

Figure S1. Flow cytometry analysis of T- and B-cell populations in spleen and MLN. On day 31, the spleens and MLNs were collected and cell suspensions were obtained. Representative CD4⁺T-cell (**A**), CD8⁺ T-cell (**B**) and B-cell (**C**) plots are shown. Percentages of regulatory T-cells (**D**), Th17 (**E**), Th1 (**F**), activated CD8⁺T-cells (**G**) were determined in the CD4⁺ or CD8⁺ lymphocyte populations in the spleen. The percentage of CD19⁺B220⁺ B-cells (**H**) in the spleen is also shown. Respectively, T- and B-cell populations for MLN are shown (**I-M**). Data are represented as mean ± SEM of sham (*n* = 3) and vaccinated mice (*n* = 9). Significant differences are show as * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001.



Figure S2. Relative mRNA expression in the colon. Relative mRNA expression of galectin-3 (**A**), -4 (**B**), -9 (**C**), ROR γ (**D**), TGF β 1 (**E**) and TGF β 3 (**F**) were measured in the colon using RT-qPCR. TGF β 1/ROR γ ratio (**G**) and TGF β 3/ROR γ ratio (**H**) were calculated to represent regulatory/Th17 balance in the colon. Data are represented as mean ± SEM of sham (*n* = 3) and vaccinated mice (*n* = 9). Significant differences are show as ** *p* < 0.01.



Figure S3. Ratio mRNA expression and serum galectins. Relative mRNA expression of galectin-3 (**A**), -4 (**B**), -9 (**C**), ROR γ (**D**), TGF β 1 (**E**) and TGF β 3 (**F**) were measured in the ileum using RT-qPCR. TGF β /ROR γ ratios were calculated to represent regulatory/Th17 balance in the colon. Data are represented as mean ± SEM of sham (*n* = 3) and vaccinated mice (*n* = 9). Significant differences are show as * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001.



Figure S4. Ratio galectin-4/galectin-3 mRNA in ileum and correlations of RORγ/TGFβ3 and serum galectins. The ratio of galectin-4/galectin-3 mRNA expression (**A**) is shown. Additionally, RORγ/TGFβ3 ratio in ileum were correlated to serum galectin3/galectin-9 ratio (**B**) and to serum galectin-4/galectin-9 ratio (**C**) using Spearman correlation. Data are represented as mean ± SEM of sham (n = 3) and vaccinated mice (n = 9).