

ONLINE SUPPLEMENTARY MATERIAL

**Identification of a PD-L1<sup>+</sup> Tim-1<sup>+</sup> iNKT subset that protects against fine particulate matter-induced airway inflammation**

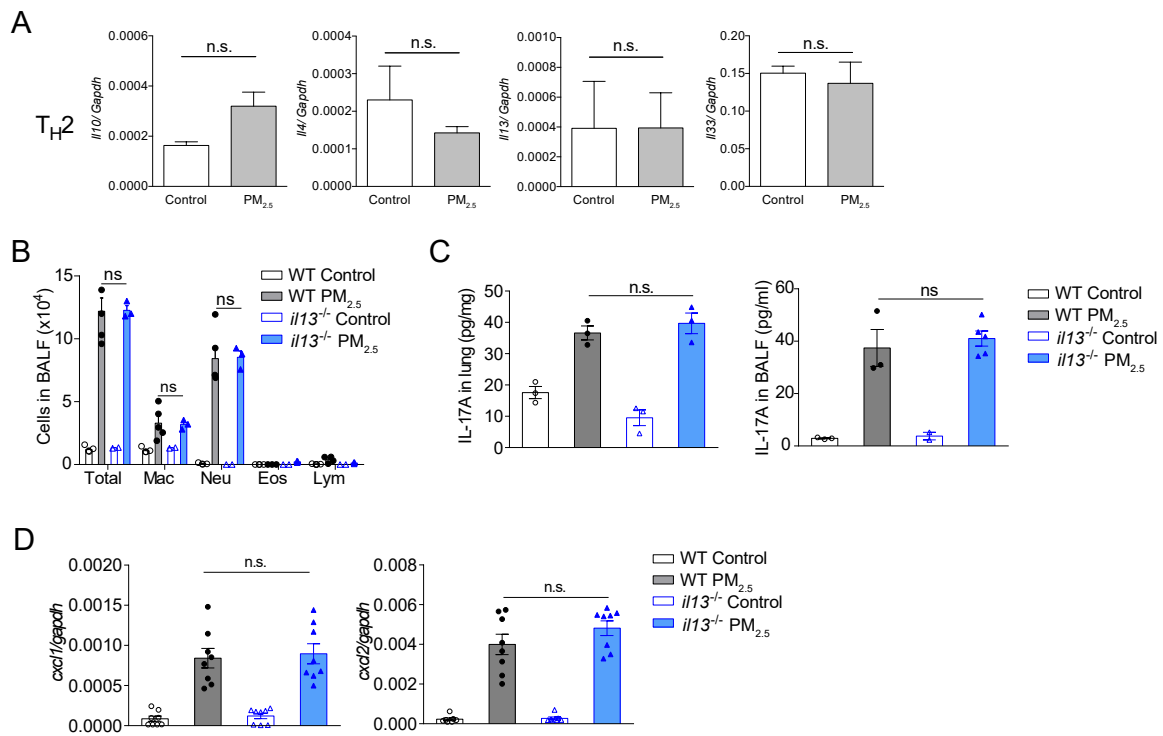
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**Affiliations:**

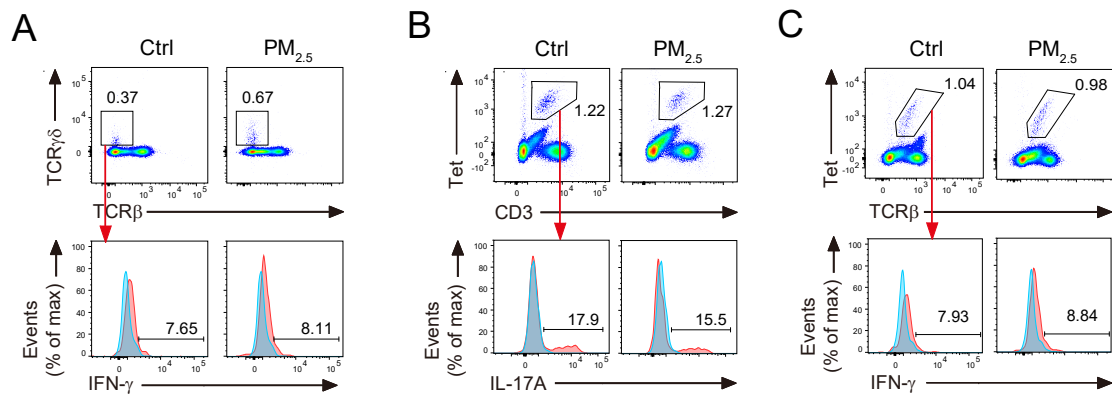
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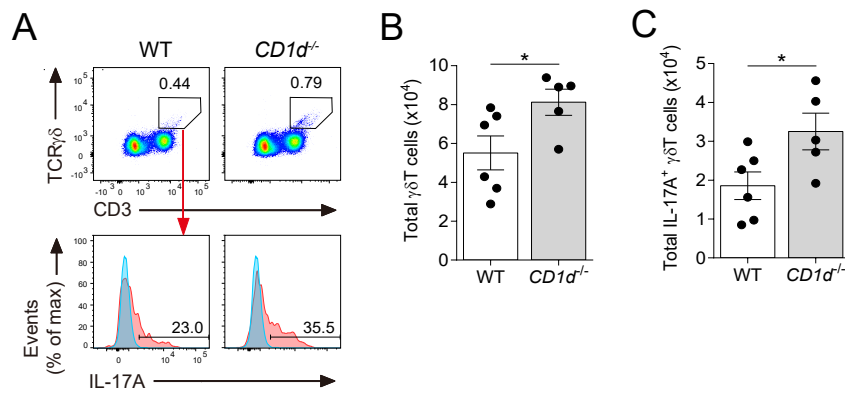
† These authors are co-first authors.



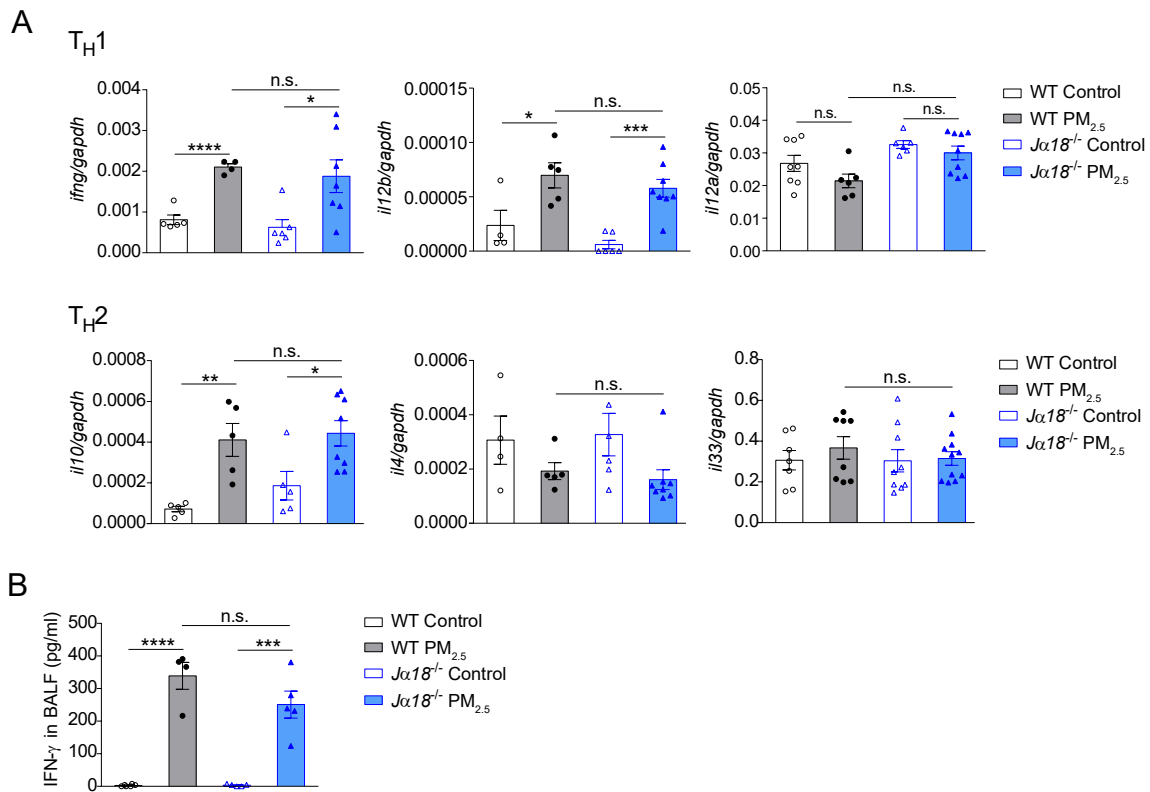
**Supplemental Figure 1. PM<sub>2.5</sub>-induced inflammation is independent of T<sub>H2</sub> response.** BALB/c (WT) (A-D) and *Il13*<sup>-/-</sup> (B-D) mice received daily intranasal exposure of PM<sub>2.5</sub> for three days and sacrificed one day after the last exposure. (A) Transcript expression levels of T<sub>H2</sub>-associated cytokines (*Il10*, *Il4*, *Il13*, and *Il33*) in the lungs. (B) Cellular composition (Mac, macrophage; Neu, neutrophil; Eos, eosinophil; Lym, lymphocyte) in BALF. (C) Levels of IL-17A in the lungs and BALF. (D) Transcript expression levels of *Cxcl1* and *Cxcl2* in the lungs. Data are expressed as mean ± SEM from 2 independent experiments (n=2-8 per group). Statistical analysis was performed using an unpaired two-tailed *t* test (A) or one-way ANOVA (B-D). n.s., Not significant.



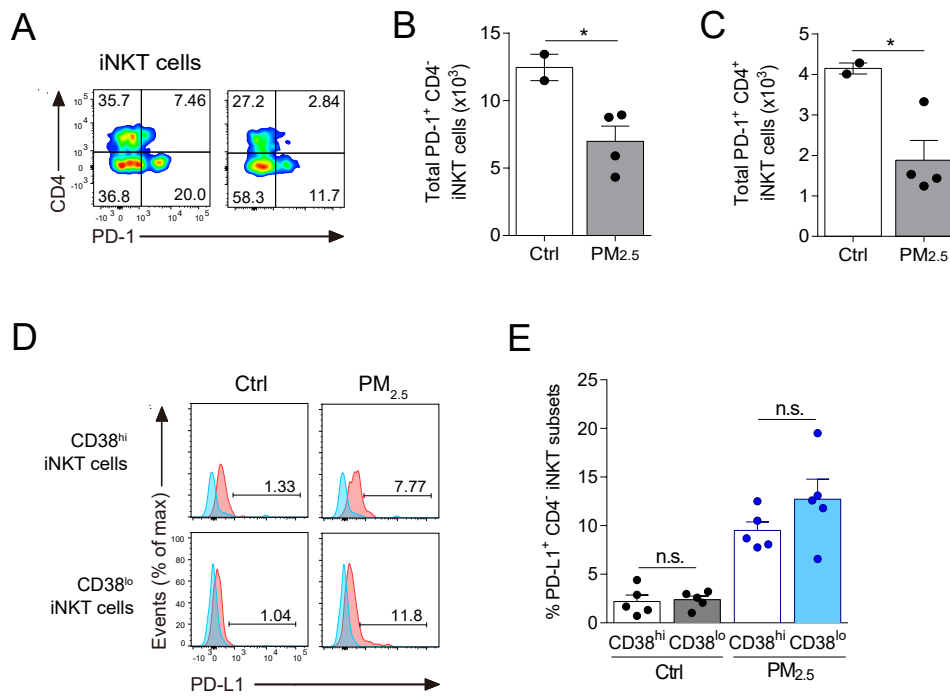
**Supplemental Figure 2. IL-17A and IFN- $\gamma$  production by  $\gamma\delta$  T and iNKT cells.** BALB/c received daily intranasal exposure of PM<sub>2.5</sub> for three days and sacrificed one day after the last exposure. **(A)** Representative flow cytometry plot showing IFN- $\gamma$  production in  $\gamma\delta$  T cells (TCR $\gamma\delta^+$  TCR $\beta^-$  cells). **(B-C)** Representative flow cytometry plot showing IL-17A **(B)** and IFN- $\gamma$  **(C)** production in iNKT cells (CD1d-tetramer $^+$  CD3 $^+$  TCR $\beta^+$  cells).



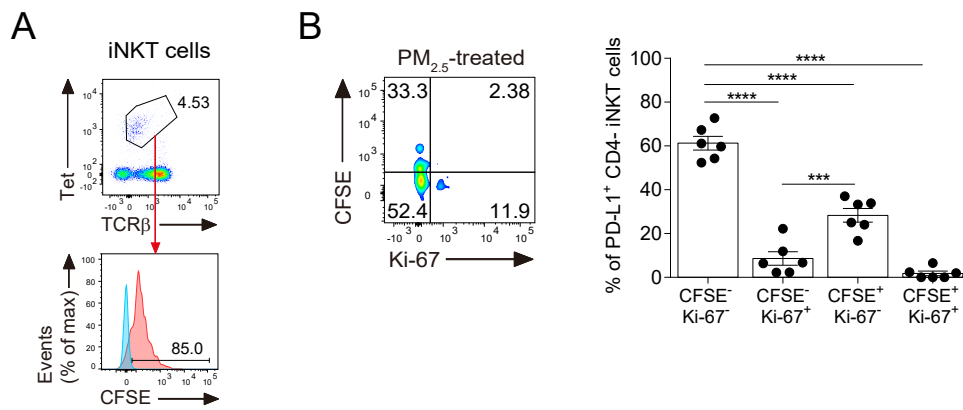
**Supplemental Figure 3. IL-17A production by  $\gamma\delta$  T cells in naïve WT and  $CD1d^{-/-}$  mice.** BALB/c (WT) and  $CD1d^{-/-}$  mice received daily intranasal exposure of  $PM_{2.5}$  for three days and sacrificed one day after the last exposure. **(A)** Representative flow cytometry plot showing IL-17A $^+$   $\gamma\delta$  T cells (CD3 $^+$  TCR $\gamma\delta^+$  cells). **(B)** Total  $\gamma\delta$  T cells and **(C)** IL-17A $^+$   $\gamma\delta$  T cells in naïve WT and  $CD1d^{-/-}$  mice, assessed as in A. Data are expressed as mean  $\pm$  SEM from 2 independent experiments (n=5-6 per group). Statistical analysis was performed using an unpaired two-tailed *t* test. \**P*<.05.



**Supplemental Figure 4. iNKT cell-deficiency does not alter T<sub>H</sub>1 and T<sub>H</sub>2 responses after PM<sub>2.5</sub> treatment.** BALB/c and *Ja18*<sup>-/-</sup> mice received daily intranasal exposure of PM<sub>2.5</sub> for three days and sacrificed one day after the last exposure. (A) Levels of T<sub>H</sub>1 (*Ifng*, *Il12a*, and *Il12b*) and T<sub>H</sub>2 (*Il10*, *Il4*, and *Il33*) mRNA in the lungs of mice. (B) Level of IFN- $\gamma$  in the BALF. Data are expressed as mean  $\pm$  SEM from 2 independent experiments (n=4-8 per group). Statistical analysis was performed using one-way ANOVA. \* $P$ <.05, \*\* $P$ <.01, \*\*\* $P$ <.001, and \*\*\*\*  $P$ <.0001. n.s., Not significant.



**Supplemental Figure 5. PD-1 and PD-L1 expression on iNKT cells.** BALB/c received daily intranasal exposure of PM<sub>2.5</sub> for three days and sacrificed one day after the last exposure. **(A)** Representative flow cytometry plot showing PD-1 expression on the CD4<sup>+</sup> and CD4<sup>-</sup> iNKT cell subsets (Gated from CD1d-tetramer<sup>+</sup> TCRβ<sup>+</sup> cells). **(B)** Total PD-1<sup>+</sup> CD4<sup>-</sup> iNKT cells and **(C)** PD-1<sup>+</sup> CD4<sup>+</sup> iNKT cells, assessed as in **A**. **(D)** Representative histogram showing PD-L1 expression on CD38<sup>hi</sup> and CD38<sup>lo</sup> iNKT cell subsets (Gated from CD1d-tetramer<sup>+</sup> TCRβ<sup>+</sup> CD4<sup>-</sup> cells). Blue solid line: Isotype-matched control; Red solid line: Antibody staining. **(E)** Frequencies of PD-L1-expressing CD38<sup>hi</sup> and CD38<sup>lo</sup> CD4<sup>-</sup> iNKT subsets, assessed as in **D**. Data are representative of one experiment (n=2-5 per group). Statistical analysis was performed using an unpaired two-tailed *t* test (**B**, **C**) or one-way ANOVA (**E**). \**P*<.05. n.s., Not significant.



**Supplemental Figure 6. Expression of CFSE and Ki-67 in PD-L1<sup>+</sup> CD4<sup>+</sup> iNKT cells.**

BALB/c mice received daily intranasal exposure of PM<sub>2.5</sub> for three days and sacrificed one day after the last exposure. CFSE (2 μg/g mouse) was administered intravenously after the second exposure to PM<sub>2.5</sub>. (A) Representative flow cytometry plot showing CFSE<sup>+</sup> iNKT cells in the blood 48 hours after CFSE labeling. (B) Representative flow cytometry plot and quantitative data showing CFSE<sup>-</sup> and Ki-67-expressing PD-L1<sup>+</sup> CD4<sup>+</sup> iNKT cell subset, gated as in **Figure 6E**. Data are shown as mean ± SEM from 2 independent experiments (n=6 per group). Statistical analysis was performed using one-way ANOVA (B). \*\*\* *P*<.001 and \*\*\*\* *P*<.0001.

## Supplemental tables

**Supplemental Table 1: List of antibodies used for CyTOF**

Antigen panel	Symbol	Mass	Antibody clone	Brand
CD90	In	113	30-H12	Biologend
TCR $\beta$	In	115	H57-597	Biologend
CD44	Cd	116	IM7	BD Biosciences
CD11b	Ce	140	M1/70	Biologend
CD69	Pr	141	H1.2F3	Biologend
CD45	Nd	142	30-F11	Biologend
CD11c	Nd	143	HL3	BD Biosciences
Gr1	Nd	144	RB6-8C5	Biologend
CD4	Nd	145	RM4-5	Fluidigm
CD3	Sm	147	17A2	Biologend
CD64	Nd	148	X54-5/7.1	Biologend
CD19	Sm	149	6D5	Fluidigm
CD27	Nd	150	LG.3A10	Fluidigm
Ly6C	Eu	151	HK1.4	Biologend
Ki-67	Sm	152	SolA15	eBioscience
CD8a	Gd	155	53-6.7	Biologend
CD1d tetramer	Gd	156		NIH
Foxp3	Gd	158	FJK-16S	eBioscience
PD-1	Tb	159	29F.1A12	Biologend
GATA3	Gd	160	TWAJ	eBioscience
Tbet	Dy	161	O4-46	Fluidigm
TCR $\gamma\delta$	Dy	162	GL3	Biologend
CD62L	Dy	164	MEL-14	Fluidigm
NK1.1	Ho	165	PK136	Fluidigm
cKit	Er	166	2B8	Biologend
NKp46	Er	167	29A1.4	Biologend
ROR $\gamma$ t	Er	168	600214	Fluidigm
F4/80	Tm	169	BM8	Biologend
CD137 (41BB)	Er	170	17B5	Biologend
CD86	Yb	172	GL-1	Biologend
FceRI	Yb	173	MAR-I	Biologend
mSiglecF	Yb	174	E50-2440	BD Biosciences
CD127	Lu	175	A7R34	Fluidigm
ST2	Yb	176	DIH9	Biologend
MHCII	Bi	209	M5/114.15.2	Fluidigm



DNA	Ir	191/193		
Cisplatin Viability	Pt	195		

**Supplemental Table 2: Gating strategy for identifying CD45<sup>+</sup> CD90<sup>+</sup> lymphocyte subsets by CyTOF analysis**

<b>Immune cell subsets</b>	<b>Gating strategy</b>
CD4 T cells	CD3 <sup>+</sup> TCRβ <sup>+</sup> CD4 <sup>+</sup> CD8 <sup>-</sup>
CD8 T cells	CD3 <sup>+</sup> TCRβ <sup>+</sup> CD8 <sup>+</sup> CD4 <sup>-</sup>
CD44 <sup>+</sup> γδ T cells	CD3 <sup>+</sup> TCRβ <sup>-</sup> TCRγδ <sup>+</sup> CD44 <sup>+</sup>
CD27 <sup>+</sup> γδ T cells	CD3 <sup>+</sup> TCRβ <sup>-</sup> TCRγδ <sup>+</sup> CD27 <sup>+</sup>
NKT cells	CD3 <sup>+</sup> TCRβ <sup>+</sup> CD1d-tetramer <sup>+</sup>
Treg	CD3 <sup>+</sup> TCRβ <sup>+</sup> Foxp3 <sup>+</sup>
NK cells	CD3 <sup>-</sup> NK1.1 <sup>+</sup> NKp46 <sup>+</sup> Tbet <sup>+</sup>
ILC2	CD3 <sup>-</sup> TCRβ <sup>-</sup> CD127 <sup>+</sup> RORγt <sup>+</sup> Tbet <sup>-</sup> GATA3 <sup>+</sup>
ILC3	CD3 <sup>-</sup> TCRβ <sup>-</sup> CD127 <sup>+</sup> GATA3 <sup>-</sup> Tbet <sup>-</sup> RORγt <sup>+</sup>

\*NKT cells: Natural killer T cells; Treg: Regulatory T cells; NK cells: Natural killer cells; ILC2: Group 2 innate lymphoid cells; ILC3: Group 3 innate lymphoid cells

**Supplemental Table 3: Lists of primers used**

<b>Gene</b>	<b>Forward</b>	<b>Reverse</b>
<i>Il17a</i>	TCCAGAAGGCCCTCAGACTA	ACACCCACCAGCATCTTCTC
<i>Ifng</i>	GGCCA TCAGC AACAA CATAA GCGT	TGGGT TGTTG ACCTC AAAC TGGC
<i>Il12a</i>	CTGTGCCTTGGTAGCATCTATG	GCAGAGTCTCGCCATTATGATTC
<i>Il12b</i>	GTCCTCAGAAGCTAACCATCTCC	CCAGAGCCTATGACTCCATGTC
<i>Il33</i>	ATTTC CCCGG CAAAG TTCAG	AACGG AGTCT CATGC AGTAG A
<i>Il10</i>	GGTTGCCAAGCCTTATCGGA	ACCTGCTCCACTGCCTTGCT
<i>Il4</i>	GGTCTCAACCCCCAGCTAGT	GCCGATGATCTCTCTCAAGTGAT
<i>Il13</i>	CCTGGCTCTTGCTTGCCTT	GGTCTTGTGTGATGTTGCTCA
<i>Il1b</i>	GAAAT GCCAC CTTTT GACAG TG	CTGGA TGCTC TCATC AGGAC A
<i>Il23a</i>	ATGCT GGATT GCAGA GCAGT A	ACGGG GCACA TTATT TTTAG TCT
<i>Cxcl1</i>	AAAAGGTGTCCCAAGTA	AAGCAGAACTGAACTACCATCG
<i>Cxcl2</i>	GGGAGAGGGTGAGTTGGG	GCACACTCCTTCATGAAAGC