Supplementary Materials

Supplementary figures



Supplementary figure 1. Modulation in cytokine production in DSS treated *II33^{-/-}* mice. Quantification of indicated cytokines in the colon explants of WT and *II33^{-/-}* mice at indicated days post DSS administration. Similar data was obtained with the sera (not shown). Data represent two independent experiments and analyzed by Mann-Whitney U test. Error bars represent mean±S.E.M with 10 mice per group per time point.

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Supplementary figure 2. Levels of IL-33 and IL-1α are increased in the colon after DSS administration. WT and *II*33^{-/-} mice were treated with DSS in drinking water for 6 days, followed by drinking water for 2 days. (A) Western blot analysis for IL-33 in the colon lysates. (B) qRT-PCR for *II*33 expression in the colon tissue. (C) qRT-PCR analysis of *II*33 and *II*1α expression from indicated cell populations from the epithelial (Ep) and colonic lamina propria (cLP) fractions of WT mice at day 8. (D) Western blot analysis for caspase-1 in colon lysates. (E) H&E staining at 40x original magnification of colon sections at day 4 post DSS. Data represent two

independent experiments and analyzed by Mann-Whitney U test. Error bars represent
 mean±S.E.M with 5 mice per group per time point.



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42 **Supplementary figure 3. IL-1α promotes colitis and associated cancer. (A)** Survival and **(B)** 43 disease Activity Index of WT and $l/1a^{-l-}$ mice administered 4% DSS in drinking water. **(C)** Body 44 weight change of WT and $l/1a^{-l-}$ mice injected with AOM on day 0 and administered 5 rounds of 45 3.5% DSS in drinking water. **(D)** and **(E)** Colon histology analysis and **(F)** proportion of mice with 46 low- or high-grade epithelial dysplasia at day 108 post AOM injection. **(G)** H&E staining at 10x 47 original magnification of the distal and middle colon sections at day 108 post AOM injection.

48	Data re	present t	two	independent	experiments	and	analyzed	by	two-way	ANOVA	followed	by
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- Holm-Sidak post test (**B** and **C**) or Mann-Whitney U test (**D** and **E**). Error bars represent mean±S.E.M.

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Supplementary figure 4. Modulation of goblet cells during DSS administration in *II33⁻¹⁻* mice. (A) Quantification of expression of mucins and goblet cell associated genes in the colons of WT and *II33⁻¹⁻* mice at indicated days post DSS administration by qRT-PCR. (B) Quantification of the number of goblet cells per crypt. (C) PAS staining at 10x original magnification of colon sections. Data was analyzed by Kruskal-Wallis test followed by Dunn's post test. Error bars represent mean±S.E.M. N= 8 each for (A), N=5 each for (B) and (C). Each dot represents an individual crypt.



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103 **Supplementary figure 5. ILC2 analysis in the colons of WT and** *II33^{-/-}* **mice. (A)** 104 Quantification of expression of type2 cytokines in the colons of WT and *II33^{-/-}* mice at indicated 105 days post DSS administration by qRT-PCR. **(B)** Gating strategy for ILC2 cells, where Lin⁻ 106 represents CD3⁻B220⁻Ly-6C⁻Ly-6G⁻CD11b⁻Ter119⁻NKp46⁻. Proportion and total number of **(C)** 107 ILC2s and **(D)** T_H2 cells in the lamina propria. Data represent two independent experiments and 108 analyzed by Kruskal-Wallis test followed by Dunn's post test. Error bars represent mean±S.E.M. 109 with 10 mice per group per time point.

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Supplementary figure 6. There is no defect in induction of anti-inflammatory and epithelium healing genes or T regs in the colons of *II33^{-/-}* mice. (A-C) Quantification of expression of indicated genes in the colons of WT and *II33^{-/-}* deficient mice at indicated days post DSS administration by qRT-PCR. (D) Quantification of T regulatory cells gated as CD19⁻ CD3⁺CD4⁺Foxp3⁺ cells in the lamina propria of the colon by flow cytometry. Data represent two independent experiments and analyzed by Kruskal-Wallis test followed by Dunn's post test. Error bars represent mean±S.E.M with 5 mice per group per time point.

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Supplementary figure 7. Increasing Akkermansia level in WT mice increases colitis susceptibility. (A) Q-PCR analysis of indicated bacteria from stool samples of WT and //33^{-/-} mice 1-week administration of water or broad-spectrum antibiotic cocktail (125 mg/l ciprofloxacin, 1 g/l bacitracin, 2 g/l streptomycin, 1.5 g/l metronidazole and 172 mg/l gentamycin) in their drinking water. (B) IL-1 α measurement in colon explants at day 4 post DSS administration (C) Body weight loss and (D) disease activity index of mice during DSS administration. (E) Colon length measurements and (F) representative colon images at day 8 post DSS administration. Data represent two independent experiments and analyzed by Kruskal-Wallis test (A) (B) and (D) or two-way ANOVA (C) followed by Dunn's post test. Error bars represent mean±S.E.M. and each dot represents an individual mouse. N=10 mice for WT and water, *I*/33^{-/-} and water group, 5 for WT and antibiotics group.



Supplementary figure 8. I/33^{-/-} littermates have decreased intestinal IgA, increased level of Akkermansia and susceptibility to colitis. (A) IgA measurement by ELISA and (B) gRT-PCR analysis of indicated bacteria from stool samples of WT, 1/33+/- and 1/33-/- mice 4 weeks after separation. (C) Body weight loss of mice during DSS administration. (D) Colon length measurements and (E) representative colon images at day 8 post DSS administration. Data is analyzed by Kruskal-Wallis test (A) (B) and (D) or two-way ANOVA (C) followed by Dunn's post test. Error bars represent mean±S.E.M. and each dot represents an individual mouse. N=11 mice for WT, 14 for $I/33^{+/-}$ and 6 for $I/33^{-/-}$ group.

182 Supplementary Methods

183 Preparation of single cell suspension from colon.

184 Single cell suspension was prepared from the colon as described previously (1). Briefly, 185 for removal of epithelial cells, the colon was washed, cut into small pieces, and then the 186 pieces were incubated with calcium- and magnesium-free HBSS supplemented with 5% 187 FBS and 5 mM EDTA (Sigma-Aldrich) at 140 rpm at 25°C for 30 min. The tissues were 188 then incubated with RPMI 1640 containing 10% FBS and 0.5 mg/ml collagenase type IV 189 for 1 hour at 37°C with shaking at 150 rpm. The liberated cells were collected by 190 passage through a 70 µm nylon mesh. The isolated cells from the EDTA (epithelial) and 191 collagenase (lamina propria) treated fractions were separated on a 40/80% 192 discontinuous Percoll gradient (GE Bioscience). The following monoclonal antibodies 193 were used in appropriate combinations: anti-CD3 (clone 145 – 2C11), anti-CD4 (clone 194 RM4-5), anti-CD19 (clone 1D3), anti-IgA (clone RMA-1), anti-Foxp3 (FJK-16s), anti-195 CD326 (clone G8.8), anti-CD16/CD32 (clone 93), anti-CD127 (clone A7R34), anti-196 GATA-3 (clone L50-823), anti-NKp46 (clone 29A1.4), anti-CD45.2 (clone 104), anti-197 CD90.2 (clone 30-H12), anti-MHCII (clone M5/114.152) and anti-mouse Lin cocktail 198 (BioLegend, Catalog #133306). For intracellular cytokine staining, cells were fixed and 199 permeabilized using fixation and permeabilization solution (eBioscience, Catalog # 00-200 5523). For intracellular cytokine staining, cells were fixed and permeabilized using 201 fixation and permeabilization solution (eBioscience). Intracellular staining for the Foxp3 202 and GATA-3 transcription factors were performed using the eBioscience Foxp3 staining 203 set according to the manufacturer's recommendations. Flow cytometry data were 204 acquired on LSRII (BD) and were analyzed with FlowJo software (TreeStar).

205 Western blotting

Proteins were extracted from colon tissues using RIPA lysis buffer supplemented proteinase and phosphatase inhibitors (Roche). Samples were resolved in 12-15% SDS-PAGE and transferred onto PVDF membranes. Blocking was performed in 5% milk for 1 hout and membranes were incubated in primary antibodies overnight at 4°C. Membranes were incubated with HRP-conjugated secondary antibody for 1 hour and proteins were visualized using ECL substrate (ThermoScientific). The primary antibodies were Caspase-1 p10 (1:500 dilution, sc-515, Santa Cruz biotechnology) GAPDH (1:10,00 dilution, clone D16H11, Cell Signaling) and IL-

- 213 33 (1:1000 dilution, Catalog # AF3626 R&D systems).
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Gene	Primer Sequences
Gapdh	Forward: 5'-CGTCCCGTAGACAAAATGGT-3'
	Reverse: 5'-TTGATGGCAACAATC TCC AC-3'
β-actin	Forward: 5'-GGCTGTATTCCC CTCCATCG-3'
	Reverse: 5'-CCAGTTGGTAACAATGCCATG T-3'
ll1α	Forward: 5'-AAAATCTCAGATTCACAACTGTTCGT-3'
	Reverse: 5'-TGGCAACTCCTTCAGCAACAC-3'
<i>II5</i>	Forward: 5'-GCAATGAGACGATGAGGCTT-3'
	Reverse: 5'-CCCACGGACAGTTTGATTCT-3'
ll13	Forward: 5'-TGTGTCTCTCCCTCTGACCC-3'
	Reverse: 5'-CACACTCCATACCATGCTGC-3'
Muc1	Forward: 5'-GCAGTCCTCAGTGGCACCTC-3'
	Reverse: 5'-CACCGTGGGCTACTGGAGAG-3'
Muc2	Forward: 5'-GCTGACGAGTGGTTGGTGAATG-3'
	Reverse: 5'-GATGAGGTGGCAGACAGGAGAC-3'
Muc3	Forward: 5'-CGTGGTCAACTGCGAGAATGG-3'
	Reverse: 5'-CGGCTCTATCTCTACGCTCTCC-3'
Muc4	Forward: 5'-CAGCAGCCAGTGGGGACAG-3'
	Reverse: 5'-CTCAGACACAGCCAGGGAACTC-3'
ll22	Forward: 5'-AGAACGTCTTCCAGGGTGAA-3'
	Reverse: 5'-CAT CGA CAT AAG TCA GCA CCA G-3'
Reg3β	Forward: 5'-ATGGCTCCTACTGCTATGCC-3'
	Reverse: 5'-GTGTCCTCCAGGCCTCTTT-3'
Reg3γ	Forward: 5'-ATGGCTCCTATTGCTATGC-3'
	Reverse: 5'-GATGTCCTGAGGGCCTCTT-3'
Lcn2	Forward: 5'-ACATTTGTTCCAAGCTCCAGGGC-3'
	Reverse: 5'-CATGGCGAACTGGTTGTAGTCCG-3'
Mptx	Forward: 5'-CCTGTTTCTCTCTGTTCTTTCAGG-3'
	Reverse: 5'-GGCCTTCATACACAGAGTGAAG-3'

Areg	Forward: 5'-GCCATTATGCAGCTGCTTTGGAGC-3'
•	Reverse: 5'-TGTTTTCTTGGGCTTAATCACCT-3'
S100A9	Forward: 5'-GGTGGAAGCACAGTTGGCA-3'
	Reverse: 5'-GTGTCCAGGTCCTCCATGATG-3'
Occludin	Forward: 5'-TTGAAAGTCCACCTCCTTACAGA -3'
	Reverse: 5'-CCGGATAAAAAGAGTACGCTGG-3'
Zo1	Forward: 5'-GCCGCTAAGAGCACAGCAA-3'
	Reverse: 5'-GCCCTCCTTTTAACACATCAGA -3'
Tff3	Forward: 5'-CCTGGTTGCTGGGTCCTCTG-3'
	Reverse: 5'-GCCACGGTTGTTACACTGCTC-3'
plgR	Forward: 5'-AAGAACTGACCAAAGGGAGGA-3'
	Reverse: 5'-AGAGTAACTTCAATTCTGCACCC-3'
Muc5ac	Forward: 5'-CTGTGACATTATCCCATAAGCCC-3'
	Reverse: 5'-AAGGGGTATAGCTGGCCTGA-3'
Eubacteria	Forward: 5'-ACTCCTACGGGAGGCAGCAGT-3'
	Reverse: 5'-ATTACCGCGGCTGCTGGC-3'
Bacteroides	Forward: 5'-GGTTCTGAGAGGAGGTCCC-3'
	Reverse: 5'-CTGCCTCCCGTAGGAGT-3'
Enterobacteriaceae	Forward: 5'-GTGCCAGCMG CCGCGGTAA-3'
	Reverse: 5'-GCCTCAAGGG CACAACCTCC AAG-3'
γ-Proteobacteria	Forward: 5'-TAACGCTTGG GAATCTGCCT RTT-3'
	Reverse: 5'-CATCTRTTAG CGCCAGGCCT TGC-3'
Fungal 18s	Forward: 5'- ATTGGAGGGCAAGTCTGGTG-3'
	Reverse: 5'-CCGATCCCTAGTCGGCATAG-3'
Prevotellacae	Forward: 5'- ATTGGAGGGCAAGTCTGGTG-3'
	Reverse: 5'-CCGATCCCTAGTCGGCATAG-3'
Akkermansia	Forward: 5'-CAGCACGTGAAGGTGGGGAC-3'
	Reverse: 5'-CCTTGCGGTTGGCTTCAGAT-3'
Anaerostipes	Forward: 5'-AAGTCGAACGAAGCACCTTG-3'
	Reverse: 5'-TCCGCCACTCAGTCACAATG-3'
Dorea	Forward: 5'-ACGGTACCTGACTAAGAAGCCC-3'
	Reverse: 5'-CCTCAACGTCAGTCATCGTCC-3'
E. rectale	Forward: 5'-ACTCCTACGGGAGGCAGC-3'
	Reverse: 5'-GCTTCTTAGTCAGGTACCGTCA-3'
Flexispira	Forward: 5'-AATACATGCAAGTCGAACGATGA-3'
	Reverse: 5'-AATCACCGTTTCCAGTGGCT-3'
Clostridia	Forward: 5'-CICAACIIGGGIGCIGCAIII-3'
– <i>v</i>	Reverse: 5'-ATTGTAGTACGTGTGTGTAGCCCC-3'
E. COll	Forward: 5'-CATGCCGCGTGTATGAAGAA-3'
	Reverse: 5'-CGGGTAACGTCAATGAGCAAA-3'
Prevotella	Forward: 5'-CACGGTAAACGATGGATGCC-3'
	Reverse: 5'-GGTUGGGTTGUAGAUU-3'
Lactobacilius	Forward: 5'-GGAAACAGATGCTAATACCG-3'
	Reverse: 5'-CAUCGUTACACATGGAG-3'
MIB	Forward: 5'-CCAGCAGCCGCGCATACTACT
9LB	
Tm7	Reverse. 3-GAUGGUAUGGATTGTTATTC-3
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	Reverse: 5 - GAGAGGATGATCAGCCAG-3

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