

Supplementary Information

Stage-specific GATA3 induction promotes ILC2 development after lineage commitment

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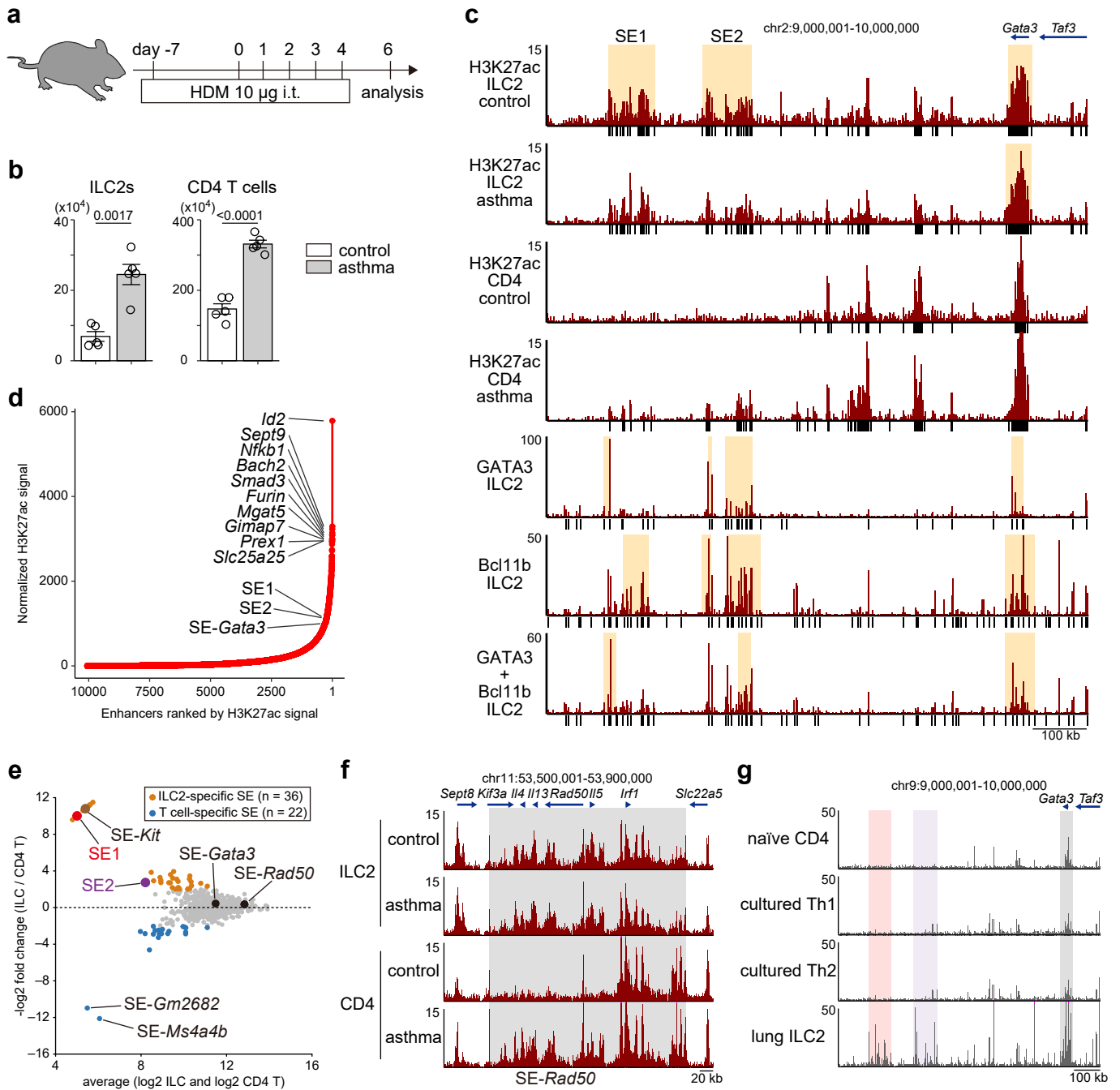
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This Supplementary Information file contains:

Supplementary Figure 1-12

Supplementary Table 1, 2

Supplementary Figure 1

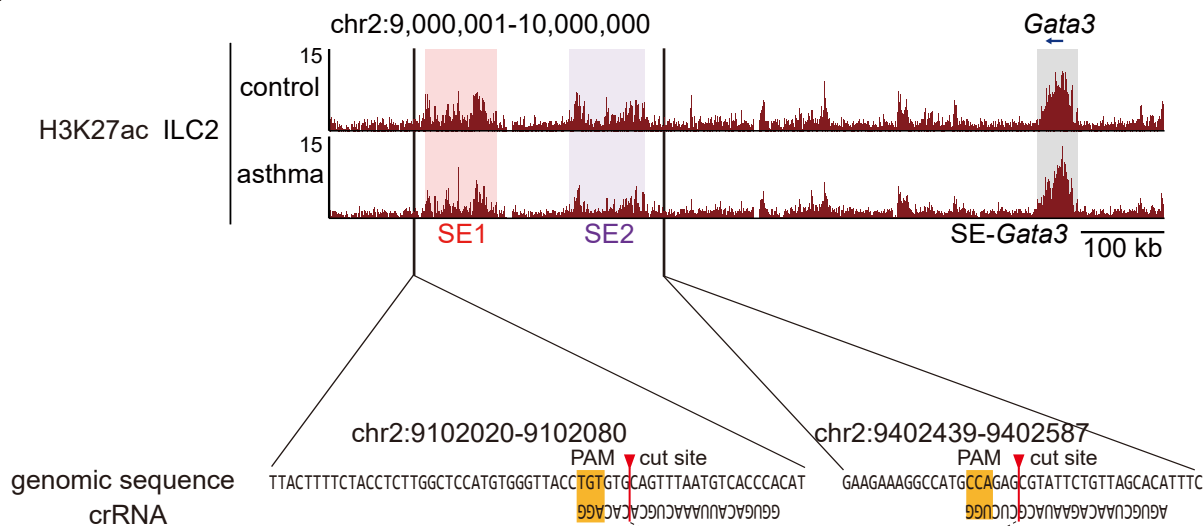


Supplementary Figure 1. ILC2-specific super-enhancers identified via H3K27ac ChIP-seq.

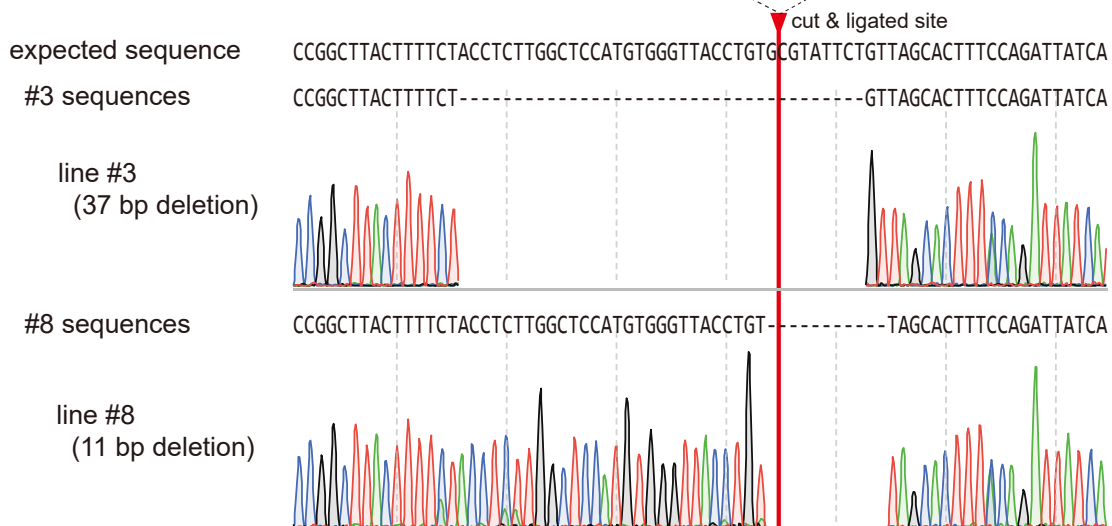
a Experimental procedure of HDM-induced allergic airway inflammation. C57BL/6J mice were intratracheally administered with house dust mites (HDM) on days -7, 0, 1, 2, 3, 4. i.t.: intratracheal. **b-f** Samples were harvested from control mice and asthma mice on day 0 and day 6. ILC2 (CD45⁺Lin1⁻Thy1.2⁺ST2⁺ cells) and CD4 T cells (CD45⁺Lin2⁻CD4⁺ cells) were isolated. Lin1: CD4, CD5, CD8a, CD11b, CD11c, CD19, CD31, CD49b, FcεR1, TCRβ, TCRγδ, Ter119; Lin2: CD8a, CD11c, CD49b, TCRγδ, EpCAM. **b** The number of ILC2s and CD4 T cells in the lung, n=5. Data are presented as mean±SE. Statistical analysis was performed using unpaired, two-sided Welch's *t*-test. *p* values are shown on the graphs. **c** ILC2s and CD4 T cells were subjected to ChIP-seq analysis of H3K27 acetylation (H3K27ac). UCSC tracks of ChIP-seq of ILC2 and CD4 T cells, and GATA3 (GSE11187), Bcl11b (GSE131082), GATA3+Bcl11b ChIP-seq data. Black bars indicate typical enhancers, and orange boxes indicate super-enhancers. **d** The plot of H3K27ac signal versus rank of stitched enhancers in control ILC2. **e** MA-plot of tag counts on the super-enhancers based on H3K27ac in ILC2 and CD4 T cells under HDM-asthmatic conditions. Total: 624 SEs. A fold change > 4 was considered specific. **f** UCSC tracks of H3K27ac ChIP-seq for ILC2 and CD4 T cells in the *Rad50-Irf1* region. Rectangles indicate super-enhancer regions. **g** UCSC tracks of ATAC-seq for naïve CD4 T cells, cultured Th1 cells, cultured Th2 cells (GSE159505), and lung ILC2 in steady-state conditions. Source data are provided as a Source Data file.

Supplementary Figure 2

a



b

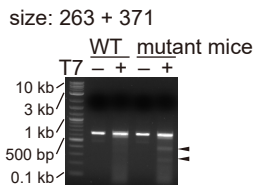


Supplementary Figure 2. Generation of G3SEKO mice.

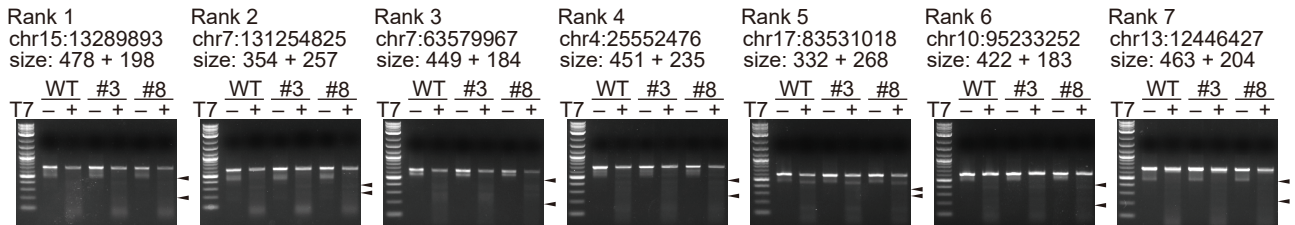
a Upper panel: UCSC genome tracks around the *Gata3* and G3SE region. Lower panel: Genomic sequence of upstream/downstream crRNA target regions. **b** DNA sequences of G3SE deleted loci from two lines of generated G3SEKO mice.

Supplementary Figure 3

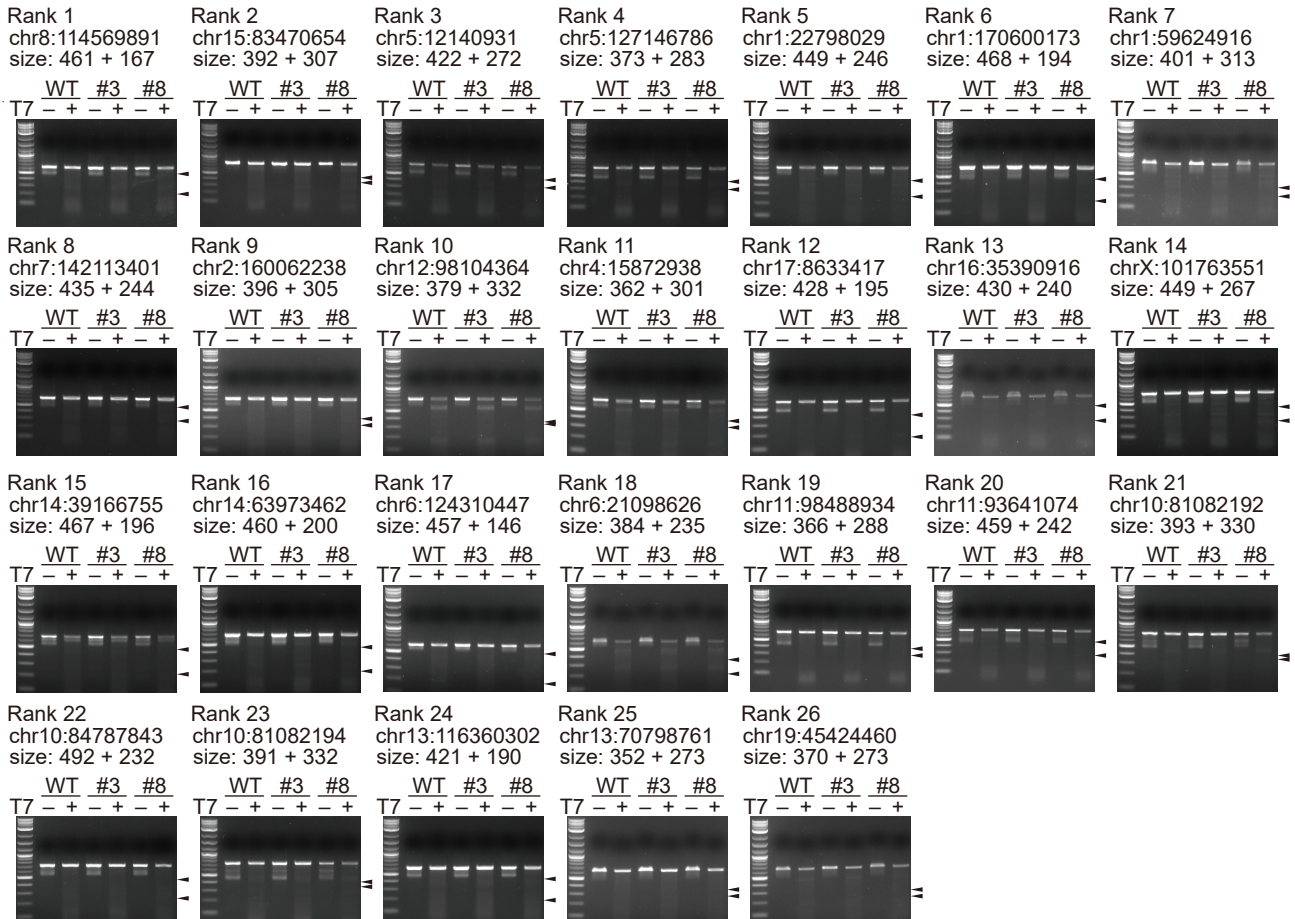
a Positive control for T7 assay



b Off-targets of upstream crRNA (AGUGCUAACAGAAUACGCUCUGG)



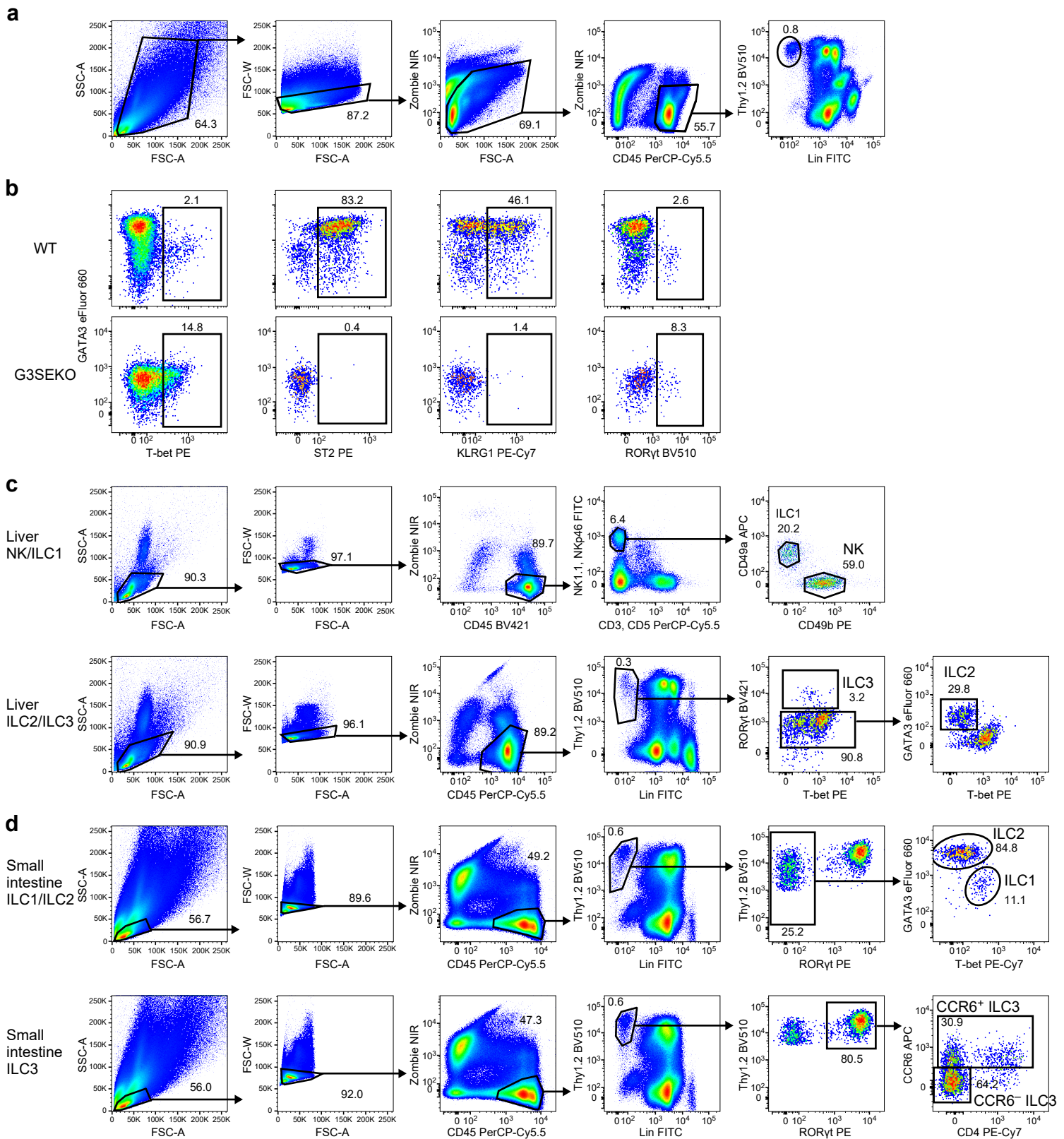
c Off-targets of upstream crRNA (GGUGACAUUAAACUGCACACAGG)



Supplementary Figure 3. Off-target analyses of crRNA-derived cleavage.

The off-target candidates induced by crRNA were identified using CAS-OFFinder. The tail DNA from wild-type and G3SEKO mice was mixed in equimolar ratios to generate PCR templates. The regions flanking the candidate sites were amplified by PCR, and the off-target insertion/deletion was determined using the T7 Endonuclease mismatch detection method. Arrowheads indicate the predicted cleavage sites. **a** The positive control of T7 endonuclease mismatch detection method. Template DNA was extracted from the tail of mutant mice generated by single crRNA. **b-c** T7 assay results for the off-target candidates of upstream (**b**) and downstream (**c**) crRNAs. Uncropped gel photos are provided as a Source Data file.

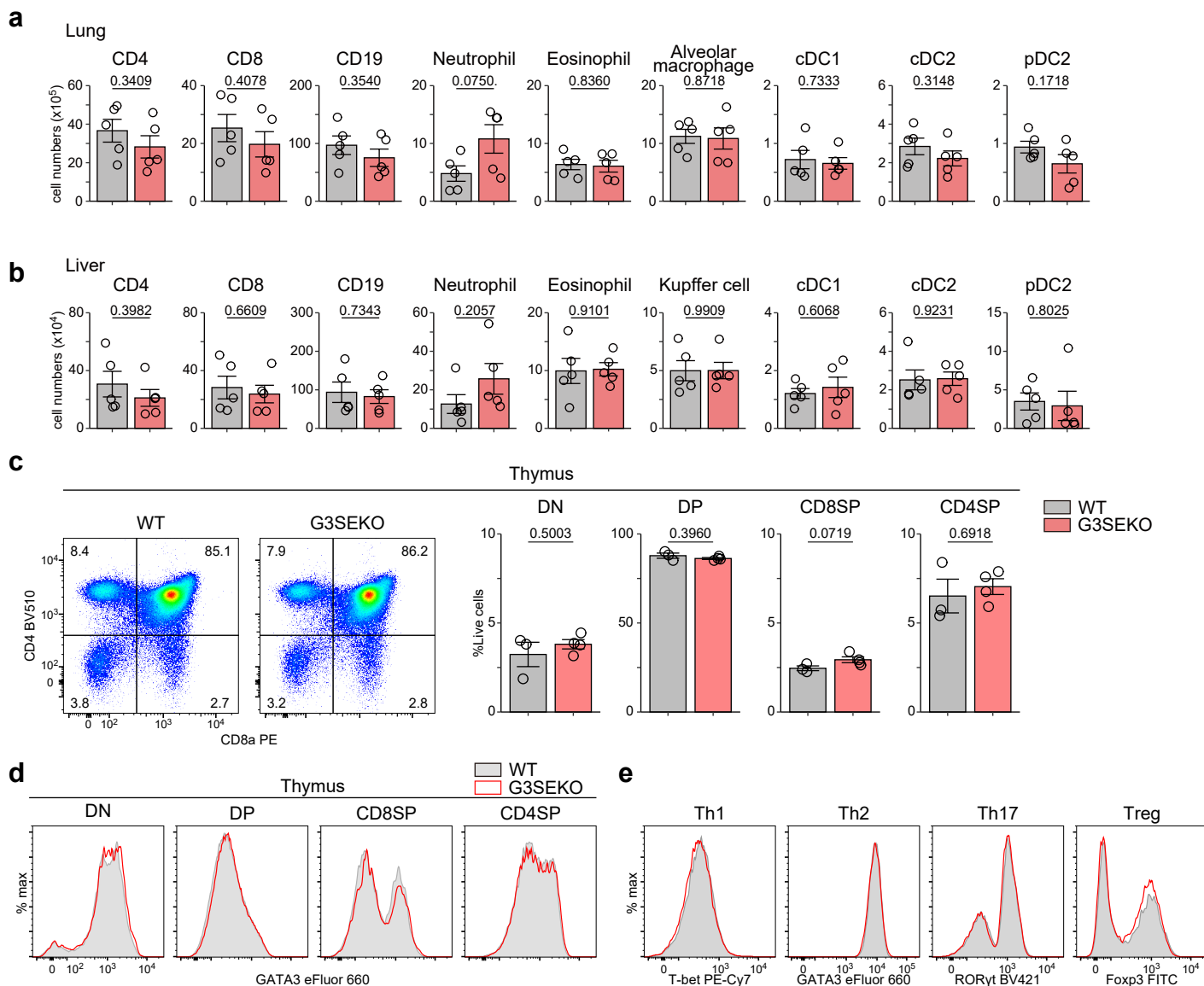
Supplementary Figure 4



Supplementary Figure 4. Gating strategies for lung, liver, and small intestinal ILCs.

a-b Gating strategy for lung ILCs (**a**) and ILC2 (**b**). Lin: CD3 ϵ , CD4, CD5, CD8a, CD11b, CD11c, CD19, CD49b, Gr1, B220, Ter119. **c** Gating strategy for liver NK cells and ILC1 (upper panels) and ILC2 and ILC3 (lower panels). Lin: CD3 ϵ , CD4, CD5, CD8a, CD11b, CD11c, CD19, Gr1, B220, Ter119. **d** Gating strategy for small intestinal ILC1/ILC2 (upper panel) and ILC3 (lower panel). Lin: CD3 ϵ , CD5, CD8a, CD11b, CD11c, CD19, Gr1, B220, Ter119.

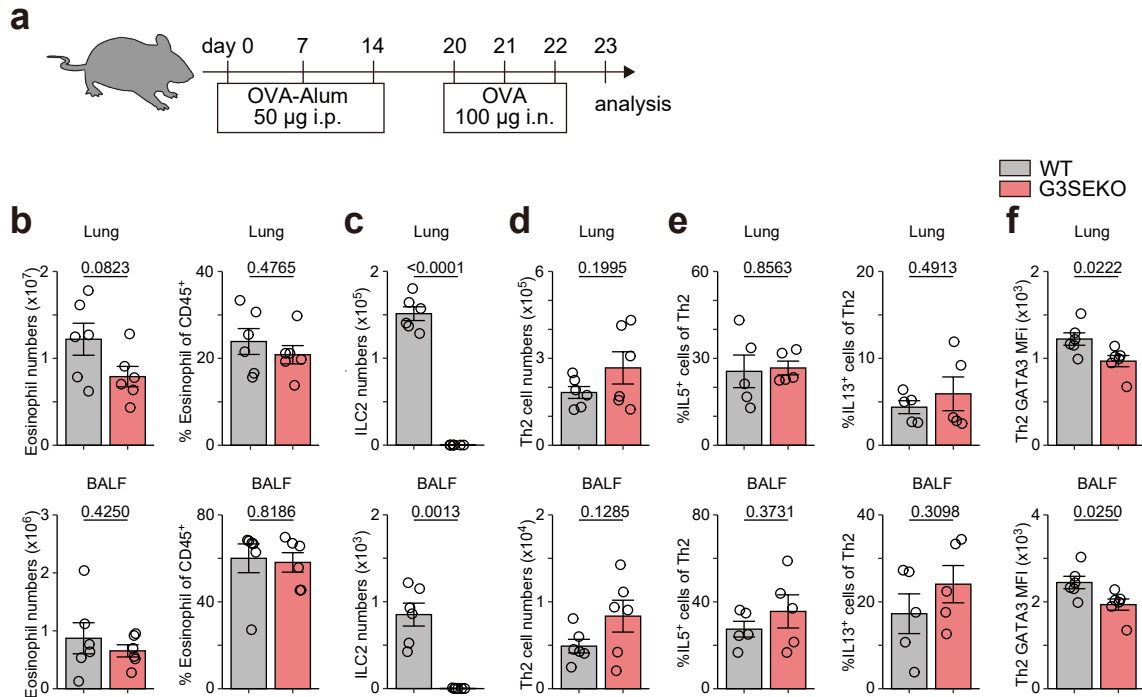
Supplementary Figure 5



Supplementary Figure 5. Immune cell development and helper T cell differentiation in G3SEKO mice.

a-b Distribution of lymphocytes and myeloid populations in the lung (**a**) and the liver (**b**) of WT and G3SEKO mice. $n=5$. **c** Flow cytometry analysis of thymocytes in WT ($n=3$) and G3SEKO ($n=4$) mice. DN: double negative, DP: double positive, SP: single positive. **d** GATA3 expression in thymocytes. **e** Naïve CD4 T cells were cultured under T cell subset-polarizing conditions for three days, and master transcription factor expression was analyzed. (**a-c**), Each data point indicates one mouse from two independent experiments. Data are presented as mean \pm SE. Statistical analysis was performed using unpaired, two-sided Welch's t-test. p values are shown on the graphs. (**d-e**) Shown are representative of six WT mice and five G3SEKO mice from two independent experiments. Source data are provided as a Source Data file.

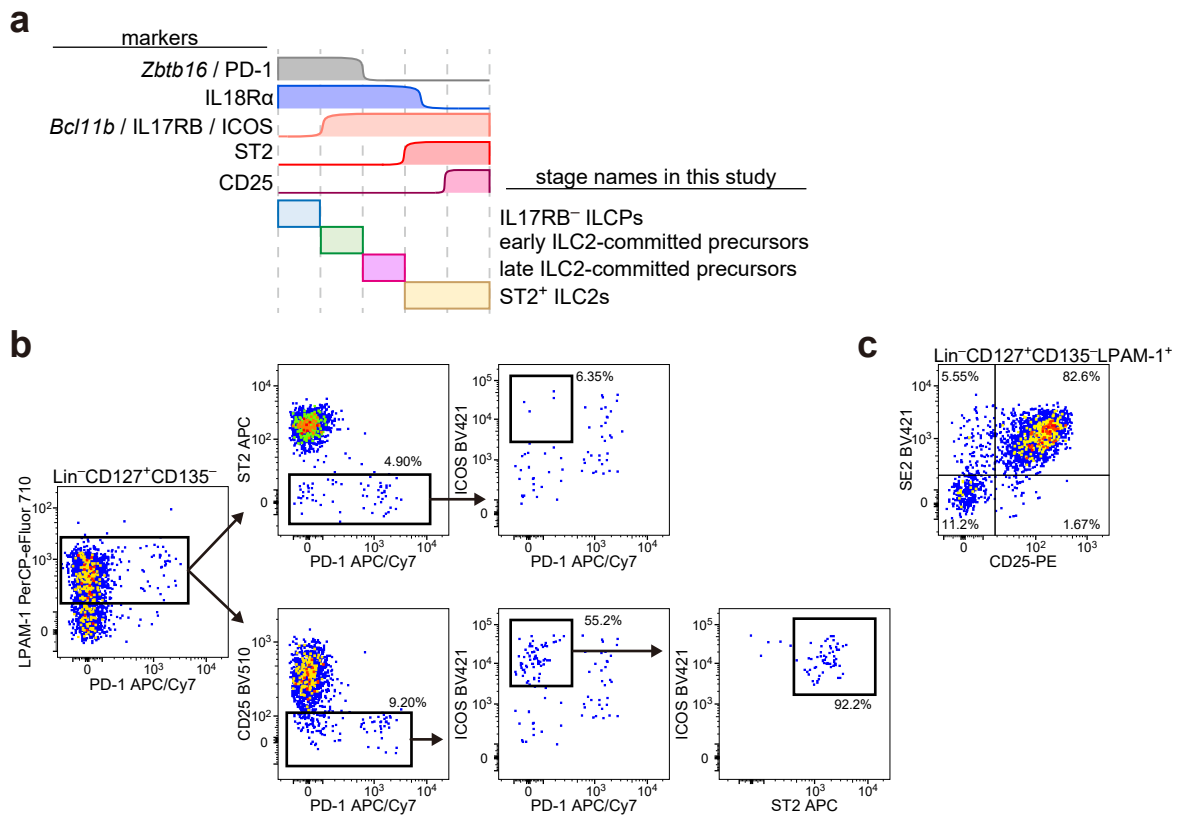
Supplementary Figure 6



Supplementary Figure 6. OVA-induced allergic airway inflammation in G3SEKO mice.

a Scheme of OVA-induced asthma model. **b** Eosinophil inflammation in lung and bronchoalveolar lavage fluid (BALF). **c** Cell numbers of ILC2s (Lin⁻Thy1.2⁺ST2⁺ cells). **d** Cell numbers of Th2 cells (CD3e⁺CD4⁺ST2⁺ cells). **e** IL-5 and IL-13 expression of Th2 cells. **f** GATA3 expression of Th2 cells. **b-f** Each data point indicates one mouse from three (**b-d, f**, n=6) or two (**e**, n=5) independent experiments. Data are presented as mean \pm SE. Statistical analysis was performed using unpaired, two-sided Welch's t-test. p values are shown on the graphs. Source data are provided as a Source Data file.

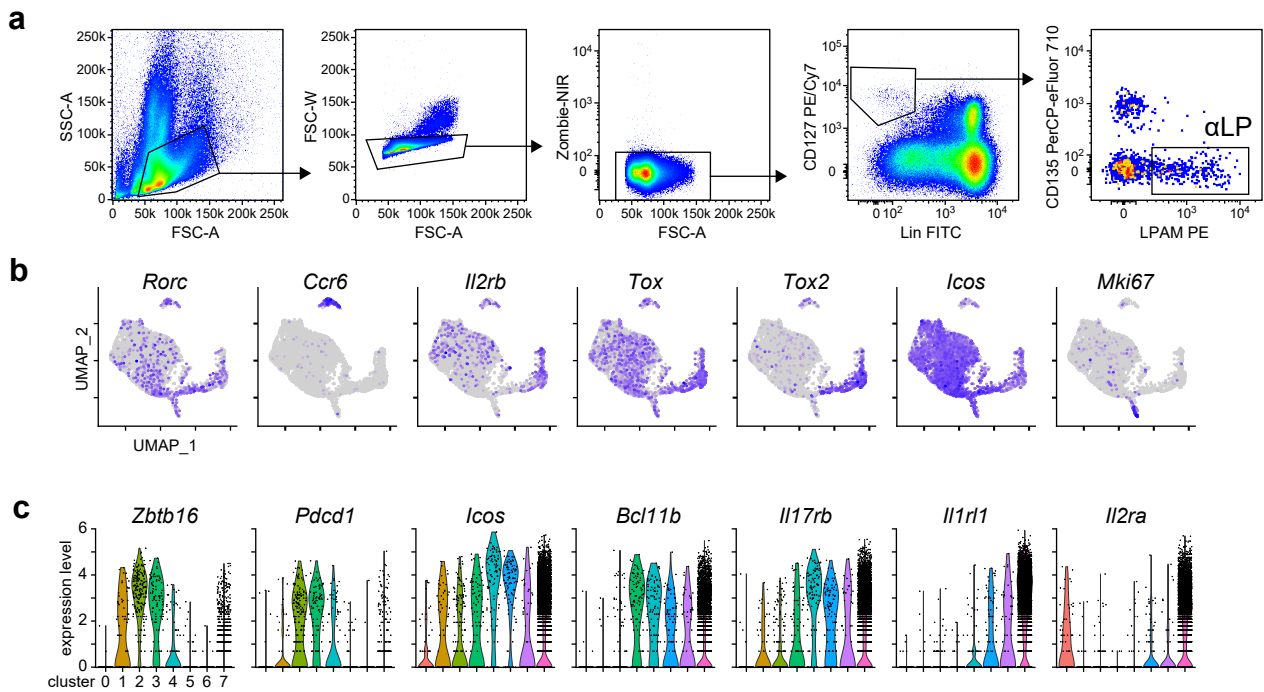
Supplementary Figure 7



Supplementary Figure 7. ILC2 lineages in the BM.

a The terms of ILC2 lineages in the BM in this study. **b** Flow cytometry analysis of WT BM cells for the late ILC2-committed precursors. **c** Flow cytometry analysis of ST2 and CD25 of WT Lin⁻CD127⁺CD135⁻LPAM-1⁺ BM cells. The plots are representative of two independent experiments (**b-c**).

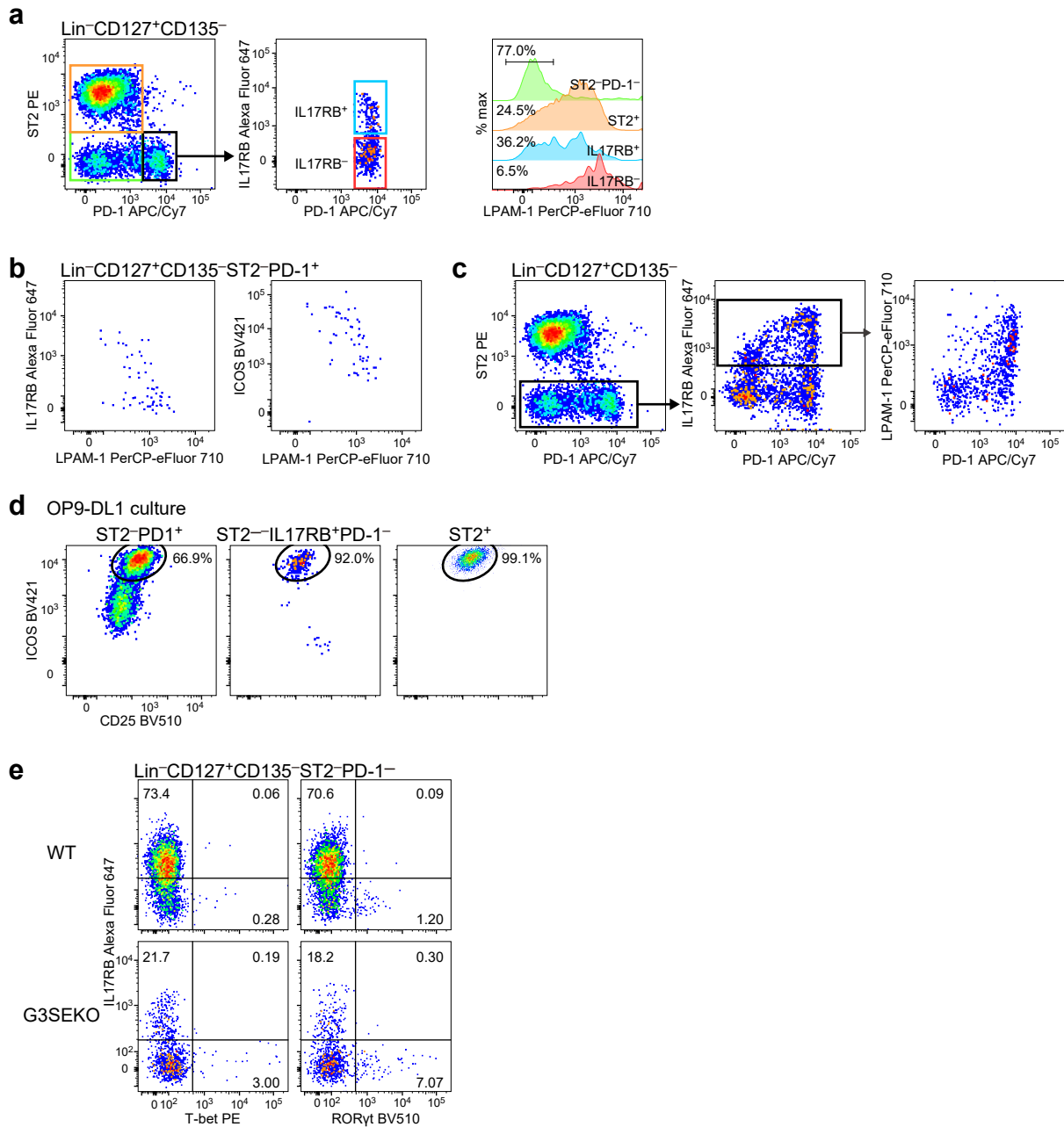
Supplementary Figure 8



Supplementary Figure 8. Gating strategies and the expression of ILC-related genes.

a Gating strategy of α LP-sorting for scRNA-seq experiments. Lin: CD3 ϵ , CD4, CD5, CD8a, CD11b, CD11c, CD19, CD49b, Gr1, B220, Ter119. **b-c** Feature plots (**b**) and violin plots (**c**) show the expression of ILC development-related genes.

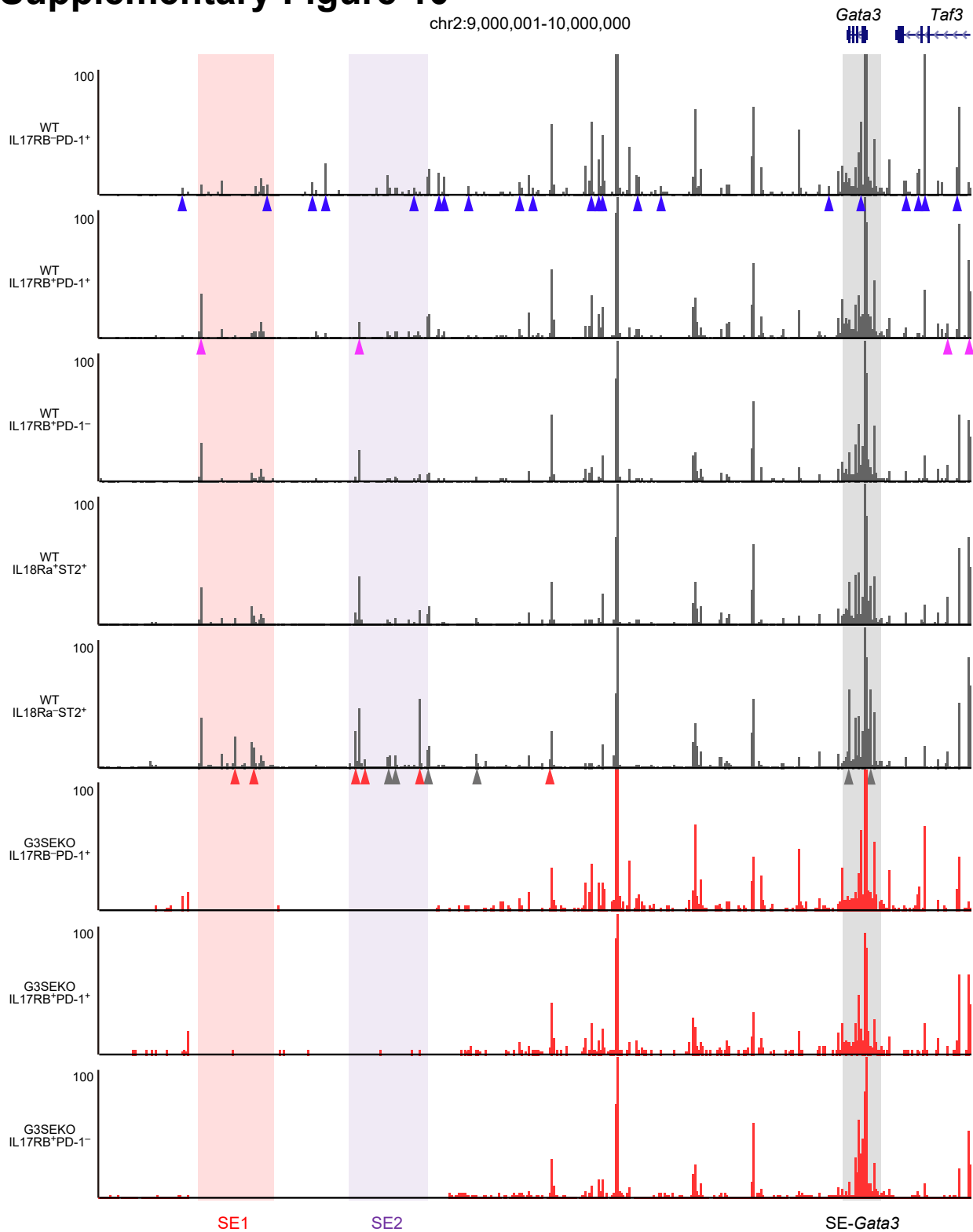
Supplementary Figure 9



Supplementary Figure 9. Analysis of late ILC2-committed precursors in the BM.

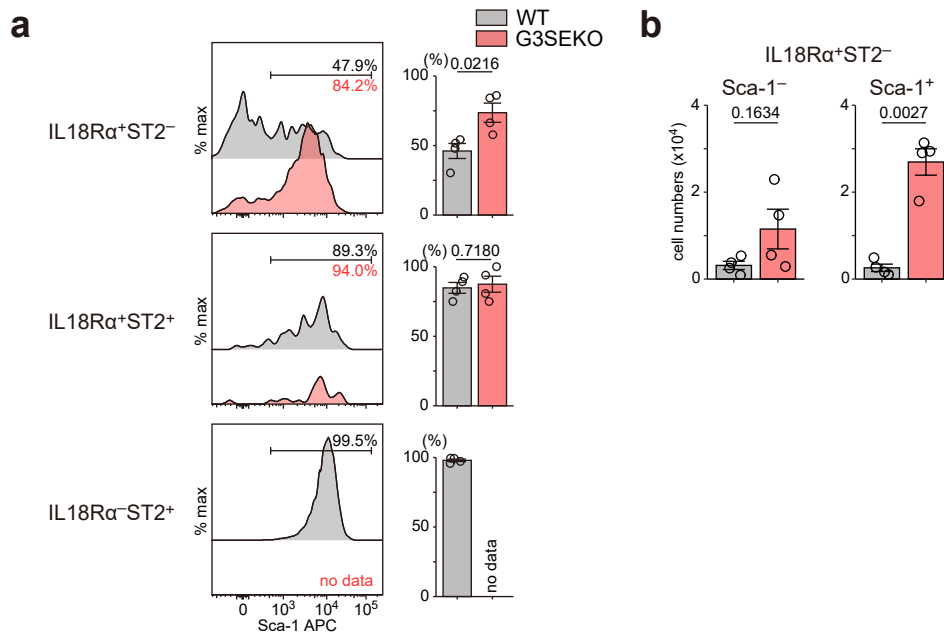
a Flow cytometry analyses of LPAM expression in Lin⁻CD127⁺CD135⁻ cells in WT BM. **b** Relationship between LPAM-1 expression and IL17RB or ICOS expression in WT PD-1⁺ ILCPs. **c** Relationship between PD-1 expression and LPAM-1 expression in WT ST2-IL17RB⁺ cells. **d** Indicated cells were sorted from the WT Lin⁻CD127⁺CD135⁻ BM cells and were cultured on OP9-DL1 cells in the presence of IL-7 and SCF for 5 days. Flow cytometry analysis of cultured cells. **e** Flow cytometry analyses of Lin⁻CD127⁺ST2⁻PD-1⁻ BM cells. Lin: CD3ε, CD4, CD5, CD8a, CD11b, CD11c, CD19, CD49b, CD135, Gr1, B220, Ter119. The plots were representative of four (**a-d**) and two (**e**) mice from two independent experiments.

Supplementary Figure 10



Supplementary Figure 10. ATAC-seq of ILC2 lineages in the BM of WT and G3SEKO mice. ATAC-seq was performed for IL17RB⁻ ILCP, early and late ILC2-committed precursors, IL18Rα⁺ST2⁺ ILC2 and IL18Rα⁻ST2⁺ ILC2 from WT BM cells and IL17RB⁻ ILCP, early and late ILC2-committed precursors from G3SEKO BM cells. Red, purple, and grey arrowheads indicate open chromatin regions; red arrow heads indicate late opened chromatin, purple arrowheads indicate early opened chromatin, and blue arrowheads indicate closed chromatin during ILC2 development.

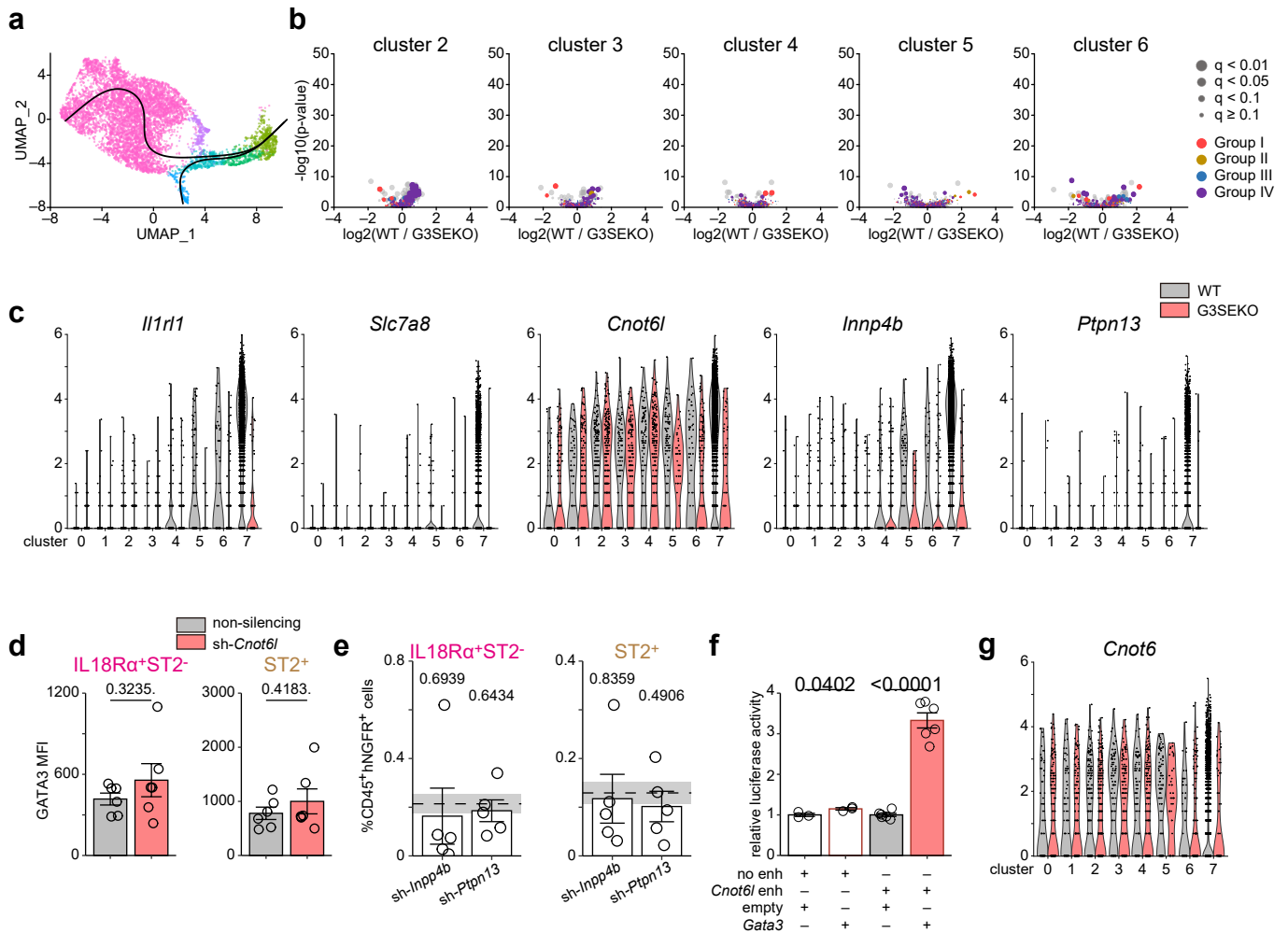
Supplementary Figure 11



Supplementary Figure 11. Sca-1 expression of lung ILC2 lineages.

a, Sca-1 expression of indicated cells in the lung. n=4. Data were representative of four mice from three independent experiments. **b**, Cell numbers of IL18Rα⁺ST2⁻Sca-1⁻ cells and IL18Rα⁺ST2⁻Sca-1⁺ cells. (**a**, **b**) Data are presented as mean±SE. Statistical analysis was performed using unpaired, two-sided Welch's t-test. p values are shown on the graphs. Source data are provided as a Source Data file.

Supplementary Figure 12



Supplementary Figure 12. Supporting data related to Fig. 5.

a Pseudotime analysis for clusters 2-7 by slingshot. **b** Differentially expressed genes between WT and G3SEKO cells in cluster 2 to cluster 6. **c** Expression of the candidate genes, related to Fig. 5f. **d** Gata3 expression of sh-*Cnot6l* transduced *IL18Rα⁺ST2⁻* cells and *ST2⁺* cells in the lung, related to Fig. 5h. n=6. **e** Lung ILC frequency (*CD45⁺hNGFR⁺Lin⁻Thy1.2⁺* cells) in *CD45⁺hNGFR⁺* cells, related to Fig. 5h. Dash lines and grey areas indicate the mean and SE of non-silencing samples, respectively. Statistical analysis were performed between indicated shRNA-transduced cells and non-silencing vector-transduced cells. n=5. **f** Dual-luciferase Reporter Assay for the *Cnot6l* enhancer region in 293T cells transiently transduced with MSCV-empty vector or MSCV *Gata3* vector. no enhancer: n=3, *Cnot6l* enhancer n=6. **g** Expression of *Cnot6* by scRNA-seq analysis. Each data point indicates one mouse from four (**d**, **e**) and two (**f**) independent experiments. (**d-f**) Data are presented as mean±SE. Statistical analysis was performed using unpaired, two-sided Welch's t-test. p values are shown on the graphs. Source data are provided as a Source Data file.

Supplementary Table 1. The list of oligonucleotides

| name | sequence |
|---------------------------------|---|
| G3SEKO 5' crRNA | Alt-R1/AGUGCUAACAGAAUACGCUC/Alt-R2 |
| G3SEKO 3' crRNA | Alt-R1/GGUGACAUUAAACUGCACAC/Alt-R2 |
| G3SEKO genotype forward primer | GACCCTTTAGTTTAGCTCCTTGC |
| G3SEKO genotype reverse primer | CTCACCATGGAACTTACAGCTC |
| G3SEWT genotype forward primer | GCTGGATAGAAGGCTGACTCATA |
| G3SEWT genotype reverse primer | ATGGGGTACAATGCTAAACAGAA |
| G3SE2KO 5' crRNA | Alt-R1/AGUGCUAACAGAAUACGCUC/Alt-R2 |
| G3SE2KO 3' crRNA | Alt-R1/GUGGAGGUGGUUACCCAAC/Alt-R2 |
| G3SE2KO genotype forward primer | TATTCTCCCTTGTGGTCAATCC |
| G3SE2KO genotype reverse primer | CAATGTCTCCACTATGCTCCAG |
| G3SE2WT genotype forward primer | GCTGGAGTGGTAAGGAGCAG |
| G3SE2WT genotype reverse primer | GGAGCCTTGAGTCCTGAGTG |
| miR-30 common forward primer | CAGAAGGCTCGAGAAGGTATATTGCTGTTGACAGTGAGCG |
| miR-30 common reverse primer | CTAAGTAGCCCCTTGAATTCGAGGCAGTAGGCA |
| shRNA targeting <i>Ptpn13</i> | TGCTGTTGACAGTGAGCGTTGCTACCAGAAATCGATACTAT AGTGAAGCCACAGATGTATAGTATCGATTTCTGGTAGCACT GCCTACTGCCTCGGA |
| shRNA targeting <i>Sptssa</i> | TGCTGTTGACAGTGAGCGCGCATTTCACGTTGTGATAAAT AGTGAAGCCACAGATGTATTTATCACACGTGCAAATGCAT GCCTACTGCCTCGGA |
| shRNA targeting <i>Cnot6l</i> | TGCTGTTGACAGTGAGCGATGGAAACAGAGCAATACTTTAT AGTGAAGCCACAGATGTATAAAGTATTGCTCTGTTTCCACT GCCTACTGCCTCGGA |
| shRNA targeting <i>Inpp4b</i> | TGCTGTTGACAGTGAGCGTAACTGCATTATCGCTAAAGTAT AGTGAAGCCACAGATGTATACTTTAGCGATAATGCAGTTCT GCCTACTGCCTCGGA |
| shRNA targeting <i>Tespa1</i> | TGCTGTTGACAGTGAGCGAATCTCGTATCCCTGCAAGATTT AGTGAAGCCACAGATGTAAATCTTGCAGGGATACGAGATGT GCCTACTGCCTCGGA |
| ATAC-seq universal primer Ad1 | AATGATACGGCGACCACCGAGATCTACACTCGTCGGCAGC GTCAGATGTG |
| ATAC-seq index primer Ad2.1 | CAAGCAGAAGACGGCATAACGAGATTCGCCTTAGTCTCGTG GGCTCGGAGATGT |
| ATAC-seq index primer Ad2.2 | CAAGCAGAAGACGGCATAACGAGATCTAGTACGGTCTCGTG GGCTCGGAGATGT |
| ATAC-seq index primer Ad2.3 | CAAGCAGAAGACGGCATAACGAGATTTCTGCCTGTCTCGTGG GCTCGGAGATGT |
| ATAC-seq index primer Ad2.4 | CAAGCAGAAGACGGCATAACGAGATGCTCAGGAGTCTCGTG GGCTCGGAGATGT |
| ATAC-seq index primer Ad2.5 | CAAGCAGAAGACGGCATAACGAGATAGGAGTCCGTCTCGTG GGCTCGGAGATGT |
| Off-target upstream Rank1_F | TTTGGAGGTTTGGCTTCCCTA |
| Off-target upstream Rank1_R | ATCAAAGTGGACCTGGTTGC |
| Off-target upstream Rank2_F | TGTCCAATGGCAGACTTCAA |
| Off-target upstream Rank2_R | GGTGGTCAGTGTCCATCAGA |
| Off-target upstream Rank3_F | CATAGGGGACTTTTCGGGATAA |
| Off-target upstream Rank3_R | TCACTGGGTGTCCCACTAA |
| Off-target upstream Rank4_F | GGCAGTAGATTCTGGGATG |
| Off-target upstream Rank4_R | TGCTACCGCTGAAAAGGTC |
| Off-target upstream Rank5_F | TGAGGTTCCCATCACTACG |
| Off-target upstream Rank5_R | GAGACGTGTTGCATCAAGGA |
| Off-target upstream Rank6_F | AGAATCGAAGTTGGCACACC |
| Off-target upstream Rank6_R | ACCCCATCTCCCAATTAAC |
| Off-target upstream Rank7_F | CAGTGCCAGTTCCAAGACAG |

| | |
|--------------------------------|-----------------------|
| Off-target upstream Rank7 R | TGTAGCCCCAAGGAATTCTG |
| Off-target downstream Rank1 F | ATGTGTGCGTGTGGATGAGT |
| Off-target downstream Rank1 R | TGGGATCTAGGCATCTGGAG |
| Off-target downstream Rank2 F | AAGCCCAAGTTCAGTCCTCA |
| Off-target downstream Rank2 R | AGGTCCCCTCTGTCACCTTT |
| Off-target downstream Rank3 F | GTGTTGGGGGAGTGGATAGA |
| Off-target downstream Rank3 R | TGTCAACAGAGCCACTTTTCG |
| Off-target downstream Rank4 F | GTGGCATGTGCCATCTGTAG |
| Off-target downstream Rank4 R | ATTTCCAGTCAGCGGTTGTC |
| Off-target downstream Rank5 F | CTTCATCAGGGACTGCTCCT |
| Off-target downstream Rank5 R | CCCCCTGCCTCAACTTATTT |
| Off-target downstream Rank6 F | GAGCTGCCTCTGCAAACCTCT |
| Off-target downstream Rank6 R | AACTCTGGGGCATCTGATTG |
| Off-target downstream Rank7 F | GGCTCACAACCACCCATAAC |
| Off-target downstream Rank7 R | AAGGCACATGGATGAAAAGG |
| Off-target downstream Rank8 F | CCTGTGTGTGTATGCGTGTG |
| Off-target downstream Rank8 R | GGGGTTCCAGTGTACAGAA |
| Off-target downstream Rank9 F | CCACTCGTCCCACGTACTION |
| Off-target downstream Rank9 R | TGGGTGCGTCTAAAGTGTGA |
| Off-target downstream Rank10 F | TTCTGTGCCCTTTTGCTCTT |
| Off-target downstream Rank10 R | CCCCAATCCACTTCTCTCA |
| Off-target downstream Rank11 F | CCCTCCCTCCCTGTTTTCTA |
| Off-target downstream Rank11 R | TTCATCCCACCTGCTTTTTTC |
| Off-target downstream Rank12 F | CGAGGGAAGGAATGTTTGAA |
| Off-target downstream Rank12 R | TATGCACGGCTAGTCAGCAC |
| Off-target downstream Rank13 F | TGGCTCAGTGCATAAAGGTG |
| Off-target downstream Rank13 R | TTTTGTCTTTCTCCCCGAAA |
| Off-target downstream Rank14 F | CCAAGTGGCAGATTCCTTA |
| Off-target downstream Rank14 R | GGGATCTGACACCGTCATCT |
| Off-target downstream Rank15 F | TCTGTGCGAGTGTGTGAGTG |
| Off-target downstream Rank15 R | GAAGCATCCTGAGCCATTA |
| Off-target downstream Rank16 F | GCTCAGCTCAGCTACCTGCT |
| Off-target downstream Rank16 R | CCCTATTTGTCCTGCTCTGC |
| Off-target downstream Rank17 F | TGTCACCGTCCTTTTAGCC |
| Off-target downstream Rank17 R | TAGCACTGTTTGGGGAGGT |
| Off-target downstream Rank18 F | TCACCCTGACCAGTGAAAAA |
| Off-target downstream Rank18 R | AATGGTGCTTGGATTGGAAG |
| Off-target downstream Rank19 F | TGTTTGATTGCTGCTGCTCT |
| Off-target downstream Rank19 R | TGCCAAGAAATGGAGAAACC |
| Off-target downstream Rank20 F | TTTGACTIONAAGCCCCAGA |
| Off-target downstream Rank20 R | AGCTGCAGGAGAAAGAGGTG |
| Off-target downstream Rank21 F | AGGTTGGCATGGACTACAGG |
| Off-target downstream Rank21 R | TGCATGTGCTCACTGTCTGA |
| Off-target downstream Rank22 F | CTTCATGGGGAGCAGTGTTT |
| Off-target downstream Rank22 R | CTCCAGGTGTATGGCATGAA |
| Off-target downstream Rank23 F | AGGTTGGCATGGACTACAGG |
| Off-target downstream Rank23 R | TGCATGTGCTCACTGTCTGA |
| Off-target downstream Rank24 F | AGGTTTCTGCAAAGCTTGA |
| Off-target downstream Rank24 R | AAAGCAATGTGTGGTGGTTTC |
| Off-target downstream Rank25 F | GGACTTGCAGAGCAGGACTC |
| Off-target downstream Rank25 R | CCACGTGCCTGTTTATGTAT |
| Off-target downstream Rank26 F | TGTGTGCAGGTGTGTGAAGA |
| Off-target downstream Rank26 R | TCCCCTACTGATCCGAATGT |

Supplementary Table 2. The list of antibodies

| Antibodies | Company |
|---|--------------------------|
| Rat anti-mouse CD3 ϵ FITC (Clone 145-2C11) | BioLegend |
| Rat anti-mouse CD3 ϵ PE (Clone 145-2C11) | BioLegend |
| Rat anti-mouse CD3 ϵ PerCP/Cyanine5.5 (Clone 145-2C11) | BioLegend |
| Rat anti-mouse CD3 ϵ Brilliant Violet 510 (Clone 145-2C11) | BioLegend |
| Rat anti-mouse CD4 FITC (Clone RM4-5) | BioLegend |
| Rat anti-mouse CD4 PE (Clone RM4-5) | BioLegend |
| Rat anti-mouse CD4 PE/Cyanine7 (Clone RM4-5) | BioLegend |
| Rat anti-mouse CD4 Brilliant Violet 510 (Clone RM4-5) | BioLegend |
| Rat anti-mouse CD5 FITC (Clone 53-7.3) | BioLegend |
| Rat anti-mouse CD5 PerCP/Cyanine5.5 (Clone 53-7.3) | BioLegend |
| Rat anti-Mouse CD8a FITC (Clone 53-6.7) | BD Bioscience |
| Rat anti-Mouse CD8a PE (Clone 53-6.7) | BD Bioscience |
| Rat anti-Mouse CD8a PE/Cyanine7 (Clone 53-6.7) | BioLegend |
| Rat anti-mouse/human CD11b FITC (Clone M1/70) | BioLegend |
| Rat anti-mouse/human CD11b Pe/Cyanine7 (Clone M1/70) | BioLegend |
| Hamster anti-mouse CD11c FITC (Clone N418) | BioLegend |
| Hamster anti-mouse CD11c PE/Cyanine7 (Clone N418) | BioLegend |
| Purified anti-mouse CD16/32 (Clone 93) | BioLegend |
| Rat anti-mouse CD19 FITC (Clone 6D5) | BioLegend |
| Rat anti-mouse CD25 PE (Clone PC61) | BioLegend |
| Rat anti-mouse CD25 Brilliant Violet 421 (Clone PC61) | BioLegend |
| Rat anti-mouse CD25 Brilliant Violet 510 (Clone PC61) | BioLegend |
| Rat anti-mouse CD44 PerCP-Cy5.5 (Clone IM7) | BD Bioscience |
| Rat anti-mouse/human CD45R/B220 FITC (Clone RA3-6B2) | BioLegend |
| Rat anti-mouse/human CD45R/B220 Brilliant Violet (Clone RA3-6B2) | BioLegend |
| Rat anti-mouse CD45 PerCP/Cyanine5.5 (Clone 30-F11) | BioLegend |
| Rat anti-mouse CD45 BV421 (Clone 30-F11) | BD Bioscience |
| Mouse anti-mouse CD45.1 FITC (Clone A20) | BioLegend |
| Mouse anti-mouse CD45.1 PerCP/Cyanine5.5 (Clone A20) | BioLegend |
| Mouse anti-mouse CD45.1 APC (Clone A20) | BioLegend |
| Mouse anti-mouse CD45.1 Brilliant Violet 605 (Clone A20) | BioLegend |
| Mouse anti-mouse CD45.2 APC/Cyanine7 (Clone 104) | BioLegend |
| Mouse anti-mouse CD45.2 Brilliant Violet 510 (Clone 104) | BioLegend |
| Mouse anti-mouse CD45.2 Brilliant Violet 785 (Clone 104) | BioLegend |
| Hamster anti-mouse CD49a APC (Clone HM α 1) | BioLegend |
| Rat anti-mouse CD49b FITC (Clone DX5) | BioLegend |
| Rat anti-mouse CD49b PE (Clone DX5) | BioLegend |
| Rat anti-mouse CD62L PE (Clone MEL-14) | Thermo Fisher Scientific |
| Rat anti-mouse CD90.2 PerCP/Cyanine5.5 (Clone 53-2.1) | BioLegend |
| Rat anti-mouse CD90.2 PE/Cyanine7 (Clone 53-2.1) | BioLegend |
| Rat anti-mouse CD90.2 Brilliant Violet 510 (Clone 53-2.1) | BioLegend |
| Rat anti-mouse CD103 PE/Cyanine7 (Clone 2E7) | BioLegend |
| Rat anti-mouse CD127 PE/Cyanine7 (Clone A7R34) | BioLegend |
| Rat anti-mouse CD127 BV421 (Clone SB/199) | BD Bioscience |
| Mouse anti-mouse CD135 PerCP-eFluor 710 (Clone A2F10) | Thermo Fisher Scientific |
| Rat anti-mouse CD135 biotin (Clone A2F10) | BioLegend |
| Hamster anti-mouse CD196 (CCR6) PE (Clone 29-2L17) | BioLegend |
| Hamster anti-mouse CD196 (CCR6) APC (Clone 29-2L17) | BioLegend |
| Rat anti-mouse CD218a (IL-18R α) PE (Clone A17071D) | BioLegend |
| Mouse anti-human CD271(NGFR) PerCP-Cy5.5 (Clone C40-1457) | BD Bioscience |

| | |
|--|--------------------------|
| Mouse anti-human CD271 (NGFR) V450 (Clone C40-1457) | BD Bioscience |
| Hamster anti-human/mouse/rat CD278 (ICOS) Brilliant Violet 421 (Clone C398.4A) | BioLegend |
| Rat anti-mouse CD279 PE/Cyanine7 (Clone 29F.1A12) | BioLegend |
| Rat anti-mouse CD279 PerCP/Cyanine5.5 Antibody (Clone 29F.1A12) | BioLegend |
| Mouse anti-mouse CD279 APC/Cyanine7 (Clone EH12.2H7) | BioLegend |
| Rat anti-mouse CD335 FITC (Clone 29A1.4) | BioLegend |
| Mouse anti-mouse/human GATA-3 eFluor 660 (Clone TWAJ) | Thermo Fisher Scientific |
| Mouse anti-mouse/human GATA-3 PE/Cyanine7 (Clone TWAJ) | Thermo Fisher Scientific |
| Mouse anti-mouse GATA3 Alexa Fluor 647 (Clone L50-823) | BD Bioscience |
| Rat anti-mouse I-A/I-E PerCP/Cyanine5.5 (Clone M5/114.15.2) | BioLegend |
| Rat anti-mouse Ly6A/E (Sca-1) PE (Clone E13-161-7) | BD Bioscience |
| Rat anti-mouse Ly6A/E (Sca-1) APC (Clone E13-161-7) | BioLegend |
| Rat anti-mouse Ly-6G/Ly-6C FITC (Clone RB6-8C5) | BioLegend |
| Rat anti-mouse Ly-6G FITC (Clone 1A8) | BioLegend |
| Rat anti-mouse Ly-6G APC (Clone 1A8) | BioLegend |
| Rat anti-mouse LPAM-1 PE (Clone DATK32) | BioLegend |
| Rat anti-mouse LPAM-1 APC (Clone DATK32) | BioLegend |
| Rat anti-mouse LPAM-1 PerCP-eFluor 710 (Clone DATK32) | Thermo Fisher Scientific |
| Hamster anti-mouse/human KLRG1 (MAFA) PE (Clone 2F1/KLRG1) | BioLegend |
| Hamster anti-mouse/human KLRG1 (MAFA) PE/Cyanine7 (Clone 2F1/KLRG1) | BioLegend |
| Rat anti-mouse/human IL-5 PE (Clone TRFK5) | BioLegend |
| Rat anti-mouse IL-13 eFluor 660 (Clone eBio13A) | Thermo Fisher Scientific |
| Rat anti-mouse IL-17RB Alexa Fluor 647 (Clone 9B10) | BioLegend |
| Rat anti-mouse IL-17RB PE (Clone 9B10) | BioLegend |
| Rat anti-mouse IL-33R α (IL1RL1, ST2) PE/Cyanine7 (Clone DIH9) | BioLegend |
| Rat anti-mouse IL-33R α (IL1RL1, ST2) APC (Clone DIH9) | BioLegend |
| Rat anti-mouse IL-33R α (IL1RL1, ST2) Brilliant Violet 421 (Clone DIH9) | BioLegend |
| Hamster anti-mouse PLZF Antibody PE (Clone 9E12) | BioLegend |
| Mouse anti-mouse NK-1.1 (Clone PK136) | BioLegend |
| Mouse anti-mouse ROR γ t PE (Clone Q31-378) | BD Bioscience |
| Mouse anti-mouse ROR γ t BV421 (Clone Q31-378) | BD Bioscience |
| Mouse anti-mouse ROR γ t BV510 (Clone Q31-378) | BD Bioscience |
| Rat anti-mouse Siglec-F PE (Clone E50-2440) | BD Bioscience |
| Mouse anti-mouse/human T-bet PE (Clone 4B10) | BioLegend |
| Mouse anti-mouse/human T-bet PE/Cyanine7 (Clone 4B10) | BioLegend |
| Hamster anti-mouse TCR β chain FITC (Clone H57-597) | BioLegend |
| Hamster anti-mouse TCR γ/δ PE (Clone UC7-13D5) | BioLegend |
| Rat anti-mouse TER-119 FITC (Clone TER-119) | BioLegend |
| TotalSeq-A0302 anti-mouse Hashtag 2 | BioLegend |
| TotalSeq-A0303 anti-mouse Hashtag 3 | BioLegend |
| TotalSeq-A0307 anti-mouse Hashtag 7 | BioLegend |
| TotalSeq-A0308 anti-mouse Hashtag 8 | BioLegend |
| TotalSeq-A0309 anti-mouse Hashtag 9 | BioLegend |
| TotalSeq-A0310 anti-mouse Hashtag 10 | BioLegend |
| TotalSeq-A0311 anti-mouse Hashtag 11 | BioLegend |
| TotalSeq-A0312 anti-mouse Hashtag 12 | BioLegend |
| Anti-mouse CD3 ϵ antibody (Clone 145-2C11) | BioXCell |
| Anti-mouse CD28 antibody (Clone 37.51) | BioXCell |
| Anti-mouse IFN γ antibody (Clone XMG1.2) | BioXCell |
| Anti-mouse IL-4 antibody (Clone 11B11) | BioXCell |
| Anti-H3K27ac antibody (Clone ab4729) | abcam |