



Title	Gut commensals suppress interleukin-2 production through microRNA-200/BCL11B and microRNA-200/ETS-1 axes in lamina propria leukocytes of murine large intestine
Author(s)	Ohsaka, Fumina; Karatsu, Yugo; Kadota, Yoshihiro et al.
Citation	Biochemical and biophysical research communications, 534, 808-814 https://doi.org/10.1016/j.bbrc.2020.10.103
Issue Date	2021-01-01
Doc URL	https://hdl.handle.net/2115/83740
Rights	©2021. This manuscript version is made available under the CC-BY-NC-ND 4.0 license http://creativecommons.org/licenses/by-nc-nd/4.0/
Rights(URL)	https://creativecommons.org/licenses/by-nc-nd/4.0/
Type	journal article
File Information	Ohsaka2020BBRC-fig-R.pdf, 



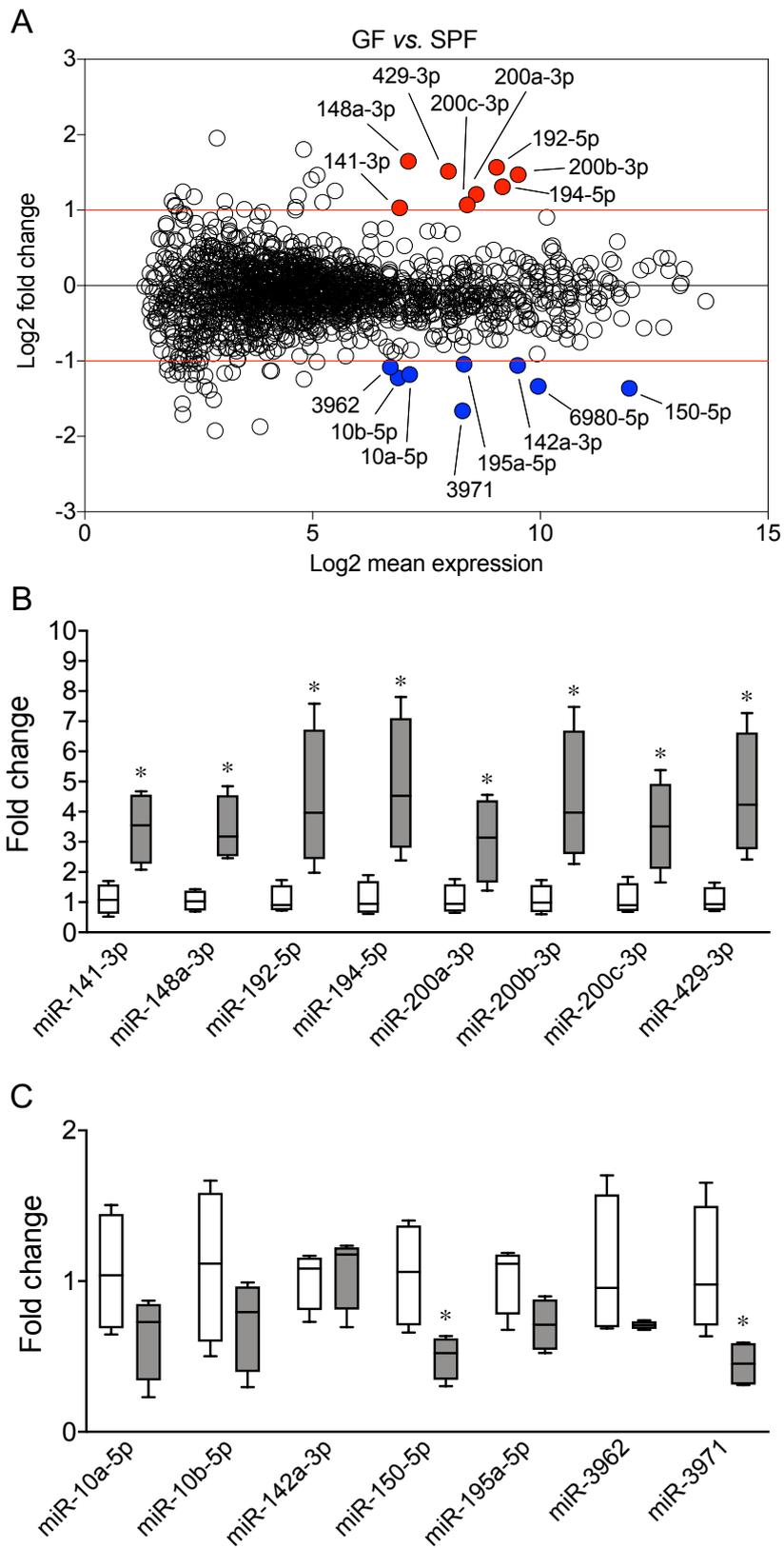


Fig. 1 Ohsaka *et al.*

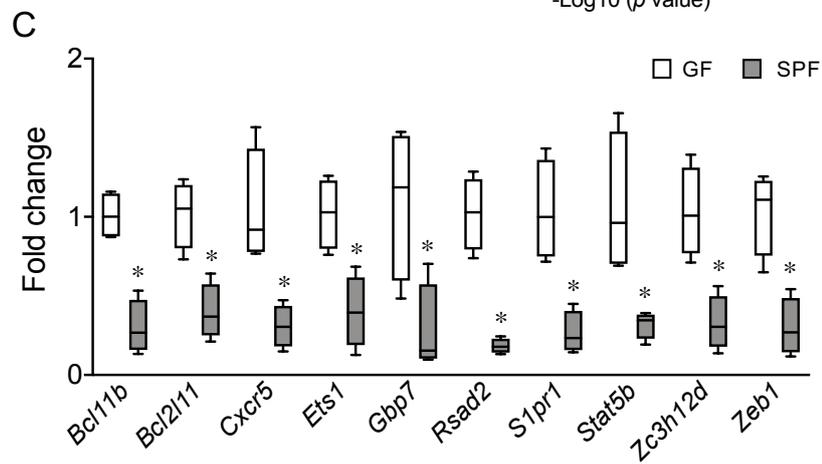
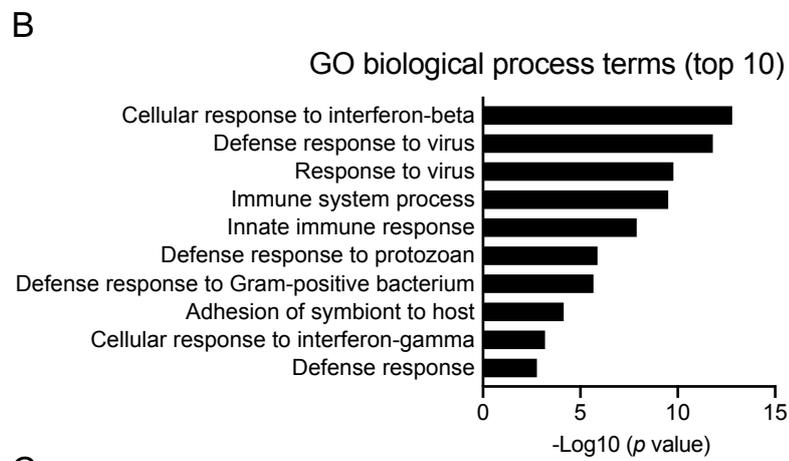
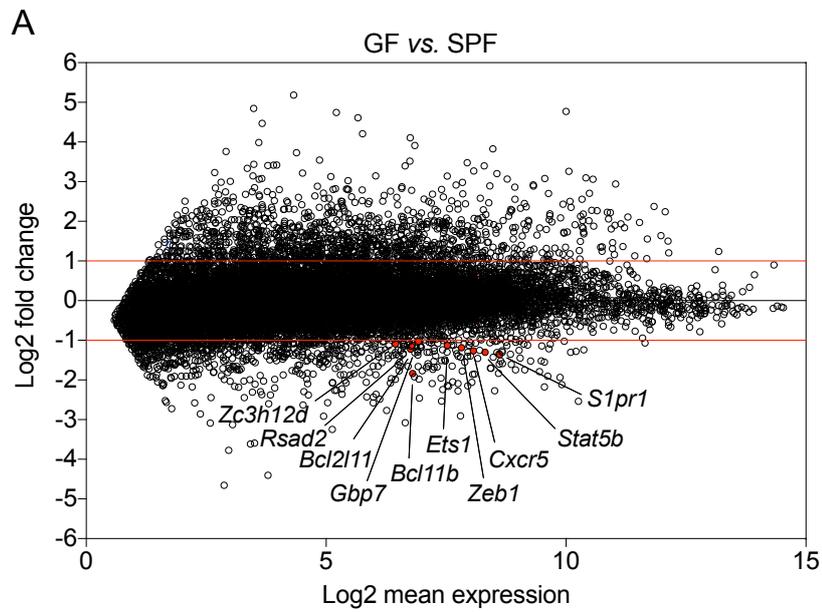


Fig. 2 Ohsaka *et al.*

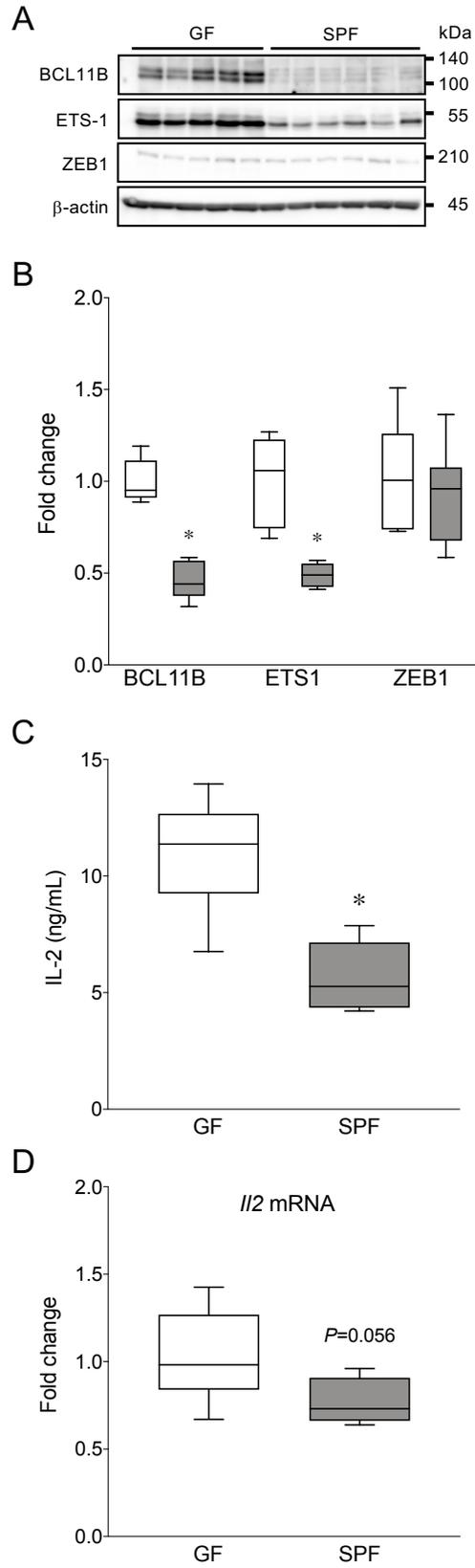


Fig. 3 Ohsaka *et al.*

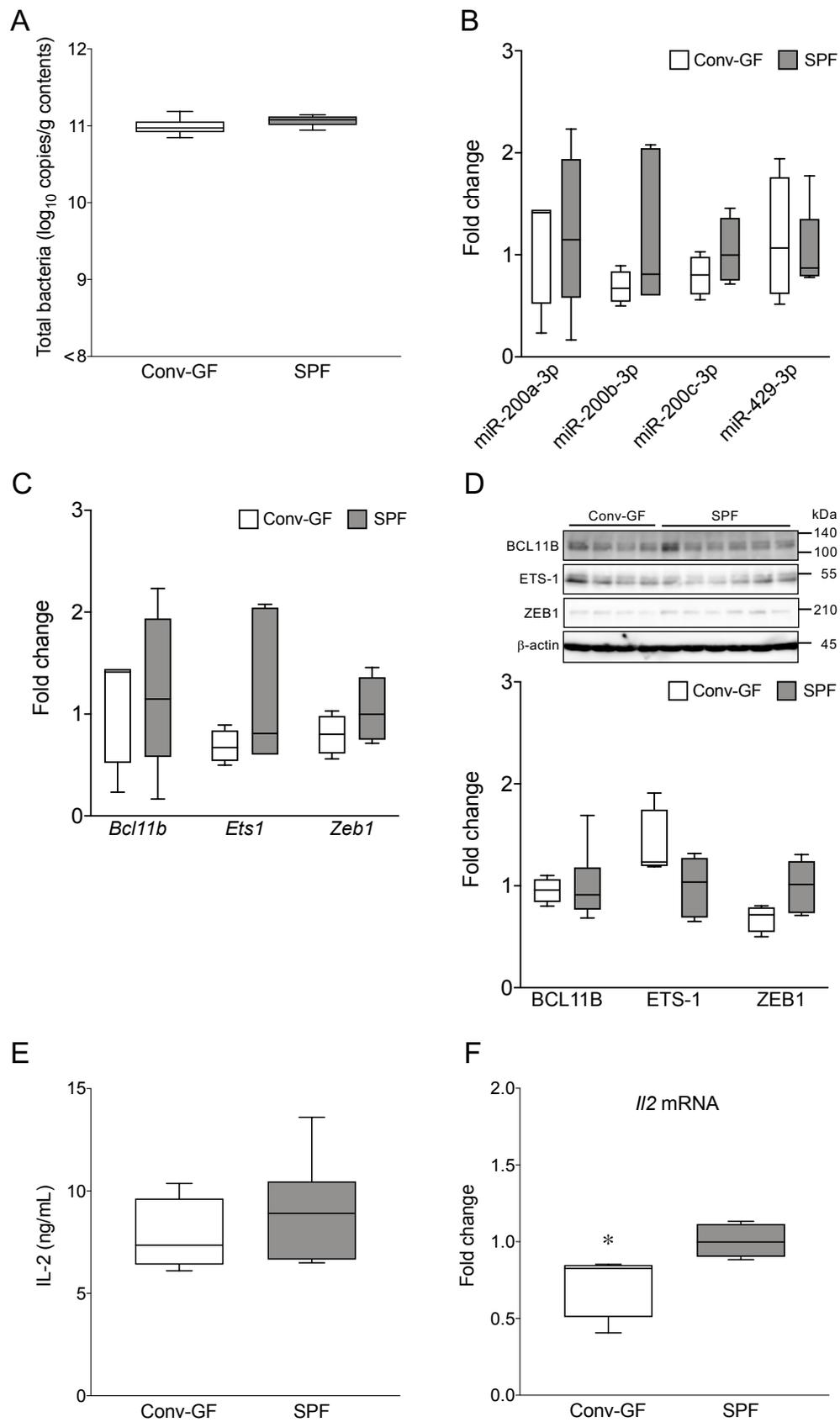


Fig. 4 Ohsaka *et al.*

Supplementary Table 1 Primer sequences for qRT-PCR of miRNA

miRNA	Forward
mmu-miR-10a-5p	TACCCTGTAGATCCGAATTTGTG
mmu-miR-10b-5p	TACCCTGTAGAACCGAATTTGTG
mmu-miR-141-3p	TAACACTGTCTGGTAAAGATGG
mmu-miR-142a-3p	CATAAAGTAGAAAGCACTACT
mmu-miR-148a-3p	TCAGTGCCTACAGAACTTTGT
mmu-miR-150-5p	TCTCCCAACCCTTGACCAGTG
mmu-miR-192-5p	CTGACCTATGAATTGACAGCC
mmu-miR-194-5p	TGTAACAGCAACTCCATGTGGA
mmu-miR-195a-5p	TAGCAGCACAGAAATATTGGC
mmu-miR-200a-3p	TAACACTGTCTGGTAACGATGT
mmu-miR-200b-3p	TAATACTGCCTGGTAATGATGA
mmu-miR-200c-3p	TAATACTGCCGGGTAATGATGGA
mmu-miR-429-3p	TAATACTGTCTGGTAATGCCGT
mmu-miR-3962	AGGTAGTAGTTTGTACATTT
mmu-miR-3971	CTCCCCACCCCTGTACCAGTGA
cel-miR-39-3p	TCACCGGGTGTAATCAGCTTG

Supplementary Table 2 Primer sequences for qRT-PCR of mRNA

Gene	Forward	Reverse
<i>Actb</i>	CTGGGACGATATGGAGAAGA	AGAGGCATACAGGGACAACA
<i>Bcl11b</i>	GGCGATGCCAGAATAGATGCCG	CCAGGCCACTTGGCTCCTCTATCTCCAGAC
<i>Bcl2l1l</i>	AATGTCTGACTCTGATTCTCGGAC	TCTCAGCAGGCTGCAATTGTCCAC
<i>Cxcr5</i>	GACTCCTTACCACAGTGCACCTT	GGAAACGGGAGGTGAACCA
<i>Ets1</i>	CGATCCAGCTGTGGCAGTT	ATCTCCTGGCCACCTCATCT
<i>Gbp7</i>	TTGAGGAAATGCCAGAGGACCAGT	GTCTCCACTATTGATAGCATCCACG
<i>Il2</i>	TGATGGACCTACAGGAGCTCCTGAG	GAGTCAAATCCAGAACATGCCGCAG
<i>Rsad2</i>	TGCTGGCTGAGAATAGCATTAGG	GCTGAGTGCTGTTCCCATCT
<i>Slpr1</i>	ATGGTGTCCACTAGCATCCC	CGATGTTCAACTTGCTGTGTAG
<i>Stat5b</i>	GGACTCCGTCCTTGATACCG	TCCATCGTGTCTTCCAGATCG
<i>Zc3h12d</i>	TGGCAGCAATGTGGCTATGA	AGCACGTGTTGCTCTCTGAT
<i>Zeb1</i>	GCTGGCAAGACAACGTGAAAG	GCCTCAGGATAAATGACGGC

Supplementary Table 3 Primary antibodies for western blot analysis

Target	Cat#	Source	Species raised in; clonality	Dilution	RRID*
BCL11B	55414-1-AP	Proteintech (Rosemont, IL)	Rabbit; poly	1:300	AB_11182609
ETS-1	sc-55581	Santa Cruz Biotechnology (Santa Cruz, CA)	Mouse; mono	1:300	AB_831289
ZEB1	NBP1-05987	Novus Biologicals (Littleton, CO)	Rabbit; poly	1:1,000	AB_2273178
β -actin	sc-47778	Santa Cruz Biotechnology	Rabbit; poly	1:200	AB_2714189

* RRID, Research Resource Identifier

Supplementary Table 4 Secondary antibodies for western blot analysis

Target	Cat#; label	Source	Species raised in; clonality	Dilution	RRID*
Rabbit IgG	HAF008; HRP	R&D Systems (Minneapolis, MN)	Goat; poly	1:1,000	AB_357235
Mouse IgG	HAF007; HRP	R&D Systems	Goat; poly	1:1,000	AB_357234

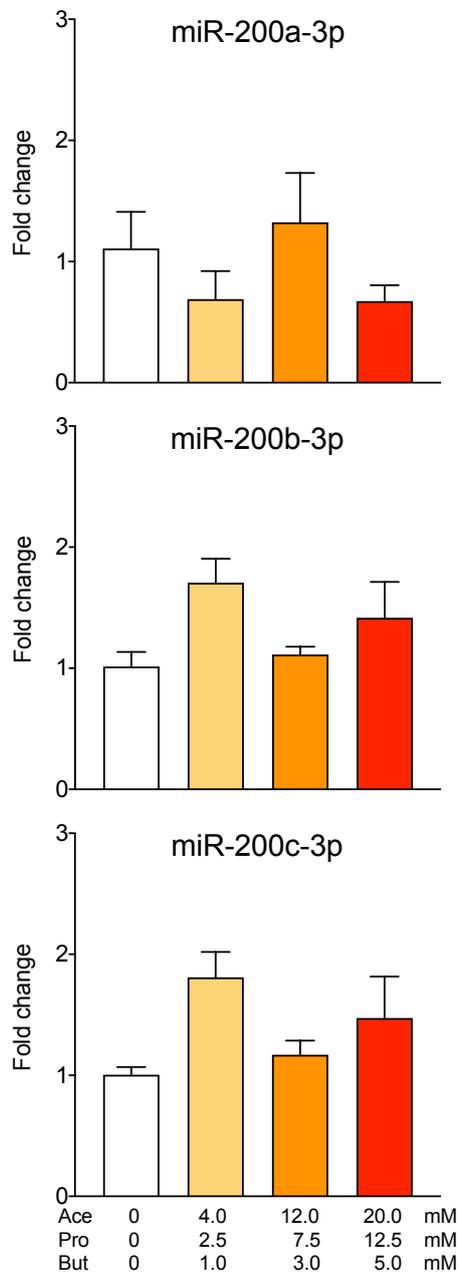
* RRID, Research Resource Identifier

Supplementary Table 5 miRNAs increased by gut microbiota and their predicted target genes

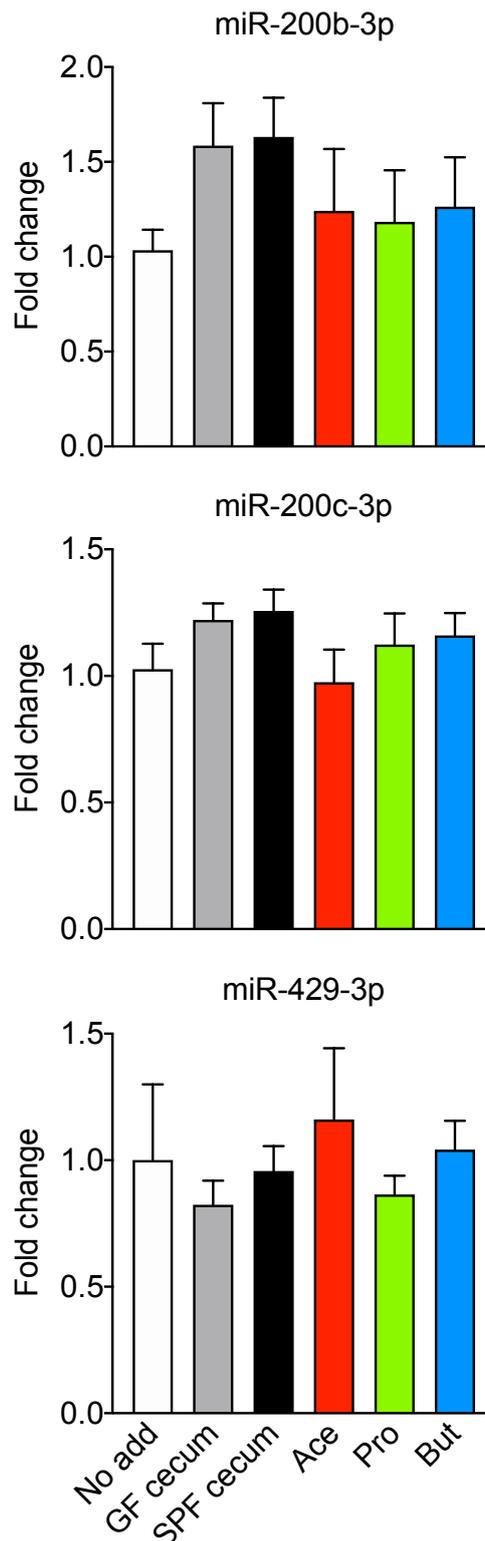
miRNA*	Target gene**			
mmu-miR-148a-3p	<i>Bcl2l1l</i>	<i>Slpr1</i>		
mmu-miR-192-5p	<i>Cxcr5</i>	<i>Rsad2</i>	<i>Slpr1</i>	
mmu-miR-194-5p	<i>Stat5b</i>	<i>Zc3h12d</i>	<i>Zeb1</i>	
mmu-miR-141-3p/200a-3p	<i>Bcl11b</i>	<i>Gbp7</i>	<i>Stat5b</i>	<i>Zeb1</i>
mmu-miR-200b-3p/200c-3p/429-3p	<i>Bcl11b</i>	<i>Ets1</i>	<i>Zeb1</i>	

* These miRNAs show higher levels in SPF mice than in GF mice as determined by qRT-PCR (**Fig. 1B**).

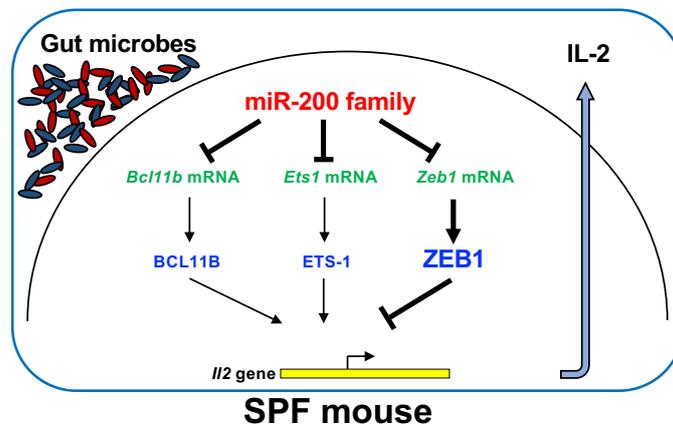
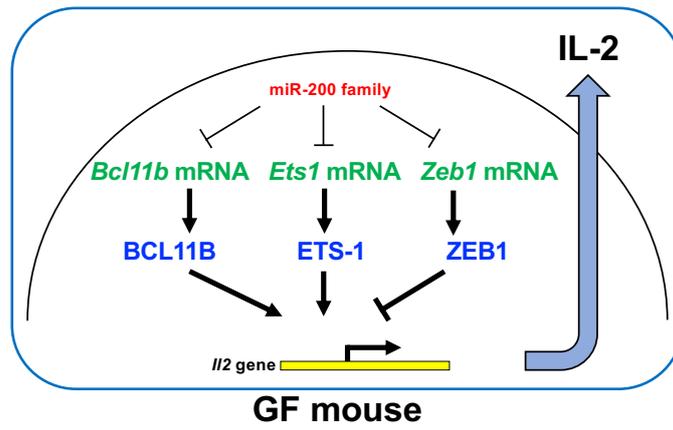
** GO terms of these genes are associated with infection defense and the immune system as shown in **Fig. 2B**, and lower levels of their mRNAs in SPF mice than in GF mice were shown by qRT-PCR (**Fig. 2C**).



Supplementary Figure 1 Expression of miRNAs in large intestinal lamina propria leukocytes cultured in the presence of short-chain fatty acids. The cells were isolated from large intestine of specific-pathogen free BALB/c mice and cultured with a mixture of different concentrations of sodium acetate, sodium propionate and sodium butyrate (Ace, Pro and But, respectively) for 24 h. Relative miRNA levels were determined by qRT-PCR. Data are expressed as means \pm SEM ($n = 3$) and are shown relative to levels in the negative control (white bar), which were set to 1.



Supplementary Figure 2 Expression of miRNAs in large intestinal lamina propria leukocytes cultured in the presence of fecal extracts or short-chain fatty acids. The cells were isolated from large intestine of germ-free BALB/c mice and cultured with fecal extracts (200 μ g protein/mL) prepared from cecal contents of germ-free (GF cecum) or specific-pathogen free (SPF cecum) mice, or 0.1 mM acetate (Ace), propionate (Pro) or butyrate (But) upon stimulation with 25 ng/mL PMA and 1 μ g/mL ionomycin for 24 h. Relative miRNA levels were determined by qRT-PCR. Data are expressed as means \pm SEM ($n = 6$) and are shown relative to levels in the negative control (No add; white bar), which were set to 1.



Graphical abstract