

## SUPPLEMENTAL MATERIAL

Rigoni et al, <http://www.jem.org/cgi/content/full/jem.20151116/DC1>

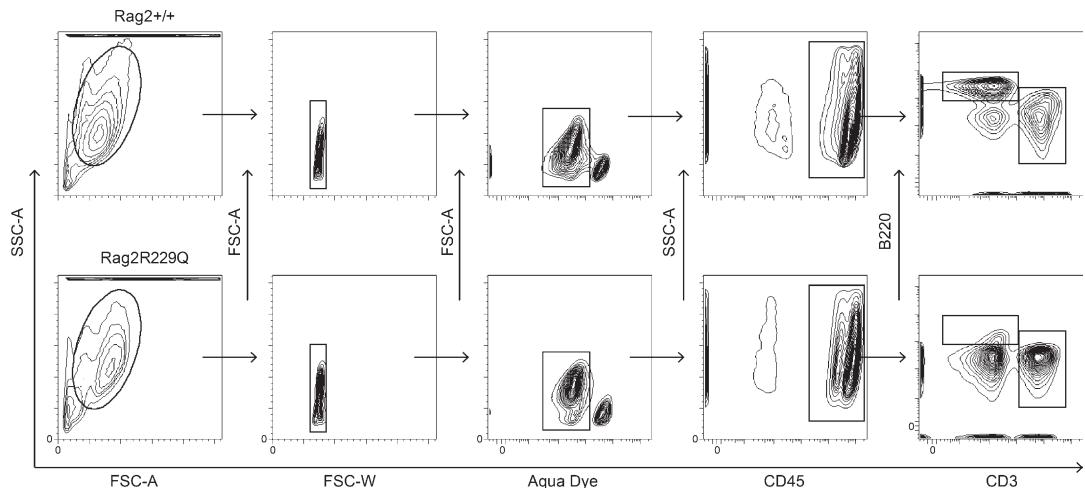
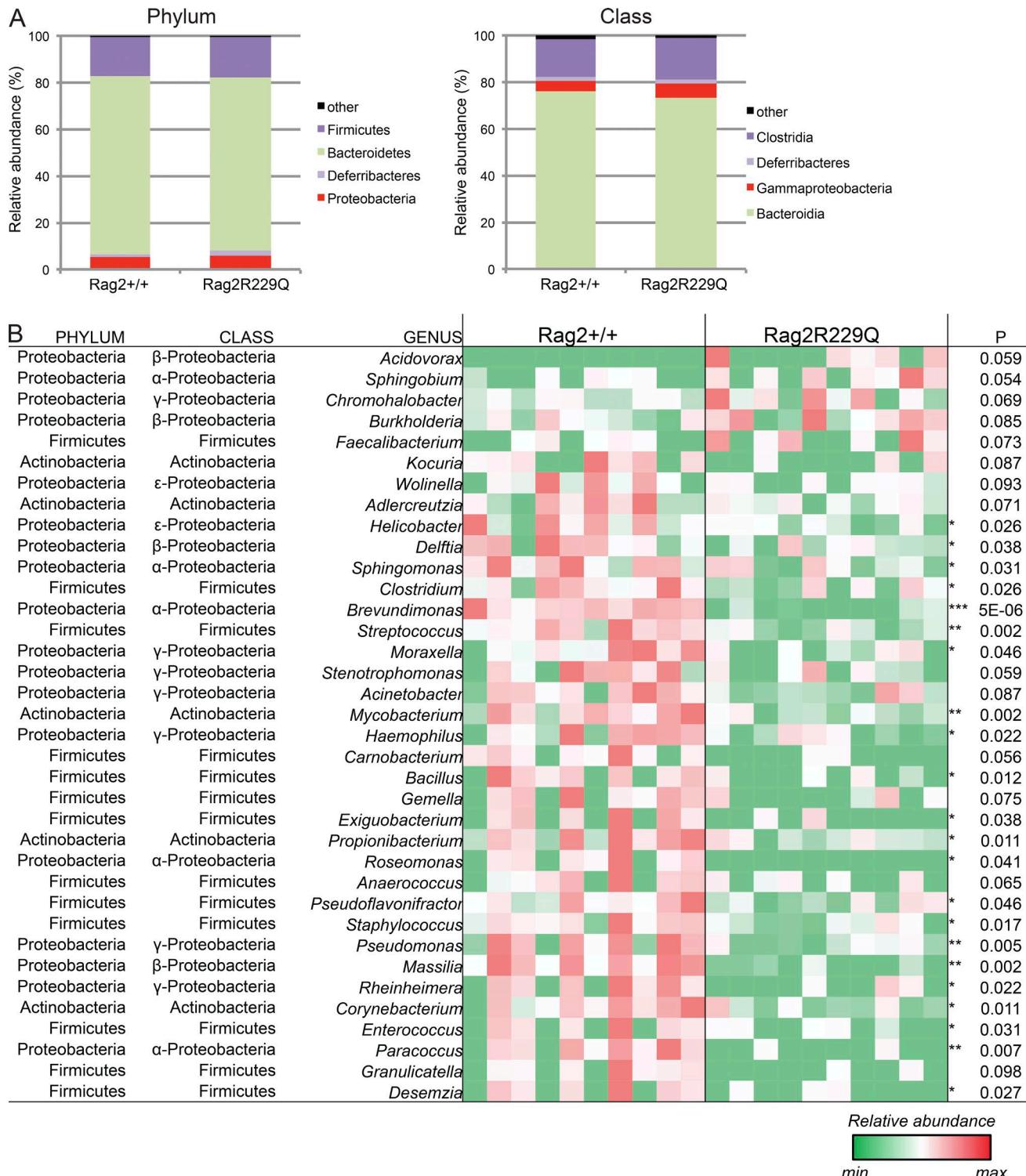


Figure S1. Representative gating strategy to identify  $\text{CD3}^+$  T cells and  $\text{B220}^+$  B cells in the intestinal LP of  $\text{Rag2}^{+/+}$  and  $\text{Rag2}^{R229Q}$ . Doublet exclusion (FSC-W vs. FSC-A) and live cell identification were performed (aqua dye vs. FSC-A). Cells were then gated on a CD45<sup>+</sup> population to identify lymphocytes.



**Figure S2. Analysis of microbiota composition.** (A) Relative abundance of intestinal microbiota at class and phylum level in *Rag2<sup>+/+</sup>* and *Rag2R229Q* mice. (B) Heat map of bacterial genera in the gut microbiota that discriminate *Rag2R229Q* from *Rag2<sup>+/+</sup>* mice with a cutoff of  $P < 0.1$ . Each line represents one genus; each column represents an individual mouse; color represents the difference between *Rag2<sup>+/+</sup>* and *Rag2R229Q* littermates, green indicating lower and red higher expression. P-values are according to two-tailed unpaired Student's t test. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ .

Table S1. List of primers used in this study

Gene	Sequence or assay number
<b>SYBR green primers</b>	
Actb for	5'-CTAAGGCCAACCGTCAAAAG-3'
Actb rev	5'-ACCAAGGGCATACAGGGACA-3'
IL-17A for	5'-TCCAGAAGGCCCTCAGACTA-3'
IL-17A rev	5'-TGAGCTTCCCAGATCACAGA-3'
IFN- $\gamma$ for	5'-TCAAGTGCATAGATGTGAAGAA-3'
IFN- $\gamma$ rev	5'-TGGCTCTGCAGGATTTCATG-3'
IL-22 for	5'-TGACGACCAGAACATCCAGA-3'
IL-22 rev	5'-AATGCCCTTGATCTCTCAC-3'
IL-10 for	5'-GGTTCCAAGCCTTATCGGA-3'
IL-10 rev	5'-ACCTGCTCCACTGCCTTGCT-3'
IL-6 for	5'-TAGCTATGGTACTCCAGAAGAC-3'
IL-6 rev	5'-ACGATGATGCACTTGCAGAA-3'
TNF for	5'-GACGTGAACTGGCAGAAGAG-3'
TNF rev	5'-TTGGTGGTTGTGACTGTGAG-3'
RegIIly for	5'-TTCCTGCTCCATGATCAAAA-3'
RegIIly rev	5'-CATCCACCTCTGTTGGTTCA-3'
IL-1 $\beta$ for	5'-GCCCATCCTCTGTGACTCAT-3'
IL-1 $\beta$ rev	5'-AGGCCACAGTATTGGTCG-3'
CCL2 for	5'-CTGGATCGGAACCAAATGAG-3'
CCL2 rev	5'-AAGGCATCACAGTCGGAGTC-3'
CCL20 for	5'-AACTGGGTGAAAAGGGCTGT-3'
CCL20 rev	5'-GTCCAATTCCATCCCCAAAAA-3'
CXCL10 for	5'-GCTGCCGTCTTTCTGC-3'
CXCL10 rev	5'-TCTCACTGCCCGTCATC-3'
CXCL9 for	5'-CTTTCTCTGGGCATCAT-3'
CXCL9 rev	5'-GCATCGTCATTCTTATCA-3'
CCL5 for	5'-GTGCCACGTCAAGGGTAT-3'
CCL5 rev	5'-CCCACTTCTCTCCTGGGTTG-3'
CCL25 for	5'-GAGTGCACCCTAGGTCACTC-3'
CCL25 rev	5'-CCAGCTGGTCTTACTCTGA-3'
16s for	5'-GTGSTGCAYGGYTGTGTCGTA-3'
16s rev	5'-ACGTCRTCCMCACCTCCTC-3'
18s for	5'-CTCAACACGGAAACCTCCTCAC-3'
18s rev	5'-CGCTCCACCAACTAAGAAGG-3'
IL-4 for	5'-CATCGCATTGAAACGAG-3'
IL-4 rev	5'-CGAGCTCACTCTCTGGTG-3'
IL-23p19 for	5'-AGCCAGTTCTGCTTCAAAGG-3'
IL-23p19 rev	5'-GGAGTTGTGAAGTTGCTCCATG-3'
TGF- $\beta$ for	5'-TGGACCAACATGTGGAACTC-3'
TGF- $\beta$ rev	5'-GTCACGCGGGTTACCA-3'
IL-12p35 for	5'-CTATCTGAGCTCCGCCCTGAAG-3'
IL-12p35 rev	5'-GGCCAAGACCACCTGACTCTTA-3'
<b>TaqMan probes</b>	
Actb	Mm00607939_s1
Rorc	Mm01261022_m1
Tbx21	Mm00450960_m1