ligand 2  | inflammatory       | monocytes and neutrophils and expressed      |
|        |       |                     | protein 2-alpha    | at sites of inflammation                     |
|        |       |                     | (MIP2α); GRO2      |  |
|        |       |                     | oncogene           |  |
|        |       |                     |                    |  |
| Cxcl3  | 0.724 | C-X-C motif         | GRO3 oncogene      | Ligand for CXCR2; attracts neutrophils       |
|        |       | chemokine ligand 3  |                    |  |
|        |       |                     |                    |  |
| Cxcl9  | 0.677 | C-X-C motif         | Humig              | Chemoattractant ligand for CXCR3; attracts   |
|        |       | chemokine ligand 9  |                    | activated T cells                            |
|        |       |                     |                    |  |
| Cxcl10 | 0.709 | C-X-C motif         | IFNγ-induced       | Macrophage, DC, T cell, and NK cell          |
|        |       | chemokine ligand 10 | protein 10 (IP-10) | chemattractant secreted by several cell      |
|        |       |                     |                    | types in response to IFNy; binds to CXCR3    |
|        |       | _                   |                    |  |
| Cxcl11 | 0.769 | C-X-C motif         |                    | Dominant ligand for CXCR3; attracts          |
|        |       | chemokine ligand 11 |                    | activated T cells; strongly induced by IFNγ  |
| Cxcr3  | 0.768 | C-X-C motif         | CD183              | Induces integrin activation, cytoskeletal    |
| CACIS  | 0.708 |                     | CD163              |  |
|        |       | chemokine receptor  |                    | remodeling, and chemotaxis; expressed by T   |
|        |       | 3                   |                    | cells and NK cells; prominently expressed in |
|        |       |                     |                    | effector and memory T cells                  |
|        | 0.740 | 0 4 6               | CD404 5 :          |  |
| Cxcr4  | 0.743 | C-X-C motif         | CD184; fusin       | Alpha-chemokine receptor specific for SDF1   |
|        |       | chemokine receptor  |                    | aka CXCL12                                   |
|        |       | 4                   |                    |  |
|        |       |                     |                    |  |

| Cxcr6  | 0.736 | C-X-C motif           | CD186              | Receptor for the C-X-C chemokine CXCL16;                    |
|--------|-------|-----------------------|--------------------|---|
|        |       | chemokine receptor    |                    | expressed in several T lymphocyte subsets                   |
|        |       | 6                     |                    | and bone marrow stromal cells                               |
|        |       | Ü                     |                    | and some marrow stromar cens                                |
| lsg15  | 0.717 | Interferon-stimulated |                    | Ubiquitin-like protein that binds                           |
|        |       | gene 15               |                    | intracellular target proteins upon activation               |
|        |       |                       |                    | by IFN $\alpha$ or $\beta$ ; can also be secreted to induce |
|        |       |                       |                    | NK cell proliferation, act as a                             |
|        |       |                       |                    | chemoattractant for neutrophils, and                        |
|        |       |                       |                    | induce IFNγ upon binding to ITGAL/ITGB2                     |
| Itgam  | 0.681 | Integrin alpha M      | CD11b              | Pairs with CD18 to form Mac-1 aka                           |
|        |       |                       |                    | complement receptor 3; mediates leukocyte                   |
|        |       |                       |                    | activation, adhesion, chemotaxis, migration,                |
|        |       |                       |                    | phagocytosis, and cell-mediated                             |
|        |       |                       |                    | cytotoxicity; serves as a macrophage marker                 |
| Pik3cd | 0.741 | Phosphatidylinositol- |                    | A subunit of PI3K; acts downstream of TLR4,                 |
|        |       | 4,5-bisphosphate 3-   |                    | TCR, BCR, and CD40; contributes to T helper                 |
|        |       | kinase catalytic      |                    | cell expansion, mast cell development, and                  |
|        |       | subunit delta isoform |                    | neutrophil chemotaxis, extravasation, and                   |
|        |       |                       |                    | respiratory burst   |
|        |       | Compler               | nent & Humoral Imm | nunity  |
|        |       |                       |                    |   |
| C3ar1  | 0.506 | Complement            |                    | GPCR that binds to C3a, activating                          |
|        |       | component 3a          |                    | chemotaxis, granule enzyme release,                         |
|        |       | receptor 1            |                    |   |
|        |       |                       |                    |   |

|       |       |  |                                       | superoxide anion production, and bacterial opsonization  |
|-------|-------|--|---------------------------------------|--|
| Cfb   | 0.358 | Complement factor B                      |                                       | Alternate complement pathway  component; when cleaved, produces a  serine protease that binds to C3b to form  C3 convertase  |
| Fcgr1 | 0.456 | Fc fragment of IgG receptor Ia           | CD64                                  | High affinity receptor for the Fc region of $\gamma$ - lgs; functions in both innate and adaptive immune responses   |
| Fcgr4 | 1.13  | Fragment crystallizable gamma receptor 4 | Fc receptor-like 3<br>(Fcrl3); CD16-2 | Putative mouse ortholog to human FcγRIIIA  |
|       |       |  | Costimulation                         |  |
| Cd40  | 0.843 | Cluster of differentiation 40            |                                       | APC-expressed costimulatory protein that binds to CD40L on CD4+T cells, causing activation of both   |
| Dpp4  | 0.388 | Dipeptidyl-peptidase  4                  | CD26                                  | Cell surface glycoprotein receptor involved in the costimulatory signal essential for TCR-mediated T cell activation; serine exopeptidase that cleaves various substrates, thereby inactivating them; involved in the migration and invasion of endothelial cells into the ECM; also |

|           |       |  |  | regulates lymphocyte-epithelial cell adhesion  |
|-----------|-------|--|--|--|
| Icos      | 0.984 | Inducible T cell costimulator                        | CD278                                      | Enhances all basic T cell responses to foreign Ag; essential both for efficient interaction between T and B cells and for normal Ab responses to T cell-dependent Ags  |
| Icosl     | 0.655 | Inducible T cell costimulator ligand                 | CD275                                      | Ligand for T cell-specific co-receptor ICOS; also induces B cell proliferation and plasma cell differentiation   |
| Ptprc     | 0.937 | Protein tyrosine  phosphatase  receptor type C       | CD45; leukocyte<br>common antigen<br>(LCA) | Delivers costimulation during T cell activation upon binding to its ligand DPP4; dephosphorylates Lyn and suppresses JAK kinases   |
| Tnfrsf13c | 2.42  | TNF receptor superfamily member                      |  | B cell-activating factor; enhances B cell survival in vitro; regulator of the peripheral B cell population   |
| Tnfrsf14  | 0.671 | Tumor necrosis factor receptor superfamily member 14 | CD270                                      | Receptor for four distinct ligands: LIGHT, lymphotoxin-α, BTLA, and CD160, altogether defining a complex stimulatory and inhibitory signaling network; signals via the TRAF2-TRAF3 E3 ligase pathway to promote immune cell survival and |

|      |       |                     |                 | differentiation which the test of the test |
|------|-------|---------------------|-----------------|--|
|      |       |                     |                 | differentiation; participates in bidirectional   |
|      |       |                     |                 | cell-cell contact signaling between APCs and   |
|      |       |                     |                 | lymphocytes; delivers costimulatory signals  |
|      |       |                     |                 | to T cells, promoting cell proliferation and   |
|      |       |                     |                 | effector functions; interacts with CD160 on  |
|      |       |                     |                 | NK cells, enhancing IFNγ production and  |
|      |       |                     |                 | antitumor immune response; upon binding  |
|      |       |                     |                 | to CD160 on activated CD4 <sup>+</sup> T cells,  |
|      |       |                     |                 | downregulates CD28 costimulatory   |
|      |       |                     |                 | signaling; participates in cis or trans  |
|      |       |                     |                 | reactions with BTLA; cis interactions seem   |
|      |       |                     |                 | to promote quiescence; trans interactions  |
|      |       |                     |                 | seem to promote survival   |
|      |       |                     | Cytokines       |  |
|      |       |                     | <b>0,10</b> 10  |  |
| II1a | 0.916 | Interleukin 1 alpha | Hematopoietin-1 | Cytokine produced by monocytes and   |
|      |       |                     |                 | macrophages in response to cell injury;  |
|      |       |                     |                 | stimulates thymocyte proliferation by  |
|      |       |                     |                 | inducing IL-2 release; also stimulates B cell  |
|      |       |                     |                 | maturation and proliferation, and fibroblast   |
|      |       |                     |                 | growth factor activity   |
| II1b | 0.694 | Interleukin 1 beta  | Catabolin       | One of the two primary inflammatory  |
|      |       |                     |                 | cytokines produced by the inflammasome   |
|      |       |                     |                 | (the other one being IL-18); induces   |
|      |       |                     |                 | neutrophil influx and activation, T cell   |
|      |       |                     |                 |  |

| II2rg   | 0.874 | Interleukin 2 receptor subunit         | Common gamma<br>chain; CD132 | activation and cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNy synthesis from T <sub>H</sub> 1 cells  Common subunit for the receptors for a variety of interleukins, including IL-2, -4, -7, and -21 |
|---------|-------|--|------------------------------|---|
| II7r    | 1.07  | gamma Interleukin 7                    | CD127                        | Receptor for IL-7   |
| "71     | 1.07  | receptor                               | CDIZI                        | Receptor for IL-7   |
| Il12rb1 | 0.83  | Interleukin 12 receptor subunit beta 1 | CD212                        | Cytokine receptor component that associates with IL12RB2 to IL23R   |
| Il12rb2 | 0.612 | Interleukin 12 receptor subunit beta 2 |                              | Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  cell proliferation; promotes T <sub>H</sub> 1 polarization  by enhancing IFNy production  |
|         |       |  | Cytotoxicity                 |   |
| Gzmb    | 1.2   | Granzyme B                             | Fragmentin 2                 | Abundant protease in the cytosolic granules of cytotoxic T and NK cells that activates caspase-mediated cell death when   |

| Gzmk   | 1.07   | Granzyme K                               | Tryptase II                           | delivered into the target cell through the immunological synapse  Granule-secreted, pro-apoptotic serine |
|--------|--------|--|---------------------------------------|--|
|        |        |  |                                       | protease found in the cytoplasmic granules  of CTLs  |
| Fcgr4  | 1.13   | Fragment crystallizable gamma receptor 4 | Fc receptor-like 3<br>(Fcrl3); CD16-2 | Putative mouse ortholog to human FcyRIIIA  |
|        |        | G  | rowth/Proliferation                   |  |
| Map2k1 | 0.181  | Dual specificity                         | MAPK/ERK kinase 1                     | Essential component of the MAP kinase  |
|        |        | mitogen-activated                        | (MEK1)                                | signal transduction pathway; participates in   |
|        |        | protein kinase kinase                    |                                       | numerous biological functions, including cell  |
|        |        | 1  |                                       | growth, adhesion, survival, differentiation,   |
|        |        |  |                                       | transcription, metabolism, and cytoskeletal  |
|        |        |  |                                       | remodeling   |
| Rps6   | 0.492  | Ribosomal protein 6                      |                                       | Component of the 40S small ribosomal   |
|        |        |  |                                       | subunit; plays an important role in  |
|        |        |  |                                       | controlling cell growth and proliferation  |
|        |        |  |                                       | through the selective translation of   |
|        |        |  |                                       | particular classes of mRNA   |
| Yy1    | 0.0857 | Yin yang 1                               |                                       | Ubiquitous factor that serves as a   |
|        |        |  |                                       | transcriptional "switch", either promoting   |
|        |        |  |                                       | or repressing the transcription of numerous  |

|              |       |                       |                | genes through the selective recruitment of         |
|--------------|-------|-----------------------|----------------|--|
|              |       |                       |                | either histone deacetylases or                     |
|              |       |                       |                | acetyltransferases; plays a fundamental role       |
|              |       |                       |                | in diverse processes, such as differentiation,     |
|              |       |                       |                | replication, and cellular proliferation            |
|              |       |                       | Hematopoiesis  |  |
| <u>Hck</u>   | 0.486 | Hematopoietic cell    |                | Src family tyrosine kinase that mediates           |
|              |       | kinase                |                | secretory lysosome mobilization,                   |
|              |       |                       |                | degranulation, and activation of NADPH             |
|              |       |                       |                | oxidase  |
| <u>lkzf1</u> | 0.994 | IKAROS family zinc    |                | Transcriptional regulator of hematopoietic         |
|              |       | finger 1              |                | cell differentiation; plays a role in T and B      |
|              |       |                       |                | cell development                                   |
| <u>lkzf2</u> | 0.383 | IKAROS family zinc    |                | Hematopoietic cell-specific TF involved in         |
|              |       | finger protein 2      |                | early hematopoietic development                    |
| Stat5b       | 0.314 | Signal transducer and |                | Carries out a dual function: signal                |
|              |       | activator of          |                | transduction and activation of transcription;      |
|              |       | transcription 5b      |                | positively regulates                               |
|              |       |                       |                | hematopoietic/erythroid differentiation.           |
|              |       |                       | Inflammation   |  |
|              |       |                       |                |  |
| Casp1        | 0.417 | Caspase 1             | Interleukin 1β | Cysteine-aspartic acid protease that               |
|              |       |                       | convertase     | mediates cleavage-based activation of IL-1 $\beta$ |

|         |       |  |                        | and IL-18; serves as the central enzymatic core of the inflammasome; also induces apoptosis   |
|---------|-------|--|------------------------|---|
| Cd38    | 1.02  | Cluster of differentiation 38          | ADP-ribosyl cyclase  1 | Synthesizes the second messengers cyclic  ADP-ribose and NADPH; appears to play a  critical role in inflammation, although its  exact immunological function(s) remain(s)  poorly defined   |
| Ctsh    | 0.363 | Cathepsin H                            |                        | Lysosomal protease; increased in macrophages in response to IFNy  |
| II1b    | 0.694 | Interleukin 1 beta                     | Catabolin              | One of the two primary inflammatory cytokines produced by the inflammasome (the other one being IL-18); induces neutrophil influx and activation, T cell activation and cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNy synthesis from T <sub>H</sub> 1 cells |
| II12rb1 | 0.83  | Interleukin 12 receptor subunit beta 1 | CD212                  | Cytokine receptor component that associates with IL12RB2 to IL23R   |

| Il12rb2 | 0.612 | Interleukin 12        |                      | Signaling component coupling to the                         |
|---------|-------|-----------------------|----------------------|---|
| 1112102 | 0.012 |                       |                      |   |
|         |       | receptor subunit      |                      | JAK2/STAT4 pathway; promotes T and NK                       |
|         |       | beta 2                |                      | cell proliferation; promotes T <sub>H</sub> 1 polarization  |
|         |       |                       |                      | by enhancing IFNγ production                                |
| Irf1    | 0.693 | Interferon regulatory |                      | Transcriptional regulator that promotes                     |
|         |       | factor 1              |                      | inflammatory innate and adaptive immune                     |
|         |       |                       |                      | responses   |
| Irf4    | 0.927 | Interferon regulatory |                      | Transcriptional activator that complexes                    |
|         |       | factor 4              |                      | with BATF and binds ISREs within the                        |
|         |       |                       |                      | promoters of multiple genes involved in                     |
|         |       |                       |                      | inflammation  |
| Irgm2   | 0.564 | Immunity-related      | Interferon-inducible | Function not fully known, but most likely                   |
|         |       | GTPase family M       | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory                    |
|         |       | member 2              |                      | cytokine production   |
| lsg15   | 0.717 | Interferon-stimulated |                      | Ubiquitin-like protein that binds                           |
|         |       | gene 15               |                      | intracellular target proteins upon activation               |
|         |       |                       |                      | by IFN $\alpha$ or $\beta$ ; can also be secreted to induce |
|         |       |                       |                      | NK cell proliferation, act as a                             |
|         |       |                       |                      | chemoattractant for neutrophils, and                        |
|         |       |                       |                      | induce IFNγ upon binding to ITGAL/ITGB2                     |
| Jak1    | 0.493 | Janus kinase 1        |                      | Essential tyrosine kinase involved signal                   |
|         |       |                       |                      | transduction in type I and II cytokines and                 |
|         |       |                       |                      | IFNs  |
|         |       |                       |                      |   |

| Lgals3        | 0.299 | Galectin 3             |           | Galactose-specific lectin that binds IgE;   |
|---------------|-------|------------------------|-----------|---|
|               |       |                        |           | involved in acute inflammatory responses,   |
|               |       |                        |           | including neutrophil activation and         |
|               |       |                        |           | adhesion, chemoattraction of                |
|               |       |                        |           | monocytes/macrophages, opsonization of      |
|               |       |                        |           | apoptotic neutrophils, and activation of    |
|               |       |                        |           | mast cells                                  |
| Nlrp3         | 0.652 | NACHT domain-,         | Cryopyrin | PRR with a wide diversity of recognized     |
|               |       | leucine-rich repeat-,  |           | targets that activates the NLRP3            |
|               |       | and PYD-containing     |           | inflammasome consisting of NLRP3,           |
|               |       | protein 3              |           | PYCARD, and caspase-1/-8                    |
| Nos2          | 1.58  | Inducible nitric oxide |           | Produces reactive oxygen species and        |
|               |       | synthase (iNOS)        |           | contributes to inflammatory cytokine        |
|               |       |                        |           | production                                  |
| <u>Pik3cd</u> | 0.741 | Phosphatidylinositol-  |           | A subunit of PI3K; acts downstream of TLR4, |
|               |       | 4,5-bisphosphate 3-    |           | TCR, BCR, and CD40; contributes to T helper |
|               |       | kinase catalytic       |           | cell expansion, mast cell development, and  |
|               |       | subunit delta isoform  |           | neutrophil chemotaxis, extravasation, and   |
|               |       |                        |           | respiratory burst                           |
| Pik3cg        | 0.387 | Phosphatidylinositol-  |           | A subunit of PI3K; modulates leukocyte      |
|               |       | 4,5-bisphosphate 3-    |           | chemotaxis to inflammatory sites and in     |
|               |       | kinase catalytic       |           | response to chemoattractant agents          |
|               |       |                        |           |   |

|            |       | subunit gamma<br>isoform      |  |   |
|------------|-------|-------------------------------|--|---|
| <u>Spn</u> | 0.812 | Sialophorin                   | Leukosialin; CD43                      | Cell surface sialoglycoprotein expressed by  T cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNy in CD4 <sup>+</sup> T  cells |
| Tbk1       | 0.338 | TANK-binding kinase           |  | Coordinates the activation of IRF3 and NFkB and induction of type I IFNs  |
| Tmem173    | 0.485 | Transmembrane protein 173     | Stimulator of interferon genes (STING) | Adaptor protein in type I IFN signaling; activates STAT6 and IRF3 through TBK1 to induce type I IFN production  |
|            |       |                               | Inhibition                             |   |
| Bcl2       | 0.418 | B cell lymphoma 2             |  | Outer mitochondrial membrane protein that inhibits apoptosis and autophagy; may attenuate inflammation by impairing inflammasome formation  |
| Btla       | 1.73  | B and T lymphocyte attenuator | CD272                                  | Inhibitory cell surface protein that inhibits T cell function by binding to B7H4 and TNFRSF14   |

|              |       | differentiation 200  |                     | surface glycoprotein; limits inflammation by     |
|--------------|-------|----------------------|---------------------|--|
|              |       | receptor 1           |                     | inhibiting the expression of                     |
|              |       |                      |                     | proinflammatory molecules including TNF $lpha$ , |
|              |       |                      |                     | IFNs, and iNOS                                   |
| Cdkn1a       | 0.478 | Cyclin dependent     | p21; CDK-           | Binds to and inhibits cyclin-dependent           |
|              |       | kinase inhibitor 1A  | interaction protein | kinase activity, preventing phosphorylation      |
|              |       |                      | 1 (CIP1)            | of critical cyclin-dependent kinase              |
|              |       |                      |                     | substrates and blocking cell cycle               |
|              |       |                      |                     | progression                                      |
| Ctla4        | 0.786 | Cytotoxic T          | CD152               | Inhibitory receptor that blocks CD28             |
|              |       | lymphocyte antigen 4 |                     | costimulation by competitively binding its       |
|              |       |                      |                     | ligands CD80 and CD86                            |
| <u>Cd274</u> | 1.1   | Cluster of           | Programmed cell     | Ubiquitously expressed ligand for co-            |
|              |       | differentiation 274  | death receptor      | inhibitory receptor PD-1; upregulated by         |
|              |       |                      | ligand 1 (PD-L1)    | tumors as an immune evasion strategy             |
| Cyld         | 0.305 | Cylindromatosis      |                     | Inhibits NFkB activation by deubiquitinating     |
|              |       | lysine 63            |                     | upstream signaling factors; inhibits Wnt         |
|              |       | deubiquitinase       |                     | signaling; restricts polyubiquitination of       |
|              |       |                      |                     | RIPK1 and -2, thereby limiting necroptosis       |
| <u>Irak3</u> | 0.515 | Interleukin-1        |                     | Adaptor protein that negatively regulates        |
|              |       | receptor-associated  |                     | TLR signaling; predominantly expressed in        |
|              |       | kinase 3             |                     | monocytes and macrophages                        |

| Foxp3         | 1.02  | Forkhead box P3      | DIETER | Master TF for Tregs; represses expression of    |
|---------------|-------|----------------------|--------|---|
|               |       |                      |        | Tnfrsf18, Il2ra, and Ctla4                      |
| <u>Nfkbia</u> | 0.553 | Nuclear factor kappa |        | Inhibits activity of REL dimers by masking of   |
|               |       | B inhibitor alpha    |        | their nuclear localization signals              |
| Nlrc5         | 0.709 | NLR family CARD      | NOD27  | Inhibits NFкВ and type I IFN signaling          |
|               |       | domain containing 5  |        | pathways; may also regulate the type II IFN     |
|               |       |                      |        | signaling pathway                               |
| Socs1         | 0.758 | Suppressor of        |        | Inhibits JAK proteins; negative regulator of    |
|               |       | cytokine signaling 1 |        | IL-6  |
| Tnfrsf14      | 0.671 | TNF receptor         | CD270  | Receptor for four distinct ligands: LIGHT,      |
|               |       | superfamily member   |        | lymphotoxin-α, BTLA, and CD160,                 |
|               |       | 14                   |        | altogether defining a complex stimulatory       |
|               |       |                      |        | and inhibitory signaling network; signals via   |
|               |       |                      |        | the TRAF2-TRAF3 E3 ligase pathway to            |
|               |       |                      |        | promote immune cell survival and                |
|               |       |                      |        | differentiation; participates in bidirectional  |
|               |       |                      |        | cell-cell contact signaling between APCs and    |
|               |       |                      |        | lymphocytes; delivers costimulatory signals     |
|               |       |                      |        | to T cells, promoting cell proliferation and    |
|               |       |                      |        | effector functions; interacts with CD160 on     |
|               |       |                      |        | NK cells, enhancing IFNγ production and         |
|               |       |                      |        | antitumor immune response; upon binding         |
|               |       |                      |        | to CD160 on activated CD4 <sup>+</sup> T cells, |

|             |       |                       |                      | downregulates CD28 costimulatory                            |
|-------------|-------|-----------------------|----------------------|---|
|             |       |                       |                      | signaling; participates in cis or trans                     |
|             |       |                       |                      | reactions with BTLA; cis interactions seem                  |
|             |       |                       |                      | to promote quiescence; trans interactions                   |
|             |       |                       |                      | seem to promote survival                                    |
|             |       |                       |                      |   |
|             |       | lı                    | nterferon Response   |   |
| <u>Ctsh</u> | 0.363 | Cathepsin H           |                      | Lysosomal protease; increased in                            |
|             |       |                       |                      | macrophages in response to IFNγ                             |
| <u>Irf1</u> | 0.693 | Interferon regulatory |                      | Transcriptional regulator that promotes                     |
|             |       | factor 1              |                      | inflammatory innate and adaptive immune                     |
|             |       |                       |                      | responses   |
|             |       |                       |                      |   |
| <u>Irf4</u> | 0.927 | Interferon regulatory |                      | Transcriptional activator that complexes                    |
|             |       | factor 4              |                      | with BATF and binds ISREs within the                        |
|             |       |                       |                      | promoters of multiple genes involved in                     |
|             |       |                       |                      | inflammation  |
| Irgm2       | 0.564 | Immunity-related      | Interferon-inducible | Function not fully known, but most likely                   |
|             |       | GTPase family M       | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory                    |
|             |       | member 2              |                      | cytokine production   |
|             |       |                       |                      | ν, σ γ. σ   |
| Isg15       | 0.717 | Interferon-stimulated |                      | Ubiquitin-like protein that binds                           |
|             |       | gene 15               |                      | intracellular target proteins upon activation               |
|             |       |                       |                      | by IFN $\alpha$ or $\beta$ ; can also be secreted to induce |
|             |       |                       |                      | NK cell proliferation, act as a                             |
|             |       |                       |                      |   |

|         |       |                      |                  | chemoattractant for neutrophils, and        |
|---------|-------|----------------------|------------------|---|
|         |       |                      |                  | induce IFNγ upon binding to ITGAL/ITGB2     |
| Jak1    | 0.493 | Janus kinase 1       |                  | Essential tyrosine kinase involved signal   |
|         |       |                      |                  | transduction in type I and II cytokines and |
|         |       |                      |                  | IFNs  |
| Mx2     | 0.932 | Myxovirus resistance |                  | IFN-induced dynamin-like GTPase with        |
|         |       | protein 2            |                  | potent antiviral activity against HIV-1     |
| Tmem173 | 0.485 | Transmembrane        | Stimulator of    | Adaptor protein in type I IFN signaling;    |
|         |       | protein 173          | interferon genes | activates STAT6 and IRF3 through TBK1 to    |
|         |       |                      | (STING)          | induce type I IFN production                |
|         |       |                      | Ion Transport    |   |
| App     | 0.157 | Amyloid-beta         |                  | Cell surface receptor and transmembrane     |
|         |       | precursor protein    |                  | precursor protein that is cleaved by        |
|         |       |                      |                  | secretases to form a number of peptides;    |
|         |       |                      |                  | involved in cell mobility, copper           |
|         |       |                      |                  | homeostasis, and oxidative stress           |
| Ms4a1   | 2.89  | Membrane spanning    | CD20; Bp35       | B cell-specific membrane protein that       |
|         |       | 4-domains A1         |                  | functions as a store-operated Ca(2+)        |
|         |       |                      |                  | channel component, promoting Ca(2+)         |
|         |       |                      |                  | influx after BCR activation                 |

| <u>Slc11a1</u> | 0.95  | Natural resistance- |                  | Macrophage-specific metal ion transporter;    |
|----------------|-------|---------------------|------------------|---|
|                |       | associated          |                  |   |
|                |       | associated          |                  | uptakes divalent metal cations to neutralize  |
|                |       | macrophage protein  |                  | ROSs  |
|                |       | 1                   |                  |   |
|                |       |                     |                  |   |
|                |       |                     | IRAKs & TRAFs    |   |
| <u>Irak2</u>   | 0.515 | Interleukin-1       |                  | Adaptor protein involved in TLR and IL-1      |
|                |       | receptor-associated |                  | signaling                                     |
|                |       | kinase 2            |                  |   |
|                |       |                     |                  |   |
| <u>Irak3</u>   | 0.734 | Interleukin-1       |                  | Adaptor protein that negatively regulates     |
|                |       | receptor-associated |                  | TLR signaling; predominantly expressed in     |
|                |       | kinase 3            |                  | monocytes and macrophages                     |
|                |       |                     | JAK-STAT Pathway |   |
| Jak1           | 0.493 | Janus kinase 1      |                  | Essential tyrosine kinase involved signal     |
|                |       |                     |                  | transduction in type I and II cytokines and   |
|                |       |                     |                  | IFNs  |
| Jak3           | 0.434 | Janus kinase 3      |                  | Non-receptor tyrosine kinase involved in      |
|                |       |                     |                  | various processes such as cell growth,        |
|                |       |                     |                  | development, or differentiation; mediates     |
|                |       |                     |                  | essential signaling events in both innate and |
|                |       |                     |                  | adaptive immunity                             |
|                |       |                     |                  |   |

| activator of transcription 1  Stat5b  0.314  Signal transducer and activator of transcription 5b  Stat6  0.187  Signal transducer and activator of transcription 6  Stat6  Atm  0.29  Ataxia telangiectasia mutated  activator of transcription 1  Carries out a dual function: signal transduction and activation of transcription 6  positively regulates hematopoietic/erythroid differential macrophage function and M2 macrophage function and M2 macrophage function and M2 macrophage function and M2 macrophage function and M3 macrophage function and M4 macrophage function and M4 macrophage function and M5 macrophage function and M6 macrophage function and M6 macrophage function and M7 macrophage function and M8 macrophage fun           |        |
|--|--------|
| Stat5b  O.314 Signal transducer and activator of transduction and activation of transcription 5b positively regulates hematopoietic/erythroid differential states  Stat6 O.187 Signal transducer and activator of transcription 6 Essential TF for TH2 CD4+ T cell and macrophage function and M2 macrophage function and M3 m           |        |
| activator of transcription 5b positively regulates hematopoietic/erythroid differentia  Stat6 0.187 Signal transducer and activator of transcription 6 Essential TF for TH2 CD4+ T cell and macrophage function and M2 macrophage function and M3 macrophage function an           |        |
| transcription 5b positively regulates hematopoietic/erythroid differential  Stat6 0.187 Signal transducer and activator of transcription 6 Essential TF for TH2 CD4+ T cell and macrophage function and M2 macrophage function and M2 macrophages  Kinases  Atm 0.29 Ataxia telangiectasia Serine/threonine protein kinase the   | otion; |
| Stat6  O.187 Signal transducer and activator of transcription 6  Kinases  Atm  O.29 Ataxia telangiectasia  hematopoietic/erythroid differential hematopoietic/e           |        |
| Stat6  O.187 Signal transducer and activator of transcription 6  Kinases  Atm  O.29 Ataxia telangiectasia  Essential TF for TH2 CD4 <sup>+</sup> T cell and macrophage function and M2 macrophage function |        |
| activator of transcription 6 macrophage function and M2 macrophage function           | ion.   |
| transcription 6 polarization  Kinases  Atm 0.29 Ataxia telangiectasia Serine/threonine protein kinase the  | k      |
| Kinases  Atm 0.29 Ataxia telangiectasia Serine/threonine protein kinase th   | hage   |
| Atm 0.29 Ataxia telangiectasia Serine/threonine protein kinase th  |        |
|  |        |
|  | at     |
|  |        |
| apoptosis, and genotoxic stresses; act   |        |
| master controller for cell cycle check   | oint   |
| signaling pathways required for the  | ONA    |
| damage response and genomic stab   |        |
| Btk 0.615 Bruton's tyrosine Crucial kinase in B cell receptor sig  | lity   |
| kinase transmission and B cell activation  |        |
| Hck 0.486 Hematopoietic cell Src family tyrosine kinase that medi  | nal    |
| kinase secretory lysosome mobilization   | nal    |

|             |       |                       |     | degranulation, and activation of NADPH  oxidase |
|-------------|-------|-----------------------|-----|---|
|             |       |                       |     |   |
| <u>Itk</u>  | 0.932 | Interleukin-2-        | LYK | Key actor in the TCR signaling cascade;         |
|             |       | inducible T cell      |     | phosphorylates PLCγ1, LAT, and LCP2             |
|             |       | kinase                |     |   |
| <u>Lck</u>  | 0.983 | Lymphocyte cell       |     | Src family tyrosine kinase that acts as one of  |
|             |       | kinase                |     | the main signaling intermediaries               |
|             |       |                       |     | downstream of the TCR; constitutively           |
|             |       |                       |     | associated with the cytoplasmic portion of      |
|             |       |                       |     | CD4   |
| <u>Lyn</u>  | 0.359 | Tyrosine-protein      |     | Src family tyrosine kinase that potentiates     |
|             |       | kinase Lyn            |     | signaling from the B cell receptor and CD40     |
| Jak1        | 0.493 | Janus kinase 1        |     | Essential tyrosine kinase involved signal       |
|             |       |                       |     | transduction in type I and II cytokines and     |
|             |       |                       |     | IFNs  |
| <u>Jak3</u> | 0.434 | Janus kinase 3        |     | Non-receptor tyrosine kinase involved in        |
|             |       |                       |     | various processes such as cell growth,          |
|             |       |                       |     | development, or differentiation; mediates       |
|             |       |                       |     | essential signaling events in both innate and   |
|             |       |                       |     | adaptive immunity                               |
| Pik3cd      | 0.741 | Phosphatidylinositol- |     | A subunit of PI3K; acts downstream of TLR4,     |
|             |       | 4,5-bisphosphate 3-   |     | TCR, BCR, and CD40; contributes to T helper     |
|             |       |                       |     | cell expansion, mast cell development, and      |

|              |       | kinase catalytic      | neutrophil chemotaxis, extravasation, and      |
|--------------|-------|-----------------------|--|
|              |       | subunit delta isoform | respiratory burst                              |
| Pik3cg       | 0.387 | Phosphatidylinositol- | A subunit of PI3K; modulates leukocyte         |
|              |       | 4,5-bisphosphate 3-   | chemotaxis to inflammatory sites and in        |
|              |       | kinase catalytic      | response to chemoattractant agents             |
|              |       | subunit gamma         |  |
|              |       | isoform               |  |
| Tbk1         | 0.338 | TANK-binding kinase   | Coordinates the activation of IRF3 and NFkB    |
|              |       | 1                     | and induction of type I IFNs                   |
| <u>Txk</u>   | 1.07  | TXK tyrosine kinase   | Regulates the development, function, and       |
|              |       |                       | differentiation of conventional T cells and    |
|              |       |                       | nonconventional NK-T cells; contributes to     |
|              |       |                       | signaling from many receptors and              |
|              |       |                       | participates in multiple downstream            |
|              |       |                       | pathways, including regulation of the actin    |
|              |       |                       | cytoskeleton; can phosphorylate PLCγ1,         |
|              |       |                       | leading to its localization in lipid rafts and |
|              |       |                       | activation, followed by subsequent cleavage    |
|              |       |                       | of its substrates                              |
| <u>Zap70</u> | 0.772 | Zeta chain of T cell  | Tyrosine kinase that plays an essential role   |
|              |       | receptor associated   | in regulation of the adaptive immune           |
|              |       | protein kinase 70     | response; regulates motility, adhesion,        |
|              |       |                       | proliferation, differentiation, and cytokine   |
|              |       |                       | expression in T cells; contributes to the      |

|             |       |                    |                     | development and activation of primary B  cells; phosphorylates at least two essential  adapter proteins: LAT and LCP2, leading in  turn to a large number of signaling  molecules being recruited |
|-------------|-------|--------------------|---------------------|---|
|             |       |                    | Lysosomal Activity  |   |
| <u>Ctsh</u> | 0.363 | Cathepsin H        |                     | Lysosomal protease; increased in  |
|             |       |                    |                     | macrophages in response to IFNγ   |
| Ctss        | 0.736 | Cathepsin S        |                     | Lysosomal protease that participates in   |
|             |       |                    |                     | processing of Ag by MHC class II  |
| <u>Hck</u>  | 0.486 | Hematopoietic cell |                     | Src family tyrosine kinase that mediates  |
|             |       | kinase             |                     | secretory lysosome mobilization,  |
|             |       |                    |                     | degranlation, and activation of NADPH   |
|             |       |                    |                     | oxidase   |
|             |       | M                  | lacrophage Function |   |
| <u>Cd14</u> | 0.268 | Cluster of         |                     | PRR that recognizes LPS; mostly found on  |
|             |       | differentiation 14 |                     | macrophages   |
| <u>Ctsh</u> | 0.363 | Cathepsin H        |                     | Lysosomal protease; increased in  |
|             |       |                    |                     | macrophages in response to IFNγ   |

| <u>Irak3</u>   | 0.515 | Interleukin-1         |                       | Adaptor protein that negatively regulates    |
|----------------|-------|-----------------------|-----------------------|--|
|                |       | receptor-associated   |                       | TLR signaling; predominantly expressed in    |
|                |       | kinase 3              |                       | monocytes and macrophages                    |
|                |       |                       |                       |  |
| <u>Itgam</u>   | 0.681 | Integrin alpha M      | CD11b                 | Pairs with CD18 to form Mac-1 aka            |
|                |       |                       |                       | complement receptor 3; mediates leukocyte    |
|                |       |                       |                       | activation, adhesion, chemotaxis, migration, |
|                |       |                       |                       | phagocytosis, and cell-mediated              |
|                |       |                       |                       | cytotoxicity; serves as a macrophage marker  |
| Marco          | 2.03  | Macrophage            |                       | A PRR that recognizes LDL                    |
| <u>iviareo</u> | 2.03  | receptor with         |                       | ATTIN that recognizes EDE                    |
|                |       |                       |                       |  |
|                |       | collagenous structure |                       |  |
| Slamf7         | 1.07  | Signaling lymphocytic |                       | A super-activator of macrophages and a       |
|                |       | activation molecule   |                       | strong promoter of phagocytosis; binds to    |
|                |       | family member 7       |                       | CD74   |
|                |       |                       |                       |  |
| Slc11a1        | 0.95  | Natural resistance-   |                       | Macrophage-specific metal ion transporter;   |
|                |       | associated            |                       | uptakes divalent metal cations to neutralize |
|                |       | macrophage protein    |                       | ROSs   |
|                |       | 1                     |                       |  |
|                |       | N                     | IAP Kinase Signaling  |  |
|                |       |                       |                       |  |
| Mapk1          | 0.444 | Mitogen-activated     | Extracellular signal- | Serine/threonine kinase that acts as an      |
|                |       | protein kinase 1      | regulated kinase 2    | essential component of the MAP kinase        |
|                |       |                       | (ERK2)                | signal transduction pathway                  |
|                |       |                       |                       |  |

| Mapk11        | 0.345 | Mitogen-activated     | Stress-activated   | Serine/threonine kinase that acts as an       |
|---------------|-------|-----------------------|--------------------|---|
|               |       | protein kinase 11     | protein kinase 2   | essential component of the MAP kinase         |
|               |       |                       | (SAPK2)            | signal transduction pathway                   |
| Map2k1        | 0.181 | Dual specificity      | MAPK/ERK kinase 1  | Essential component of the MAP kinase         |
|               |       | mitogen-activated     | (MEK1)             | signal transduction pathway; participates in  |
|               |       | protein kinase kinase |                    | numerous biological functions, including cell |
|               |       | 1                     |                    | growth, adhesion, survival, differentiation,  |
|               |       |                       |                    | transcription, metabolism, and cytoskeletal   |
|               |       |                       |                    | remodeling                                    |
| Map4k2        | 0.536 | Mitogen-activated     |                    | Essential component of the MAP kinase         |
|               |       | protein kinase kinase |                    | signal transduction pathway downstream of     |
|               |       | kinase kinase 2       |                    | TRAF6; upstream activator of the SAP/JNK      |
|               |       |                       |                    | signaling pathway;                            |
|               |       |                       | Mast Cell Function |   |
| <u>Lgals3</u> | 0.299 | Galectin 3            |                    | Galactose-specific lectin that binds IgE;     |
|               |       |                       |                    | involved in acute inflammatory responses,     |
|               |       |                       |                    | including neutrophil activation and           |
|               |       |                       |                    | adhesion, chemoattraction of monocytes        |
|               |       |                       |                    | macrophages, opsonization of apoptotic        |
|               |       |                       |                    | neutrophils, and activation of mast cells     |
| <u>Pik3cd</u> | 0.741 | Phosphatidylinositol- |                    | A subunit of PI3K; acts downstream of TLR4,   |
|               |       | 4,5-bisphosphate 3-   |                    | TCR, BCR, and CD40; contributes to T helper   |
|               |       |                       |                    | cell expansion, mast cell development, and    |

|             |       | kinase catalytic      |                    | neutrophil chemotaxis, extravasation, and     |
|-------------|-------|-----------------------|--------------------|---|
|             |       | ·                     |                    |   |
|             |       | subunit delta isoform |                    | respiratory burst                             |
| Pik3cg      | 0.387 | Phosphatidylinositol- |                    | A subunit of PI3K; modulates leukocyte        |
|             |       | 4,5-bisphosphate 3-   |                    | chemotaxis to inflammatory sites and in       |
|             |       | kinase catalytic      |                    | response to chemoattractant agents            |
|             |       | subunit gamma         |                    |   |
|             |       |                       |                    |   |
|             |       | isoform               |                    |   |
|             |       |                       | Metabolism         |   |
|             |       |                       |                    |   |
| Abcg1       | 1.01  | ATP-binding cassette  |                    | Membrane-associated cholesterol efflux        |
|             |       | transporter G1        |                    | pump  |
|             |       |                       |                    |   |
| <u>Cd36</u> | 0.93  | Cluster of            | Fatty acid         | Class B scavenger receptor that mediates      |
|             |       | differentiation 36    | translocase (FAT)  | fatty acid uptake                             |
|             |       |                       |                    |   |
| Map2k1      | 0.181 | Dual specificity      | MAPK/ERK kinase 1  | Essential component of the MAP kinase         |
|             |       | mitogen-activated     | (MEK1)             | signal transduction pathway; participates in  |
|             |       | protein kinase kinase |                    | numerous biological functions, including cell |
|             |       | 1                     |                    | growth, adhesion, survival, differentiation,  |
|             |       |                       |                    | transcription, metabolism, and cytoskeletal   |
|             |       |                       |                    | remodeling                                    |
|             |       |                       |                    |   |
|             |       |                       | Migration/Motility |   |
| <u>App</u>  | 0.157 | Amyloid-beta          |                    | Cell surface receptor and transmembrane       |
|             |       | precursor protein     |                    | precursor protein that is cleaved by          |
|             |       |                       |                    | secretases to form a number of peptides;      |
|             | 1     | 1                     | I                  |   |

|              |       |                               |                     | involved in cell mobility, copper homeostasis, and oxidative stress  |
|--------------|-------|-------------------------------|---------------------|--|
| Cd97         | 0.29  | Cluster of differentiation 97 | BL-Ac[F2]           | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells   |
| <u>Dpp4</u>  | 0.388 | Dipeptidyl-peptidase 4        | CD26                | Cell surface glycoprotein receptor involved in the costimulatory signal essential for TCR-mediated T cell activation; serine exopeptidase that cleaves various substrates, thereby inactivating them; involved in the migration and invasion of endothelial cells into the ECM; also regulates lymphocyte-epithelial cell adhesion |
| <u>Itgam</u> | 0.681 | Integrin alpha M              | CD11b               | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker   |
|              |       | Ŋ                             | ieutropnii Function |  |

| Fpr2   | 1.04  | Formyl peptide        | Lipoxin A4 receptor  | Low affinity receptor for N-formyl-         |
|--------|-------|-----------------------|----------------------|---|
|        |       | receptor 2            |                      | methionyl peptides; activates neutrophils   |
| Ncf4   | 0.545 | Neutrophil cytosolic  | SH3 and PX domain-   | Cytosolic regulatory component of the       |
|        |       | factor 4              | containing protein 4 | superoxide-producing phagocyte NADPH-       |
|        |       |                       | (SH3PXD4)            | oxidase, a multicomponent enzyme system     |
|        |       |                       |                      | important for host defense                  |
| Pik3cd | 0.741 | Phosphatidylinositol- |                      | A subunit of PI3K; acts downstream of TLR4, |
|        |       | 4,5-bisphosphate 3-   |                      | TCR, BCR, and CD40; contributes to T helper |
|        |       | kinase catalytic      |                      | cell expansion, mast cell development, and  |
|        |       | subunit delta isoform |                      | neutrophil chemotaxis, extravasation, and   |
|        |       |                       |                      | respiratory burst                           |
| Cxcl3  | 0.724 | C-X-C motif           | GRO3 oncogene        | Ligand for CXCR2; attracts neutrophils      |
|        |       | chemokine ligand 3    |                      |   |
|        |       |                       | NFkB Signaling       |   |
| Dol10  | 0.137 | D. coll               |                      | Activates NEVD via ubiquitination of IVV    |
| Bcl10  | 0.137 | B cell                |                      | Activates NFκB via ubiquitination of ΙΚΚγ   |
|        |       | lymphoma/leukemia     |                      |   |
|        |       | 10                    |                      |   |
| Card11 | 1.24  | Caspase recruitment   | Bcl10-interacting    | Adapter protein that plays a key role in    |
|        |       | domain family         | MAGUK protein 3      | adaptive immune response by transducing     |
|        |       | member 11             | (BIMP3)              | the activation of NFkB downstream of TCR    |
|        |       |                       |                      | and BCR engagement; upon activation in      |
|        |       |                       |                      | response to TCR or BCR triggering,          |
|        |       |                       |                      | homooligomerizes to form a nucleating       |

|              |       |  |       | helical template that recruits BCL10 via  CARD-CARD interaction, thereby promoting  polymerization of BCL10 and subsequent  recruitment of MALT1, leading to IKK  phosphorylation and degradation and  release of NFkB proteins for nuclear  translocation |
|--------------|-------|--|-------|--|
| Cxcr3        | 0.768 | C-X-C motif chemokine receptor                             | CD183 | Induces integrin activation, cytoskeletal remodeling, and chemotaxis; expressed by T cells and NK cells; prominently expressed in effector and memory T cells  |
| <u>Ikbke</u> | 0.868 | Inhibitor of nuclear factor kappa B kinase subunit epsilon |       | Noncanonical IKB kinase; plays an essential role in regulating inflammatory through the activation of the type I IFN, NFKB, and STAT signaling   |
| <u>Rel</u>   | 0.623 | Avian reticuloendotheliosis viral oncogene homolog         | c-Rel | One of the NFkB family TFs; important for B  cell and Treg development   |
| Relb         | 0.332 | Avian reticuloendotheliosis viral oncogene homolog B       |       | One of the NFkB family TFs; controls  lymphoid development, DC biology, and  noncanonical NFkB signaling   |

| Dielo          | 0.274 | Document interesting |                  | DID kinned that national actions is and                    |
|----------------|-------|----------------------|------------------|--|
| Ripk2          | 0.374 | Receptor-interacting |                  | RIP kinase that potentiates signals                        |
|                |       | serine/threonine-    |                  | downstream of NOD1 and -2, leading to                      |
|                |       | protein kinase 2     |                  | NFkB activation; promotes BCL10                            |
|                |       |                      |                  | phosphorylation and subsequent NFкВ                        |
|                |       |                      |                  | activation following TCR engagement                        |
|                |       |                      | NK Cell Function |  |
|                |       |                      |                  |  |
| <u>Cd69</u>    | 0.997 | Cluster of           | C-type lectin    | Signal transmitting receptor in lymphocytes,               |
|                |       | differentiation 69   | domain family 2, | NK cells, and platelets; induced upon T cell               |
|                |       |                      | member C         | activation; involved in lymphocyte                         |
|                |       |                      |                  | proliferation  |
|                |       |                      |                  |  |
| <u>Gzmb</u>    | 1.2   | Granzyme B           | Fragmentin 2     | Abundant protease in the cytosolic granules                |
|                |       |                      |                  | of cytotoxic T and NK cells that activates                 |
|                |       |                      |                  | caspase-mediated cell death when                           |
|                |       |                      |                  | delivered into the target cell through the                 |
|                |       |                      |                  | immunological synapse                                      |
| <u>Gzmk</u>    | 1.07  | Granzyme K           | Tryptase II      | Granule-secreted, pro-apoptotic serine                     |
| <u> </u>       | 1.07  | Granzyme K           | mypease m        | protease found in the cytoplasmic granules                 |
|                |       |                      |                  |  |
|                |       |                      |                  | of CTLs  |
| <u>ll12rb2</u> | 0.612 | Interleukin 12       |                  | Signaling component coupling to the                        |
|                |       | receptor subunit     |                  | JAK2/STAT4 pathway; promotes T and NK                      |
|                |       | beta 2               |                  | cell proliferation; promotes T <sub>H</sub> 1 polarization |
|                |       |                      |                  | by enhancing IFNγ production                               |
|                |       |                      |                  |  |

| Pik3cd      | 0.741                         | Phosphatidylinositol-  |       | A subunit of PI3K; acts downstream of TLR4, |  |  |
|-------------|-------------------------------|------------------------|-------|---|--|--|
|             |                               | 4,5-bisphosphate 3-    |       | TCR, BCR, and CD40; contributes to T helper |  |  |
|             |                               | kinase catalytic       |       | cell expansion, mast cell development, and  |  |  |
|             |                               | subunit delta isoform  |       | neutrophil chemotaxis, extravasation, and   |  |  |
|             |                               |                        |       | respiratory burst                           |  |  |
| Pik3cg      | 0.387                         | Phosphatidylinositol-  |       | A subunit of PI3K; modulates leukocyte      |  |  |
|             |                               | 4,5-bisphosphate 3-    |       | chemotaxis to inflammatory sites and in     |  |  |
|             |                               | kinase catalytic       |       | response to chemoattractant agents          |  |  |
|             |                               | subunit gamma          |       |   |  |  |
|             |                               | isoform                |       |   |  |  |
| <u>Pvr</u>  | 0.319                         | Poliovirus receptor    | CD155 | Mediates NK cell adhesion and triggers NK   |  |  |
|             |                               |                        |       | cell effector functions; binds CD96 and     |  |  |
|             |                               |                        |       | CD226, leading to the formation of a        |  |  |
|             |                               |                        |       | mature immunological synapse between NK     |  |  |
|             |                               |                        |       | cell and target cell                        |  |  |
| Tcf7        | 1.24                          | Transcription factor 7 |       | HMG box TF predominantly expressed by T     |  |  |
|             |                               |                        |       | cells that drives their development,        |  |  |
|             |                               |                        |       | although also involved in NK cell           |  |  |
|             |                               |                        |       | development; activates transcription        |  |  |
|             |                               |                        |       | through a Wnt/β-catenin signaling pathway   |  |  |
|             | Pattern Recognition Receptors |                        |       |   |  |  |
| <u>Cd14</u> | 0.268                         | Cluster of             |       | PRR that recognizes LPS; mostly found on    |  |  |
|             |                               | differentiation 14     |       | macrophages                                 |  |  |

| <u>Cd180</u>  | 1.2   | Cluster of            |                     | Heterodimeric binding partner of Ly86 that   |
|---------------|-------|-----------------------|---------------------|--|
|               |       | differentiation 180   |                     | participates in LPS binding in APCs          |
| Clec7a        | 0.519 | C-Type lectin domain  | Dectin-1            | PRR specific for β-1,3- and β-1,6-linked     |
|               |       | family 7, member a    |                     | glucans from fungi and plants; necessary for |
|               |       |                       |                     | the TLR2-mediated inflammatory response      |
|               |       |                       |                     | and for TLR2-mediated activation of NF-кВ    |
| Fpr2          | 1.04  | Formyl peptide        | Lipoxin A4 receptor | Low affinity receptor for N-formyl-          |
|               |       | receptor 2            |                     | methionyl peptides; activates neutrophils    |
| <u>Ly96</u>   | 0.285 | Lymphocyte antigen    | Myeloid             | Heterodimeric binding partner of TLR4 that   |
|               |       | 96                    | differentiation     | participates in LPS binding                  |
|               |       |                       | factor 2 (MD-2)     |  |
| Marco         | 2.03  | Macrophage            |                     | A PRR that recognizes LDL                    |
|               |       | receptor with         |                     |  |
|               |       | collagenous structure |                     |  |
| Nlrp3         | 0.652 | NACHT domain-,        | Cryopyrin           | PRR with a wide diversity of recognized      |
|               |       | leucine-rich repeat-, |                     | targets that activates the NLRP3             |
|               |       | and PYD-containing    |                     | inflammasome consisting of NLRP3,            |
|               |       | protein 3             |                     | PYCARD, and caspase-1/-8                     |
| <u>Ticam2</u> | 0.445 | TIR domain-           |                     | Sorting adapter in various innate immune     |
|               |       | containing adaptor    |                     | signaling cascades; bridges TLR2 and MyD88   |
|               |       | molecule 2            |                     |  |

| Tlr1                        | 0.561 | Toll-like receptor 1  Toll-like receptor 7                | CD281   | Ubiquitously expressed surface PRR that recognizes diacylated and triacylated lipopeptides; pairs with TLR2 and CD14  Endosomic PRR that recognizes ssRNA  |  |
|-----------------------------|-------|---|---|--|--|
|                             |       |   |   |  |  |
|                             |       |   | Phagocytosis  |  |  |
| <u>Itgam</u>                | 0.681 | Integrin alpha M  | CD11b   | Pairs with CD18 to form Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated cytotoxicity; serves as a macrophage marker |  |
| Ncf4                        | 0.545 | Neutrophil cytosolic factor 4                             | SH3 and PX domain-<br>containing protein 4<br>(SH3PXD4) | Cytosolic regulatory component of the superoxide-producing phagocyte NADPH-oxidase, a multicomponent enzyme system important for host defense  |  |
| <u>Slamf7</u>               | 1.07  | Signaling lymphocytic activation molecule family member 7 |   | A super-activator of macrophages and a strong promoter of phagocytosis; binds to CD74  |  |
| ROS Generation & Protection |       |   |   |  |  |
| <u>Cybb</u>                 | 0.516 | Cytochrome b-245<br>heavy chain                           | Nox2  | Part of the NADPH oxidase process; generates superoxides   |  |

| Ncf4         | 0.545 | Neutrophil cytosolic   | SH3 and PX domain-   | Cytosolic regulatory component of the        |
|--------------|-------|------------------------|----------------------|--|
| IVELT        | 0.545 |                        |                      |  |
|              |       | factor 4               | containing protein 4 | superoxide-producing phagocyte NADPH-        |
|              |       |                        | (SH3PXD4)            | oxidase, a multicomponent enzyme system      |
|              |       |                        |                      | important for host defense                   |
| Nos2         | 1.58  | Inducible nitric oxide |                      | Produces reactive oxygen species and         |
|              |       | synthase (iNOS)        |                      | contributes to inflammatory cytokine         |
|              |       |                        |                      | production                                   |
|              |       |                        |                      |  |
| <u>Txnip</u> | 0.364 | Thioredoxin            |                      | Thiol-oxidoreductase; protects cells from    |
|              |       | interacting protein    |                      | oxidative stress by inhibiting thioredoxin   |
|              |       |                        | Stress Response      |  |
|              |       |                        | ·                    |  |
| Арр          | 0.157 | Amyloid-beta           |                      | Cell surface receptor and transmembrane      |
|              |       | precursor protein      |                      | precursor protein that is cleaved by         |
|              |       |                        |                      | secretases to form a number of peptides;     |
|              |       |                        |                      | involved in cell mobility, copper            |
|              |       |                        |                      | homeostasis, and oxidative stress            |
| <u>Atm</u>   | 0.29  | Ataxia telangiectasia  |                      | Serine/threonine protein kinase that         |
|              |       | mutated                |                      | activates checkpoint signaling upon DSBs,    |
|              |       |                        |                      | apoptosis, and genotoxic stresses; acts as a |
|              |       |                        |                      | master controller for cell cycle checkpoint  |
|              |       |                        |                      | signaling pathways required for the DNA      |
|              |       |                        |                      | damage response and genomic stability        |
|              |       |                        | T Cell Function      |  |
|              |       |                        |                      |  |

| Cd2         | 1.09  | Cluster of        | Leukocyte            | Interacts with LFA-3 and CD48 to mediate                  |
|-------------|-------|-------------------|----------------------|---|
|             |       | differentiation 2 | functional antigen 2 | adhesion between T cells and other cell                   |
|             |       |                   | (LFA-2)              | types   |
| <u>Cd3d</u> | 1.16  | Cluster of        |                      | Component of the TCR-CD3 complex; upon                    |
|             |       | differentiation 3 |                      | phosphorylation by Lck, serves as a docking               |
|             |       | delta             |                      | station for downstream TCR signaling                      |
|             |       |                   |                      | adaptors  |
| <u>Cd3e</u> | 0.69  | Cluster of        |                      | Component of the TCR-CD3 complex;                         |
|             |       | differentiation 3 |                      | initiates the TCR-CD3 complex assembly by                 |
|             |       | epsilon           |                      | forming the two heterodimers CD3 $\delta$ /CD3 $\epsilon$ |
|             |       |                   |                      | and CD3γ/CD3ε; also participates in                       |
|             |       |                   |                      | internalization and cell surface down-                    |
|             |       |                   |                      | regulation of TCR-CD3 complexes via                       |
|             |       |                   |                      | endocytosis sequences present in CD3ε                     |
|             |       |                   |                      | cytosolic region  |
| Cd3g        | 0.672 | Cluster of        |                      | Component of the TCR-CD3 complex; plays                   |
|             |       | differentiation 3 |                      | an essential role in the dynamic regulation               |
|             |       | gamma             |                      | of TCR expression at the cell surface                     |
| <u>Cd4</u>  | 1.1   | Cluster of        |                      | Signature helper T cell marker; binds to                  |
|             |       | differentiation 4 |                      | MHC class II and provides necessary                       |
|             |       |                   |                      | costimulation for T cell activation                       |
|             |       |                   | <u> </u>             |   |

| <u>Cd5</u>   | 0.972 | Cluster of                        | LEU1   | Type-I transmembrane glycoprotein found  |
|--------------|-------|-----------------------------------|--|--|
|              |       | differentiation 5                 |  | on the surface of T and B cells; may act as a  |
|              |       |                                   |  | receptor in regulating T cell proliferation  |
| <u>Cd8b1</u> | 0.82  | Cluster of differentiation 8 beta |  | Beta chain of the CD8 coreceptor, which binds to MHC class I   |
|              |       | 1                                 |  |  |
| <u>Cd37</u>  | 1.12  | Cluster of differentiation 37     | Tetraspanin-26                                   | Cell surface glycoprotein known to complex with integrins and other transmembrane 4 superfamily proteins; may play a role in T cell-B cell interactions                              |
| <u>Cd69</u>  | 0.997 | Cluster of differentiation 69     | C-type lectin  domain family 2,  member C        | Signal transmitting receptor in lymphocytes,  NK cells, and platelets; induced upon T cell  activation; involved in lymphocyte  proliferation  |
| <u>Cd97</u>  | 0.29  | Cluster of differentiation 97     | BL-Ac[F2]  | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells |
| <u>Cd247</u> | 1.37  | Cluster of differentiation 247    | T cell surface<br>glycoprotein CD3<br>zeta chain | Central intracellular signaling chain of the  TCR, to which downstream signaling  adaptors dock  |

| - 6: -       |       |                       |                     |   |
|--------------|-------|-----------------------|---------------------|---|
| Cyfip2       | 0.877 | Cytoplasmic FMR1-     |                     | Involved in T cell adhesion and p53/TP53-               |
|              |       | interacting protein 2 |                     | dependent induction of apoptosis                        |
| Cxcr3        | 0.768 | C-X-C motif           | CD183               | Induces integrin activation, cytoskeletal               |
|              |       | chemokine receptor    |                     | remodeling, and chemotaxis; expressed by T              |
|              |       | 3                     |                     | cells and NK cells; prominently expressed in            |
|              |       |                       |                     | effector and memory T cells                             |
| Dpp4         | 0.388 | Dipeptidyl-peptidase  | CD26                | Cell surface glycoprotein receptor involved             |
|              |       | 4                     |                     | in the costimulatory signal essential for TCR-          |
|              |       |                       |                     | mediated T cell activation; serine                      |
|              |       |                       |                     | exopeptidase that cleaves various                       |
|              |       |                       |                     | substrates, thereby inactivating them;                  |
|              |       |                       |                     | involved in the migration and invasion of               |
|              |       |                       |                     | endothelial cells into the ECM; also                    |
|              |       |                       |                     | regulates lymphocyte-epithelial cell                    |
|              |       |                       |                     | adhesion  |
| <u>Eomes</u> | 0.488 | Eomesodermin          | T-box brain protein | Transcriptional activator critical for                  |
|              |       |                       | 2 (TBR2)            | development; involved in CD8 <sup>+</sup> T cell        |
|              |       |                       |                     | differentiation   |
| Gata3        | 0.763 | GATA binding protein  |                     | Transcriptional activator that binds to the             |
|              |       | 3                     |                     | enhancer of the TCR $\alpha$ and $\delta$ genes;        |
|              |       |                       |                     | required for T <sub>H</sub> 2 differentiation following |
|              |       |                       |                     | immune and inflammatory responses                       |

| Gzmb         | 1.2   | Granzyme B   | Fragmentin 2 | Abundant protease in the cytosolic granules  |
|--------------|-------|--|--------------|--|
| CEITIO       | ±.£   | S. G. Lyme B   |              |  |
|              |       |  |              | of cytotoxic T and NK cells that activates   |
|              |       |  |              | caspase-mediated cell death when   |
|              |       |  |              | delivered into the target cell through the   |
|              |       |  |              | immunological synapse  |
| <u>Gzmk</u>  | 1.07  | Granzyme K   | Tryptase II  | Granule-secreted, pro-apoptotic serine   |
|              |       |  |              | protease found in the cytoplasmic granules   |
|              |       |  |              | of CTLs  |
| lcos         | 0.984 | Inducible T cell   | CD278        | Enhances all basic T cell responses to   |
|              |       | costimulator   |              | foreign Ag; essential both for efficient   |
|              |       |  |              | interaction between T and B cells and for  |
|              |       |  |              | normal Ab responses to T cell-dependent  |
|              |       |  |              | Ags  |
|              |       |  |              |  |
| <u>lkzf1</u> | 0.994 | IKAROS family zinc   |              | Transcriptional regulator of hematopoietic   |
| <u>lkzf1</u> | 0.994 | IKAROS family zinc   |              | Transcriptional regulator of hematopoietic cell differentiation; plays a role in T and B   |
| <u>lkzf1</u> | 0.994 | ·  |              |  |
| ll12rb2      | 0.994 | ·  |              | cell differentiation; plays a role in T and B  |
|              |       | finger 1   |              | cell differentiation; plays a role in T and B  cell development  |
|              |       | finger 1  Interleukin 12   |              | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the   |
|              |       | finger 1  Interleukin 12  receptor subunit                         |              | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  |
|              |       | finger 1  Interleukin 12  receptor subunit                         |              | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  cell proliferation; promotes T <sub>H</sub> 1 polarization  |
|              |       | finger 1  Interleukin 12  receptor subunit                         | LYK          | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  cell proliferation; promotes T <sub>H</sub> 1 polarization  |
| ll12rb2      | 0.612 | finger 1  Interleukin 12  receptor subunit  beta 2                 | LYK          | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  cell proliferation; promotes T <sub>H</sub> 1 polarization  by enhancing IFNγ production  |
| ll12rb2      | 0.612 | finger 1  Interleukin 12  receptor subunit  beta 2  Interleukin-2- | LYK          | cell differentiation; plays a role in T and B  cell development  Signaling component coupling to the  JAK2/STAT4 pathway; promotes T and NK  cell proliferation; promotes T <sub>H</sub> 1 polarization  by enhancing IFNy production  Key actor in the TCR signaling cascade; |

| <u>Lck</u> | 0.983 | Lymphocyte cell kinase  |           | Src family tyrosine kinase that acts as one of the main signaling intermediaries downstream of the TCR; constitutively   |
|------------|-------|---|-----------|--|
|            |       |   |           | associated with the cytoplasmic portion of   |
| Lcp1       | 0.504 | Lymphocyte cytosolic protein 1  | Plastin-2 | Actin-binding protein that promotes T cell activation in response to costimulation through TCR/CD3 and CD2 or CD28; assists with IL2RA transport to the cell surface                           |
| Nfatc1     | 0.435 | Nuclear factor of activated T cells, cytoplasmic 1  |           | Inducible nuclear component of the NFAT  TF complex; mediates induction of IL-2 and  IL-4 in T cells   |
| Nfatc2     | 0.602 | Nuclear factor of activated T cells, cytoplasmic 2  |           | Cytosolic component of the NFAT TF complex; mediates induction of IL-2, IL-3, IL-4, TNF $\alpha$ , and GM-CSF  |
| Pik3cd     | 0.741 | Phosphatidylinositol-<br>4,5-bisphosphate 3-<br>kinase catalytic<br>subunit delta isoform |           | A subunit of PI3K; acts downstream of TLR4, TCR, BCR, and CD40; contributes to T helper cell expansion, mast cell development, and neutrophil chemotaxis, extravasation, and respiratory burst |
| Pik3cg     | 0.387 | Phosphatidylinositol- 4,5-bisphosphate 3- kinase catalytic                                |           | A subunit of PI3K; modulates leukocyte chemotaxis to inflammatory sites and in response to chemoattractant agents  |

|       |       | subunit gamma<br>isoform                       |                                      |   |
|-------|-------|--|--------------------------------------|---|
| Ptprc | 0.937 | Protein tyrosine  phosphatase  receptor type C | CD45; leukocyte common antigen (LCA) | Delivers costimulation during T cell activation upon binding to its ligand DPP4; dephosphorylates Lyn and suppresses JAK kinases  |
| Spn   | 0.812 | Sialophorin                                    | Leukosialin; CD43                    | T cells, B cells, monocytes, and granulocytes; promotes lymph node localization in T cells; shunts T cells away from the T <sub>H</sub> 2 phenotype and towards T <sub>H</sub> 1; promotes the expression of IFNγ in CD4 <sup>+</sup> T cells |
| Tbx21 | 0.624 | T-box transcription<br>factor 21               |                                      | Initiates $T_H1$ lineage development from naïve $T_H$ precursor cells both by activating $T_H1$ genetic programs and by repressing the opposing $T_H2$ and $T_H17$ genetic programs   |
| Tcf7  | 1.24  | Transcription factor 7                         |                                      | HMG box TF predominantly expressed by T cells that drives their development, although also involved in NK cell development; activates transcription through a Wnt/β-catenin signaling pathway   |

| Txk          | 1.07                                 | TXK tyrosine kinase  | Regulates the development, function, and       |  |  |
|--------------|--------------------------------------|----------------------|--|--|--|
| TAK          | 1.07                                 | TAR CYTOSITIC KITUSC |  |  |  |
|              |                                      |                      | differentiation of conventional T cells and    |  |  |
|              |                                      |                      | nonconventional NK-T cells; contributes to     |  |  |
|              |                                      |                      | signaling from many receptors and              |  |  |
|              |                                      |                      | participates in multiple downstream            |  |  |
|              |                                      |                      | pathways, including regulation of the actin    |  |  |
|              |                                      |                      | cytoskeleton; can phosphorylate PLCγ1,         |  |  |
|              |                                      |                      | leading to its localization in lipid rafts and |  |  |
|              |                                      |                      | activation, followed by subsequent cleavage    |  |  |
|              |                                      |                      | of its substrates                              |  |  |
| <u>Txnip</u> | 0.364                                | Thioredoxin          | Thiol-oxidoreductase; protects cells from      |  |  |
|              |                                      | interacting protein  | oxidative stress by inhibiting thioredoxin     |  |  |
| <u>Zap70</u> | 0.772                                | Zeta chain of T cell | Tyrosine kinase that plays an essential role   |  |  |
|              |                                      | receptor associated  | in regulation of the adaptive immune           |  |  |
|              |                                      | protein kinase 70    | response; regulates motility, adhesion,        |  |  |
|              |                                      |                      | proliferation, differentiation, and cytokine   |  |  |
|              |                                      |                      | expression in T cells; contributes to the      |  |  |
|              |                                      |                      | development and activation of primary B        |  |  |
|              |                                      |                      | cells; phosphorylates at least two essential   |  |  |
|              |                                      |                      | adapter proteins: LAT and LCP2, leading in     |  |  |
|              |                                      |                      | turn to a large number of signaling            |  |  |
|              |                                      |                      | molecules being recruited                      |  |  |
|              |                                      | _                    |  |  |  |
|              | Transcription Factors & Coactivators |                      |  |  |  |

| Foxp3        | 1.02  | Forkhead box P3       | DIETER              | Master TF for Tregs; represses expression of            |
|--------------|-------|-----------------------|---------------------|---|
|              |       |                       |                     | II2 and Ifng; activates expression of                   |
|              |       |                       |                     | Tnfrsf18, Il2ra, and Ctla4                              |
| Eomes        | 0.488 | Eomesodermin          | T-box brain protein | Transcriptional activator critical for                  |
|              |       |                       | 2 (TBR2)            | development; involved in CD8 <sup>+</sup> T cell        |
|              |       |                       |                     | differentiation   |
| Gata3        | 0.763 | GATA binding protein  |                     | Transcriptional activator that binds to the             |
|              |       | 3                     |                     | enhancer of the TCR $\alpha$ and $\delta$ genes;        |
|              |       |                       |                     | required for T <sub>H</sub> 2 differentiation following |
|              |       |                       |                     | immune and inflammatory responses                       |
| <u>Irf1</u>  | 0.693 | Interferon regulatory |                     | Transcriptional regulator that promotes                 |
|              |       | factor 1              |                     | inflammatory innate and adaptive immune                 |
|              |       |                       |                     | responses   |
| <u>Irf4</u>  | 0.927 | Interferon regulatory |                     | Transcriptional activator that complexes                |
|              |       | factor 4              |                     | with BATF and binds ISREs within the                    |
|              |       |                       |                     | promoters of multiple genes involved in                 |
|              |       |                       |                     | inflammation  |
| <u>lkzf1</u> | 0.994 | IKAROS family zinc    |                     | Transcriptional regulator of hematopoietic              |
|              |       | finger 1              |                     | cell differentiation; plays a role in T and B           |
|              |       |                       |                     | cell development  |
| <u>lkzf2</u> | 0.383 | IKAROS family zinc    |                     | Hematopoietic cell-specific TF involved in              |
|              |       | finger protein 2      |                     | early hematopoietic development                         |

| Nfatc1     | 0.435  | Nuclear factor of      | Inducible nuclear component of the NFAT                          |
|------------|--------|------------------------|--|
|            |        | activated T cells,     | TF complex; mediates induction of IL-2 and                       |
|            |        | cytoplasmic 1          | IL-4 in T cells  |
| Nfatc2     | 0.602  | Nuclear factor of      | Cytosolic component of the NFAT TF                               |
|            |        | activated T cells,     | complex; mediates induction of IL-2, IL-3, IL-                   |
|            |        | cytoplasmic 2          | 4, TNFα, and GM-CSF  |
| Stat1      | 0.675  | Signal transducer and  | Transcriptional activator that mediates                          |
|            |        | activator of           | cellular responses to IFNs, cytokines, and                       |
|            |        | transcription 1        | other growth factors   |
| Stat6      | 0.187  | Signal transducer and  | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and    |
|            |        | activator of           | macrophage function and M2 macrophage                            |
|            |        | transcription 6        | polarization   |
| Tbx21      | 0.624  | T-box transcription    | Initiates T <sub>H</sub> 1 lineage development from              |
|            |        | factor 21              | naïve T <sub>H</sub> precursor cells both by activating          |
|            |        |                        | T <sub>H</sub> 1 genetic programs and by repressing the          |
|            |        |                        | opposing T <sub>H</sub> 2 and T <sub>H</sub> 17 genetic programs |
| Tcf7       | 1.24   | Transcription factor 7 | HMG box TF predominantly expressed by T                          |
|            |        |                        | cells that drives their development,                             |
|            |        |                        | although also involved in NK cell                                |
|            |        |                        | development; activates transcription                             |
|            |        |                        | through a Wnt/β-catenin signaling pathway                        |
| <u>Yy1</u> | 0.0857 | Yin yang 1             | Ubiquitous factor that serves as a                               |
|            |        |                        | transcriptional "switch", either promoting                       |

|                |       |                         |                            | or repressing the transcription of numerous genes through the selective recruitment of either histone deacetylases or acetyltransferases; plays a fundamental role in diverse processes, such as differentiation, replication, and cellular proliferation |
|----------------|-------|-------------------------|----------------------------|---|
| Zbp1           | 0.822 | Z-DNA binding protein 1 | Tumor stroma and activated | Key innate sensor that recognizes and binds  Z-RNA structures, which are produced by a  |
|                |       |                         | macrophage protein         | number of viruses, and induces type-I IFN   |
|                |       |                         | DLM-1                      | production; key activator of cellular   |
|                |       |                         |                            | necroptosis; promotes IL-1α induction in an   |
|                |       |                         |                            | NLRP3-inflammasome-independent manner   |
|                |       | l                       | <br>  Jbiquitin Regulation |   |
|                |       |                         |                            |   |
| <u>Bcl10</u>   | 0.137 | B cell                  |                            | Activates NFκB via ubiquitination of ΙΚΚγ   |
|                |       | lymphoma/leukemia       |                            |   |
|                |       | 10                      |                            |   |
| Cyld           | 0.305 | Cylindromatosis         |                            | Inhibits NFkB activation by deubiquitinating  |
|                |       | lysine 63               |                            | upstream signaling factors; inhibits Wnt  |
|                |       | deubiquitinase          |                            | signaling; restricts polyubiquitination of  |
|                |       |                         |                            | RIPK1 and -2, thereby limiting necroptosis  |
|                |       |                         |                            |   |
| <u>Tnfaip3</u> | 0.387 | Tumor necrosis          |                            | Ubiquitin-editing enzyme that complexes   |
|                |       | factor, alpha-induced   |                            | with ITCH to degrade inflammatory   |
|                |       | protein 3               |                            | signaling components in the TNF, IL1, and   |
|                |       |                         |                            |   |

|  |  | TLR pathways; targets TRAF2, TRAF6, and |
|--|--|---|
|  |  | IKK                                     |
|  |  |   |

|      | NBTXR3+PBT+αPD1 vs PBT+αPD1       |                            |                      |  |  |
|------|-----------------------------------|----------------------------|----------------------|--|--|
| Gene | Log2 fold                         | Full Name                  | Notable Aliases      | Function   |  |
|      |                                   | А                          | cute Phase Response  |  |  |
| Cma1 | 0.453                             | Chymase 1                  | Mast cell protease 1 | Serine protease with pro-inflammatory peptidolytic activation predominantly secreted by mast cells                         |  |
| Dpp4 | 0.585                             | Dipeptidyl-<br>peptidase 4 | CD26                 | Serine exopeptidase that cleaves various substrates, thereby inactivating them   |  |
| Cfd  | 1.33                              | Complement factor D        | Adipsin              | Chymotrypsin-family peptidase that cleaves factor B when the latter is complexed with factor C3b, activating C3 convertase |  |
|      | Adhesion & Cell-Cell Interactions |                            |                      |  |  |
| Sell | 0.962                             | L-selectin                 |                      | Mediates cell adhesion by binding to glycoproteins on neighboring cells  |  |
|      |                                   | Antigen                    | Processing & Present | ation  |  |

| <u>Cd1d1</u> | 0.7   | Cluster of differentiation 1 D1 |                       | Murine non-classical class I MHC; primarily presents lipid and glycolipid Ags |  |  |
|--------------|-------|---------------------------------|-----------------------|---|--|--|
|              |       |                                 | Apoptosis             |   |  |  |
| Tof          |       |                                 |                       |   |  |  |
| <u>Tnf</u>   | 1.12  | Tumor necrosis                  | Cachectin             | Inflammatory cytokine mainly produced by                                      |  |  |
|              |       | factor                          |                       | macrophages; binds to TNFRSF1A/TNFR1  |  |  |
|              |       |                                 |                       | and TNFRSF1B/TNFBR; capable of inducing                                       |  |  |
|              |       |                                 |                       | cell death in certain tumor cell lines  |  |  |
| Nlrp3        | 0.865 | NACHT domain-,                  | Cryopyrin             | PRR with a wide diversity of recognized                                       |  |  |
|              |       | leucine-rich repeat-,           |                       | targets that activates the NLRP3  |  |  |
|              |       | and PYD-containing              |                       | inflammasome consisting of NLRP3,   |  |  |
|              |       | protein 3                       |                       | PYCARD, and caspase-1/-8  |  |  |
|              |       | В                               | Cell-associated Genes |   |  |  |
| Dink         | 1.02  | D call linker                   | Src hamalagy 1        | Functions as a control linker protein   |  |  |
| <u>Blnk</u>  | 1.02  | B cell linker                   | Src homology 1        | Functions as a central linker protein   |  |  |
|              |       |                                 | domain-containing     | downstream of the B cell receptor, bridging                                   |  |  |
|              |       |                                 | leukocyte protein of  | SYK kinase to a multitude of signaling  |  |  |
|              |       |                                 | 65 kDa (SLP-65);      | pathways and regulating biological  |  |  |
|              |       |                                 | Ly57                  | outcomes of B cell function and   |  |  |
|              |       |                                 |                       | development; plays a role in the activation                                   |  |  |
|              |       |                                 |                       | of ERK/EPHB2, MAP kinase p38 and JNK;   |  |  |
|              |       |                                 |                       | modulates AP1 activation; important for the                                   |  |  |
|              |       |                                 |                       | activation of NFкВ and NFAT   |  |  |
| <u>Btk</u>   | 0.885 | Bruton's tyrosine               |                       | Crucial kinase in B cell receptor signal                                      |  |  |

|             |       | kinase                        |   | transmission and B cell activation  |
|-------------|-------|-------------------------------|---|---|
| <u>Btla</u> | 1.68  | B and T lymphocyte attenuator | CD272   | Inhibitory cell surface protein that inhibits T  cell function by binding to B7H4 and  TNFRSF14   |
| <u>Cd19</u> | 2.14  | Cluster of differentiation 19 | B-lymphocyte<br>surface antigen B4  | BCR coreceptor; activates signaling pathways that lead to the activation of PI3K and Ca(2+) flux  |
| <u>Cd22</u> | 1.67  | Cluster of differentiation 22 | Sialic acid-binding Ig-<br>like lectin 2<br>(SIGLEC2)<br>B lymphocyte cell<br>adhesion molecule<br>(BL-CAM) | Mediates interactions between B cells; binds CD45   |
| Cd40lg      | 1.48  | CD40 ligand                   | CD154; TNF-related activation protein (TRAP) T-B cell activating molecule (T-BAM)                           | Costimulates T cell proliferation and cytokine production; enhances IL-4 and IL- 10 production in conjunction with TCR/CD3 ligation and CD28 costimulation; induces  NFkB activation; mediates B cell proliferation in the absence of costimulation; ligand for integrins ITGA5:ITGB1 and ITGAV:ITGB3 |
| <u>Cd69</u> | 0.867 | Cluster of differentiation 69 | C-type lectin domain family 2, member C   | Signal transmitting receptor in lymphocytes,  NK cells, and platelets; induced upon T cell  |

|              |               |   |            | activation; involved in lymphocyte proliferation   |  |  |
|--------------|---------------|---|------------|--|--|--|
| Cd79b        | 2.56          | Cluster of differentiation 79b              | B29        | One of the two flanking proteins that initiate signaling downstream of the BCR   |  |  |
| <u>lcosl</u> | 0.897         | Inducible T cell costimulator ligand        | CD275      | Ligand for T cell-specific co-receptor ICOS; also induces B cell proliferation and plasma cell differentiation                             |  |  |
| <u>lkzf1</u> | 1.05          | IKAROS family zinc finger 1                 |            | Transcriptional regulator of hematopoietic cell differentiation; plays a role in T and B cell development                                  |  |  |
| <u>Ms4a1</u> | 2.07          | Membrane spanning 4-domains A1              | CD20; Bp35 | B cell-specific membrane protein that functions as a store-operated Ca(2+) channel component, promoting Ca(2+) influx after BCR activation |  |  |
| Pou2f2       | 1.12          | POU domain class 2,<br>transcription factor |            | TF that regulates Ab and IL-6 expression in  B cells   |  |  |
| Tnfrsf13c    | 2.38          | TNF receptor superfamily member 13C         |            | B cell-activating factor; enhances B cell survival in vitro; regulator of the peripheral B cell population                                 |  |  |
|              | Costimulation |   |            |  |  |  |

| Cd40lg       | 1.48  | CD40 ligand         | CD154; TNF-related  | Costimulates T cell proliferation and        |
|--------------|-------|---------------------|---------------------|--|
|              |       |                     | activation protein  | cytokine production; enhances IL-4 and IL-   |
|              |       |                     | (TRAP)              | 10 production in conjunction with TCR/CD3    |
|              |       |                     | T-B cell activating | ligation and CD28 costimulation; induces     |
|              |       |                     | molecule (T-BAM)    | NFκB activation; mediates B cell             |
|              |       |                     |                     | proliferation in the absence of              |
|              |       |                     |                     | costimulation; ligand for integrins          |
|              |       |                     |                     | ITGA5:ITGB1 and ITGAV:ITGB3                  |
| <u>lcosl</u> | 0.897 | Inducible T cell    | CD275               | Ligand for T cell-specific co-receptor ICOS; |
|              |       | costimulator ligand |                     | also induces B cell proliferation and plasma |
|              |       |                     |                     | cell differentiation                         |
|              |       |                     | Cytokines           |  |
|              |       |                     | Cytokiiics          |  |
| <u>II1b</u>  | 0.765 | Interleukin 1 beta  | Catabolin           | One of the two primary inflammatory          |
|              |       |                     |                     | cytokines produced by the inflammasome       |
|              |       |                     |                     | (the other one being IL-18); induces         |
|              |       |                     |                     | neutrophil influx and activation, T cell     |
|              |       |                     |                     | activation and cytokine production, B cell   |
|              |       |                     |                     | activation and Ab production, fibroblast     |
|              |       |                     |                     | proliferation, and collagen production;      |
|              |       |                     |                     | synergizes with IL-12 to induce IFNγ         |
|              |       |                     |                     | synthesis from T <sub>H</sub> 1 cells        |
| <u>II7r</u>  | 1.02  | Interleukin 7       | CD127               | Receptor for IL-7                            |
|              |       | receptor            |                     |  |
|              |       |                     |                     |  |

| _           |       |                     |                     |  |
|-------------|-------|---------------------|---------------------|--|
| <u>Tnf</u>  | 1.12  | Tumor necrosis      | Cachectin           | Inflammatory cytokine mainly produced by     |
|             |       | factor              |                     | macrophages; binds to TNFRSF1A/TNFR1         |
|             |       |                     |                     | and TNFRSF1B/TNFBR; capable of inducing      |
|             |       |                     |                     | cell death in certain tumor cell lines       |
|             |       | G                   | rowth/Proliferation |  |
|             |       |                     | •                   |  |
| Rps6        | 0.606 | Ribosomal protein 6 |                     | Component of the 40S small ribosomal         |
|             |       |                     |                     | subunit; plays an important role in          |
|             |       |                     |                     | controlling cell growth and proliferation    |
|             |       |                     |                     | through the selective translation of         |
|             |       |                     |                     | particular classes of mRNA                   |
| Tyk2        | 0.74  | Tyrosine kinase 2   | JTK1                | Plays both structural and catalytic roles in |
|             |       |                     |                     | numerous cytokines and interferons           |
|             |       |                     |                     | signaling; associates with cytokine and      |
|             |       |                     |                     | growth factor receptors and activate STAT    |
|             |       |                     |                     | family members including STAT1, STAT3,       |
|             |       |                     |                     | STAT4, or STAT6                              |
|             |       |                     | Inflammation        |  |
|             |       |                     |                     |  |
| <u>II1b</u> | 0.765 | Interleukin 1 beta  | Catabolin           | One of the two primary inflammatory          |
|             |       |                     |                     | cytokines produced by the inflammasome       |
|             |       |                     |                     | (the other one being IL-18); induces         |
|             |       |                     |                     | neutrophil influx and activation,T cell      |
|             |       |                     |                     | activation and cytokine production, B cell   |
|             |       |                     |                     | activation and Ab production, fibroblast     |
|             |       | 1                   |                     |  |

| <u>Irf4</u> | 0.828 | Interferon   |            | proliferation, and collagen production; synergizes with IL-12 to induce IFN $\gamma$ synthesis from $T_H1$ cells   |
|-------------|-------|--|------------|--|
|             |       | regulatory factor 4  |            | with BATF and binds ISREs within the promoters of multiple genes involved in inflammation  |
| Nlrp3       | 0.865 | NACHT domain-,<br>leucine-rich repeat-,<br>and PYD-containing<br>protein 3 | Cryopyrin  | PRR with a wide diversity of recognized targets that activates the NLRP3 inflammasome consisting of NLRP3, PYCARD, and caspase-1/-8                          |
| <u>Tnf</u>  | 1.12  | Tumor necrosis<br>factor   | Cachectin  | Inflammatory cytokine mainly produced by macrophages; binds to TNFRSF1A/TNFR1 and TNFRSF1B/TNFBR; capable of inducing cell death in certain tumor cell lines |
|             |       |  | Inhibition |  |
| <u>Btla</u> | 1.68  | B and T lymphocyte attenuator  | CD272      | Inhibitory cell surface protein that inhibits T cell function by binding to B7H4 and TNFRSF14  |
| Foxp3       | 0.544 | Forkhead box P3  | DIETER     | Master TF for Tregs; represses expression of  II2 and Ifng; activates expression of  Tnfrsf18, II2ra, and Ctla4  |

| 0.442 | Nuclear factor      | Inhibits activity of REL dimers by masking of                         |
|-------|---------------------|---|
|       | kappa B inhibitor   | their nuclear localization signals                                    |
|       | alpha               |   |
|       |                     |   |
| 0.6   | Src homology 2B     | Adapter protein for several members of the                            |
|       | adaptor protein 2   | tyrosine kinase receptor family; involved in                          |
|       |                     | multiple signaling pathways; may be                                   |
|       |                     | involved in coupling from immunoreceptor                              |
|       |                     | to Ras signaling; acts as a negative regulator                        |
|       |                     | of cytokine signaling in collaboration with                           |
|       |                     | CBL; may induce cytoskeletal reorganization                           |
|       |                     | via interaction with Vav3   |
|       |                     |   |
|       |                     | Kinases   |
| 0.885 | Bruton's tyrosine   | Crucial kinase in B cell receptor signal                              |
|       | kinase              | transmission and B cell activation                                    |
| 0.93  | TXK tyrosine kinase | Regulates the development, function, and                              |
|       |                     | differentiation of conventional T cells and                           |
|       |                     | nonconventional NK-T cells; contributes to                            |
|       |                     | signaling from many receptors and                                     |
|       |                     | participates in multiple downstream                                   |
|       |                     | pathways, including regulation of the actin                           |
|       |                     | cytoskeleton; can phosphorylate PLCγ1,                                |
|       |                     | leading to its localization in lipid rafts and                        |
|       |                     | activation, followed by subsequent cleavage                           |
|       | 0.6                 | 0.6 Src homology 2B adaptor protein 2  0.885 Bruton's tyrosine kinase |

|               |       |   |                     | of its substrates  |
|---------------|-------|---|---------------------|--|
| Tyk2          | 0.74  | Tyrosine kinase 2   | JTK1                | Plays both structural and catalytic roles in numerous cytokines and interferons signaling; associates with cytokine and growth factor receptors and activate STAT family members including STAT1, STAT3, STAT4, or STAT6 |
|               |       | N   | lacrophage Function |  |
| Marco Slc11a1 | 0.667 | Macrophage receptor with collagenous structure  Natural resistance- associated macrophage protein |                     | A PRR that recognizes LDL  Macrophage-specific metal ion transporter; uptakes divalent metal cations to neutralize  ROSs   |
|               |       | 1   |                     |  |
|               |       |   | Metabolism          |  |
| Abcg1         | 1.05  | ATP-binding cassette transporter G1   |                     | Membrane-associated cholesterol efflux pump  |
|               |       |   | Migration/Motility  |  |

| Ccl3   | 0.512 | C-C motif           | Macrophage                         | Chemoattractant ligand for CCR1, -4, and -5                 |
|--------|-------|---------------------|------------------------------------|---|
| 33.0   | 0.011 |                     |                                    | gana ia sana, iyana a                                       |
|        |       | chemokine ligand 3  | inflammatory                       |   |
|        |       |                     | protein $1\alpha$ (MIP $1\alpha$ ) |   |
|        |       |                     |                                    |   |
| Ccr2   | 0.694 | C-C motif           | CD192                              | Receptor for CCL2, a monocyte-specific                      |
|        |       | chemokine receptor  |                                    | chemokine   |
|        |       | 2                   |                                    |   |
|        |       |                     |                                    |   |
| Ccr6   | 1.74  | C-C motif           | CD196                              | Receptor for CCL20 and β-defensins; signals                 |
|        |       | chemokine receptor  |                                    | through Ca(2+) flux; controls chemotaxis of                 |
|        |       | 6                   |                                    | DCs and effector/memory T cells and B                       |
|        |       |                     |                                    | cells; essential for recruitment of T <sub>H</sub> 17 cells |
|        |       |                     |                                    | and Tregs   |
|        |       |                     |                                    |   |
| Cxcr5  | 1.83  | C-X-C motif         | CD185; Burkitt's                   | Cytokine receptor that binds to B                           |
|        |       | chemokine receptor  | lymphoma receptor                  | lymphocyte chemoattractant (BLC);                           |
|        |       | 5                   | 1 (BLR1)                           | involved in B cell migration into splenic                   |
|        |       |                     |                                    | follicles and Peyer's patches                               |
|        |       |                     |                                    |   |
|        |       |                     | NFkB Signaling                     |   |
| Card11 | 1.33  | Caspase recruitment | Bcl10-interacting                  | Adapter protein that plays a key role in                    |
| 001011 | 1.33  |                     |                                    |   |
|        |       | domain family       | MAGUK protein 3                    | adaptive immune response by transducing                     |
|        |       | member 11           | (BIMP3)                            | the activation of NFkB downstream of TCR                    |
|        |       |                     |                                    | and BCR engagement; upon activation in                      |
|        |       |                     |                                    | response to TCR or BCR triggering,                          |
|        |       |                     |                                    | homooligomerizes to form a nucleating                       |
|        |       |                     |                                    | helical template that recruits BCL10 via                    |
|        |       |                     |                                    |   |

| Ikbke        | 0.776 | Inhibitor of nuclear factor kappa B kinase subunit epsilon        |                       | CARD-CARD interaction, thereby promoting polymerization of BCL10 and subsequent recruitment of MALT1, leading to IKK phosphorylation and degradation and release of NFκB proteins for nuclear translocation  Noncanonical IκB kinase; plays an essential role in regulating inflammatory through the activation of the type I IFN, NFκB, and STAT signaling |
|--------------|-------|---|-----------------------|---|
|              |       | Patte   | rn Recognition Recept | tors  |
| <u>Cd180</u> | 1.33  | Cluster of differentiation 180                                    |                       | Heterodimeric binding partner of Ly86 that participates in LPS binding in APCs  |
| <u>Marco</u> | 1.37  | Macrophage receptor with collagenous structure                    |                       | A PRR that recognizes LDL   |
| Nlrp3        | 0.865 | NACHT domain-, leucine-rich repeat-, and PYD-containing protein 3 | Cryopyrin             | PRR with a wide diversity of recognized targets that activates the NLRP3 inflammasome consisting of NLRP3,  PYCARD, and caspase-1/-8  |
| <u>Tlr7</u>  | 0.876 | Toll-like receptor 7  | CD287                 | Endosomic PRR that recognizes ssRNA   |

|              | T Cell Function |                     |                      |  |
|--------------|-----------------|---------------------|----------------------|--|
| <u>Cd69</u>  | 0.867           | Cluster of          | C-type lectin domain | Signal transmitting receptor in lymphocytes,   |
|              |                 | differentiation 69  | family 2, member C   | NK cells, and platelets; induced upon T cell   |
|              |                 |                     |                      | activation; involved in lymphocyte             |
|              |                 |                     |                      | proliferation                                  |
| <u>lkzf1</u> | 1.05            | IKAROS family zinc  |                      | Transcriptional regulator of hematopoietic     |
|              |                 | finger 1            |                      | cell differentiation; plays a role in T and B  |
|              |                 |                     |                      | cell development                               |
| <u>Txk</u>   | 0.93            | TXK tyrosine kinase |                      | Regulates the development, function, and       |
|              |                 |                     |                      | differentiation of conventional T cells and    |
|              |                 |                     |                      | nonconventional NK-T cells; contributes to     |
|              |                 |                     |                      | signaling from many receptors and              |
|              |                 |                     |                      | participates in multiple downstream            |
|              |                 |                     |                      | pathways, including regulation of the actin    |
|              |                 |                     |                      | cytoskeleton; can phosphorylate PLCγ1,         |
|              |                 |                     |                      | leading to its localization in lipid rafts and |
|              |                 |                     |                      | activation, followed by subsequent cleavage    |
|              |                 |                     |                      | of its substrates                              |

## Supplemental Table 3: Differentially expressed genes in various immune cells

| Markers | Cluster     | Markers | Cluster   | Markers | Cluster           |
|---------|-------------|---------|-----------|---------|-------------------|
| Mrc1    | Macrophages | Ms4a4c  | Monocytes | Cd8a    | Cytotoxic T cells |
| Sdc3    | Macrophages | Ms4a6c  | Monocytes | Cd8b1   | Cytotoxic T cells |
| Msr1    | Macrophages | Plac8   | Monocytes | Lag3    | Cytotoxic T cells |
| Lrp1    | Macrophages | Tgfbi   | Monocytes | Nkg7    | Cytotoxic T cells |

| Curl4C | NA           | DA-4-4- | N.A              | Count    | O 4-4-3-7         |
|--------|--------------|---------|------------------|----------|-------------------|
| Cxcl16 | Macrophages  | Ms4a4a  | Monocytes        | Gzmb     | Cytotoxic T cells |
| Grn    | Macrophages  | Gm9733  | Monocytes        | Gzmk     | Cytotoxic T cells |
| Lgmn   | Macrophages  | Plbd1   | Monocytes        | Klrc1    | Cytotoxic T cells |
| Dab2   | Macrophages  | Ifitm3  | Monocytes        | Prf1     | Cytotoxic T cells |
| Trem2  | Macrophages  | Lrp1    | Monocytes        | Cxcr6    | Cytotoxic T cells |
| Csf1r  | Macrophages  | H2-DMb1 | Monocytes        | Pdcd1    | Cytotoxic T cells |
| Cd4    | CD4+ T cells | Cd8a    | CD8+ T cells     | Klra4    | NK cells          |
| Tcf7   | CD4+ T cells | Cd8b1   | CD8+ T cells     | Ncr1     | NK cells          |
| Nsg2   | CD4+ T cells | Lat     | CD8+ T cells     | Klra7    | NK cells          |
| S1pr1  | CD4+ T cells | Fcer1g  | CD8+ T cells     | Klra1    | NK cells          |
| Ccr7   | CD4+ T cells | Rps7    | CD8+ T cells     | Trdc     | NK cells          |
| Tnfsf8 | CD4+ T cells | Rps5    | CD8+ T cells     | Gzma     | NK cells          |
| Tbc1d4 | CD4+ T cells | Rpsa    | CD8+ T cells     | Car2     | NK cells          |
| Ets1   | CD4+ T cells | Lck     | CD8+ T cells     | Nrarp    | NK cells          |
| Lrig1  | CD4+ T cells | Rpl30   | CD8+ T cells     | Klrb1f   | NK cells          |
| Lck    | CD4+ T cells | Rps15a  | CD8+ T cells     | Atp1b1   | NK cells          |
| Rpl7   | Neutrophils  | Flt3    | DC               | Kif11    | Tgd               |
| Cxcr2  | Neutrophils  | 3-Sep   | DC               | Cd8b1    | Tgd               |
| G0s2   | Neutrophils  | Clec9a  | DC               | Cd8a     | Tgd               |
| Rpl7a  | Neutrophils  | Gcsam   | DC               | Cd3e     | Tgd               |
| Hdc    | Neutrophils  | Mycl    | DC               | Lat      | Tgd               |
| Cebpb  | Neutrophils  | Tspan33 | DC               | Thy1     | Tgd               |
| Rps3a1 | Neutrophils  | Timd4   | DC               | Cd3d     | Tgd               |
| S100a8 | Neutrophils  | Cbfa2t3 | DC               | Trbc2    | Tgd               |
| S100a9 | Neutrophils  | Hmgn3   | DC               | 1-Sep    | Tgd               |
| Mcl1   | Neutrophils  | Gnb4    | DC               | Cd3g     | Tgd               |
| Rps20  | Neutrophils  | Nkg7    | ILC              | Foxp3    | Tregs             |
| Rps8   | Neutrophils  | Klrd1   | ILC              | Cd4      | Tregs             |
| Csf3r  | Neutrophils  | II2rb   | ILC              | Rln3     | Tregs             |
| Rpl11  | Neutrophils  | Klrc1   | ILC              | lzumo1r  | Tregs             |
| Cxcl2  | Neutrophils  | Ctsw    | ILC              | Tnfrsf4  | Tregs             |
| Rplp0  | Neutrophils  | Ctla2a  | ILC              | lkzf2    | Tregs             |
| Rpl6   | Neutrophils  | Thy1    | ILC              | Tnfrsf18 | Tregs             |
| Tmsb10 | Neutrophils  | Klrk1   | ILC              | Cd2      | Tregs             |
|        | ·            |         |                  |          |                   |
| Rpl32  | Neutrophils  | Ms4a4b  | ILC              | lgsf23   | Tregs             |
| Rpl28  | Neutrophils  | Klre1   | ILC              | Neb      | Tregs             |
| Rps5   | Neutrophils  | Ccr7    | CD4-CD8- T cells | Cd79a    | B cells           |
| Rps19  | Neutrophils  | S1pr1   | CD4-CD8- T cells | Ebf1     | B cells           |
| Rps11  | Neutrophils  | Tcf7    | CD4-CD8- T cells | Ly6d     | B cells           |
| Rpl18  | Neutrophils  | Lat     | CD4-CD8- T cells | Iglc3    | B cells           |
| Rps3   | Neutrophils  | Ets1    | CD4-CD8- T cells | Iglc2    | B cells           |

| Rps15a Neutrophils Plek CD4-CD8-T cells Cd79b B cells Rpl13 Neutrophils Lgals3 CD4-CD8-T cells Spib B cells Eef1a1 Neutrophils lifitm2 CD4-CD8-T cells Igkc B cells Rpsa Neutrophils Ctsd CD4-CD8-T cells Snn B cells Rpsa Neutrophils Ctsd CD4-CD8-T cells Snn B cells Rpl14 Neutrophils Cd3e NKT Fcer1a Mast cells Srgn Neutrophils Cd3d NKT Cpa3 Mast cells Eef2 Neutrophils Cd3g NKT Car8 Mast cells Ppia Neutrophils Icos NKT Atp8b5 Mast cells Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Cd3e NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Ptprcap NKT Cd200r3 Mast cells Rps23 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Neutrophils Rps8 Neutrophils Rps9 Neutrophils Rps9 Neutrophils Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps11 Neutrophils Rps11 Neutrophils Rps12 Neutrophils Rps2 Neutrophils Rps3 Neutrophils Rps10 Neutrophils Rps11 Neutrophils Rps2 Neutrophils Rps12 Neutrophils Rps13 Neutrophils Rps14 Neutrophils Rps15 Neutrophils Rps2 Neutrophils Rps2 Neutrophils Rps3 Neutrophils Rps4 Neutrophils Rps5 Neutrophils Rps6 Neutrophils Rps7 Neutrophils Rps7 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps11 Neutrophils Rps2 Neutrophils Rps2 Neutrophils Rps3 Neutrophils Rps4 Neutrophils Rps4 Neutrophils Rps5 Neutrophils Rps6 Neutrophils Rps7 Neutrophils Rps7 Neutrophils Rps8 Neutrophils Rps8 Neutrophils Rps9 Neutrophils | Rpl27a   | Neutrophils | Emilin2 | CD4-CD8- T cells | Mzb1    | B cells    |
|---|----------|-------------|---------|------------------|---------|------------|
| Eef1a1       Neutrophils       Ifitm2       CD4-CD8-T cells       lgkc       B cells         Rpsa       Neutrophils       Ctsd       CD4-CD8-T cells       Snn       B cells         Rp114       Neutrophils       Cd3e       NKT       Fcer1a       Mast cells         Srgn       Neutrophils       Cd3d       NKT       Cpa3       Mast cells         Eef2       Neutrophils       Cd3g       NKT       Car8       Mast cells         Ppia       Neutrophils       Icos       NKT       Atp8b5       Mast cells         Rack1       Neutrophils       Trac       NKT       Atp8b5       Mast cells         Rpl23       Neutrophils       Ctla4       NKT       Tph1       Mast cells         Rpl19       Neutrophils       Ptprcap       NKT       Cyp11a1       Mast cells         Rps7       Neutrophils       Ptprcap       NKT       Cd0       Mast cells         Rps23       Neutrophils       Cd6       NKT       Cd200r3       Mast cells         Rps15       Neutrophils       H2-Q7       NKT       Madd number cells         Rps24       Neutrophils       Neutrophils         Rps10       Neutrophils       Neutrophils  | Rps15a   | Neutrophils | Plek    | CD4-CD8- T cells | Cd79b   | B cells    |
| Rpsa Neutrophils Ctsd CD4-CD8-T cells Snn B cells Rpl14 Neutrophils Cd3e NKT Fcer1a Mast cells Srgn Neutrophils Cd3d NKT Cpa3 Mast cells Eef2 Neutrophils Cd3g NKT Car8 Mast cells Ppia Neutrophils Icos NKT Atp8b5 Mast cells Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Ctla4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rps24 Neutrophils H2-Q7 NKT Maob Mast cells Rps84 Neutrophils Neutrophils Rps9 Neutrophils Neutrophils Rps10 Neutrophils Rps40 Neutrophils Rps40 Neutrophils Rps40 Neutrophils Rps40 Neutrophils Rps40 Neutrophils  | Rpl13    | Neutrophils | Lgals3  | CD4-CD8- T cells | Spib    | B cells    |
| Rpl14 Neutrophils Cd3e NKT Fcer1a Mast cells Srgn Neutrophils Cd3d NKT Cpa3 Mast cells Eef2 Neutrophils Cd3g NKT Car8 Mast cells Ppia Neutrophils Icos NKT Atp8b5 Mast cells Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Ct1a4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cyp11a1 Mast cells Rps87 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rps15 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils H2-Q7 NKT Maob Mast cells Rps8 Neutrophils Rpl8 Neutrophils Rpl8 Neutrophils Rps10 Neutrophils Rps4x Neutrophils Rps4x Neutrophils   | Eef1a1   | Neutrophils | Ifitm2  | CD4-CD8- T cells | Igkc    | B cells    |
| Srgn Neutrophils Cd3d NKT Cpa3 Mast cells  Eef2 Neutrophils Cd3g NKT Car8 Mast cells  Ppia Neutrophils Icos NKT Atp8b5 Mast cells  Rack1 Neutrophils Trac NKT Gata2 Mast cells  Rpl23 Neutrophils Ct1a4 NKT Tph1 Mast cells  Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells  Rps7 Neutrophils Ptprcap NKT Cobl Mast cells  Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells  Rps24 Neutrophils H2-Q7 NKT Maob Mast cells  Rps84 Neutrophils  Rpl8 Neutrophils  Rps95 Neutrophils  Rpl8 Neutrophils  Rps10 Neutrophils  Rps110 Neutrophils  Rps110 Neutrophils  Rps12 Neutrophils  Rps13 Neutrophils  Rps14 Neutrophils  Rps44 Neutrophils  Rps45 Neutrophils  Rps47 Neutrophils  | Rpsa     | Neutrophils | Ctsd    | CD4-CD8- T cells | Snn     | B cells    |
| Eef2 Neutrophils Cd3g NKT Car8 Mast cells Ppia Neutrophils Icos NKT Atp8b5 Mast cells Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Ct1a4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rps24 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Neutrophils Rps8 Neutrophils Rps8 Neutrophils Rps9 Neutrophils Rps1 Neutrophils Rps4 Neutrophils Rps4 Neutrophils Rps4x Neutrophils  | Rpl14    | Neutrophils | Cd3e    | NKT              | Fcer1a  | Mast cells |
| Ppia Neutrophils Icos NKT Atp8b5 Mast cells Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Ctla4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rps24 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils H2-Q7 NKT Maob Mast cells Rps15 Neutrophils H2-Q7 NKT Maob Mast cells Rps16 Neutrophils Rps17 Neutrophils Neutrophils Rps18 Neutrophils Rps10 Neutrophils  | Srgn     | Neutrophils | Cd3d    | NKT              | Cpa3    | Mast cells |
| Rack1 Neutrophils Trac NKT Gata2 Mast cells Rpl23 Neutrophils Ctla4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rps15 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Rpl8 Neutrophils Rpl8 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps10 Neutrophils  | Eef2     | Neutrophils | Cd3g    | NKT              | Car8    | Mast cells |
| Rpl23 Neutrophils Ctla4 NKT Tph1 Mast cells Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rpl5 Neutrophils H2-Q7 NKT Maob Mast cells Rpl8 Neutrophils Rpl8 Neutrophils Rpl8 Neutrophils Rpl9 Neutrophils Rps2 Neutrophils Rps1 Neutrophils Rps4 Neutrophils Rps4 Neutrophils Rps4 Neutrophils   | Ppia     | Neutrophils | Icos    | NKT              | Atp8b5  | Mast cells |
| Rpl19 Neutrophils Cd2 NKT Cyp11a1 Mast cells Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rpl15 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Rpl8 Neutrophils Rpl8 Neutrophils Rps2 Neutrophils Rps1 Neutrophils Rps1 Neutrophils Rps1 Neutrophils Rps1 Neutrophils Rps1 Neutrophils Rps1 Neutrophils Rps10 Neutrophils Rps10 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps4x Neutrophils Rps4x Neutrophils Rps4x Neutrophils   | Rack1    | Neutrophils | Trac    | NKT              | Gata2   | Mast cells |
| Rps7 Neutrophils Ptprcap NKT Cobl Mast cells Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rpl15 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Rpl8 Neutrophils Rpl3 Neutrophils Rps2 Neutrophils Rps10 Neutrophils  | Rpl23    | Neutrophils | Ctla4   | NKT              | Tph1    | Mast cells |
| Rps23 Neutrophils Cd6 NKT Cd200r3 Mast cells Rpl15 Neutrophils H2-Q7 NKT Maob Mast cells Rps24 Neutrophils Rpl8 Neutrophils Rpl3 Neutrophils Rps2 Neutrophils Rps10 Neutrophils Rpl10a Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps18 Neutrophils Rps4x Neutrophils Rps4x Neutrophils Neutrophils  | Rpl19    | Neutrophils | Cd2     | NKT              | Cyp11a1 | Mast cells |
| Rp15 Neutrophils H2-Q7 NKT Maob Mast cells  Rps24 Neutrophils  Rpl8 Neutrophils  Rpl3 Neutrophils  Rps2 Neutrophils  Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils  | Rps7     | Neutrophils | Ptprcap | NKT              | Cobl    | Mast cells |
| Rps24 Neutrophils  Rpl8 Neutrophils  Rpl3 Neutrophils  Rps2 Neutrophils  Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rps23    | Neutrophils | Cd6     | NKT              | Cd200r3 | Mast cells |
| Rpl8 Neutrophils  Rpl3 Neutrophils  Rps2 Neutrophils  Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rpl15    | Neutrophils | H2-Q7   | NKT              | Maob    | Mast cells |
| Rps2 Neutrophils  Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rps24    | Neutrophils |         |                  |         |            |
| Rps2 Neutrophils  Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rpl8     | Neutrophils |         |                  |         |            |
| Rps10 Neutrophils  Rpl10a Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rpl3     | Neutrophils |         |                  |         |            |
| Rpl10a Neutrophils  Rps18 Neutrophils  Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils  | Rps2     | Neutrophils |         |                  |         |            |
| Rps18 Neutrophils Hsp90ab1 Neutrophils Rps4x Neutrophils mt-Co1 Neutrophils   | Rps10    | Neutrophils |         |                  |         |            |
| Hsp90ab1 Neutrophils  Rps4x Neutrophils  mt-Co1 Neutrophils   | Rpl10a   | Neutrophils |         |                  |         |            |
| Rps4x Neutrophils mt-Co1 Neutrophils  | Rps18    | Neutrophils |         |                  |         |            |
| mt-Co1 Neutrophils  | Hsp90ab1 | Neutrophils |         |                  |         |            |
| ·   | Rps4x    | Neutrophils |         |                  |         |            |
| mt-Co2 Neutrophils  | mt-Co1   | Neutrophils |         |                  |         |            |
|   | mt-Co2   | Neutrophils |         |                  |         |            |
| mt-Atp6 Neutrophils   | mt-Atp6  | Neutrophils |         |                  |         |            |
| mt-Co3 Neutrophils  | mt-Co3   | Neutrophils |         |                  |         |            |
| mt-Cytb Neutrophils   | mt-Cytb  | Neutrophils |         |                  |         |            |

## Supplemental Table 4: Genes Differentially Upregulated in Survivor Mice Prior to Rechallenge

| Gene | Log2 fold | Full Name | Notable Aliases | Function |
|------|-----------|-----------|-----------------|----------|
|      | change    |           |                 |          |
|      |           |           |                 |          |

|      |      | A                  | Acute Phase Response     |  |
|------|------|--------------------|--------------------------|--|
| Арр  | 2.46 | Amyloid-beta       |                          | Cell surface receptor and transmembrane      |
|      |      | precursor protein  |                          | precursor protein that is cleaved by         |
|      |      |                    |                          | secretases to form a number of peptides;     |
|      |      |                    |                          | involved in cell mobility, copper            |
|      |      |                    |                          | homeostasis, and oxidative stress            |
| Clu  | 1.62 | Clusterin          | Apolipoprotein J;        | Extracellular chaperone that promotes        |
|      |      |                    | Ku70-binding protein     | clearance of inflammation and injury-        |
|      |      |                    | 1 (KUB1)                 | induced immune complexes; protects cells     |
|      |      |                    |                          | against apoptosis and against cytolysis by   |
|      |      |                    |                          | complement                                   |
| Lcn2 | 2.98 | Lipocalin 2        | Neutrophil               | Neutrophil-secreted factor that sequesters   |
|      |      |                    | gelatinase-associated    | iron-containing siderophores; also           |
|      |      |                    | lipocalin (NGAL)         | functions as a growth factor                 |
|      |      | Adhes              | ion & Cell-Cell Interact | ions   |
|      |      |                    |                          |  |
| Cd97 | 1.35 | Cluster of         | BL-Ac[F2]                | GPCR that promotes granulocyte adhesion      |
|      |      | differentiation 97 |                          | and migration; activates T cells via binding |
|      |      |                    |                          | to CD55; stimulates angiogenesis through     |
|      |      |                    |                          | binding integrin counterreceptors on         |
|      |      |                    |                          | endothelial cells                            |

| lfih1  | 1.09  | Interferon induced | Helicard; melanoma   | PRR for cytoplasmic dsRNA; upon target               |
|--------|-------|--------------------|----------------------|--|
|        |       | with helicase C    | differentiation-     | recognition, associates with MAVS to                 |
|        |       | domain 1           | associated protein 5 | activate TNK1 and IKBKE, which                       |
|        |       |                    | (MDA5)               | phosphorylate IRF3 and -7, which, in turn,           |
|        |       |                    |                      | activate transcription of IFN $\alpha$ and - $\beta$ |
| Itgam  | 1.77  | Integrin alpha M   | CD11b                | Pairs with CD18 to forms Mac-1 aka                   |
|        |       |                    |                      | complement receptor 3; mediates                      |
|        |       |                    |                      | leukocyte activation, adhesion, chemotaxis,          |
|        |       |                    |                      | migration, phagocytosis, and cell-mediated           |
|        |       |                    |                      | cytotoxicity; serves as a macrophage                 |
|        |       |                    |                      | marker   |
| Lgals3 | 1.99  | Galectin 3         |                      | Galactose-specific lectin that binds IgE;            |
|        |       |                    |                      | involved in acute inflammatory responses,            |
|        |       |                    |                      | including neutrophil activation and                  |
|        |       |                    |                      | adhesion, chemoattraction of monocytes               |
|        |       |                    |                      | macrophages, opsonization of apoptotic               |
|        |       |                    |                      | neutrophils, and activation of mast cells            |
| Map2k1 | 0.824 | Dual specificity   | MAPK/ERK kinase 1    | Essential component of the MAP kinase                |
|        |       | mitogen-activated  | (MEK1)               | signal transduction pathway; participates in         |
|        |       | protein kinase     |                      | numerous biological functions, including             |
|        |       | kinase 1           |                      | cell growth, adhesion, survival,                     |
|        |       |                    |                      | differentiation, transcription, metabolism,          |
|        |       |                    |                      | and cytoskeletal remodeling                          |
|        |       |                    |                      |  |

| S100a8 | 2.82 | S100 calcium-      | Calgranulin A | Calcium- and zinc-binding protein involved   |
|--------|------|--------------------|---------------|--|
|        |      | binding protein A8 |               | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |               | scavenging and apoptosis-inducing            |
|        |      |                    |               | activities; can induce neutrophil            |
|        |      |                    |               | chemotaxis, adhesion, phagocytosis, and      |
|        |      |                    |               | degranulation; predominantly found as        |
|        |      |                    |               | calprotectin (S100A8/A9) which has a wide    |
|        |      |                    |               | plethora of intra- and extracellular         |
|        |      |                    |               | functions, including adhesion, apoptosis,    |
|        |      |                    |               | autophagy, cytoskeletal remodeling,          |
|        |      |                    |               | cytokine production, chemotaxis,             |
|        |      |                    |               | migration, inflammation, arachidonic acid    |
|        |      |                    |               | metabolism, oxidant-scavenging, and PRR      |
|        |      |                    |               | signaling                                    |
| Thbs1  | 1.44 |                    |               | Adhesive glycoprotein that mediates cell-    |
|        |      |                    |               | to-cell and cell-to-matrix interactions;     |
|        |      | Thrombospondin 1   |               | ligand for CD36                              |
|        |      |                    | Angiogenesis  |  |
|        |      |                    |               |  |
| Angpt1 | 1.72 | Angiopoietin 1     |               | Secreted glycoprotein that binds and         |
|        |      |                    |               | activates the TEK/TIE2 receptor by inducing  |
|        |      |                    |               | its dimerization and tyrosine                |
|        |      |                    |               | phosphorylation; involved in angiogenesis,   |
|        |      |                    |               | endothelial cell survival, proliferation,    |

| Cd97    | 1.35  | Cluster of<br>differentiation 97              | BL-Ac[F2]                | migration, adhesion, cell spreading, and reorganization of the actin cytoskeleton  GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through  |
|---------|-------|---|--------------------------|---|
|         |       |   |                          | binding integrin counterreceptors on endothelial cells  |
| II1b    | 2.64  | Interleukin 1 beta                            | Catabolin                | One of the two primary inflammatory cytokines produced by the inflammasome (the other one being IL-18); induces neutrophil influx and activation,T cell activation and cytokine production, B cell activation and Ab production, fibroblast proliferation, and collagen production; synergizes with IL-12 to induce IFNy synthesis from TH1 cells |
|         |       | Antiger                                       | n Processing & Present   | ation   |
| Cd74    | 0.853 | Cluster of differentiation 74                 | MHC class II gamma chain | Stabilizes peptide-free class II αβ heterodimers during MHC-Ag complex formation  |
| Clec4a2 | 2.18  | C-type lectin<br>domain family 4<br>member A2 |                          | PRR that, upon binding mannose or fucose, is endocytosed and processed in the Ag presentation pathway   |

| Ctss    | 1.42 | Cathepsin S          |                      | Lysosomal protease that participates in     |
|---------|------|----------------------|----------------------|---|
|         |      |                      |                      | processing of Ag by MHC class II            |
|         |      |                      |                      | ,   |
| H2-Q10  | 1.98 | Histocompatibility   |                      | MHC class I molecule; presents Ags to CD8+  |
|         |      | 2, Q region locus 10 |                      | T cells                                     |
|         |      |                      |                      |   |
| lcam1   | 1.22 | Intracellular        | CD54                 | Cell surface glycoprotein that serves as    |
|         |      | adhesion molecule    |                      | strong adhesive ligand for LFA-1; important |
|         |      | 1                    |                      | for leukocyte mobility and costimulation    |
|         |      |                      | Anti-Inflammatory    |   |
|         |      |                      | Anti-initianimatory  |   |
| Bcl2    | 1.12 | B cell lymphoma 2    |                      | Outer mitochondrial membrane protein        |
|         |      |                      |                      | that inhibits apoptosis and autophagy; may  |
|         |      |                      |                      | attenuate inflammation by impairing         |
|         |      |                      |                      | inflammasome formation                      |
|         |      |                      |                      |   |
| Clu     | 1.62 | Clusterin            | Apolipoprotein J;    | Extracellular chaperone that promotes       |
|         |      |                      | Ku70-binding protein | clearance of inflammation and injury-       |
|         |      |                      | 1 (KUB1)             | induced immune complexes; protects cells    |
|         |      |                      |                      | against apoptosis and against cytolysis by  |
|         |      |                      |                      | complement                                  |
|         |      |                      |                      |   |
| Tnfaip3 | 1.38 | Tumor necrosis       |                      | Ubiquitin-editing enzyme that complexes     |
|         |      | factor, alpha-       |                      | with ITCH to degrade inflammatory           |
|         |      | induced protein 3    |                      | signaling components in the TNF, IL1, and   |
|         |      |                      |                      | TLR pathways; targets TRAF2, TRAF6, and     |
|         |      |                      |                      | IKK   |
|         |      |                      |                      |   |

| Tollip | 1.32  | Toll interacting protein                   |   | Inhibitory adaptor protein; recruits IRAK1  to the IL-1 receptor complex and inhibitively phosphorylates it   |
|--------|-------|--|---|---|
|        |       |  | Apoptosis                                 |   |
| Bid    | 1.4   | BH3 interacting domain death agonist       | Desmocollin type 4, apoptic death agonist | Induces caspases and apoptosis; counters the protective effect of BCL2, allowing release of cytochrome C  |
| Casp1  | 1.66  | Caspase 1                                  | Interleukin 1β convertase                 | Cysteine-aspartic acid protease that mediates cleavage-based activation of IL-1β and IL-18; serves as the central enzymatic core of the inflammasome; also induces apoptosis  |
| Casp8  | 0.914 | Caspase 8                                  |   | Cysteine-aspartic acid protease; cleaves and activates effector caspases CASP3, -4, -6, -7, -9, and -10; initiates pyroptosis through cleavage of gasdermin-D; mediates noncanonical cleavage of IL-1β in DCs and macrophages |
| Ifitm2 | 2.63  | Interferon-induced transmembrane protein 2 |   | IFN-induced antiviral protein which inhibits the entry of viruses to the host cell cytoplasm; induces cell cycle arrest and mediates p53-independent apoptosis through caspase activation                                     |

| S100a8 | 2.82 | S100 calcium-      | Calgranulin A        | Calcium- and zinc-binding protein involved   |
|--------|------|--------------------|----------------------|--|
|        |      | binding protein A8 |                      | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |                      | scavenging and apoptosis-inducing            |
|        |      |                    |                      | activities; can induce neutrophil            |
|        |      |                    |                      | chemotaxis, adhesion, phagocytosis, and      |
|        |      |                    |                      | degranulation; predominantly found as        |
|        |      |                    |                      | calprotectin (S100A8/A9) which has a wide    |
|        |      |                    |                      | plethora of intra- and extracellular         |
|        |      |                    |                      | functions, including adhesion, apoptosis,    |
|        |      |                    |                      | autophagy, cytoskeletal remodeling,          |
|        |      |                    |                      | cytokine production, chemotaxis,             |
|        |      |                    |                      | migration, inflammation, arachidonic acid    |
|        |      |                    |                      | metabolism, oxidant-scavenging, and PRR      |
|        |      |                    |                      | signaling                                    |
|        |      |                    | Autophagy            |  |
| Atg5   | 1.26 | Autophagy related  |                      | Pairs with ATG12 to promote the extension    |
|        |      | 5                  |                      | of the phagophoric membrane in               |
|        |      |                    |                      | autophagic vesicles                          |
| Irgm2  | 1.39 | Immunity-related   | Interferon-inducible | Function not fully known, but most likely    |
|        |      | GTPase family M    | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory     |
|        |      | member 2           |                      | cytokine production                          |
| S100a8 | 2.82 | S100 calcium-      | Calgranulin A        | Calcium- and zinc-binding protein involved   |
|        |      | binding protein A8 |                      | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |                      | scavenging and apoptosis-inducing            |

|                                |      |   | activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |
|--------------------------------|------|---|---|
| Ubc                            | 0.99 | Polyubiquitin C   | Serves various roles, including immate immunity, DNA repair, and stimulation of autophagy and the proteasomal response  |
| B Cell-<br>associated<br>Genes |      |   |   |
| Lyn                            | 1.88 | Lck/Yes-related novel kinase                              | Src family tyrosine kinase that potentiates signaling from the B cell receptor and CD40   |
| Pik3cg                         | 1.44 | Phosphatidylinosito I-4,5-bisphosphate 3-kinase catalytic | A subunit of PI3K; modulates leukocyte chemotaxis to inflammatory sites and in response to chemoattractant agents   |

|            |        | aub=:+ ==          |                      |   |  |  |  |  |
|------------|--------|--------------------|----------------------|---|--|--|--|--|
|            |        | subunit gamma      |                      |   |  |  |  |  |
|            |        | isoform            |                      |   |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
| Chemotaxis |        |                    |                      |   |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
| Ccl3       | 1.01   | C-C motif          | Macrophage           | Chemoattractant ligand for CCR1, -4, and -5 |  |  |  |  |
|            |        | chemokine ligand 3 | inflammatory protein |   |  |  |  |  |
|            |        |                    | 1α (MIP1α)           |   |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
| Ccr1       | 1.85   | C-C motif          | MIP1α receptor       | Receptor for CCL3, -5, -7, and -23          |  |  |  |  |
|            |        | chemokine          |                      |   |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
|            |        | receptor 1         |                      |   |  |  |  |  |
| Ccr2       | 2.01   | C-C motif          | CD192                | December for CCI2 a managita specific       |  |  |  |  |
| CCIZ       | 2.01   |                    | CD192                | Receptor for CCL2, a monocyte-specific      |  |  |  |  |
|            |        | chemokine          |                      | chemokine                                   |  |  |  |  |
|            |        | receptor 2         |                      |   |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
| Ccr3       | -0.901 | C-C motif          | CD193                | Receptor for a variety of chemokines,       |  |  |  |  |
|            |        | chemokine          |                      | including CCL11, CCL26, CCL7, CCL13, CCL5   |  |  |  |  |
|            |        | receptor 3         |                      | (RANTES), and CCL15; signals through        |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
|            |        |                    |                      | Ca(2+) flux                                 |  |  |  |  |
| Ccr5       | 1.18   | C-C motif          | CD195                | Receptor for a number of inflammatory CC-   |  |  |  |  |
|            |        | chemokine          |                      | chemokines, including CCL3/MIP1α,           |  |  |  |  |
|            |        |                    |                      | -   |  |  |  |  |
|            |        | receptor 5         |                      | CCL4/MIP1β, and RANTES; signals via         |  |  |  |  |
|            |        |                    |                      | Ca(2+) flux                                 |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |
| Cklf       | 1.14   | Chemokine-like     |                      | Chemoattractant for monocytes,              |  |  |  |  |
|            |        | factor             |                      | neutrophils, and lymphocytes                |  |  |  |  |
|            |        |                    |                      |   |  |  |  |  |

| Cxcr2  | 2.38 | C-X-C motif        | CD182; IL-8 receptor | Receptor for IL-8 and CXCL3; powerful        |
|--------|------|--------------------|----------------------|--|
|        |      | chemokine          | В                    | chemoattractant for neutrophils              |
|        |      | receptor 2         |                      |  |
|        |      |                    |                      |  |
| Cxcr3  | 1.55 | C-X-C motif        | CD183                | Induces integrin activation, cytoskeletal    |
|        |      | chemokine          |                      | remodeling, and chemotaxis; expressed by     |
|        |      | receptor 3         |                      | T cells and NK cells; prominently expressed  |
|        |      |                    |                      | in effector and memory T cells               |
| Itgam  | 1.77 | Integrin alpha M   | CD11b                | Pairs with CD18 to forms Mac-1 aka           |
|        |      |                    |                      | complement receptor 3; mediates              |
|        |      |                    |                      | leukocyte activation, adhesion, chemotaxis,  |
|        |      |                    |                      | migration, phagocytosis, and cell-mediated   |
|        |      |                    |                      | cytotoxicity; serves as a macrophage         |
|        |      |                    |                      | marker                                       |
| Lgals3 | 1.99 | Galectin 3         |                      | Galactose-specific lectin that binds IgE;    |
|        |      |                    |                      | involved in acute inflammatory responses,    |
|        |      |                    |                      | including neutrophil activation and          |
|        |      |                    |                      | adhesion, chemoattraction of monocytes       |
|        |      |                    |                      | macrophages, opsonization of apoptotic       |
|        |      |                    |                      | neutrophils, and activation of mast cells    |
| S100a8 | 2.82 | S100 calcium-      | Calgranulin A        | Calcium- and zinc-binding protein involved   |
|        |      | binding protein A8 |                      | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |                      | scavenging and apoptosis-inducing            |
|        |      |                    |                      | activities; can induce neutrophil            |
|        |      |                    |                      | chemotaxis, adhesion, phagocytosis, and      |

|                               |      |                    | calprotections plethon functions autoph cytoki | ation; predominantly found as  n (S100A8/A9) which has a wide ra of intra- and extracellular , including adhesion, apoptosis, agy, cytoskeletal remodeling, ne production, chemotaxis, inflammation, arachidonic acid m, oxidant-scavenging, and PRR signaling |  |  |  |
|-------------------------------|------|--------------------|--|--|--|--|--|
| Chromatin                     |      |                    |  |  |  |  |  |
| Remodeling                    |      |                    |  |  |  |  |  |
| Ep300                         | 1.1  | Adenovirus early   | Histone a                                      | cetyltransferase; participates in  |  |  |  |
|                               |      | region 1A-         | chromati                                       | n remodeling to facilitate gene  |  |  |  |
|                               |      | associated protein |  | accessibility  |  |  |  |
|                               |      | p300               |  |  |  |  |  |
| Hmgb1                         | 1.19 | High-mobility      | Remodels                                       | s chromatin to make DNA more   |  |  |  |
|                               |      | group box 1        | av   | vailable for transcription   |  |  |  |
| Complement & Humoral Immunity |      |                    |  |  |  |  |  |
| C3                            | 2    | Complement         | Cleaved by                                     | / C3 convertase to form C3a and  |  |  |  |
|                               |      | component 3        | C3b, an a                                      | naphalotoxin and an opsonizing   |  |  |  |
|                               |      |                    |  | agent, respectively  |  |  |  |

| Cfh    | 2.77 | Complement factor   | Soluble glycoprotein that regulates the      |
|--------|------|---------------------|--|
|        |      | Н                   | alternate pathway by accelerating decay of   |
|        |      |                     | C3 convertase                                |
|        |      |                     |  |
| Cfp    | 1.82 | Complement factor   | Alternate complement pathway                 |
|        |      | properdin           | component; when cleaved, produces a          |
|        |      |                     | serine protease that binds to C3b to form    |
|        |      |                     | C3 convertase                                |
| Fcer1a | 1.7  | Fragment            | High affinity receptor for IgE; responsible  |
|        |      | crystallizable of   | for initiating the allergic response         |
|        |      | immunoglobulin      |  |
|        |      | epsilon receptor 1a |  |
|        |      |                     |  |
| Lgals3 | 1.99 | Galectin 3          | Galactose-specific lectin that binds IgE;    |
|        |      |                     | involved in acute inflammatory responses,    |
|        |      |                     | including neutrophil activation and          |
|        |      |                     | adhesion, chemoattraction of monocytes       |
|        |      |                     | macrophages, opsonization of apoptotic       |
|        |      |                     | neutrophils, and activation of mast cells    |
|        |      |                     | Costimulation                                |
|        |      |                     |  |
| Cd28   | 1.04 | Cluster of          | Essential T cell co-receptor that enhances T |
|        |      | differentiation 28  | cell activation, proliferation, cytokine     |
|        |      |                     | production, and survival; binds to CD80 and  |
|        |      |                     | CD86   |
|        |      |                     |  |

| lcam1  | 1.22 | Intracellular      | CD54      | Cell surface glycoprotein that serves as    |
|--------|------|--------------------|-----------|---|
|        |      | adhesion molecule  |           | strong adhesive ligand for LFA-1; important |
|        |      | 1                  |           | for leukocyte mobility and costimulation    |
|        |      |                    | Cytokines |   |
|        |      |                    | Cytokiics |   |
| Csf1   | 1.75 | Macrophage         |           | Cytokine that promote activation and        |
|        |      | colony-stimulating |           | survival of monocytes                       |
|        |      | factor 1           |           |   |
| Csf1r  | 1.94 | Macrophage         | CD115     | Receptor for CSF1; promotes release of      |
|        |      | colony-stimulating |           | inflammatory cytokines in response to IL-34 |
|        |      | factor 1 receptor  |           | and CSF1                                    |
| Ifnar1 | 1.12 | Interferon-        |           | Component of the receptor for type I IFNs,  |
|        |      | alpha/beta         |           | binding of which activates the JAK-STAT     |
|        |      | receptor alpha     |           | pathway                                     |
|        |      | chain              |           |   |
| lfngr1 | 1.45 | Interferon gamma   | CD119     | One of the two components of the IFNy       |
|        |      | receptor 1         |           | receptor; stimulates activation of the      |
|        |      |                    |           | JAK/STAT signaling pathway                  |
| II1b   | 2.64 | Interleukin 1 beta | Catabolin | One of the two primary inflammatory         |
|        |      |                    |           | cytokines produced by the inflammasome      |
|        |      |                    |           | (the other one being IL-18); induces        |
|        |      |                    |           | neutrophil influx and activation,T cell     |
|        |      |                    |           | activation and cytokine production, B cell  |
|        |      |                    |           | activation and Ab production, fibroblast    |

|        |       |   |                                    | proliferation, and collagen production; synergizes with IL-12 to induce IFN $\gamma$ synthesis from $T_H1$ cells  |
|--------|-------|---|------------------------------------|---|
| Il1rap | 2.32  | Interleukin 1 receptor accessory protein          |                                    | Co-receptor for several ligands, including IL-<br>1R1 in the IL-1 pathway, IL-RL1 in the IL-33<br>pathway, IL-1RL2 in the IL-36 pathway;<br>signaling involves Tollip, MyD88, IRAK1, and<br>IRAK2   |
| ll1rl1 | 1.63  | Interleukin 1 receptor-like 1                     |                                    | Receptor for IL-33; recruits MyD88, IRAK1, IRAK4, and TRAF6; activates ERK1, ERK2, and MAPK14   |
| II6st  | 0.913 | Interleukin 6  cytokine family  signal transducer | Glycoprotein 130<br>(Gp130); CD130 | Transmembrane protein that acts a component in several cytokine receptors, including IL-6   |
| II18   | 1.29  | Interleukin 18                                    | Iboctadekin                        | One of the two primary inflammatory cytokines produced by the inflammasome (the other one being IL-1 $\beta$ ); upon binding to IL18R1 and IL18RAP, forms a signaling ternary complex which activates NF $\kappa$ B, triggering synthesis of inflammatory mediators |
| II18r1 | 1.21  | Interleukin 18 receptor 1                         | CD218a                             | Receptor for IL-18  |

| II18rap | 1.83  | Interleukin 18 receptor accessory protein                  | CD218b                    | Accessory subunit of the heterodimeric IL-  18 receptor involved in IL18-dependent signal transduction, leading to NFkB and JNK activation  |
|---------|-------|--|---------------------------|---|
|         |       | Су   | rtoskeletal Remodeling    |   |
| Angpt1  | 1.72  | Angiopoietin 1   |                           | Secreted glycoprotein that binds and activates the TEK/TIE2 receptor by inducing its dimerization and tyrosine phosphorylation; involved in angiogenesis, endothelial cell survival, proliferation, migration, adhesion, cell spreading, and reorganization of the actin cytoskeleton |
| Map2k1  | 0.824 | Dual specificity mitogen-activated protein kinase kinase 1 | MAPK/ERK kinase 1  (MEK1) | Essential component of the MAP kinase signal transduction pathway; participates in numerous biological functions, including cell growth, adhesion, survival, differentiation, transcription, metabolism, and cytoskeletal remodeling  |
|         |       |  | Cytotoxicity              |   |
| Itgam   | 1.77  | Integrin alpha M   | CD11b                     | Pairs with CD18 to forms Mac-1 aka complement receptor 3; mediates leukocyte activation, adhesion, chemotaxis, migration, phagocytosis, and cell-mediated   |

|        |      |                                |                      | cytotoxicity; serves as a macrophage marker  |
|--------|------|--------------------------------|----------------------|--|
| S100a8 | 2.82 | S100 calciumbinding protein A8 | Calgranulin A        | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |
|        |      | •                              | Growth/Proliferation |  |
| Angpt1 | 1.72 | Angiopoietin 1                 |                      | Secreted glycoprotein that binds and activates the TEK/TIE2 receptor by inducing its dimerization and tyrosine phosphorylation; involved in angiogenesis, endothelial cell survival, proliferation, migration, adhesion, cell spreading, and reorganization of the actin cytoskeleton  |

| Atf2   | 1.13  | Activating           | Cyclic AMP-         | Regulates transcription of various genes     |
|--------|-------|----------------------|---------------------|--|
|        |       | transcription factor | responsive element- | involved in anti-apoptosis, cell growth, and |
|        |       | 2                    | binding protein 2   | DNA damage response; in the nucleus,         |
|        |       |                      | (CREB2)             | contributes to global transcription and the  |
|        |       |                      |                     | DNA damage response, in addition to          |
|        |       |                      |                     | specific transcriptional activities that are |
|        |       |                      |                     | related to cell development, proliferation   |
|        |       |                      |                     | and death; in the cytoplasm, impairs         |
|        |       |                      |                     | mitochondrial membrane potential,            |
|        |       |                      |                     | inducing mitochondrial leakage and           |
|        |       |                      |                     | promoting cell death; phosphorylated form    |
|        |       |                      |                     | (mediated by ATM) plays a role in the DNA    |
|        |       |                      |                     | damage response                              |
| Map2k1 | 0.824 | Dual specificity     | MAPK/ERK kinase 1   | Essential component of the MAP kinase        |
|        |       | mitogen-activated    | (MEK1)              | signal transduction pathway; participates in |
|        |       | protein kinase       |                     | numerous biological functions, including     |
|        |       | kinase 1             |                     | cell growth, adhesion, survival,             |
|        |       |                      |                     | differentiation, transcription, metabolism,  |
|        |       |                      |                     | and cytoskeletal remodeling                  |
| Tyk2   | 1.54  | Tyrosine kinase 2    | JTK1                | Plays both structural and catalytic roles in |
|        |       |                      |                     | numerous cytokines and interferons           |
|        |       |                      |                     | signaling; associates with cytokine and      |
|        |       |                      |                     | growth factor receptors and activate STAT    |

|         |       |  |                              | family members including STAT1, STAT3,  STAT4, or STAT6   |
|---------|-------|--|------------------------------|---|
|         |       |  | Ion Transport                |   |
| Арр     | 2.46  | Amyloid-beta precursor protein                               |                              | Cell surface receptor and transmembrane precursor protein that is cleaved by secretases to form a number of peptides; involved in cell mobility, copper homeostasis, and oxidative stress |
| Slc11a1 | 2.01  | Natural resistance-<br>associated<br>macrophage<br>protein 1 | Inflammation                 | Macrophage-specific metal ion transporter; uptakes divalent metal cations to neutralize ROSs  |
|         |       |  |                              |   |
| Bst2    | 0.822 | Bone marrow stromal cell antigen 2                           | Tethrin; CD317               | IFN-induced antiviral factor that blocks budding of enveloped viruses by directly tethering nascent virions to the membranes of infected cells  |
| Casp1   | 1.66  | Caspase 1  | Interleukin 1β<br>convertase | Cysteine-aspartic acid protease that mediates cleavage-based activation of IL-1β and IL-18; serves as the central enzymatic core of the inflammasome; also induces apoptosis              |

| Casp8  | 0.914 | Caspase 8          |                      | Cysteine-aspartic acid protease; cleaves     |
|--------|-------|--------------------|----------------------|--|
|        |       |                    |                      | and activates effector caspases CASP3, -4, - |
|        |       |                    |                      | 6, -7, -9, and -10; initiates pyroptosis     |
|        |       |                    |                      | through cleavage of gasdermin-D; mediates    |
|        |       |                    |                      | noncanonical cleavage of IL-1β in DCs and    |
|        |       |                    |                      | macrophages                                  |
| Cebpb  | 2.13  | CCAAT/enhancer-    |                      | Critical macrophage TF that promotes         |
|        |       | binding protein    |                      | expression of several acute-phase and        |
|        |       | beta               |                      | inflammatory cytokine genes, including II6   |
| Clu    | 1.62  | Clusterin          | Apolipoprotein J;    | Extracellular chaperone that promotes        |
|        |       |                    | Ku70-binding protein | clearance of inflammation and injury-        |
|        |       |                    | 1 (KUB1)             | induced immune complexes; protects cells     |
|        |       |                    |                      | against apoptosis and against cytolysis by   |
|        |       |                    |                      | complement                                   |
| Csf1r  | 1.94  | Macrophage         | CD115                | Receptor for CSF1; promotes release of       |
|        |       | colony-stimulating |                      | inflammatory cytokines in response to IL-34  |
|        |       | factor 1 receptor  |                      | and CSF1                                     |
| Ifnar1 | 1.12  | Interferon-        |                      | Component of the receptor for type I IFNs,   |
|        |       | alpha/beta         |                      | binding of which activates the JAK-STAT      |
|        |       | receptor alpha     |                      | pathway                                      |
|        |       | chain              |                      |  |
|        |       |                    |                      |  |

| Ifngr1 | 1.45  | Interferon gamma   | CD119            | One of the two components of the IFNy        |
|--------|-------|--------------------|------------------|--|
|        |       | receptor 1         |                  | receptor; stimulates activation of the       |
|        |       |                    |                  | JAK/STAT signaling pathway                   |
|        |       |                    |                  |  |
| II1b   | 2.64  | Interleukin 1 beta | Catabolin        | One of the two primary inflammatory          |
|        |       |                    |                  | cytokines produced by the inflammasome       |
|        |       |                    |                  | (the other one being IL-18); induces         |
|        |       |                    |                  | neutrophil influx and activation,T cell      |
|        |       |                    |                  | activation and cytokine production, B cell   |
|        |       |                    |                  | activation and Ab production, fibroblast     |
|        |       |                    |                  | proliferation, and collagen production;      |
|        |       |                    |                  | synergizes with IL-12 to induce IFNγ         |
|        |       |                    |                  | synthesis from T <sub>H</sub> 1 cells        |
|        |       |                    |                  |  |
| Il6st  | 0.913 | Interleukin 6      | Glycoprotein 130 | Transmembrane protein that acts a            |
|        |       | cytokine family    | (Gp130); CD130   | component in several cytokine receptors,     |
|        |       | signal transducer  |                  | including IL-6                               |
| II18   | 1.29  | Interleukin 18     | Iboctadekin      | One of the two primary inflammatory          |
|        |       |                    |                  | cytokines produced by the inflammasome       |
|        |       |                    |                  | (the other one being IL-1β); upon binding to |
|        |       |                    |                  | IL18R1 and IL18RAP, forms a signaling        |
|        |       |                    |                  | ternary complex which activates NFĸB,        |
|        |       |                    |                  | triggering synthesis of inflammatory         |
|        |       |                    |                  | mediators                                    |
|        |       |                    |                  |  |

| ll18r1  | 1.21 | Interleukin 18 receptor 1                  | CD218a                                | Receptor for IL-18  |
|---------|------|--|---------------------------------------|---|
| Il18rap | 1.83 | Interleukin 18 receptor accessory protein  | CD218b                                | Accessory subunit of the heterodimeric IL-  18 receptor involved in IL18-dependent  signal transduction, leading to NFkB and  JNK activation  |
| Irf7    | 1.22 | Interferon regulatory factor 7             |                                       | Key transcriptional regulator of type I IFN-dependent immune responses; promotes $transcription\ of\ IFN\alpha\ and\ -\beta$  |
| Irgm2   | 1.39 | Immunity-related GTPase family M member 2  | Interferon-inducible protein 1 (IFI1) | Function not fully known, but most likely regulates autophagy and pro-inflammatory cytokine production  |
| Lgals3  | 1.99 | Galectin 3                                 |                                       | Galactose-specific lectin that binds IgE; involved in acute inflammatory responses, including neutrophil activation and adhesion, chemoattraction of monocytes macrophages, opsonization of apoptotic neutrophils, and activation of mast cells |
| Lilra5  | 2.23 | Leukocyte immunoglobulin- like receptor A5 | CD85                                  | Function poorly understood; may play a role in triggering innate immune responses through triggering Ca(2+) influx and proinflammatory cytokine release   |

| Mavs   | 0.866 | Mitochondrial       | IFNβ promoter        | Intermediary protein involved in the        |
|--------|-------|---------------------|----------------------|---|
|        |       | antiviral signaling | stimulator protein 1 | nonclassical inflammasome pathway; acts     |
|        |       | protein             | (ISP-1)              | downstream of DDX58 and IFIH1, leading to   |
|        |       |                     |                      | the activation of NFkB, IRF3, and IRF7, and |
|        |       |                     |                      | the subsequent induction of IFNβ and        |
|        |       |                     |                      | RANTES                                      |
| Mefv   | 2.88  | Mediterranean       | Marenostrin; pyrin   | Involved in the regulation of innate        |
|        |       | fever               |                      | immunity and the inflammatory response in   |
|        |       |                     |                      | response to IFNγ; both stimulates and       |
|        |       |                     |                      | restrains the inflammasome; also acts as a  |
|        |       |                     |                      | mediator of pyroptosis                      |
| Mif    | 1.07  | Macrophage          | L-dopachrome         | Pro-inflammatory cytokine that promotes     |
|        |       | migration           | tautomerase          | macrophage function through suppression     |
|        |       | inhibitory factor   |                      | of anti-inflammatory effects of             |
|        |       |                     |                      | glucocorticoids                             |
| Nlrp3  | 2.42  | NACHT domain-,      | Cryopyrin            | PRR with a wide diversity of recognized     |
|        |       | leucine-rich        |                      | targets that activates the NLRP3            |
|        |       | repeat-, and PYD-   |                      | inflammasome consisting of NLRP3,           |
|        |       | containing protein  |                      | PYCARD, and caspase-1/-8                    |
|        |       | 3                   |                      |   |
| Pik3cg | 1.44  | Phosphatidylinosito |                      | A subunit of PI3K; modulates leukocyte      |
|        |       | I-4,5-bisphosphate  |                      | chemotaxis to inflammatory sites and in     |
|        |       | 3-kinase catalytic  |                      | response to chemoattractant agents          |

|         |       | subunit gamma<br>isoform |                  |  |
|---------|-------|--------------------------|------------------|--|
| S100a8  | 2.82  | S100 calcium-            | Calgranulin A    | Calcium- and zinc-binding protein involved   |
|         |       | binding protein A8       |                  | in pro-inflammatory, antimicrobial, oxidant- |
|         |       |                          |                  | scavenging and apoptosis-inducing            |
|         |       |                          |                  | activities; can induce neutrophil            |
|         |       |                          |                  | chemotaxis, adhesion, phagocytosis, and      |
|         |       |                          |                  | degranulation; predominantly found as        |
|         |       |                          |                  | calprotectin (S100A8/A9) which has a wide    |
|         |       |                          |                  | plethora of intra- and extracellular         |
|         |       |                          |                  | functions, including adhesion, apoptosis,    |
|         |       |                          |                  | autophagy, cytoskeletal remodeling,          |
|         |       |                          |                  | cytokine production, chemotaxis,             |
|         |       |                          |                  | migration, inflammation, arachidonic acid    |
|         |       |                          |                  | metabolism, oxidant-scavenging, and PRR      |
|         |       |                          |                  | signaling                                    |
| Tbk1    | 1.22  | TANK-binding             |                  | Coordinates the activation of IRF3 and NFkB  |
|         |       | kinase 1                 |                  | and induction of type I IFNs                 |
| Tmem173 | 0.811 | Transmembrane            | Stimulator of    | Adaptor protein in type I IFN signaling;     |
|         |       | protein 173              | interferon genes | activates STAT6 and IRF3 through TBK1 to     |
|         |       |                          | (STING)          | induce type I IFN production                 |

| Tollip | 1.32  | Toll interacting  |                      | Inhibitory adaptor protein; recruits IRAK1   |
|--------|-------|-------------------|----------------------|--|
|        |       | protein           |                      | to the IL-1 receptor complex and             |
|        |       |                   |                      | inhibitively phosphorylates it               |
|        |       |                   |                      |  |
|        |       |                   | Inhibition           |  |
| Bcl2   | 1.12  | B cell lymphoma 2 |                      | Outer mitochondrial membrane protein         |
|        |       |                   |                      | that inhibits apoptosis and autophagy; may   |
|        |       |                   |                      | attenuate inflammation by impairing          |
|        |       |                   |                      | inflammasome formation                       |
| Chuk   | 0.868 | Conserved helix-  | Inhibitor of NFκB    | Part of the IKK complex that inhibits IκBα   |
|        |       | loop-helix        | kinase subunit alpha | and permits NFkB nuclear localization        |
|        |       | ubiquitous kinase | (ΙΚΚα)               |  |
|        |       |                   |                      |  |
| Clu    | 1.62  | Clusterin         | Apolipoprotein J;    | Extracellular chaperone that promotes        |
|        |       |                   | Ku70-binding protein | clearance of inflammation and injury-        |
|        |       |                   | 1 (KUB1)             | induced immune complexes; protects cells     |
|        |       |                   |                      | against apoptosis and against cytolysis by   |
|        |       |                   |                      | complement                                   |
| Cyld   | 0.997 | Cylindromatosis   |                      | Inhibits NFkB activation by deubiquitinating |
|        |       | lysine 63         |                      | upstream signaling factors; inhibits Wnt     |
|        |       | deubiquitinase    |                      | signaling; restricts polyubiquitination of   |
|        |       |                   |                      | RIPK1 and -2, thereby limiting necroptosis   |
| Dusp6  | 1.46  | Dual specificity  |                      | Inhibitively phosphorylates ERK1 and 2       |
|        |       | phosphatase 6     |                      |  |

| Foxp3 | 1.5   | Forkhead box P3                              | DIETER                              | Master TF for Tregs; represses expression of II2 and Ifng; activates expression of Tnfrsf18, II2ra, and Ctla4                                  |
|-------|-------|--|-------------------------------------|--|
| Irak3 | 1.31  | Interleukin-1 receptor- associated kinase 3  |                                     | Adaptor protein that negatively regulates  TLR signaling; predominantly expressed in  monocytes and macrophages                                |
| Nfkb1 | 0.928 | Nuclear factor<br>kappa B subunit 1          | p105/p50                            | One of the NFκB family TFs; inhibits inflammation  |
| NIrc5 | 0.845 | NLR family CARD domain containing 5          | NOD27                               | Inhibits NFkB and type I IFN signaling pathways; may also regulate the type II IFN signaling pathway   |
| Tank  | 0.916 | TRAF family member-associated NFkB activator |                                     | Inhibitory protein that sequesters TRAFs in the cytoplasm, constitutively binds TBK1, and serves as a negative regulator of NFkB               |
|       |       |  | Interferon Response                 |  |
| Bst2  | 0.822 | Bone marrow<br>stromal cell antigen<br>2     | Tethrin; CD317                      | IFN-induced antiviral factor that blocks budding of enveloped viruses by directly tethering nascent virions to the membranes of infected cells |
| Ifih1 | 1.09  | Interferon induced with helicase C domain 1  | Helicard; melanoma differentiation- | PRR for cytoplasmic dsRNA; upon target recognition, associates with MAVS to activate TNK1 and IKBKE, which                                     |

|        |      |                     | associated protein 5 | phosphorylate IRF3 and -7, which, in turn,           |
|--------|------|---------------------|----------------------|--|
|        |      |                     | (MDA5)               | activate transcription of IFN $\alpha$ and - $\beta$ |
| lfitm1 | 3.2  | Interferon-induced  | CD225                | IFN-induced antiviral protein implicated in          |
|        |      | transmembrane       |                      | cell adhesion and control of cell growth and         |
|        |      | protein 1           |                      | migration  |
| Ifitm2 | 2.63 | Interferon-induced  |                      | IFN-induced antiviral protein which inhibits         |
|        |      | transmembrane       |                      | the entry of viruses to the host cell                |
|        |      | protein 2           |                      | cytoplasm; induces cell cycle arrest and             |
|        |      |                     |                      | mediates p53-independent apoptosis                   |
|        |      |                     |                      | through caspase activation                           |
| Ifnar1 | 1.12 | Interferon-         |                      | Component of the receptor for type I IFNs,           |
|        |      | alpha/beta          |                      | binding of which activates the JAK-STAT              |
|        |      | receptor alpha      |                      | pathway  |
|        |      | chain               |                      |  |
| Ifngr1 | 1.45 | Interferon gamma    | CD119                | One of the two components of the IFNy                |
|        |      | receptor 1          |                      | receptor; stimulates activation of the               |
|        |      |                     |                      | JAK/STAT signaling pathway                           |
| Irf7   | 1.22 | Interferon          |                      | Key transcriptional regulator of type I IFN-         |
|        |      | regulatory factor 7 |                      | dependent immune responses; promotes                 |
|        |      |                     |                      | transcription of IFNα and -β                         |

| Irgm2   | 1.39  | Immunity-related  | Interferon-inducible | Function not fully known, but most likely    |  |  |
|---------|-------|-------------------|----------------------|--|--|--|
|         |       | GTPase family M   | protein 1 (IFI1)     | regulates autophagy and pro-inflammatory     |  |  |
|         |       | member 2          | , , ,                | cytokine production                          |  |  |
|         |       | member 2          |                      | cytokine production                          |  |  |
| Mefv    | 2.88  | Mediterranean     | Marenostrin; pyrin   | Involved in the regulation of innate         |  |  |
|         |       | fever             |                      | immunity and the inflammatory response in    |  |  |
|         |       |                   |                      | response to IFNγ; both stimulates and        |  |  |
|         |       |                   |                      | restrains the inflammasome; also acts as a   |  |  |
|         |       |                   |                      | mediator of pyroptosis                       |  |  |
| Tbk1    | 1.22  | TANK-binding      |                      | Coordinates the activation of IRF3 and NFkB  |  |  |
|         |       | kinase 1          |                      | and induction of type I IFNs                 |  |  |
| Tmem173 | 0.811 | Transmembrane     | Stimulator of        | Adaptor protein in type I IFN signaling;     |  |  |
|         |       | protein 173       | interferon genes     | activates STAT6 and IRF3 through TBK1 to     |  |  |
|         |       |                   | (STING)              | induce type I IFN production                 |  |  |
| Tyk2    | 1.54  | Tyrosine kinase 2 | JTK1                 | Plays both structural and catalytic roles in |  |  |
|         |       |                   |                      | numerous cytokines and interferons           |  |  |
|         |       |                   |                      | signaling; associates with cytokine and      |  |  |
|         |       |                   |                      | growth factor receptors and activate STAT    |  |  |
|         |       |                   |                      | family members including STAT1, STAT3,       |  |  |
|         |       |                   |                      | STAT4, or STAT6                              |  |  |
|         |       |                   | IRAKs & TRAFs        |  |  |  |
|         |       |                   |                      |  |  |  |

| Irak1 | 1.2   | Interleukin-1       | Adaptor protein involved in TLR and IL-1  |
|-------|-------|---------------------|---|
|       |       | receptor-           | signaling; recruited to TLRs by MyD88 and |
|       |       | associated kinase 1 | phosphorylated by IRAK4; promotes the     |
|       |       |                     | degradation of TIRAP                      |
| Irak2 | 0.791 | Interleukin-1       | Adaptor protein involved in TLR and IL-1  |
|       |       | receptor-           | signaling                                 |
|       |       | associated kinase 2 |   |
| Irak3 | 1.31  | Interleukin-1       | Adaptor protein that negatively regulates |
|       |       | receptor-           | TLR signaling; predominantly expressed in |
|       |       | associated kinase 3 | monocytes and macrophages                 |
| Irak4 | 1.33  | Interleukin-1       | The primary IRAK family member in         |
|       |       | receptor-           | mammalian TLR and IL-1 signaling; joins   |
|       |       | associated kinase 4 | with IRAK2 and MyD88 to form the          |
|       |       |                     | myddosome complex to activate IRAK1       |
| Traf2 | 0.8   | Tumor necrosis      | Adaptor protein required for TNFα-        |
|       |       | factor receptor-    | mediated activation of JNK and NFκB       |
|       |       | associated factor 2 |   |
| Traf3 | 0.939 | Tumor necrosis      | Adaptor protein that acts in the CD40     |
|       |       | factor receptor-    | signaling cascade; induces NFkB and MAPK  |
|       |       | associated factor 3 | activation                                |

| Traf6 | 0.707 | Tumor necrosis      |                   | Adaptor protein that acts in the CD40                         |
|-------|-------|---------------------|-------------------|---|
|       |       | factor receptor-    |                   | signaling cascade; promotes inflammation,                     |
|       |       | associated factor 6 |                   | IL-6, and TNFα  |
|       |       |                     | JAK-STAT Pathway  |   |
|       |       |                     | JAK-STAT Falliway |   |
| Jak1  | 0.834 | Janus kinase 1      |                   | Essential tyrosine kinase involved signal                     |
|       |       |                     |                   | transduction in type I and II cytokines and                   |
|       |       |                     |                   | IFNs  |
| Jak2  | 1.33  | Janus kinase 2      |                   | Tyrosine kinase that participates in IFN and                  |
|       |       |                     |                   | IL6ST signaling cascades                                      |
| Stat1 | 1.24  | Signal transducer   |                   | Transcriptional activator that mediates                       |
|       |       | and activator of    |                   | cellular responses to IFNs, cytokines, and                    |
|       |       | transcription 1     |                   | other growth factors  |
| Stat3 | 1.26  | Signal transducer   |                   | Transcriptional activator of genes involved                   |
|       |       | and activator of    |                   | in cell growth and apoptosis; activated by                    |
|       |       | transcription 3     |                   | JAKs  |
| Stat6 | 1.22  | Signal transducer   |                   | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and |
|       |       | and activator of    |                   | macrophage function and M2 macrophage                         |
|       |       | transcription 6     |                   | polarization  |
|       |       |                     | Kinases           |   |
| 200   | 4.5.5 |                     |                   |   |
| Btk   | 1.24  | Bruton's tyrosine   |                   | Crucial kinase in B cell receptor signal                      |
|       |       | kinase              |                   | transmission and B cell activation                            |

| Hck    | 1.62  | Hematopoietic cell  | Src family tyrosine kinase that mediates     |
|--------|-------|---------------------|--|
|        |       | kinase              | secretory lysosome mobilization,             |
|        |       |                     | degranulation, and activation of NADPH       |
|        |       |                     | oxidase                                      |
| Jak1   | 0.834 | Janus kinase 1      | Essential tyrosine kinase involved signal    |
|        |       |                     | transduction in type I and II cytokines and  |
|        |       |                     | IFNs   |
| Jak2   | 1.33  | Janus kinase 2      | Tyrosine kinase that participates in IFN and |
|        |       |                     | IL6ST signaling cascades                     |
| Lyn    | 1.88  | Lck/Yes-related     | Src family tyrosine kinase that potentiates  |
|        |       | novel kinase        | signaling from the B cell receptor and CD40  |
| Pik3cg | 1.44  | Phosphatidylinosito | A subunit of PI3K; modulates leukocyte       |
|        |       | I-4,5-bisphosphate  | chemotaxis to inflammatory sites and in      |
|        |       | 3-kinase catalytic  | response to chemoattractant agents           |
|        |       | subunit gamma       |  |
|        |       | isoform             |  |
| Ripk2  | 1.37  | Receptor-           | RIP kinase that potentiates signals          |
|        |       | interacting         | downstream of NOD1 and -2, leading to        |
|        |       | serine/threonine-   | NFkB activation; promotes BCL10              |
|        |       | protein kinase 2    | phosphorylation and subsequent NFкВ          |
|        |       |                     | activation following TCR engagement          |
|        |       |                     |  |

| Syk   | 1.23  | Spleen-associated  |                     | Critical kinase that transmits signals from  |
|-------|-------|--------------------|---------------------|--|
|       |       | tyrosine kinase    |                     | the TCR and BCR                              |
| Tbk1  | 1.22  | TANK-binding       |                     | Coordinates the activation of IRF3 and NFkB  |
|       |       | kinase 1           |                     | and induction of type I IFNs                 |
| Tyk2  | 1.54  | Tyrosine kinase 2  | JTK1                | Plays both structural and catalytic roles in |
|       |       |                    |                     | numerous cytokines and interferons           |
|       |       |                    |                     | signaling; associates with cytokine and      |
|       |       |                    |                     | growth factor receptors and activate STAT    |
|       |       |                    |                     | family members including STAT1, STAT3,       |
|       |       |                    |                     | STAT4, or STAT6                              |
|       |       |                    | Lysosomal Activity  |  |
|       |       |                    |                     |  |
| Ctss  | 1.42  | Cathepsin S        |                     | Lysosomal protease that participates in      |
|       |       |                    |                     | processing of Ag by MHC class II             |
| Hck   | 1.62  | Hematopoietic cell |                     | Src family tyrosine kinase that mediates     |
|       |       | kinase             |                     | secretory lysosome mobilization,             |
|       |       |                    |                     | degranulation, and activation of NADPH       |
|       |       |                    |                     | oxidase                                      |
|       |       | <u> </u>           | Macrophage Function |  |
|       |       |                    |                     |  |
| Casp8 | 0.914 | Caspase 8          |                     | Cysteine-aspartic acid protease; cleaves     |
|       |       |                    |                     | and activates effector caspases CASP3, -4, - |
|       |       |                    |                     | 6, -7, -9, and -10; initiates pyroptosis     |
|       |       |                    |                     | through cleavage of gasdermin-D; mediates    |

|        |      |  |   | noncanonical cleavage of IL-1β in DCs and macrophages   |
|--------|------|--|---|---|
| Ccl3   | 1.01 | C-C motif<br>chemokine ligand 3            | Macrophage inflammatory protein $1\alpha \ (\text{MIP1}\alpha)$ | Chemoattractant ligand for CCR1, -4, and -5   |
| Ccr2   | 2.01 | C-C motif chemokine receptor 2             | CD192   | Receptor for CCL2, a monocyte-specific chemokine  |
| Cd14   | 2.78 | Cluster of differentiation 14              |   | PRR that recognizes LPS; mostly found on macrophages  |
| Cebpb  | 2.13 | CCAAT/enhancer-<br>binding protein<br>beta |   | Critical macrophage TF that promotes expression of several acute-phase and inflammatory cytokine genes, including II6 |
| CkIf   | 1.14 | Chemokine-like<br>factor                   |   | Chemoattractant for monocytes, neutrophils, and lymphocytes   |
| Clec5a | 2.43 | C-Type lectin  domain family 5,  member a  | Myeloid DAP12-<br>associating lectin-1                          | Critical macrophage receptor for dengue virus serotypes 1-4; positive regulator of osteoclastogenesis                 |
| Csf1   | 1.75 | Macrophage colony-stimulating factor 1     |   | Cytokine that promote activation and survival of monocytes  |

| Csf1r   | 1.94 | Macrophage          | CD115              | Receptor for CSF1; promotes release of                        |
|---------|------|---------------------|--------------------|---|
|         |      | colony-stimulating  |                    | inflammatory cytokines in response to IL-34                   |
|         |      | factor 1 receptor   |                    | and CSF1  |
|         |      |                     |                    |   |
| Irak3   | 1.31 | Interleukin-1       |                    | Adaptor protein that negatively regulates                     |
|         |      | receptor-           |                    | TLR signaling; predominantly expressed in                     |
|         |      | associated kinase 3 |                    | monocytes and macrophages                                     |
| Mif     | 1.07 | Macrophage          | L-dopachrome       | Pro-inflammatory cytokine that promotes                       |
|         |      | migration           | tautomerase        | macrophage function through suppression                       |
|         |      | inhibitory factor   |                    | of anti-inflammatory effects of                               |
|         |      |                     |                    | glucocorticoids   |
| Slc11a1 | 2.01 | Natural resistance- |                    | Macrophage-specific metal ion transporter;                    |
|         |      | associated          |                    | uptakes divalent metal cations to neutralize                  |
|         |      | macrophage          |                    | ROSs  |
|         |      | protein 1           |                    |   |
|         |      | protein 1           |                    |   |
| Stat6   | 1.22 | Signal transducer   |                    | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and |
|         |      | and activator of    |                    | macrophage function and M2 macrophage                         |
|         |      | transcription 6     |                    | polarization  |
|         |      |                     | Mast Cell Function |   |
|         |      |                     |                    |   |
| Lgals3  | 1.99 | Galectin 3          |                    | Galactose-specific lectin that binds IgE;                     |
|         |      |                     |                    | involved in acute inflammatory responses,                     |
|         |      |                     |                    | including neutrophil activation and                           |
|         |      |                     |                    | adhesion, chemoattraction of monocytes                        |

|       |       |                   |                       | macrophages, opsonization of apoptotic         |
|-------|-------|-------------------|-----------------------|--|
|       |       |                   |                       | neutrophils, and activation of mast cells      |
| Ms4a2 | 1.57  | Membrane          | Fc fragment of IgE,   | High affinity receptor that binds to the Fc    |
|       |       | spanning 4-       | high affinity I,      | region of IgE; required for the full mast cell |
|       |       | domains A2        | receptor subunit beta | response; also mediates the secretion of       |
|       |       |                   | (FCER1B)              | important lymphokines                          |
|       |       |                   | Immunoglobulin E      |  |
|       |       |                   | receptor (IGER)       |  |
|       |       |                   | MAP Kinase Signaling  |  |
|       |       |                   |                       |  |
| Mapk1 | 0.938 | Mitogen-activated | Extracellular signal- | Serine/threonine kinase that acts as an        |
|       |       | protein kinase 1  | regulated kinase 2    | essential component of the MAP kinase          |
|       |       |                   | (ERK2)                | signal transduction pathway                    |
| Mapk3 | 1.57  | Mitogen-activated | Extracellular signal- | Serine/threonine kinase that acts as an        |
|       |       | protein kinase 3  | regulated kinase 1    | essential component of the MAP kinase          |
|       |       |                   | (ERK1)                | signal transduction pathway                    |
| Mapk8 | 0.903 | Mitogen-activated | c-Jun N-terminal      | Serine/threonine-protein kinase involved in    |
|       |       | protein kinase 8  | kinase 1 (JNK1);      | various processes such as cell proliferation,  |
|       |       |                   | Stress-activated      | differentiation, migration, transformation     |
|       |       |                   | protein kinase 1c     | and programmed cell death;                     |
|       |       |                   | (SAPK1)               | phosphorylates a number of TFs, primarily      |
|       |       |                   |                       | components of AP-1 such as JUN, JDP2, and      |
|       |       |                   |                       | ATF2, thus regulating AP-1 transcriptional     |
|       |       |                   |                       | activity; promotes stressed cell apoptosis     |

|        |       |                   |                   | by phosphorylating key regulatory factors    |
|--------|-------|-------------------|-------------------|--|
|        |       |                   |                   | including p53/TP53 and Yes-associates        |
|        |       |                   |                   | protein YAP1; required for T <sub>H</sub> 1  |
|        |       |                   |                   | differentiation                              |
| Mapk14 | 1.42  | Mitogen-activated |                   | One of the four p38 MAPKs; key kinase in     |
|        |       | protein kinase 14 |                   | the cascades of cellular responses evoked    |
|        |       |                   |                   | by extracellular stimuli such as             |
|        |       |                   |                   | proinflammatory cytokines                    |
| Map2k1 | 0.824 | Dual specificity  | MAPK/ERK kinase 1 | Essential component of the MAP kinase        |
|        |       | mitogen-activated | (MEK1)            | signal transduction pathway; participates in |
|        |       | protein kinase    |                   | numerous biological functions, including     |
|        |       | kinase 1          |                   | cell growth, adhesion, survival,             |
|        |       |                   |                   | differentiation, transcription, metabolism,  |
|        |       |                   |                   | and cytoskeletal remodeling                  |
| Map2k2 | 1.22  | Mitogen-activated | MAPK/ERK kinase 2 | Catalyzes the concomitant phosphorylation    |
|        |       | protein kinase    | (MEK2)            | of a threonine and a tyrosine residue in a   |
|        |       | kinase 2          |                   | TQY sequence located in MAP kinases;         |
|        |       |                   |                   | activates ERK1 and -2                        |
| Map2k4 | 1.03  | Mitogen-activated | MAPK/ERK kinase 4 | Dual specificity protein kinase that acts as |
|        |       | protein kinase    | (MEK4); c-Jun N-  | an essential component of the stress-        |
|        |       | kinase 4          | terminal kinase   | activated protein/c-Jun N-terminal kinase    |
|        |       |                   | kinase 1 (JNKK1)  | (SAP/JNK) signaling pathway                  |

| Map3k1   | 0.853 | Mitogen-activated protein kinase kinase kinase 1 |                                      | Serine/threonine kinase that activates the ERK and JNK kinase pathways by phosphorylation of MAP2K1 and MAP2K4; also activates CHUK and IKBKB, the central protein kinases of the NFKB pathway  |
|----------|-------|--|--------------------------------------|---|
| Map3k5   | 1.54  | Mitogen-activated protein kinase kinase kinase 5 | Apoptosis signal-regulating kinase 1 | Essential component of the MAP kinase signal transduction pathway; mediates signaling for determination of cell fate such as differentiation and survival; plays a crucial role in the apoptosis signal transduction pathway through mitochondria-dependent caspase activation; required for the innate immune response; mediates signal transduction of receptor-mediated inflammatory signals, such as TNF or LPS |
| Map3k7   | 0.844 | Mitogen-activated protein kinase kinase kinase 7 | TGFβ-activated kinase (TAK1)         | Signal transducer downstream of TGFβ and BMP; controls a variety of cell functions, including transcription regulation and apoptosis  |
| Mapkapk2 | 1.49  | MAP kinase-<br>activated protein<br>kinase 2     |                                      | Serine/threonine-protein kinase involved in cytokine production, endocytosis, reorganization of the cytoskeleton, cell migration, cell cycle control, chromatin   |

|        |       |                    |                        | anno della e DNA de                          |
|--------|-------|--------------------|------------------------|--|
|        |       |                    |                        | remodeling, DNA damage response, and         |
|        |       |                    |                        | transcriptional regulation                   |
| Tab1   | 0.88  | TGF-beta activated |                        | Activates TAK1 kinase; mediates various      |
|        |       | kinase 1 binding   |                        | intracellular signaling pathways, such as    |
|        |       | protein 1          |                        | those induced by TGFβ, IL-1, and WNT-1       |
|        |       |                    | Metabolism             |  |
|        |       |                    |                        |  |
| Cd36   | 2.24  | Cluster of         | Fatty acid translocase | Class B scavenger receptor that mediates     |
|        |       | differentiation 36 | (FAT)                  | fatty acid uptake                            |
| Map2k1 | 0.824 | Dual specificity   | MAPK/ERK kinase 1      | Essential component of the MAP kinase        |
|        |       | mitogen-activated  | (MEK1)                 | signal transduction pathway; participates in |
|        |       | protein kinase     |                        | numerous biological functions, including     |
|        |       | kinase 1           |                        | cell growth, adhesion, survival,             |
|        |       |                    |                        | differentiation, transcription, metabolism,  |
|        |       |                    |                        | and cytoskeletal remodeling                  |
| S100a8 | 2.82  | S100 calcium-      | Calgranulin A          | Calcium- and zinc-binding protein involved   |
|        |       | binding protein A8 |                        | in pro-inflammatory, antimicrobial, oxidant- |
|        |       |                    |                        | scavenging and apoptosis-inducing            |
|        |       |                    |                        | activities; can induce neutrophil            |
|        |       |                    |                        | chemotaxis, adhesion, phagocytosis, and      |
|        |       |                    |                        | degranulation; predominantly found as        |
|        |       |                    |                        | calprotectin (S100A8/A9) which has a wide    |
|        |       |                    |                        | plethora of intra- and extracellular         |
|        |       |                    |                        | functions, including adhesion, apoptosis,    |
|        |       |                    |                        |  |

|        |      |                                |                    | autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid  |
|--------|------|--------------------------------|--------------------|---|
|        |      |                                |                    | metabolism, oxidant-scavenging, and PRR signaling   |
|        |      |                                | Migration/Motility |   |
| Angpt1 | 1.72 | Angiopoietin 1                 |                    | Secreted glycoprotein that binds and activates the TEK/TIE2 receptor by inducing its dimerization and tyrosine phosphorylation; involved in angiogenesis, endothelial cell survival, proliferation, migration, adhesion, cell spreading, and reorganization of the actin cytoskeleton |
| Арр    | 2.46 | Amyloid-beta precursor protein |                    | Cell surface receptor and transmembrane  precursor protein that is cleaved by  secretases to form a number of peptides;  involved in cell mobility, copper  homeostasis, and oxidative stress   |
| Cd97   | 1.35 | Cluster of differentiation 97  | BL-Ac[F2]          | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells  |

| S100a8 | 2.82 | S100 calcium-      | Calgranulin A        | Calcium- and zinc-binding protein involved   |
|--------|------|--------------------|----------------------|--|
|        |      | binding protein A8 |                      | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |                      | scavenging and apoptosis-inducing            |
|        |      |                    |                      | activities; can induce neutrophil            |
|        |      |                    |                      | chemotaxis, adhesion, phagocytosis, and      |
|        |      |                    |                      | degranulation; predominantly found as        |
|        |      |                    |                      | calprotectin (S100A8/A9) which has a wide    |
|        |      |                    |                      | plethora of intra- and extracellular         |
|        |      |                    |                      | functions, including adhesion, apoptosis,    |
|        |      |                    |                      | autophagy, cytoskeletal remodeling,          |
|        |      |                    |                      | cytokine production, chemotaxis,             |
|        |      |                    |                      | migration, inflammation, arachidonic acid    |
|        |      |                    |                      | metabolism, oxidant-scavenging, and PRR      |
|        |      |                    |                      | signaling                                    |
|        |      |                    | Neutrophil Function  |  |
|        |      |                    |                      |  |
| Cklf   | 1.14 | Chemokine-like     |                      | Chemoattractant for monocytes,               |
|        |      | factor             |                      | neutrophils, and lymphocytes                 |
| Cxcr2  | 2.38 | C-X-C motif        | CD182; IL-8 receptor | Receptor for IL-8 and CXCL3; powerful        |
|        |      | chemokine          | В                    | chemoattractant for neutrophils              |
|        |      | receptor 2         |                      |  |
| Fpr2   | 2.91 | Formyl peptide     | Lipoxin A4 receptor  | Low affinity receptor for N-formyl-          |
|        |      | receptor 2         |                      | methionyl peptides; activates neutrophils    |
|        |      |                    |                      |  |

| Lcn2   | 2.98 | Lipocalin 2        | Neutrophil            | Neutrophil-secreted factor that sequesters   |
|--------|------|--------------------|-----------------------|--|
|        |      |                    | gelatinase-associated | iron-containing siderophores; also           |
|        |      |                    | lipocalin (NGAL)      | functions as a growth factor                 |
|        |      |                    | , ,                   |  |
| Lgals3 | 1.99 | Galectin 3         |                       | Galactose-specific lectin that binds IgE;    |
|        |      |                    |                       | involved in acute inflammatory responses,    |
|        |      |                    |                       | including neutrophil activation and          |
|        |      |                    |                       | adhesion, chemoattraction of monocytes       |
|        |      |                    |                       | macrophages, opsonization of apoptotic       |
|        |      |                    |                       | neutrophils, and activation of mast cells    |
|        |      |                    |                       |  |
| Ncf4   | 1.98 | Neutrophil         | SH3 and PX domain-    | Cytosolic regulatory component of the        |
|        |      | cytosolic factor 4 | containing protein 4  | superoxide-producing phagocyte NADPH-        |
|        |      |                    | (SH3PXD4)             | oxidase, a multicomponent enzyme system      |
|        |      |                    |                       | important for host defense                   |
|        |      |                    |                       |  |
| S100a8 | 2.82 | S100 calcium-      | Calgranulin A         | Calcium- and zinc-binding protein involved   |
|        |      | binding protein A8 |                       | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |                       | scavenging and apoptosis-inducing            |
|        |      |                    |                       | activities; can induce neutrophil            |
|        |      |                    |                       | chemotaxis, adhesion, phagocytosis, and      |
|        |      |                    |                       | degranulation; predominantly found as        |
|        |      |                    |                       | calprotectin (S100A8/A9) which has a wide    |
|        |      |                    |                       | plethora of intra- and extracellular         |
|        |      |                    |                       | functions, including adhesion, apoptosis,    |
|        |      |                    |                       | autophagy, cytoskeletal remodeling,          |
|        |      |                    |                       | cytokine production, chemotaxis,             |
|        |      |                    |                       | -,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,      |

|        |       |                      |                | migration, inflammation, arachidonic acid     |
|--------|-------|----------------------|----------------|---|
|        |       |                      |                | metabolism, oxidant-scavenging, and PRR       |
|        |       |                      |                | signaling                                     |
|        |       |                      |                |   |
|        |       |                      | NFkB Signaling |   |
|        |       |                      |                |   |
| Bcl10  | 1.26  | B cell               |                | Activates NFκB via ubiquitination of ΙΚΚγ     |
|        |       | lymphoma/leukemi     |                |   |
|        |       | a 10                 |                |   |
|        |       |                      |                |   |
| Ikbkb  | 1.02  | Inhibitor of nuclear |                | Part of the IKK complex that inhibits IκBα    |
|        |       | factor kappa B       |                | and permits NFkB nuclear localization         |
|        |       | kinase subunit beta  |                |   |
|        |       |                      |                |   |
| Nfkb1  | 0.928 | Nuclear factor       | p105/p50       | One of the NFkB family TFs; inhibits          |
|        |       | kappa B subunit 1    |                | inflammation                                  |
|        |       |                      |                |   |
| Nfkb2  | 0.857 | Nuclear factor       | p100/p52       | One of the NFkB family TFs; major driver of   |
|        |       | kappa B subunit 2    |                | inflammation                                  |
|        |       |                      |                |   |
| Nfkbia | 1.19  | Nuclear factor       |                | Inhibits activity of REL dimers by masking of |
|        |       | kappa B inhibitor    |                | their nuclear localization signals            |
|        |       | alpha                |                |   |
|        |       |                      |                |   |
| Rela   | 1     | Avian                | p65            | One of the NFkB family TFs; major driver of   |
|        |       | reticuloendothelios  |                | inflammation                                  |
|        |       | is viral oncogene    |                |   |
|        |       | _                    |                |   |
|        |       | homolog A            |                |   |
|        |       |                      |                |   |

| Ripk2   | 1.37 | Receptor-           |                        | RIP kinase that potentiates signals        |
|---------|------|---------------------|------------------------|--|
|         |      | interacting         |                        | downstream of NOD1 and -2, leading to      |
|         |      | serine/threonine-   |                        | NFkB activation; promotes BCL10            |
|         |      | protein kinase 2    |                        | phosphorylation and subsequent NFкВ        |
|         |      |                     |                        | activation following TCR engagement        |
|         |      | Patte               | ern Recognition Recept | ors  |
|         |      |                     |                        |  |
| Cd14    | 2.78 | Cluster of          |                        | PRR that recognizes LPS; mostly found on   |
|         |      | differentiation 14  |                        | macrophages                                |
| Cd180   | 0.93 | Cluster of          |                        | Heterodimeric binding partner of Ly86 that |
|         |      | differentiation 180 |                        | participates in LPS binding in APCs        |
| Clec4a2 | 2.18 | C-type lectin       |                        | PRR that, upon binding mannose or fucose,  |
|         |      | domain family 4     |                        | is endocytosed and processed in the Ag     |
|         |      | member A2           |                        | presentation pathway                       |
| Clec5a  | 2.43 | C-Type lectin       | Myeloid DAP12-         | Critical macrophage receptor for dengue    |
|         |      | domain family 5,    | associating lectin-1   | virus serotypes 1-4; positive regulator of |
|         |      | member a            |                        | osteoclastogenesis                         |
| Clec7a  | 2.29 | C-Type lectin       | Dectin-1               | PRR specific for β-1,3- and β-1,6-linked   |
|         |      | domain family 7,    |                        | glucans from fungi and plants; necessary   |
|         |      | member a            |                        | for the TLR2-mediated inflammatory         |
|         |      |                     |                        | response and for TLR2-mediated activation  |
|         |      |                     |                        | of NF-κB                                   |
|         |      |                     |                        |  |

| Ddx58 | 0.956 | DExD/H-box         | Retinoic acid-         | Cytoplasmic PRR that recognizes dsRNA;               |
|-------|-------|--------------------|------------------------|--|
|       |       | helicase 58        | inducible gene I (RIG- | can promote T cell-independent B cell                |
|       |       |                    | I)                     | activation; uses MAVS as an adaptor                  |
| Ifih1 | 1.09  | Interferon induced | Helicard; melanoma     | PRR for cytoplasmic dsRNA; upon target               |
|       |       | with helicase C    | differentiation-       | recognition, associates with MAVS to                 |
|       |       | domain 1           | associated protein 5   | activate TNK1 and IKBKE, which                       |
|       |       |                    | (MDA5)                 | phosphorylate IRF3 and -7, which, in turn,           |
|       |       |                    |                        | activate transcription of IFN $\alpha$ and - $\beta$ |
| Ly86  | 1.15  | Lymphocyte         | Myeloid                | Heterodimeric binding partner of CD180               |
|       |       | antigen 86         | differentiation factor | that participates in LPS binding in APCs             |
|       |       |                    | 1 (MD-1)               |  |
| Ly96  | 2.07  | Lymphocyte         | Myeloid                | Heterodimeric binding partner of TLR4 that           |
|       |       | antigen 96         | differentiation factor | participates in LPS binding                          |
|       |       |                    | 2 (MD-2)               |  |
| Myd88 | 1.97  | Myeloid            |                        | Key adaptor in the TLR signaling pathways;           |
|       |       | differentiation    |                        | interacts with all TLRs except TLR3;                 |
|       |       | primary response   |                        | activates NFкB and IRFs                              |
|       |       | 88                 |                        |  |
| Nod2  | 1.2   | Nucleotide-binding |                        | PRR specific for muramyl dipeptide (MDP);            |
|       |       | oligomerization    |                        | upon binding to its ligand, recruits RIPK2           |
|       |       | domain containing  |                        | and triggers MAPK and NFkB signaling                 |
|       |       | 2                  |                        |  |
|       |       |                    |                        |  |

| Nlrp3  | 2.42 | NACHT domain-,     | Cryopyrin     | PRR with a wide diversity of recognized      |
|--------|------|--------------------|---------------|--|
|        |      | leucine-rich       |               | targets that activates the NLRP3             |
|        |      | repeat-, and PYD-  |               | inflammasome consisting of NLRP3,            |
|        |      | containing protein |               | PYCARD, and caspase-1/-8                     |
|        |      | 3                  |               |  |
| S100a8 | 2.82 | S100 calcium-      | Calgranulin A | Calcium- and zinc-binding protein involved   |
|        |      | binding protein A8 |               | in pro-inflammatory, antimicrobial, oxidant- |
|        |      |                    |               | scavenging and apoptosis-inducing            |
|        |      |                    |               | activities; can induce neutrophil            |
|        |      |                    |               | chemotaxis, adhesion, phagocytosis, and      |
|        |      |                    |               | degranulation; predominantly found as        |
|        |      |                    |               | calprotectin (S100A8/A9) which has a wide    |
|        |      |                    |               | plethora of intra- and extracellular         |
|        |      |                    |               | functions, including adhesion, apoptosis,    |
|        |      |                    |               | autophagy, cytoskeletal remodeling,          |
|        |      |                    |               | cytokine production, chemotaxis,             |
|        |      |                    |               | migration, inflammation, arachidonic acid    |
|        |      |                    |               | metabolism, oxidant-scavenging, and PRR      |
|        |      |                    |               | signaling                                    |
| Ticam2 | 1.87 | TIR domain-        |               | Sorting adapter in various innate immune     |
|        |      | containing adaptor |               | signaling cascades; bridges TLR2 and         |
|        |      | molecule 2         |               | MyD88  |

| Tirap | 2    | TIR domain-<br>containing adaptor<br>protein |              | Adaptor protein involved in TLR2 and TLR4 signaling; acts via IRAK2 and TRAF6, leading to the activation of NFκB, MAPK1, MAPK3 and JNK, and resulting in cytokine secretion and the inflammatory response; positively regulates the production of TNFα and IL-6.                              |
|-------|------|--|--------------|---|
| Tlr2  | 2.27 | Toll-like receptor 2                         | CD282        | Surface PRR that binds to various lipid-<br>containing PAMPs  |
| Tlr4  | 2.38 | Toll-like receptor 4                         | CD284        | Surface PRR that recognizes LPS; pairs with LY96 and CD14; acts via MYD88, TIRAP, and TRAF6, leading to NFκB activation, cytokine secretion, and the inflammatory response; in complex with TLR6, promotes sterile inflammation in monocytes/macrophages in response to oxLDL or amyloid-β 42 |
| Tlr6  | 2.43 | Toll like receptor 6                         | CD286        | Surface PRR that recognizes diacylated and triacylated lipopeptides   |
| Tlr7  | 2.49 | Toll-like receptor 7                         | CD287        | Endosomic PRR that recognizes ssRNA   |
| Tlr8  | 2.26 | Toll-like receptor 8                         | CD288        | Endosomic PRR that recognizes ssRNA   |
|       |      | ·  | Phagocytosis |   |
| Itgam | 1.77 | Integrin alpha M                             | CD11b        | Pairs with CD18 to forms Mac-1 aka complement receptor 3; mediates  |

| Ncf4   | 1.98 | Neutrophil cytosolic factor 4  | SH3 and PX domain-<br>containing protein 4<br>(SH3PXD4) | cytotoxicity; serves as a macrophage  marker  Cytosolic regulatory component of the  superoxide-producing phagocyte NADPH- oxidase, a multicomponent enzyme system  important for host defense   |
|--------|------|--------------------------------|---|--|
| S100a8 | 2.82 | S100 calciumbinding protein A8 | Calgranulin A   | Calcium- and zinc-binding protein involved in pro-inflammatory, antimicrobial, oxidant-scavenging and apoptosis-inducing activities; can induce neutrophil chemotaxis, adhesion, phagocytosis, and degranulation; predominantly found as calprotectin (S100A8/A9) which has a wide plethora of intra- and extracellular functions, including adhesion, apoptosis, autophagy, cytoskeletal remodeling, cytokine production, chemotaxis, migration, inflammation, arachidonic acid metabolism, oxidant-scavenging, and PRR signaling |

**ROS Generation & Protection** 

| Cybb  | 1.8   | Cytochrome b-245    | Nox2                 | Part of the NADPH oxidase process;            |
|-------|-------|---------------------|----------------------|---|
|       |       | heavy chain         |                      | generates superoxides                         |
| Ncf4  | 1.98  | Neutrophil          | SH3 and PX domain-   | Cytosolic regulatory component of the         |
|       |       | cytosolic factor 4  | containing protein 4 | superoxide-producing phagocyte NADPH-         |
|       |       |                     | (SH3PXD4)            | oxidase, a multicomponent enzyme system       |
|       |       |                     |                      | important for host defense                    |
| Txnip | 1.1   | Thioredoxin         |                      | Thiol-oxidoreductase; protects cells from     |
|       |       | interacting protein |                      | oxidative stress by inhibiting thioredoxin    |
|       |       |                     | Stress Response      |   |
| Арр   | 2.46  | Amyloid-beta        |                      | Cell surface receptor and transmembrane       |
|       |       | precursor protein   |                      | precursor protein that is cleaved by          |
|       |       |                     |                      | secretases to form a number of peptides;      |
|       |       |                     |                      | involved in cell mobility, copper             |
|       |       |                     |                      | homeostasis, and oxidative stress             |
| Mapk8 | 0.903 | Mitogen-activated   | c-Jun N-terminal     | Serine/threonine-protein kinase involved in   |
|       |       | protein kinase 8    | kinase 1 (JNK1);     | various processes such as cell proliferation, |
|       |       |                     | Stress-activated     | differentiation, migration, transformation    |
|       |       |                     | protein kinase 1c    | and programmed cell death;                    |
|       |       |                     | (SAPK1)              | phosphorylates a number of TFs, primarily     |
|       |       |                     |                      | components of AP-1 such as JUN, JDP2, and     |
|       |       |                     |                      | ATF2, thus regulating AP-1 transcriptional    |
|       |       |                     |                      | activity; promotes stressed cell apoptosis    |
|       |       |                     |                      | by phosphorylating key regulatory factors     |

| Map2k4 | 1.03            | Mitogen-activated               | MAPK/ERK kinase 4                                       | including p53/TP53 and Yes-associates  protein YAP1; required for TH1  differentiation  Dual specificity protein kinase that acts as   |  |
|--------|-----------------|---------------------------------|---|--|--|
| ·      |                 | protein kinase<br>kinase 4      | (MEK4); c-Jun N-<br>terminal kinase<br>kinase 1 (JNKK1) | an essential component of the stress-<br>activated protein/c-Jun N-terminal kinase<br>(SAP/JNK) signaling pathway  |  |
| Txnip  | 1.1             | Thioredoxin interacting protein |   | Thiol-oxidoreductase; protects cells from oxidative stress by inhibiting thioredoxin   |  |
|        | T Cell Function |                                 |   |  |  |
| Cd28   | 1.04            | Cluster of differentiation 28   |   | Essential T cell co-receptor that enhances T cell activation, proliferation, cytokine production, and survival; binds to CD80 and CD86   |  |
| Cd4    | 0.937           | Cluster of differentiation 4    |   | Signature helper T cell marker; binds to  MHC class II and provides necessary  costimulation for T cell activation   |  |
| Cd97   | 1.35            | Cluster of differentiation 97   | BL-Ac[F2]   | GPCR that promotes granulocyte adhesion and migration; activates T cells via binding to CD55; stimulates angiogenesis through binding integrin counterreceptors on endothelial cells |  |

| Pik3cg | 1.44  | Phosphatidylinosito  |                        | A subunit of PI3K; modulates leukocyte       |
|--------|-------|----------------------|------------------------|--|
|        |       | I-4,5-bisphosphate   |                        | chemotaxis to inflammatory sites and in      |
|        |       | 3-kinase catalytic   |                        | response to chemoattractant agents           |
|        |       | subunit gamma        |                        |  |
|        |       | isoform              |                        |  |
|        |       |                      |                        |  |
|        |       | Transcri             | ption Factors & Coacti | vators                                       |
|        |       |                      |                        |  |
| Atf1   | 0.973 | Activating           |                        | Basic leucine zipper TF regulating the       |
|        |       | transcription factor |                        | expression of genes related to growth and    |
|        |       | 1                    |                        | survival                                     |
|        |       |                      |                        |  |
| Atf2   | 1.13  | Activating           | Cyclic AMP-            | Regulates transcription of various genes     |
|        |       | transcription factor | responsive element-    | involved in anti-apoptosis, cell growth, and |
|        |       | 2                    | binding protein 2      | DNA damage response; in the nucleus,         |
|        |       |                      | (CREB2)                | contributes to global transcription and the  |
|        |       |                      |                        | DNA damage response, in addition to          |
|        |       |                      |                        | specific transcriptional activities that are |
|        |       |                      |                        | related to cell development, proliferation   |
|        |       |                      |                        | and death; in the cytoplasm, impairs         |
|        |       |                      |                        | mitochondrial membrane potential,            |
|        |       |                      |                        | inducing mitochondrial leakage and           |
|        |       |                      |                        | promoting cell death; phosphorylated form    |
|        |       |                      |                        | (mediated by ATM) plays a role in the DNA    |
|        |       |                      |                        | damage response                              |
|        |       |                      |                        |  |

| Cebpb  | 2.13  | CCAAT/enhancer-binding protein |                      | Critical macrophage TF that promotes expression of several acute-phase and |
|--------|-------|--------------------------------|----------------------|--|
|        |       | beta                           |                      | inflammatory cytokine genes, including II6                                 |
| Crebbp | 0.949 | CREB binding                   | Lysine               | Binds specifically to phosphorylated CREB                                  |
|        |       | protein                        | acetyltransferase 3A | and enhances its transcriptional activity                                  |
|        |       |                                | (KAT3A)              | toward cAMP-responsive genes; also   |
|        |       |                                |                      | acetylates histones, giving a specific tag for                             |
|        |       |                                |                      | transcriptional activation   |
| Fos    | 2.49  | Finkel-Biskel-                 |                      | Basic leucine zipper TF that dimerizes with                                |
|        |       | Jenkins                        |                      | JUN proteins to form the AP-1 TF complex                                   |
|        |       | osteosarcoma                   |                      |  |
|        |       | proto-oncogene                 |                      |  |
|        |       | homolog                        |                      |  |
| Foxp3  | 1.5   | Forkhead box P3                | DIETER               | Master TF for Tregs; represses expression                                  |
|        |       |                                |                      | of II2 and Ifng; activates expression of                                   |
|        |       |                                |                      | Tnfrsf18, Il2ra, and Ctla4   |
| Nfkb1  | 0.928 | Nuclear factor                 | p105/p50             | One of the NFкВ family TFs; inhibits                                       |
|        |       | kappa B subunit 1              |                      | inflammation   |
| Nfkb2  | 0.857 | Nuclear factor                 | p100/p52             | One of the NFkB family TFs; major driver of                                |
|        |       | kappa B subunit 2              |                      | inflammation   |
| Rela   | 1     | Avian                          | p65                  | One of the NFkB family TFs; major driver of                                |
|        |       | reticuloendothelios            |                      | inflammation   |

|       |      | is viral oncogene                                  |   |   |
|-------|------|--|---|---|
| Stat1 | 1.24 | Signal transducer and activator of transcription 1 |   | Transcriptional activator that mediates  cellular responses to IFNs, cytokines, and  other growth factors   |
| Stat3 | 1.26 | Signal transducer and activator of transcription 3 |   | Transcriptional activator of genes involved in cell growth and apoptosis; activated by JAKs   |
| Stat6 | 1.22 | Signal transducer and activator of transcription 6 |   | Essential TF for T <sub>H</sub> 2 CD4 <sup>+</sup> T cell and macrophage function and M2 macrophage polarization  |
| Zbp1  | 1.05 | Z-DNA binding protein 1                            | Tumor stroma and activated macrophage protein DLM-1 | Key innate sensor that recognizes and binds Z-RNA structures, which are produced by a number of viruses, and induces type-I IFN production; key activator of cellular necroptosis; promotes IL-1α induction in an NLRP3-inflammasome-independent manner |
|       |      |  | Ubiquitin Regulation                                |   |
| Bcl10 | 1.26 | B cell<br>lymphoma/leukemi<br>a 10                 |   | Activates NFκB via ubiquitination of ΙΚΚγ   |

| Cyld    | 0.997 | Cylindromatosis lysine 63 deubiquitinase         | Inhibits NFkB activation by deubiquitinating upstream signaling factors; inhibits Wnt signaling; restricts polyubiquitination of RIPK1 and -2, thereby limiting necroptosis |
|---------|-------|--|---|
| Itch    | 0.947 | Itchy E3 ubiquitin protein ligase                | Participates with TNFAIP3 in a ubiquitin- editing complex that marks components of inflammatory signaling pathways such as JUNB and CXCR4 for degradation                   |
| Tnfaip3 | 1.38  | Tumor necrosis  factor, alpha- induced protein 3 | Ubiquitin-editing enzyme that complexes  with ITCH to degrade inflammatory  signaling components in the TNF, IL1, and  TLR pathways; targets TRAF2, TRAF6, and  IKK         |
| Ubc     | 0.99  | Polyubiquitin C                                  | Serves various roles, including immate immunity, DNA repair, and stimulation of autophagy and the proteasomal response  |
|         |       |  | Vesicular Trafficking   |
| Atg5    | 1.26  | Autophagy related 5                              | Pairs with ATG12 to promote the extension of the phagophoric membrane in autophagic vesicles  |