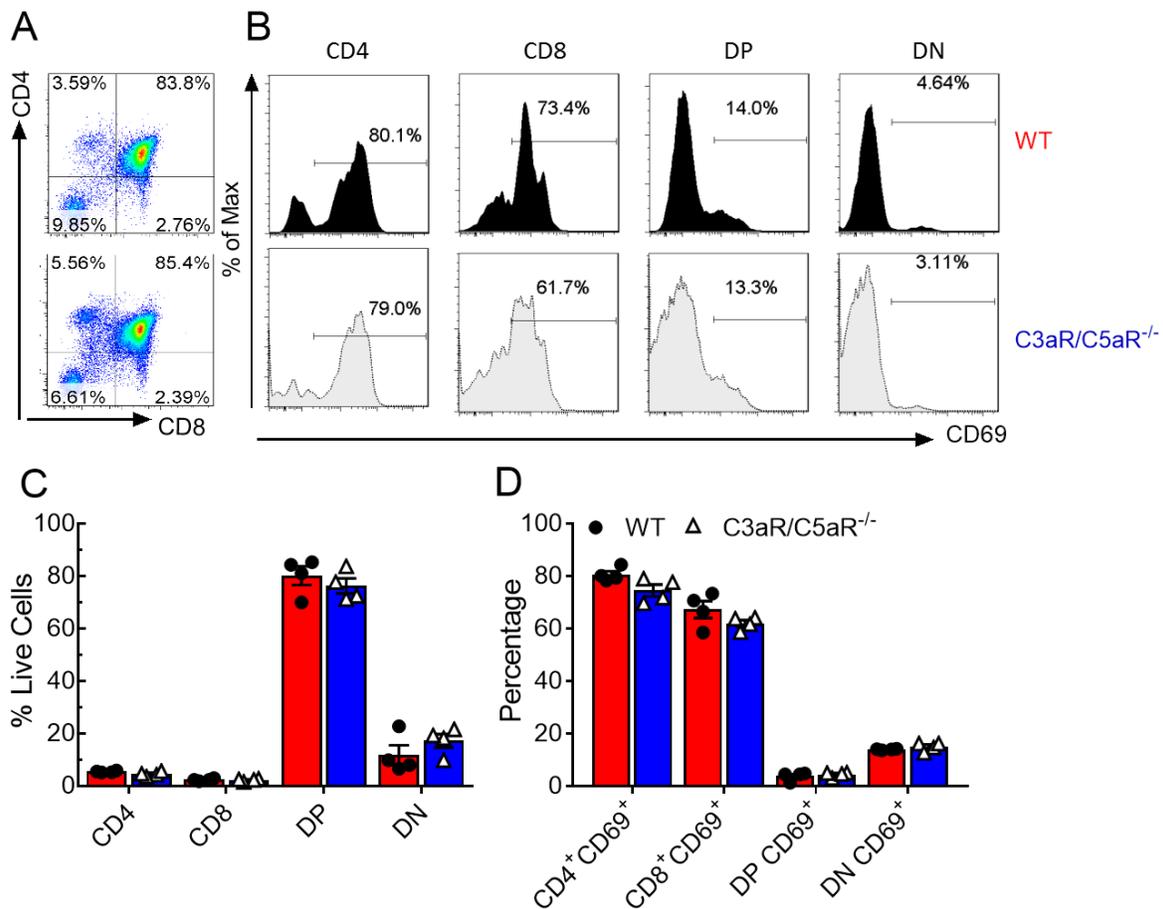


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732 **Fig. S1. (A-E) Deficiency of C3aR/C5aR increases ceramide-triggered mitophagy in recipient DCs after HCT. (A)** WT BALB/c  
 733 mice were lethally irradiated at 700 cGy ( $n=3$ ). Representative histograms for the C5aR expressed on MHCII<sup>+</sup>CD11c<sup>+</sup> DCs 24hr after  
 734 TBI. **(B and C)** WT and C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> mice ( $n=5$ ) were lethally irradiated at 700 cGy. Representative histograms or summary bar  
 735 graph for the expression of annexin V and Fas **(B)** are shown. **(D and E)** Irradiated BM derived-DCs were matured with 20 ng/mL  
 736 LPS. The relative ratios of amounts of sphingosine ceramide **(D)** and glucosyl/galactosyl ceramide **(E)** in DCs differentiated from  
 737 bone marrow of WT and C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> mice ( $n=3$ ) are shown. **(F and G)** Lethally irradiated WT or C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> recipients ( $n=5$ )  
 738 were transplanted with CFSE-labelled T-cells. The splenic cells were analyzed 4d after cell transfer. Representative flow cytometry  
 739 histogram and summary bar graphs for the expression of FasL of recipient DCs **(C)**; and bcl-2 **(F)** and DR-5 **(G)** of donor H2b<sup>+</sup> CD4<sup>+</sup>  
 740 or CD8<sup>+</sup> Tc are shown. Unpaired two-tail *t*-test was used to compare between groups. Data are representative of two independent  
 741 experiments and presented as mean  $\pm$  S.E.M., and \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

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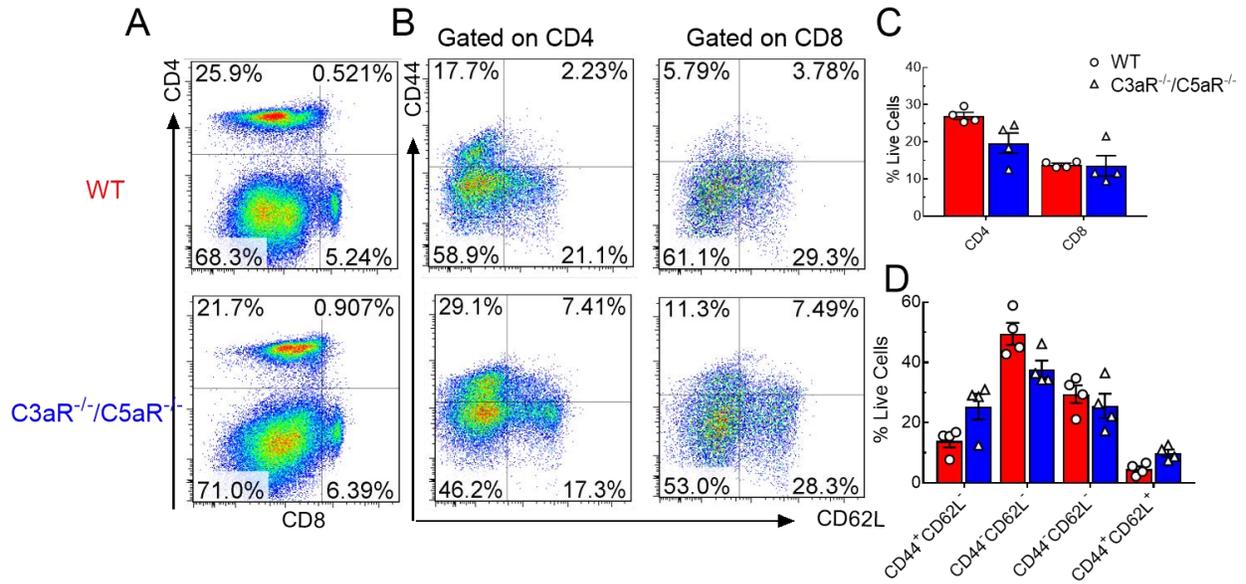
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746 **Fig. S2. C3aR/C5aR deficiency does not affect T-cells development and maturation in**  
747 **thymus.** WT or C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> BALB/c mice (*n*=4) were euthanized and thymus were isolated  
748 and analyzed by flow cytometry. Flow color contour plots present the frequencies of CD4<sup>+</sup>, CD8<sup>+</sup>,  
749 CD4<sup>+</sup>CD8<sup>+</sup> (DP), CD4<sup>+</sup>CD8<sup>-</sup> (DN). Flow histograms illustrate the fluorescence intensity of  
750 CD69<sup>+</sup>CD4<sup>+</sup>, CD69<sup>+</sup>CD8<sup>+</sup>, DP and DN cells in the thymus. **The result is representative of two**  
751 **independent experiments.** Unpaired two-tail *t*-test was used to compare between groups.

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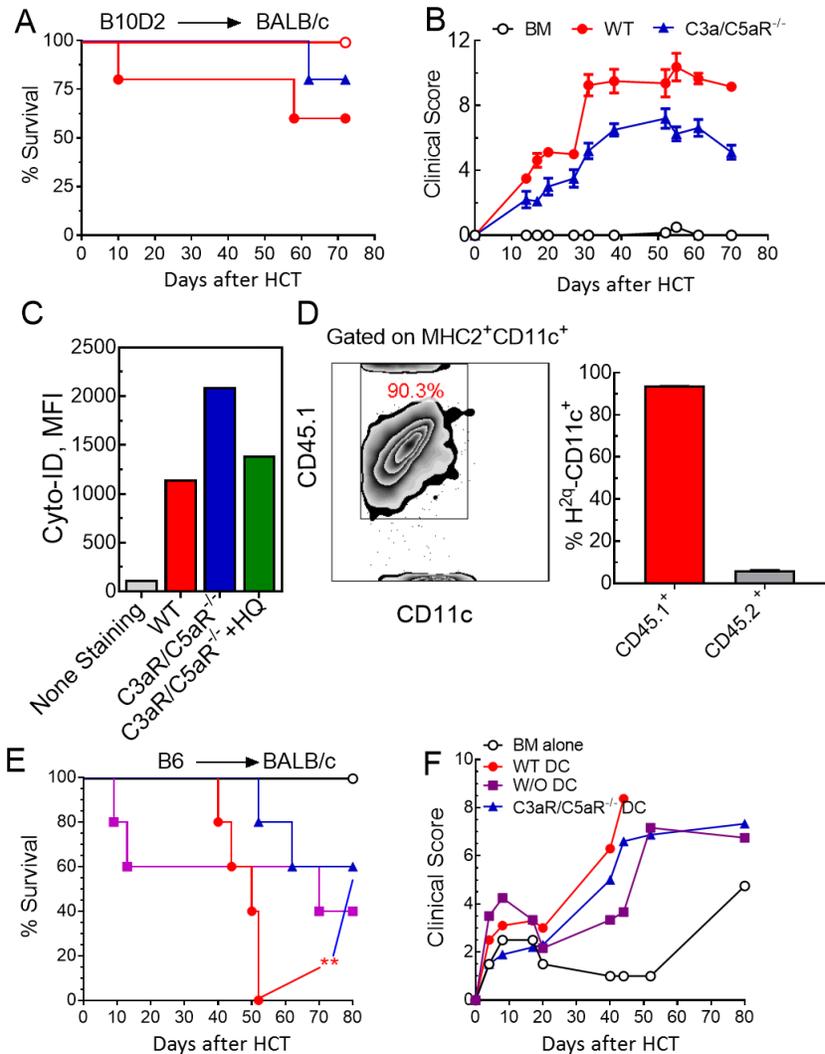


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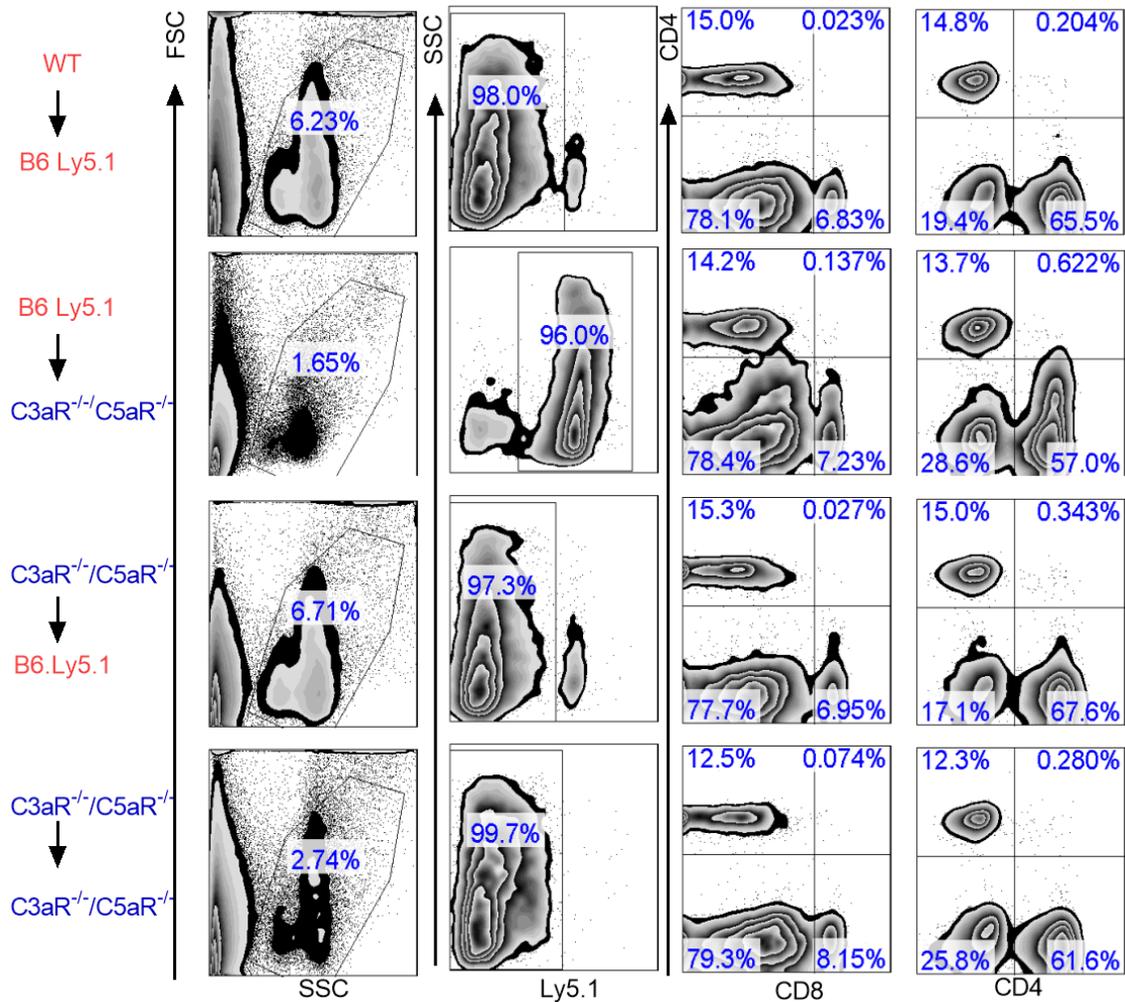
755 **Fig. S3. C3aR/C5aR deficiency does not affect mature T-cells in spleen.** WT or C3aR<sup>-/-</sup>  
 756 /C5aR<sup>-/-</sup> BALB/c mice (n=4) were euthanized and spleen were isolated and analyzed by flow  
 757 cytometry. Flow color contour plots present the frequencies and of CD4<sup>+</sup> or CD8<sup>+</sup> in the spleen  
 758 (A); memory effector phenotype CD44<sup>+</sup>CD62L<sup>+</sup> gated on splenic CD4<sup>+</sup> or CD8<sup>+</sup> T-cells (B), the  
 759 bar graphs summarize the frequencies of CD4<sup>+</sup> or CD8<sup>+</sup> (C) and memory/effector phenotype of  
 760 splenic lymphocytes (D). The result is representative of two independent experiments. Unpaired  
 761 two-tail t-test was used to compare between groups.

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764  
765 **Fig. S4. (A-E) Deficiency C3aR<sup>-1</sup>/C5aR<sup>-1</sup> in recipient increases mitophagy in host DCs and**  
766 **reduces GVHD development. (A-B)** Lethally irradiated BALB/c mice (*n*=6-7) were transplanted  
767 with BM alone (5×10<sup>6</sup>/mouse) or plus splenocytes (5×10<sup>6</sup>/mouse) from B10.D2 donor mice. The  
768 survival (A) and GVHD clinical score (B) were monitored throughout experiment (C) Irradiated  
769 BM derived-DCs were stimulated with 20 ng/mL LPS and are stained with Cyto-ID green. The  
770 summary bar for mean of fluorescence of cyto-ID expression in WT, C3aR<sup>-1</sup>/C5aR<sup>-1</sup> and HQ-  
771 treated C3aR<sup>-1</sup>/C5aR<sup>-1</sup> DCs are shown. (D) Lethally irradiated B6 recipients were transplanted  
772 with BM alone (3×10<sup>6</sup>/mouse) or plus purified T-cells (1×10<sup>6</sup>/mouse) from FVB donors.  
773 Recipients also received 2×10<sup>6</sup> WT or C3aR<sup>-1</sup>/C5aR<sup>-1</sup> B6 BM-DCs. The representative zebra  
774 color and the summary bar for percentage of co-transplanted DCs (CD45.1<sup>+</sup>CD11c<sup>+</sup>) in  
775 transplanted recipient peripheral blood are shown. (E) Lethally irradiated BALB/c recipients  
776 (*n*=5) were transplanted with BM alone (3×10<sup>6</sup>/mouse) or plus purified T-cells (1×10<sup>6</sup>/mouse)  
777 from B6 donors. Recipients also received 2×10<sup>6</sup> WT or C3aR<sup>-1</sup>/C5aR<sup>-1</sup> BALB/c BM-DCs. The  
778 survival (E) and clinical score (F) were monitored throughout the experiment. **Wilcoxon rank-**  
779 **sum test (A,E) and nonparametric Mann-Whitne U test (B,F) were used to compare between**  
780 **groups.** Data are representative of two independent experiments and presented as mean ±  
781 S.E.M., and \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001.



782

783 **Fig. S5. Generation of BM chimeras.** Lethally irradiated WT or C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> B6 mice (*n*=10)  
 784 were reconstituted with TCD-BM from WT or C3aR/C5aR donor. In order to distinguish donor  
 785 vs. host, B6 Ly5.1 congenic mice were used either as donor or recipients when possible. After  
 786 three months, the peripheral blood was collected for the analysis of donor cell reconstitution  
 787 reflecting by expression of Ly5.1, CD4 and CD8. The representative flow cytometric zebra blots  
 788 are shown.

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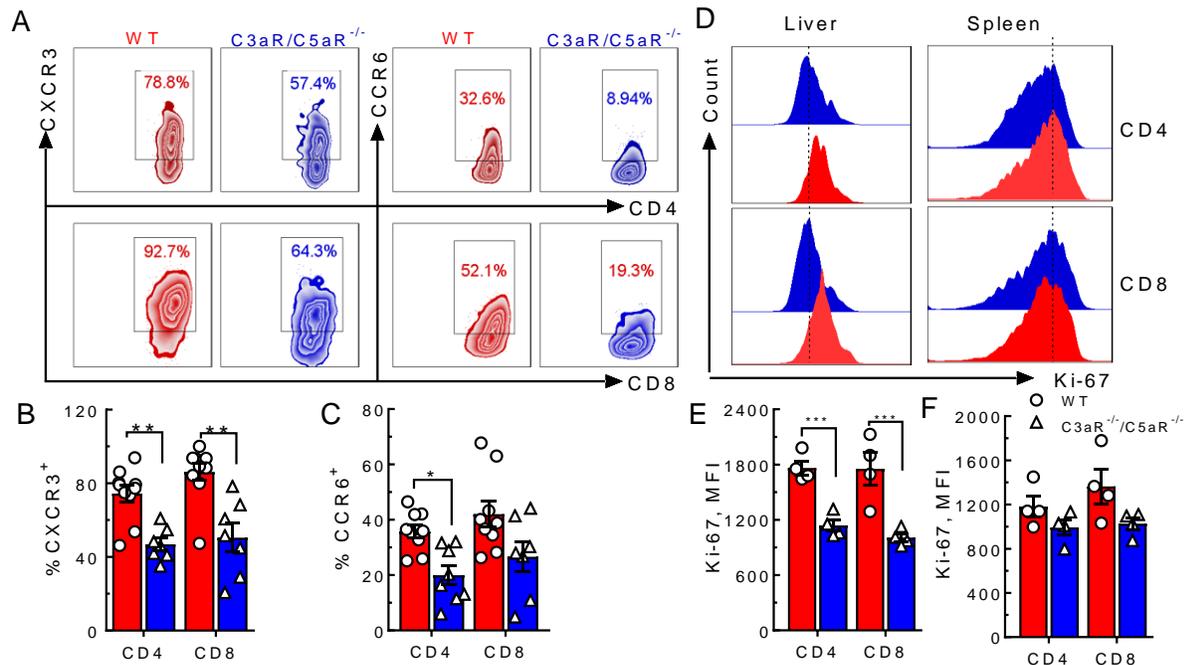
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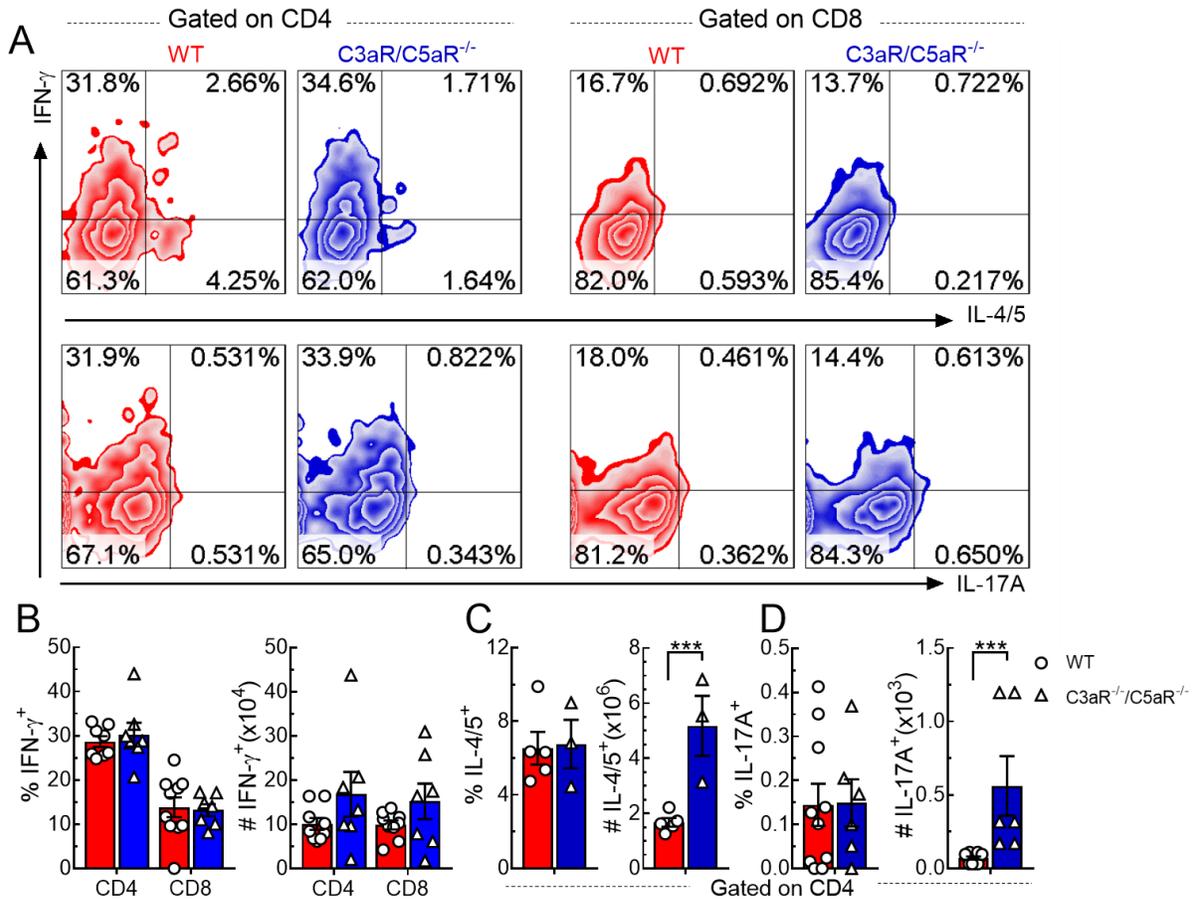
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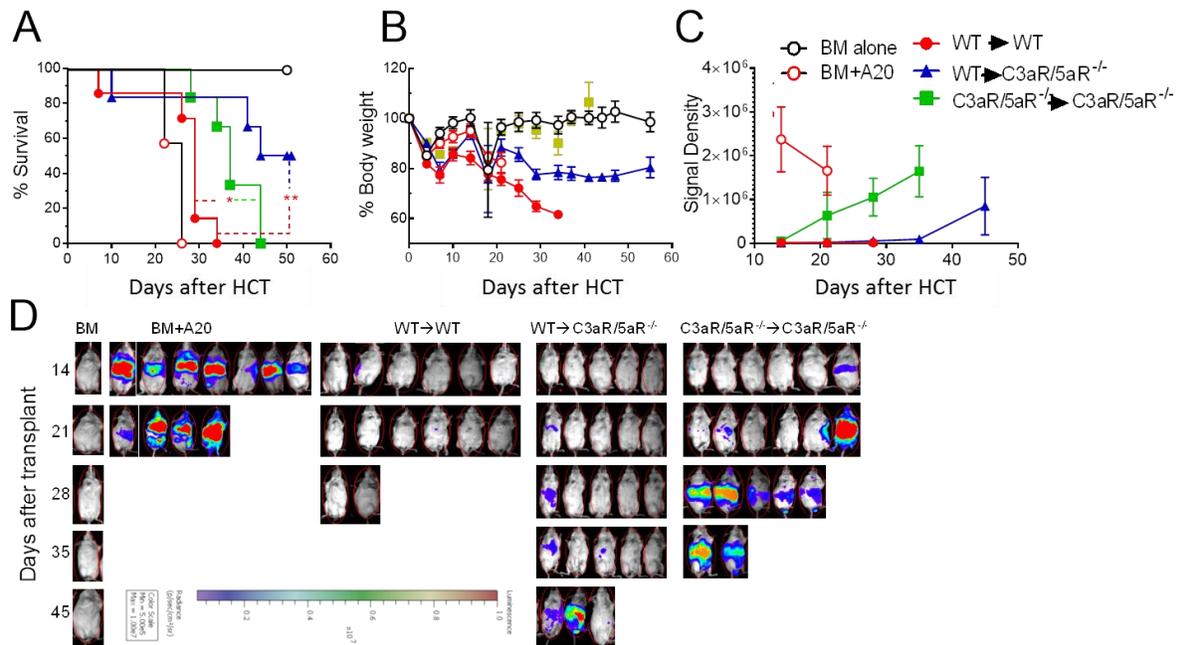
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798 **Fig. S6. Host C3aR/C5aR deficiency decreases the migration of donor T-cells toward**  
 799 **target organs.** HCT was performed as in Fig. 4. Recipient spleen and liver cells were isolated  
 800 14 post-transplant and stained for CD4, CD8, H2<sup>b</sup>, CXCR3, and CCR6. Representative zebra  
 801 plots of CXCR3<sup>+</sup> and CCR6<sup>+</sup> lymphocytes in the spleens (A). Graphs summarize frequencies of  
 802 CXCR3<sup>+</sup> (B)/CCR6<sup>+</sup> (C) donor lymphocytes ( $n=8$ ). Representative histogram and summary  
 803 graphs for MFI of ki-67 on H2<sup>b</sup>CD4<sup>+</sup>/CD8<sup>+</sup> in the liver (D, E) and spleen (D, F) ( $n=4$ ). The result  
 804 is combined of two independent experiments. Unpaired two-tail *t*-test was used to compare  
 805 between groups. Data are presented as mean  $\pm$  S.E.M. and \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .  
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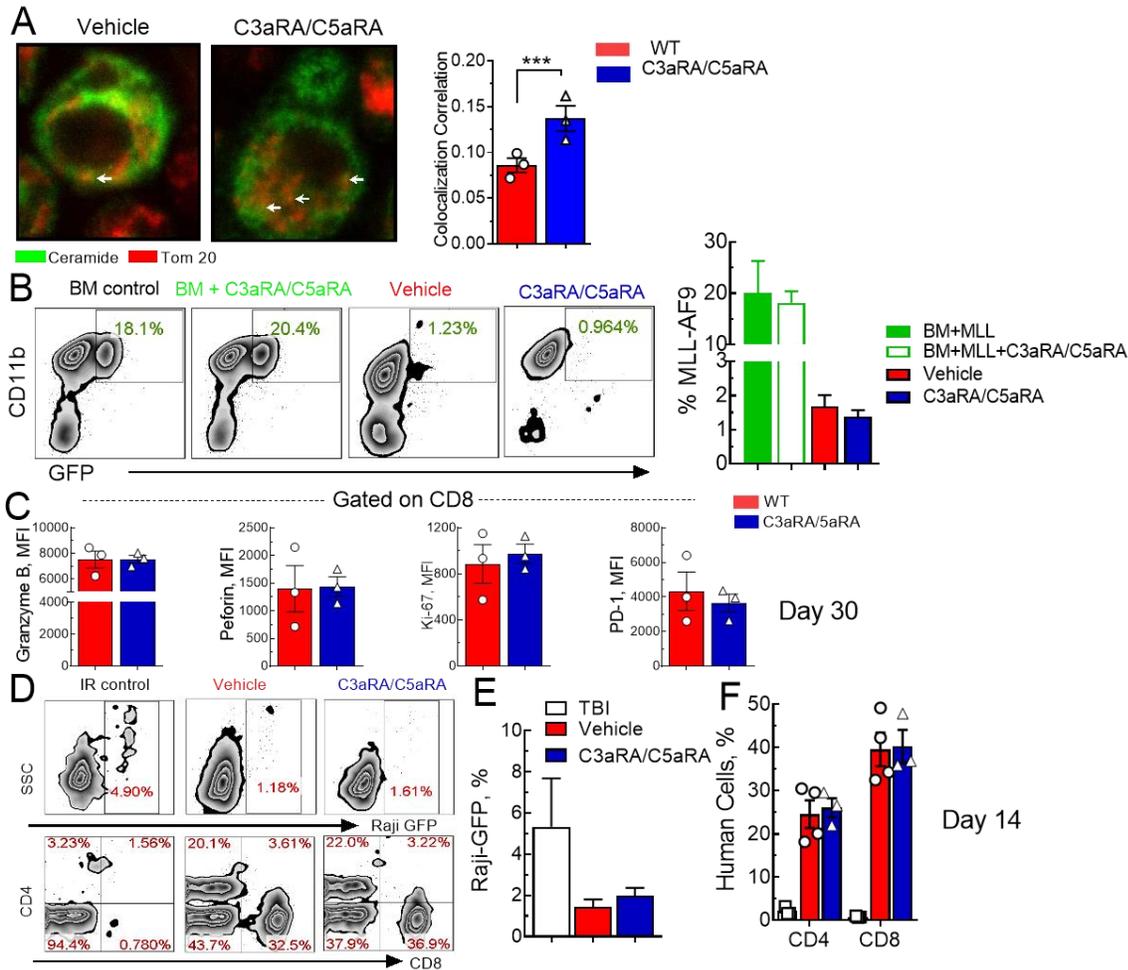
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808 **Fig. S7. Recipient C3aR/C5aR is required for Th1 differentiation while suppressing Treg**  
 809 **generation after HCT.** HCT was performed as described in Fig. 5 A and B. On day 14 after  
 810 transplantation, recipient spleen cells were stained with H2<sup>b</sup>, live/death, CD4, CD8 for surface  
 811 and intracellularly with IFN- $\gamma$ , IL-4/5, IL-17A, and Foxp3. The flow zebra plots of IFN- $\gamma$ <sup>+</sup>, IL-4/5<sup>+</sup>,  
 812 IL-17<sup>+</sup> donor cells from liver (A) are displayed. Bar graphs summarize the percentage and  
 813 absolute number of H2<sup>b</sup>IFN- $\gamma$ <sup>+</sup> (B) ( $n=8$ ), H2<sup>b</sup>IL-4/5<sup>+</sup> (C) ( $n=3-4$ ), and H2<sup>b</sup>IL-17A<sup>+</sup> (D) ( $n=8$ )  
 814 cells gated on CD4. Data were combined from two independent experiments. **Unpaired two-tail**  
 815 ***t*-test was used to compare between groups.** Results are presented as mean  $\pm$  S.E.M. and \* $p <$   
 816  $0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .



817

818 **Fig. S8. Recipient C3aR/C5aR is essential for optimal GVL activity.** Lethally WT or C3aR<sup>-/-</sup>  
 819 /C5aR<sup>-/-</sup> BALB/c recipients (n=7-10) were transplanted with TCD-BM (5×10<sup>6</sup>/mouse) with or  
 820 without purified T cells from WT or C3aR<sup>-/-</sup>/C5aR<sup>-/-</sup> B6 donors (1.0×10<sup>6</sup>/mouse). In addition,  
 821 recipients received 2×10<sup>3</sup> A20-luc at the time of transplant. Recipients were monitored for  
 822 survival (**A**), GVHD related body weight loss (**B**), and tumor growth determined by whole-body  
 823 BLI (**C, D**). The log-rank (Mantel-Cox) test (**A**) and nonparametric Mann-Whitney U test (**B, C**)  
 824 were used to compare between groups. Data are presented as mean SEM; \**p* < 0.05, \*\**p* < 0.01,  
 825 \*\*\**p* < 0.001.



826

827 **Fig.S9. C3aRA/C5aRA induced mitophagy in DCs.** (A) Irradiated BM derived-DCs were  
 828 cultured with or without C3aRA (2 mg/mL) and C5aRA (1mg/mL) and stimulated with 20 ng/mL  
 829 LPS following by dual staining with ceramide antibody and autophagosomal marker Tom 20 and  
 830 visualized using confocal microscopy. White arrows indicate colocalization. Data were combined  
 831 from two independent experiments ( $n=3$ ) and are presented as mean  $\pm$  S.D. and  $***p < 0.001$ .  
 832 (B-C) HCT ( $n=8$ ) was performed and C3RA/C5aRA was administered as described above. At  
 833 the time of HCT recipients also received MLL-AF9 GFP tumor cells. The tumor growths were  
 834 indicated by the frequencies of GFP<sup>+</sup>/CD11b<sup>+</sup> double positive cells. The representative flow  
 835 zebra plots of MLL-AF9 GFP tumor cells and the summary bar graphs for the frequency of MLL-  
 836 AF9 GFP are shown (B). On day 30 post-transplant, peripheral blood cells were stained for H<sup>2b</sup>,  
 837 CD4, CD8 and intracellular staining for granzyme B (grzm B), perforin, and CD107a. The bar  
 838 columns for the mean of fluorescence of grzm B, perforin, Ki-67 and PD-1 are shown (C) ( $n=3$ ).  
 839 (D-F) NSG-A2 mice ( $n=5-7$ ) were irradiated at the dose of 280 cGy, treated with a combination  
 840 of C3RA/C5aRA and transfused with HLA-A2 negative human PBMCs ( $15 \times 10^6$  i.v.). eGFP<sup>+</sup> Raji  
 841 cells ( $1 \times 10^6$ /mouse) were injected i.v. on day 0. Tumor burden was determined by measuring  
 842 GFP<sup>+</sup> cells in peripheral blood of transplant recipients. The engraftment was determined by  
 843 monitoring frequencies of human donor CD4<sup>+</sup> and CD8<sup>+</sup> cells in peripheral blood. The  
 844 representative flow zebra plots and summary bar graph of frequencies of MLL-AF9 GFP tumor  
 845 cells (D-E) and CD4<sup>+</sup> or CD8<sup>+</sup> (D-F) are shown ( $n=4-5$ ). Unpaired two-tail *t*-test was used to  
 846 compare between groups.  $***p < 0.001$ .