

Supplementary Information

FFAR2/3 dysfunction aggravates IL-33-induced eosinophilic airway inflammation via ST2⁺CD4⁺ T cell activation

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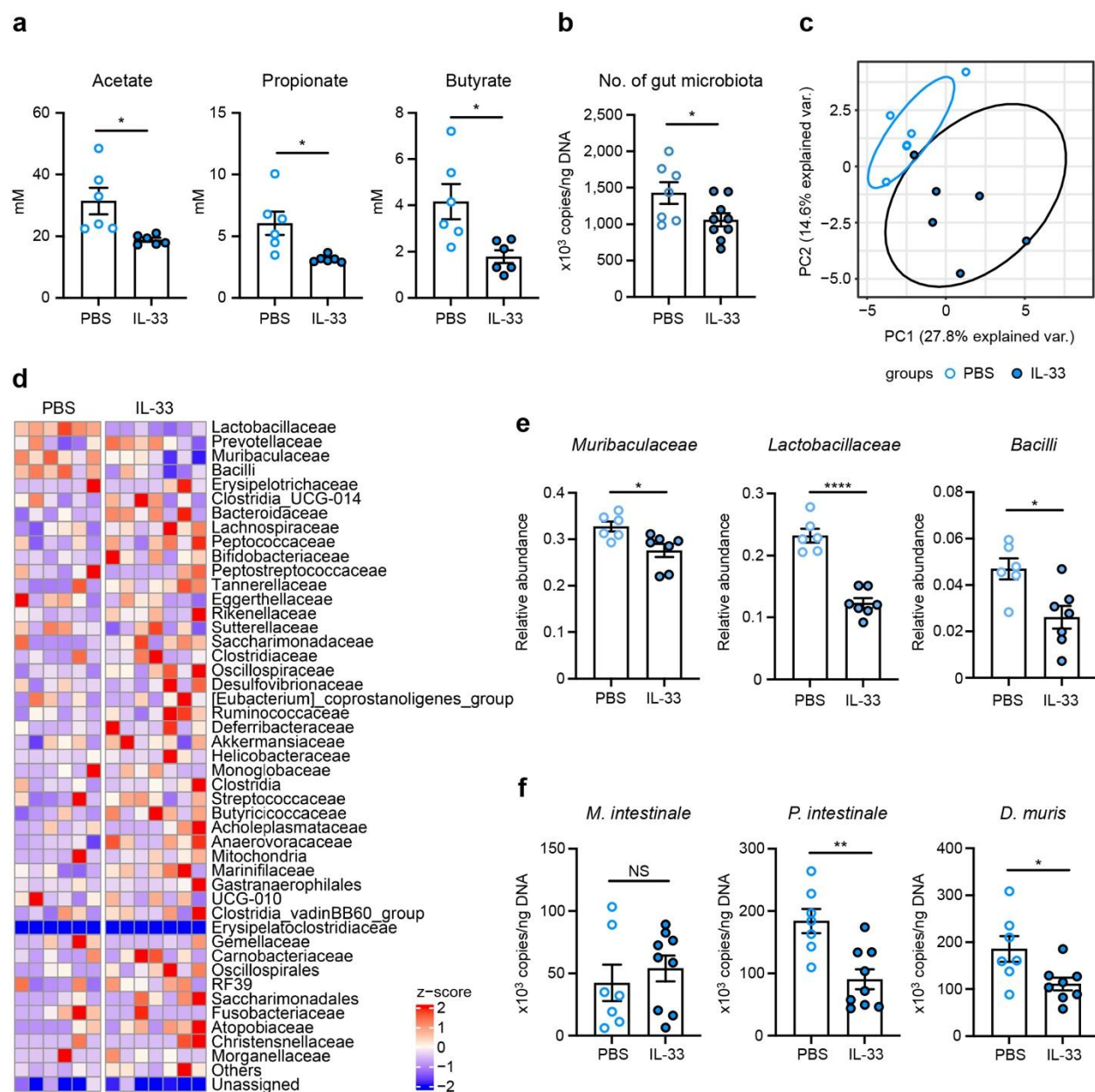
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This supplementary file includes:

Supplementary Figures 1 to 8

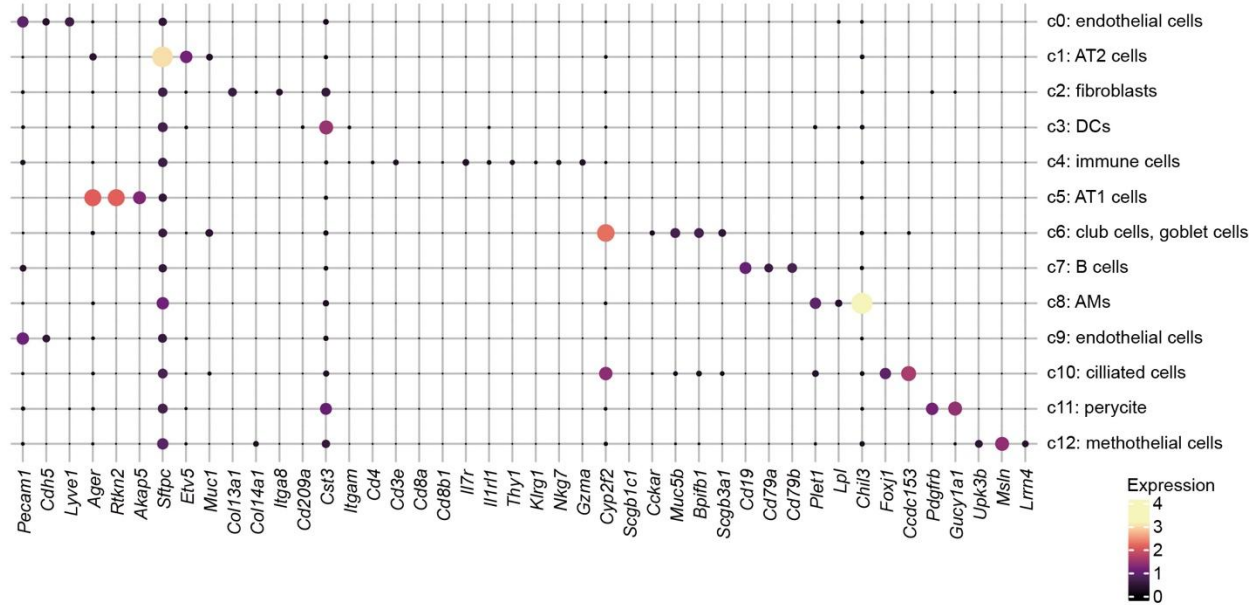
Supplementary Tables 1 to 3



Supplementary Fig. 1. Fecal short-chain fatty acid concentration and gut microbiome analysis of IL-33-induced eosinophilic airway inflammation model. **a** Concentration of short-chain fatty acids in feces. **b** Total bacterial load. **c** Principal coordinates analysis. **d** Heatmap showing the gut microbial composition at the family level. **e** The abundance of *Muribaculaceae*, *Lactobacillaceae*, and *Bacilli* in the fecal samples. **f** The abundances of *Muribaculaceae* species.

Mean $\pm$ s.e.m., n = 6–9 mice per group. * P < 0.05, ** P < 0.01, **** P < 0.0001 (Student's t -test).

a

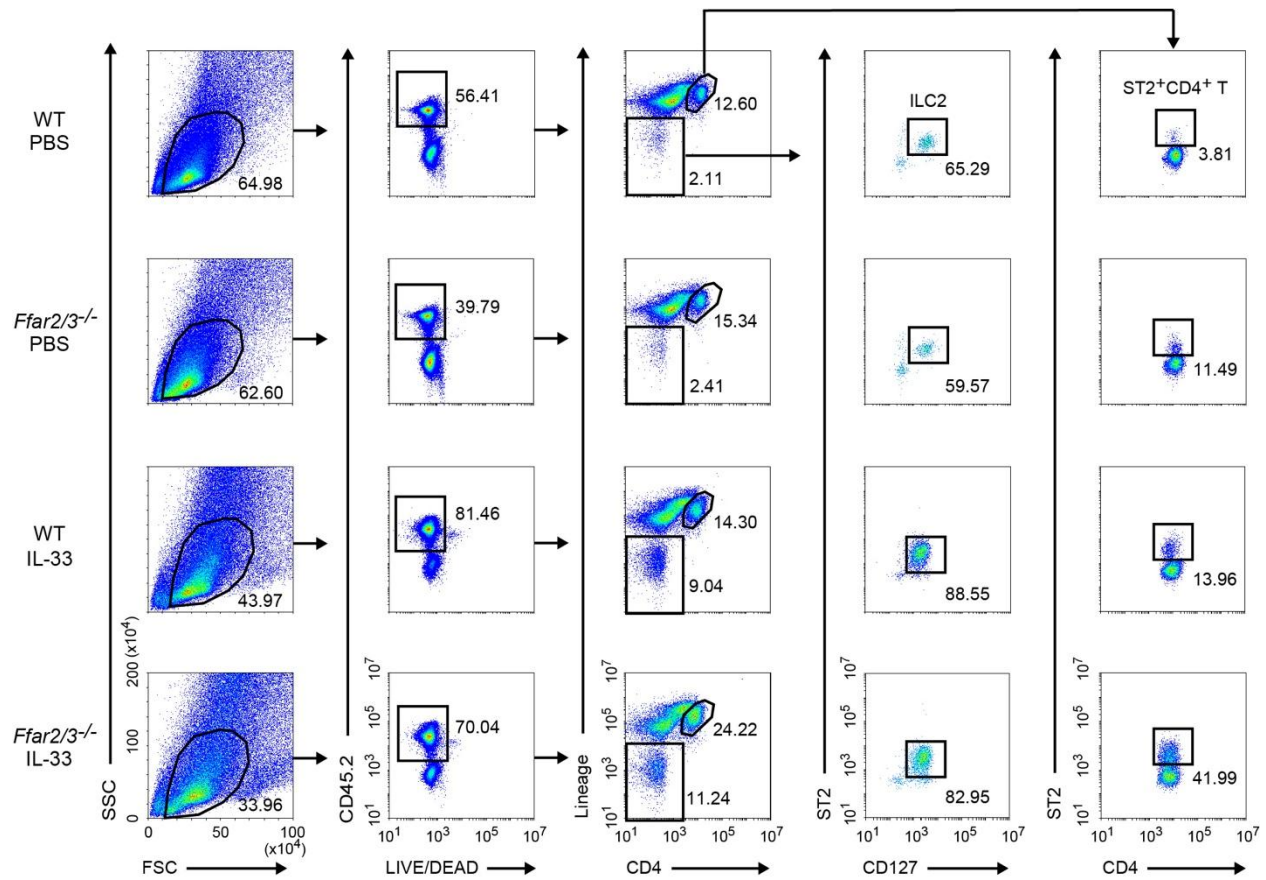


b

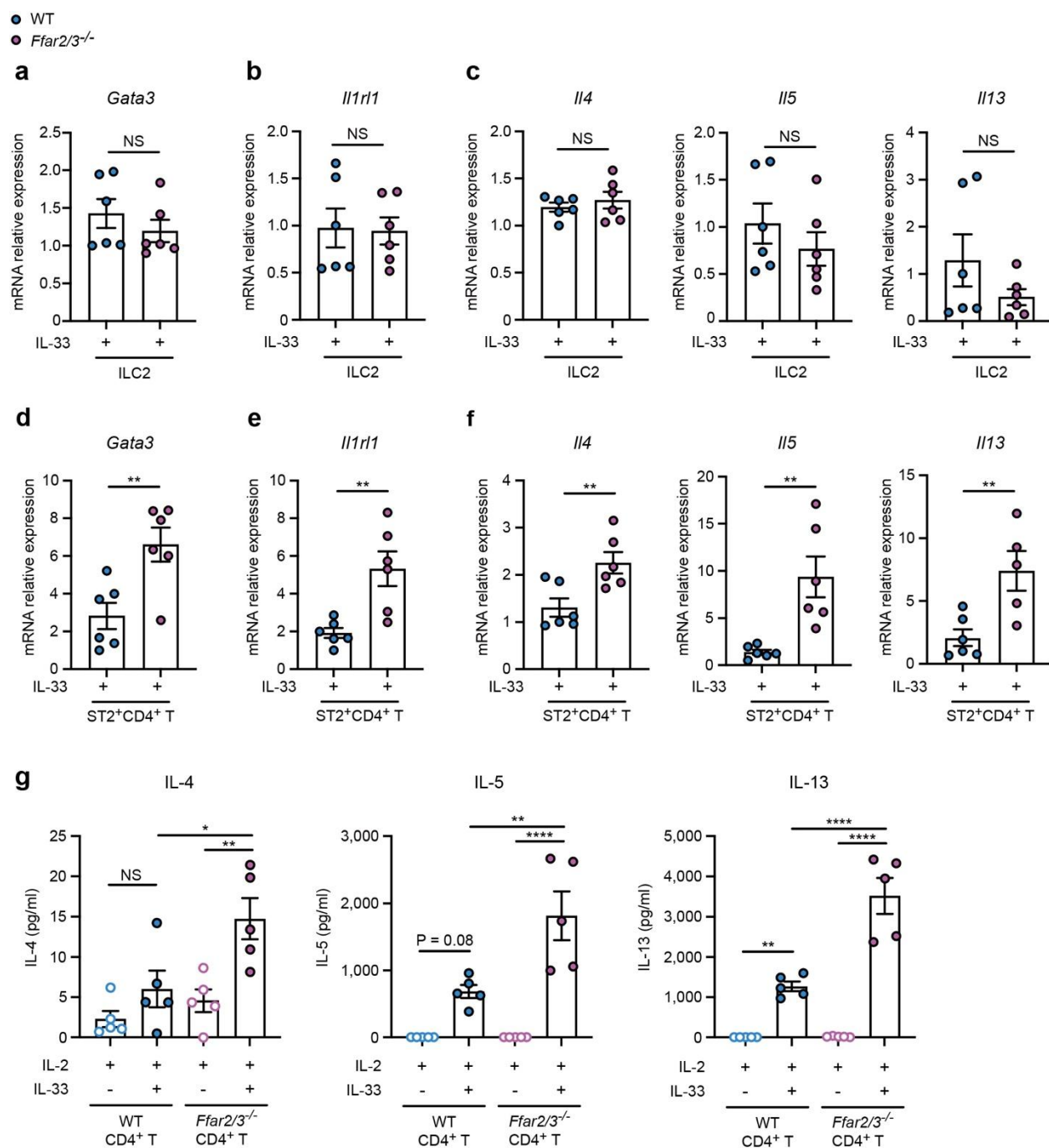
	Subcluster 0	Subcluster 1	Subcluster 2	Subcluster 3	Subcluster 4	Subcluster 5	Subcluster 6	Subcluster 7	Subcluster 8
	Naive T cells	ILC2s	CD4 ⁺ T cells	NK cells	B cells	Proliferated cells	$\gamma\delta$ T cells	Endothelial cells	Plasma cells
marker1	<i>Dgka</i>	<i>Stab2</i>	<i>Cxcr3</i>	<i>Gzma</i>	<i>Ncf1</i>	<i>Ankle1</i>	<i>Rorc</i>	<i>Arhgef15</i>	<i>Jchain</i>
marker2	<i>Gramd3</i>	<i>Il17rb</i>	<i>Ctla4</i>	<i>Adamts14</i>	<i>Siglecg</i>	<i>Rrm2</i>	<i>Blk</i>	<i>Myzap</i>	<i>Eaf2</i>
marker3	<i>Sell</i>	<i>Il1r1</i>	<i>Zfp683</i>	<i>Klre1</i>	<i>Ms4a1</i>	<i>Top2a</i>	<i>Il23r</i>	<i>Cyyr1</i>	<i>Derl3</i>
marker4	<i>Lef1</i>	<i>Arg1</i>	<i>Zfp36l2</i>	<i>Ccl5</i>	<i>Pirb</i>	<i>Ncapg</i>	<i>Cd163l1</i>	<i>Shank3</i>	<i>Cacna1h</i>
marker5	<i>Satb1</i>	<i>Hlf</i>	<i>Slamf1</i>	<i>Nkg7</i>	<i>Cd19</i>	<i>Klf1</i>	<i>Il1r1</i>	<i>Klf4</i>	<i>Igkc</i>
marker6	<i>Actn1</i>	<i>Grin2d</i>	<i>Cd4</i>	<i>Tyrbp</i>	<i>Tbxas1</i>	<i>Pimreg</i>	<i>Lingo4</i>	<i>Palmd</i>	<i>Pou2af1</i>
marker7	<i>Klf2</i>	<i>Slc7a8</i>	<i>Tmem154</i>	<i>Klrl1</i>	<i>Cd74</i>	<i>Ska1</i>	<i>Zbtb16</i>	<i>Sema3g</i>	<i>Irf4</i>
marker8	<i>Ccr7</i>	<i>Gata3</i>	<i>P2rx7</i>	<i>Ncr1</i>	<i>Cd79b</i>	<i>Ncapd2</i>	<i>Kcnk1</i>	<i>Smad6</i>	<i>Mzb1</i>
marker9	<i>Plk3ip1</i>	<i>Adam8</i>	<i>Atp2b4</i>	<i>Klrd1</i>	<i>Ebf1</i>	<i>Hist1h1b</i>	<i>Il17re</i>	<i>Efnb2</i>	<i>Igfc2</i>
marker10	<i>Evl</i>	<i>Znrf1</i>	<i>Trac</i>	<i>Cd244a</i>	<i>Unc93b1</i>	<i>Mxd3</i>	<i>Kcnc1</i>	<i>Ushbp1</i>	<i>Igha</i>
marker11	<i>Als2cl</i>	<i>Pcsk1</i>	<i>Maf</i>	<i>Klra8</i>	<i>Rassf4</i>	<i>Spag5</i>	<i>Adgrg5</i>	<i>Ecscr</i>	<i>Dennd5b</i>
marker12	<i>Tdrp</i>	<i>Rasgrp4</i>	<i>Lck</i>	<i>Aoah</i>	<i>Ppfbp2</i>	<i>Anln</i>	<i>Igf1r</i>	<i>Meis2</i>	<i>Igfc1</i>
marker13	<i>Nsg2</i>	<i>Aff4</i>	<i>Hnmp1l</i>	<i>Prf1</i>	<i>H2-Eb1</i>	<i>Tpx2</i>	<i>Smox</i>	<i>Cldn5</i>	<i>Txndc5</i>
marker14	<i>Ikbke</i>	<i>Syne1</i>	<i>Fgl2</i>	<i>Samd3</i>	<i>H2-Ab1</i>	<i>Ncapg2</i>	<i>Lmo4</i>	<i>Ccm2l</i>	<i>Rhobtb1</i>
marker15	<i>Slamf6</i>	<i>Ern1</i>	<i>Sntb1</i>	<i>Gsap</i>	<i>Sftpd</i>	<i>Ndc80</i>	<i>Il18r1</i>	<i>Cd36</i>	<i>Tnfrsf13b</i>
marker16	<i>Gimap4</i>	<i>Ccr8</i>	<i>Foxp3</i>	<i>Zeb2</i>	<i>Snhg11</i>	<i>Clsn</i>	<i>8-Sep</i>	<i>Ace</i>	<i>Tnfrsf17</i>
marker17	<i>Atp1b3</i>	<i>Il5</i>	<i>Cd3e</i>	<i>Itgam</i>	<i>Ptprj</i>	<i>Hmmr</i>	<i>Trdc</i>	<i>Sema3c</i>	<i>Prg2</i>
marker18	<i>Tcf7</i>	<i>Ccr2</i>	<i>Crybg2</i>	<i>Gzmb</i>	<i>Rilp</i>	<i>Dlgap5</i>	<i>Sox13</i>	<i>Egfl7</i>	<i>Txndc11</i>
marker19	<i>Usp28</i>	<i>Itm2b</i>	<i>Il21r</i>	<i>Klrb1f</i>	<i>Map3k9</i>	<i>Klf23</i>	<i>Capg</i>	<i>Eng</i>	<i>Ckap4</i>
marker20	<i>Rlnb</i>	<i>Furin</i>	<i>Il12rb1</i>	<i>Fcer1g</i>	<i>Il33</i>	<i>Mki67</i>	<i>Mmp25</i>	<i>Tie1</i>	<i>Ell2</i>

Supplementary Fig. 2. Single-cell RNA-seq analysis of the lung tissue from IL-33-treated wild-type and *Ffar2/Ffar3*-deficient mice. a Cell-type annotation of each cluster in single-cell

RNA-seq analysis based on gene expression pattern. **b** The list of top 20 genes in each subcluster population within Cluster 4 (immune cells).

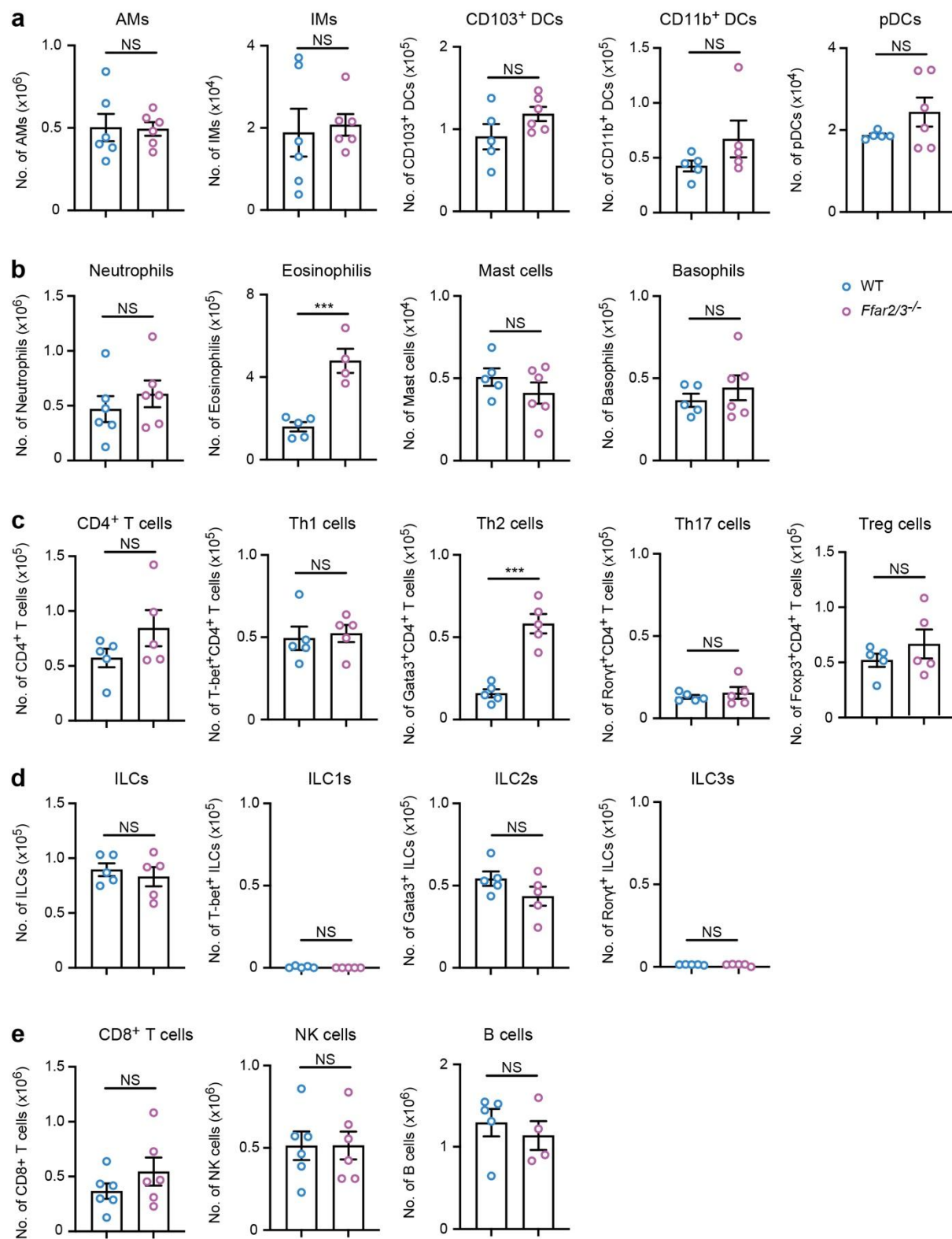


Supplementary Fig. 3. Gating strategy for flow cytometry. Flow cytometry analysis of lung ILC2s and ST2⁺CD4⁺ T cells in wild-type and *Ffar2/Ffar3*-deficient mice treated with PBS- or IL-33-treated condition.

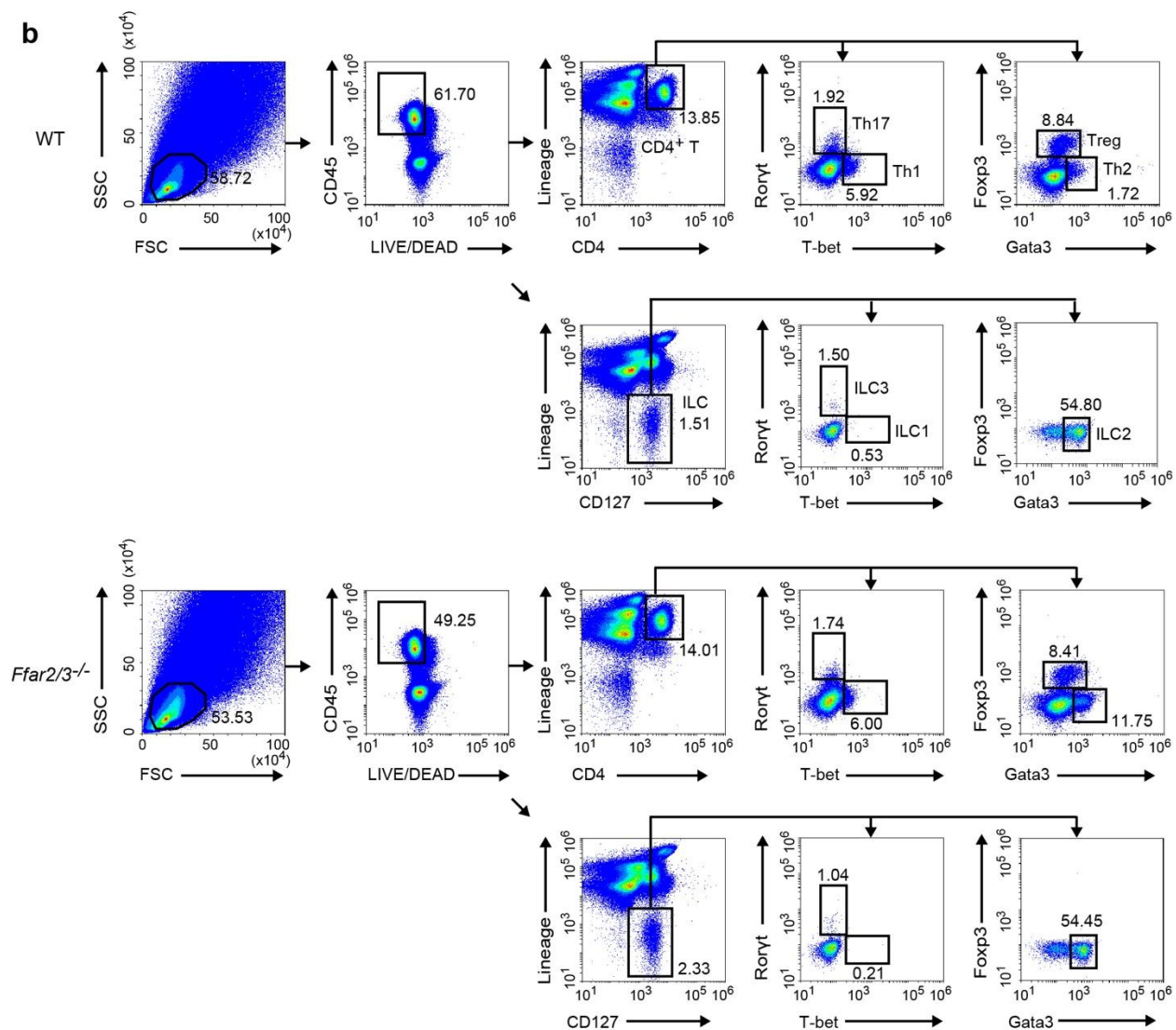
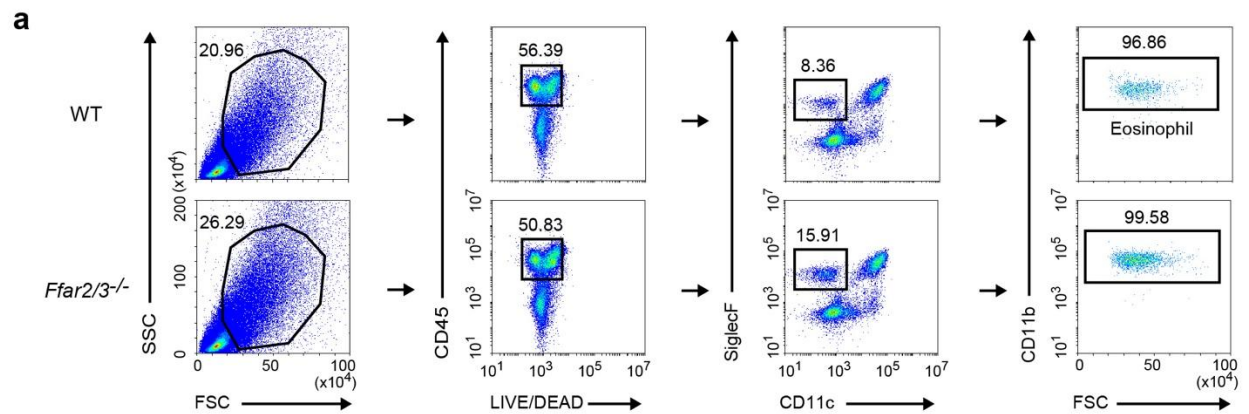


Supplementary Fig. 4. Comparison of gene expression in ILC2s and ST2⁺CD4⁺ T cells isolated from lung tissue of IL-33-treated mice and cytokine production in CD4⁺ T cells derived from spleen under IL-33 stimulation. The mRNA expression levels of *Gata3* (a, d), *Il1rl1* (b, e), type 2 cytokines (*Il4*, *Il5*, and *Il13*) (c, f) in ILC2 (a–c) and ST2⁺CD4⁺ T cells (d–f)

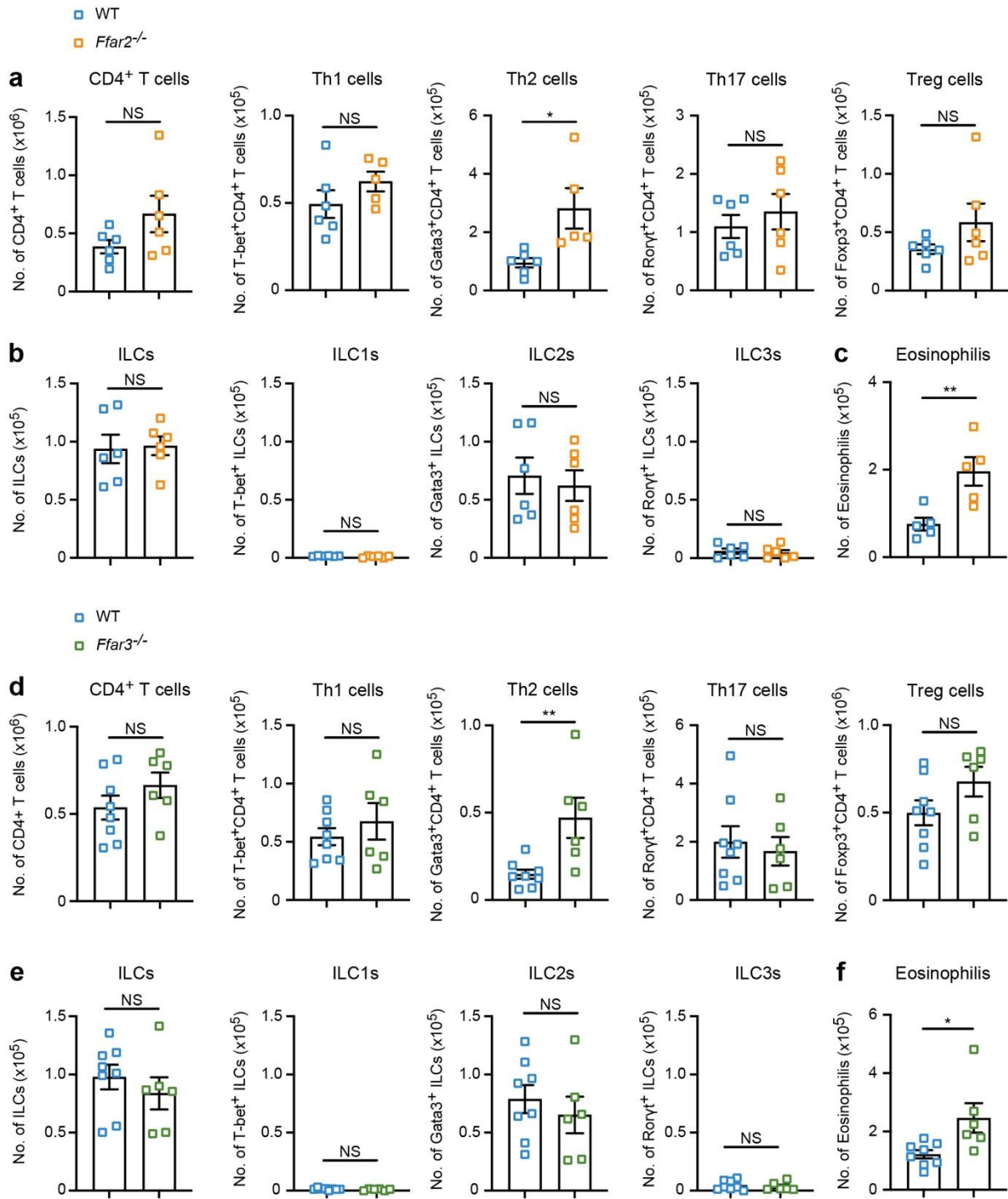
isolated from the lung tissue of wild-type and *Ffar2/Ffar3*-deficient mice in the IL-33-induced eosinophilic airway inflammation model. **g** CD4⁺ T cells (2.0×10^5 cells/well) isolated from the spleens of wild-type and *Ffar2/Ffar3*-deficient mice were stimulated with IL-2 (10 ng/ml) and IL-33 (10 ng/ml) for 72 h. IL-4, IL-5, and IL-13 concentrations in the culture supernatants. Mean $\pm$ s.e.m., n = 5-6 mice per group. **P* < 0.05, ***P* < 0.01, *****P* < 0.0001 (Student's *t*-test [**a–f**], one-way analysis of variance (ANOVA) and Tukey's post hoc test [**g**]). NS: not significant.



Supplementary Fig. 5. Comprehensive analysis of lung immune cell populations in a healthy state. Numbers of macrophage and dendritic cell subsets **(a)**, granulocyte subsets **(b)**, CD4⁺ T cell subsets **(c)**, ILC subsets **(d)**, and other lymphoid cells **(e)** isolated from the lung tissue of wild-type and *Ffar2/Ffar3*-deficient mice. Mean $\pm$ s.e.m., n = 4–6 mice per group. *** P < 0.001, (Student's *t*-test). NS: not significant. AMs: alveolar macrophages. IMs: Interstitial macrophages. DCs; dendritic cells. NK cells, natural killer cells.

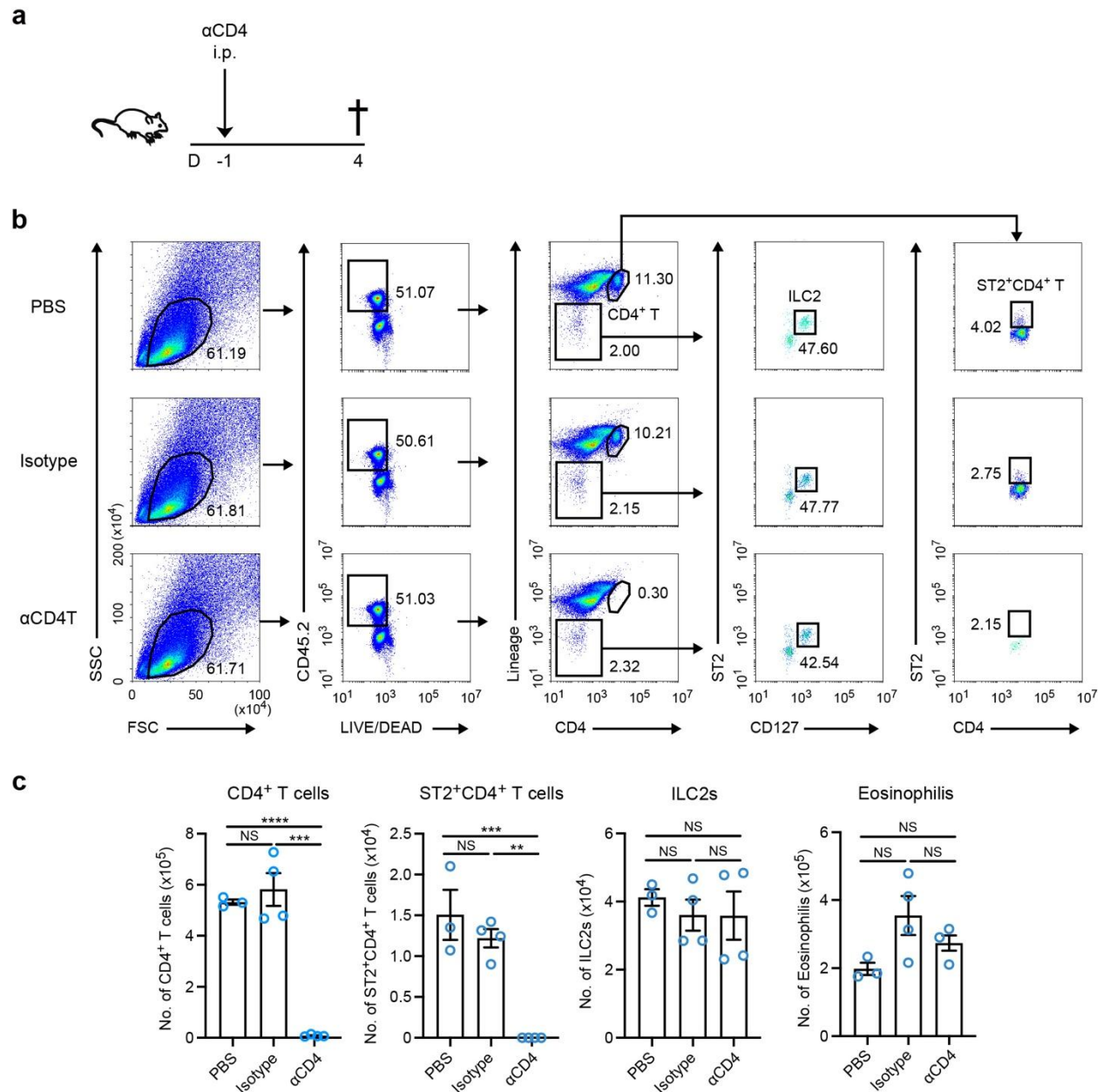


Supplementary Fig. 6. Gating strategy for flow cytometry. Flow cytometric analysis of lung eosinophils **(a)**, CD4⁺ T cell subsets, and ILC subsets **(b)** in wild-type and *Ffar2/Ffar3*-deficient mice in a healthy state.



Supplementary Fig. 7. Comparison of CD4⁺ T cell subset, ILC subsets, and eosinophils of lung tissue of *Ffar2*-deficient and *Ffar3*-deficient mice in a healthy state. CD4⁺ T cell subsets (a, d), ILC subsets (b, e), and eosinophils (c, f) in wild-type, *Ffar2*-deficient (a–c), and

Ffar3-deficient (**d–f**) mice in a healthy state. Mean $\pm$ s.e.m., n = 5–8 mice per group. * $P < 0.05$, ** $P < 0.01$ (Student's *t*-test). NS: not significant.



Supplementary Fig. 8. CD4⁺ T cell depletion. **a** The experimental schemes of CD4⁺ T cell depletion. Wild-type mice were intraperitoneally administered PBS or Isotype control (100 µg/mouse) or anti-CD4 antibodies (100 µg/mouse) once (day -1) and evaluated on day 4. **b** Flow cytometric analysis of lung ILC2s and ST2⁺CD4⁺ T cells. **c** The number of CD4⁺ T cells, ST2⁺CD4⁺ T cells, ILC2s, and Eosinophils. Mean ± s.e.m., n = 3–4 mice per group. ***P* < 0.01,

*** $P < 0.001$, **** $P < 0.0001$ (one-way analysis of variance (ANOVA) and Tukey's post hoc test). NS; not significant.

Supplementary Table 1. Antibodies and reagents

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Biotin-conjugated anti-CD3 (17A2)	BioLegend	Cat# 100244
Biotin-conjugated anti-CD19 (6D5)	BioLegend	Cat# 115503
Biotin-conjugated anti-B220 (RA3-6B2)	BioLegend	Cat# 103204
Biotin-conjugated anti-CD11c (N418)	BioLegend	Cat# 117304
Biotin-conjugated anti-CD11b (M1/70)	BioLegend	Cat# 101204
Biotin-conjugated anti-Gr-1 (RB6-8C5)	BioLegend	Cat# 108404
Biotin-conjugated anti-CD5 (53-7.3)	eBioscience	Cat# 13-0051-85
Biotin-conjugated anti-CD8a (53-6.7)	eBioscience	Cat# 553029
Biotin-conjugated anti-NK1.1 (PK136)	eBioscience	Cat# 553163
Biotin-conjugated anti-FcεRI (MAR-1)	eBioscience	Cat# 13-5898-82
Biotin-conjugated anti-TER119	eBioscience	Cat# 13-5921-85
FITC-conjugated anti-CD45.2 (104)	BioLegend	Cat# 109805

FITC-conjugated anti-MHCII (M5/114.15.2)	BioLegend	Cat# 107605
FITC-conjugated anti-CD11c (N418)	BioLegend	Cat# 117305
FITC-conjugated anti-CD49b (DX5)	BioLegend	Cat# 108906
FITC-conjugated anti-CD3 (17A2)	BioLegend	Cat# 100203
FITC-conjugated anti-Foxp3 (FJK-16s)	eBioscience	Cat# 11-5773-82
PE-conjugated anti-FcεRIa (MAR-1)	BioLegend	Cat# 134307
PE-conjugated anti-NK1.1 (PK136)	BioLegend	Cat# 108708
PE-conjugated anti-Roryt (AFKJS-9)	eBioscience	Cat# 12-6988-80
PE-conjugated anti-SiglecF (1RNM44N)	eBioscience	Cat# 12-1702-82
PE-conjugated anti-ST2 (U29-93)	BD Pharmingen	Cat# 566311
PerCP Cy5.5-conjugated anti-CD45 (30-F11)	BioLegend	Cat# 103132
PerCP Cy5.5-conjugated anti-CD4 (30-GK1.5)	BioLegend	Cat# 100434
PECy7-conjugated anti-Gr-1 (RB6-8C5)	BioLegend	Cat# 108415

PECy7-conjugated anti-CD103 (2E7)	BioLegend	Cat# 121425
PECy7-conjugated anti-CD64 (X54-5/7.1)	BioLegend	Cat# 139313
PECy7-conjugated anti-CD127 (A7R34)	BioLegend	Cat# 135013
PECy7-conjugated anti-streptavidin	BD Pharmingen	Cat# 557598
APC-conjugated anti-CD317 (927)	BioLegend	Cat# 127015
APC-conjugated anti-CD127 (SB/199)	BioLegend	Cat# 121121
APC-conjugated anti-CD8a (53-6.7)	BioLegend	Cat# 100712
APC-conjugated anti-SiglecF (S17007L)	BioLegend	Cat# 155507
APC-conjugated anti-CD24 (M1/69)	BioLegend	Cat# 101813
Alexa Fluor 647-conjugated anti-Gata3 (L50-823)	BD Pharmingen	Cat# 560078
Alexa Fluor 700-conjugated anti-CD45 (30-F11)	BioLegend	Cat# 103127
APC Cy7-conjugated anti-CD4 (GK1.5)	BioLegend	Cat# 100413
APC Cy7-conjugated anti-CD11b (M1/70)	BioLegend	Cat# 101226
APC Cy7-conjugated anti-CD19 (6D5)	BioLegend	Cat# 115530

APC Cy7-conjugated anti-streptavidin	BioLegend	Cat# 405208
BV421-conjugated anti-CD11b (M1/70)	BioLegend	Cat# 101235
BV421-conjugated anti-CD11c (N418)	BioLegend	Cat# 117329
BV421-conjugated anti-T-bet (O4-46)	BD Pharmingen	Cat# 644815
CD16/32 (2.4G2) (93)	BioLegend	Cat# 101302
LIVE/DEAD Fixable Aqua Dead Cell Stain Kit for 405 nm excitation	Thermo Fisher Scientific	Cat# L34965
Chemicals, peptides, and recombinant proteins		
Papain, Carica papaya	Roche Diagnostics GmbH	Cat# 10108014001
Alternaria. Alternata (Alternaria tenuis), Lyophilized Form	Greer Laboratories	Cat#
Recombinant mouse IL-2	R&D Systems	Cat#
Recombinant mouse IL-33	R&D Systems	Cat#
Critical commercial assays		
RNeasy Mini Kit	Qiagen	Cat# 74106
SYBR Premix Ex Taq II	TAKARA	Cat# RR820B

Foxp3 / Transcription Factor Staining Buffer Set	eBioscience	Cat# 00-5523-00
Sodium propionate	Fujifilm Wako	Cat# 198-03015
InVivoMAb anti-mouse CD4 (clone YTS 191)	Bio X Cell	Cat# BE0119
InVivoMAb rat IgG2b isotype control, anti-keyhole limpet hemocyanin	Bio X Cell	Cat# BE0090
Quantikine ELISA Kits Mouse IL-4 Immunoassay	R&D Systems	Cat# M4000B-1
Quantikine ELISA Kits Mouse IL-5 Immunoassay	R&D Systems	Cat# M5000
Quantikine ELISA Kits Mouse IL-13 Immunoassay	R&D Systems	Cat# M1300CB
Naive CD4+ T cell isolation Kit, mouse	Miltenyi Biotec	Cat# 130-104-453

Supplementary Table 2. Primers used for qRT-PCR

Gene	Forward 5'- 3'	Reverse 5'- 3'
<i>mGapdh</i>	CGTCCCGTAGACAAAATGGT	TTGATGGCAACAATCTCCAC
<i>mIl4</i>	GGTCTCAACCCCCAGCTAGT	GCCGATGATCTCTCTCAAGTGAT
<i>mIl5</i>	GCAATGAGACGATGAGGCTT	CCCACGGACAGTTTGATTCT
<i>mIl13</i>	TGTGTCTCTCCCTCTGACCC	CACACTCCATACCATGCTGC
<i>mIl1rl1</i>	TGTATTTGACAGTTACGGAGGGC	ACTTCAGACGATCTCTTGAGACA
<i>mGata3</i>	CTCGGCCATTCGTACATGGAA	GGATACCTCTGCACCGTAGC
<i>mTnfrs4</i>	TACCTACCCCAGTGGTCACAA	ACGGATGACATAGAGTATCCCTG
<i>mMaf</i>	CTGCCGCTTCAAGAGGGTGCAGC	GATCTCCTGCTTGAGGTGGTC
<i>mCcr3</i>	TCGAGCCCGAACTGTGACT	CCTCTGGATAGCGAGGACTG

Supplementary Table 3. Gating strategies for flowcytometry

Cell strain	Flowcytometry gating
Alveolar macrophages (AMs)	CD45 ⁺ CD11b ^{int} CD11c ⁺ SiglecF ⁺ CD64 ⁺
Interstitial macrophages (IMs)	CD45 ⁺ CD11b ⁺ CD11c ⁻ SiglecF ⁻ CD64 ⁺
CD103 ⁺ dendritic cells (CD103 ⁺ DCs)	CD45 ⁺ CD11c ⁺ MHCII ⁺ CD24 ⁺ CD103 ⁺
CD11b ⁺ dendritic cells (CD11b ⁺ DCs)	CD45 ⁺ CD11c ⁺ MHCII ⁺ CD24 ⁺ CD11b ⁺
Plasmacytoid dendritic cells (pDCs)	CD45 ⁺ CD11b ⁻ MHCII ⁺ CD317 ⁺
Neutrophils	CD45 ⁺ CD11b ⁺ Gr-1 ⁺
Eosinophils	CD45 ⁺ CD11c ⁻ SiglecF ⁺ CD11b ⁺
Basophils	CD45 ⁺ FcεRI ⁺ CD49b ⁺
Mast cells	CD45 ⁺ FcεRI ⁺ c-kit ⁺
CD4 ⁺ T cells	CD45 ⁺ CD4 ⁺

Th1 cells	CD45 ⁺ CD4 ⁺ T-bet ⁺
Th2 cells	CD45 ⁺ CD4 ⁺ Gata3 ⁺
ST2 ⁺ CD4 ⁺ T cells	CD45 ⁺ CD4 ⁺ ST2 ⁺
Th17 cells	CD45 ⁺ CD4 ⁺ Rory ⁺
Regulatory T cells (Treg cells)	CD45 ⁺ CD4 ⁺ Foxp3 ⁺
ILCs	CD45 ⁺ Lineage ⁻ CD4 ⁻ CD127 ⁺
ILC1s	CD45 ⁺ Lineage ⁻ CD4 ⁻ CD127 ⁺ T-bet ⁺
ILC2s	CD45 ⁺ Lineage ⁻ CD4 ⁻ CD127 ⁺ Gata3 ⁺ or CD45 ⁺ Lineage ⁻ CD4 ⁻ CD127 ⁺ ST2 ⁺
ILC3s	CD45 ⁺ Lineage ⁻ CD4 ⁻ CD127 ⁺ Rory ⁺
CD8 ⁺ T cells	CD45 ⁺ CD3 ⁺ CD8 ⁺
NK cells	CD45 ⁺ NK1.1 ⁺

B cells	CD45 ⁺ CD19 ⁺
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