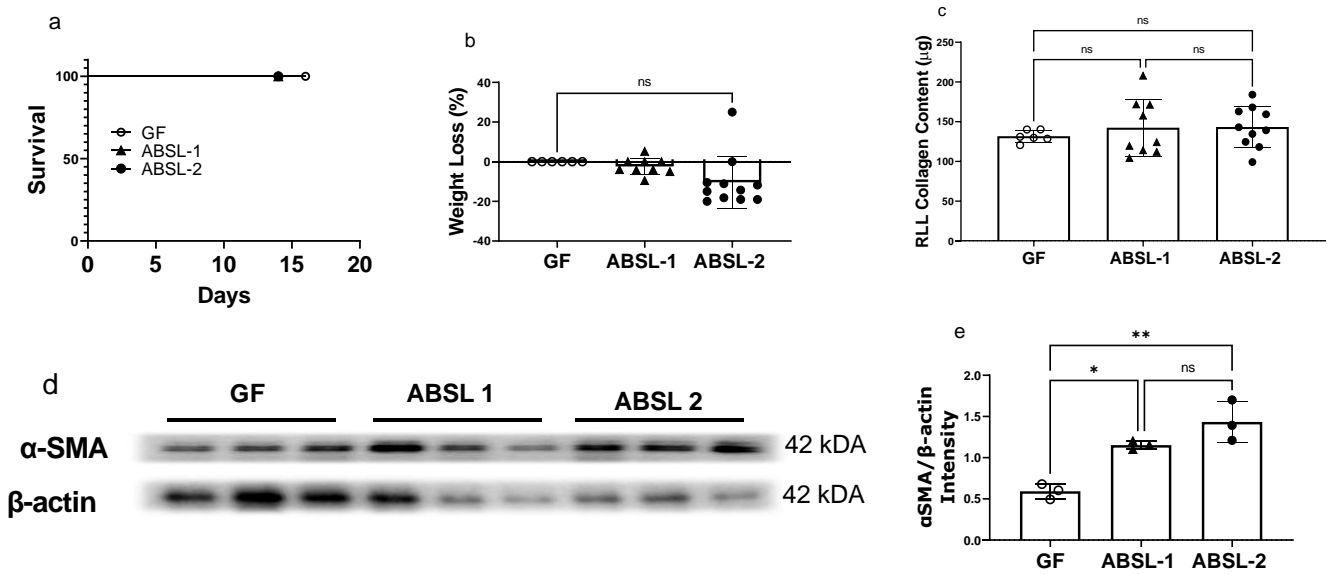


Supplementary Figures

Gut microbiota modulates lung fibrosis severity following acute lung injury in mice

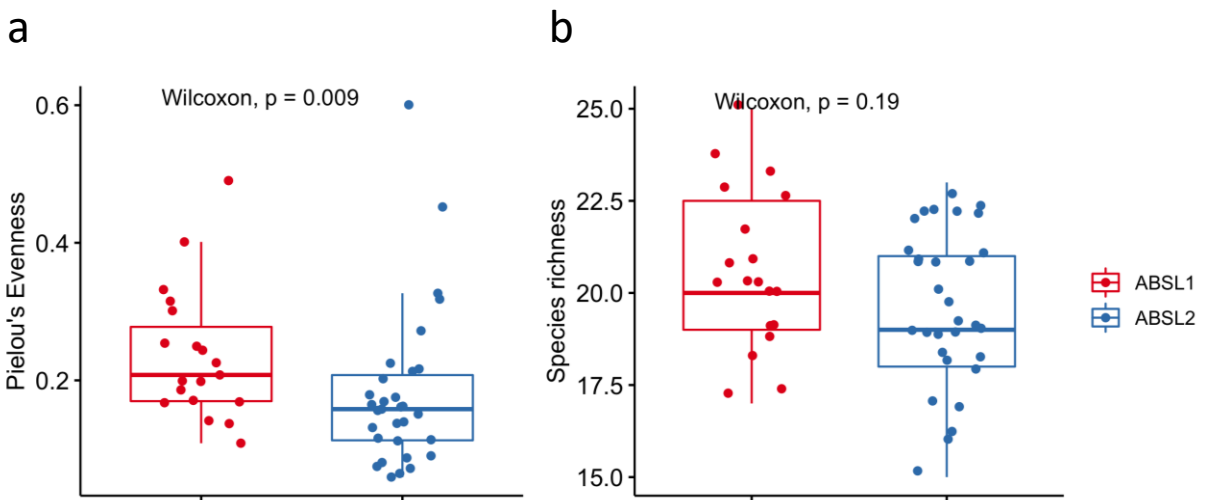
Ozioma S. Chioma¹, Elizabeth K. Mallott, Austin Chapman¹, Joseph C. Van Amburg¹, Hongmei Wu¹, Binal Shah-Gandhi¹, Nandita Dey¹, Marina E. Kirkland¹, M. Blanca Piazuelo⁵; Joyce Johnson⁴, Gordon R. Bernard⁶, Sobha R. Bodduluri⁷, Steven Davison⁸, Bodduluri Haribabu⁷, Seth R. Bordenstein^{2, 3, 4}, Wonder P. Drake^{*1, 4}

Supplemental Fig. 1: Housing conditions does not impact morbidity and mortality during saline treatment.



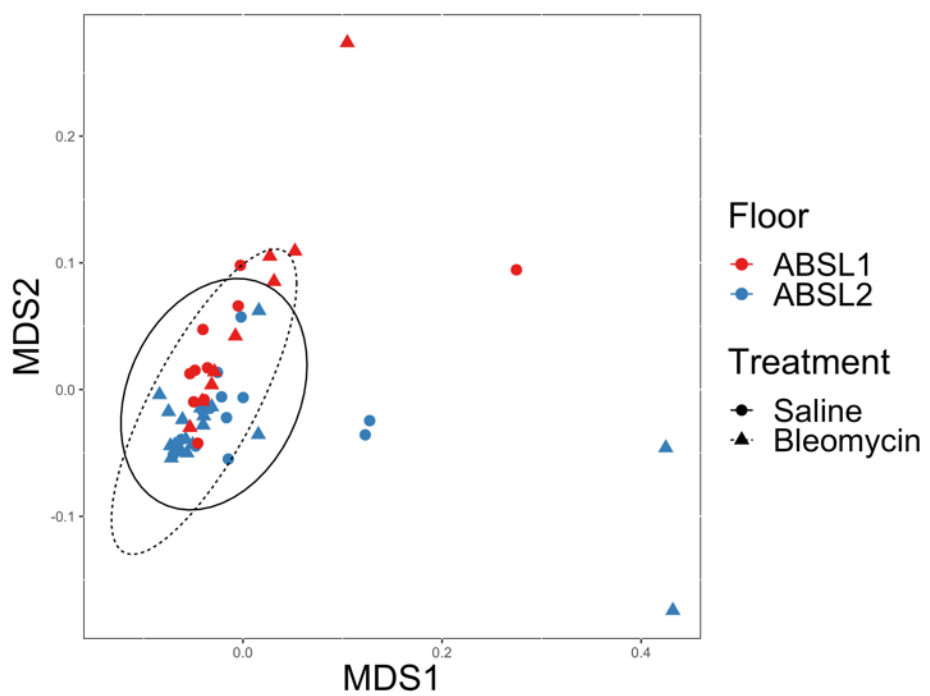
(a) Survival curve of mice housed in 3 separate facilities and intranasally inoculated with saline. **(b)** Weight loss of mice in the saline treatment cohorts; **(c)** Sircol assay of mice housed in 3 separate facilities and treated with saline (control group); **(d)** immunoblot of mice lung samples with antibodies against alpha smooth muscle actin and beta actin in bleomycin treated cohort; **(e)** quantification of immunoblot analysis; Data are reported as mean \pm SD with each dot representing an individual mouse. Statistical significance was assessed using a one-way ANOVA with Sidak's multiple comparison's test for b, c; subpanel e was analyzed using Bonferroni's test due to an n=3 for each group. *P<0.05 **P < 0.01, ****P < 0.0001; ns = no significance; RLL: right lower lobe. n= 3-11 mice

Supplementary Fig. 2: Species evenness contributes to alpha diversity in ABSL-1 housed mice.



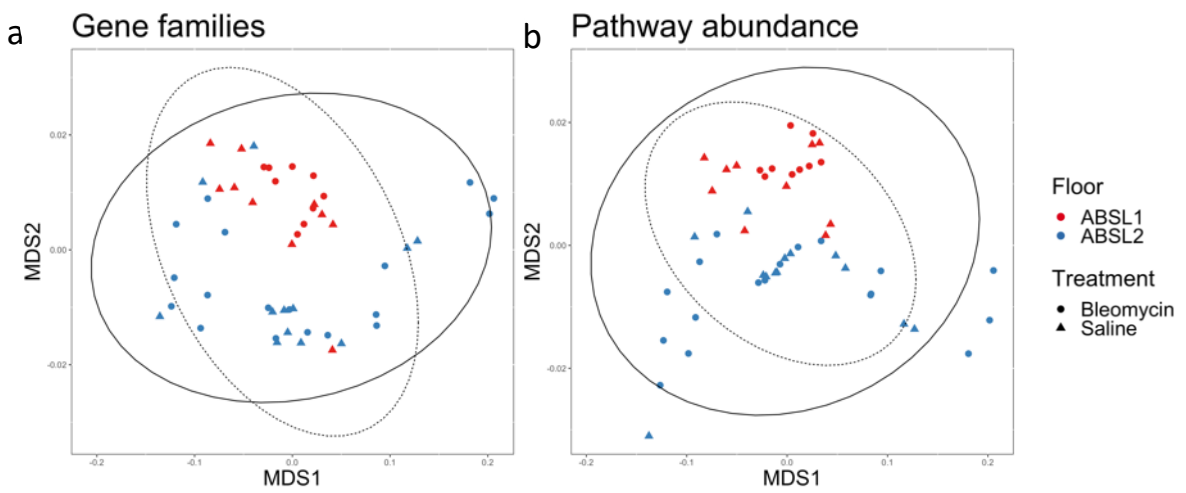
(a) Pielou's evenness and **(b)** species richness were assessed in ABSL-1 and ABSL-2 reared mice. Each dot represents an individual mouse, and median and interquartile range are indicated in the figure.

Supplemental Fig. 3: Nonlinear multidimensional scaling (MDS) plot showing differences in microbial taxonomic composition.



Nonlinear multidimensional scaling (MDS) plot showing differences in microbial taxonomic composition of the gut microbiome based on Bray-Curtis index in ABSL1 or ABSL2 mice). $n = 13-27$ mice per cohort. The experiment was independently conducted once.

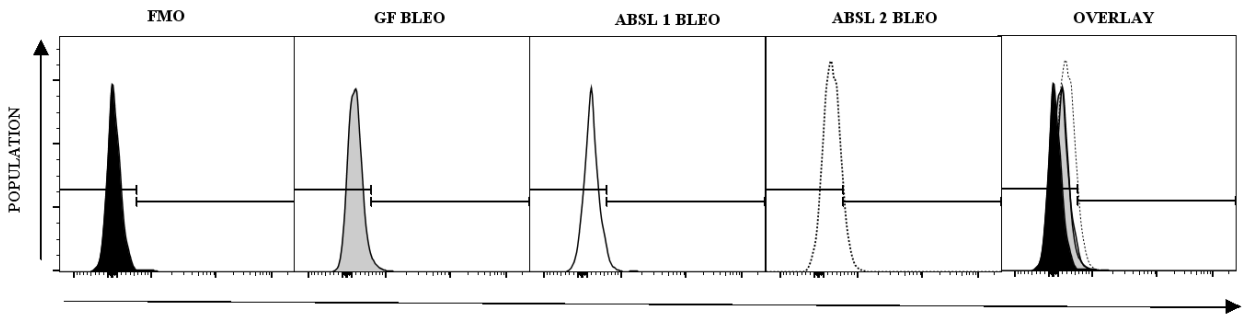
Supplemental Fig. 4: Functional profiles are not different between floor or treatment when using Bray-Curtis dissimilarity matrices.



(a) Gene family abundance and **(b)** MetaCyc reaction pathway abundance profiles based on Bray-Curtis dissimilarity matrices.

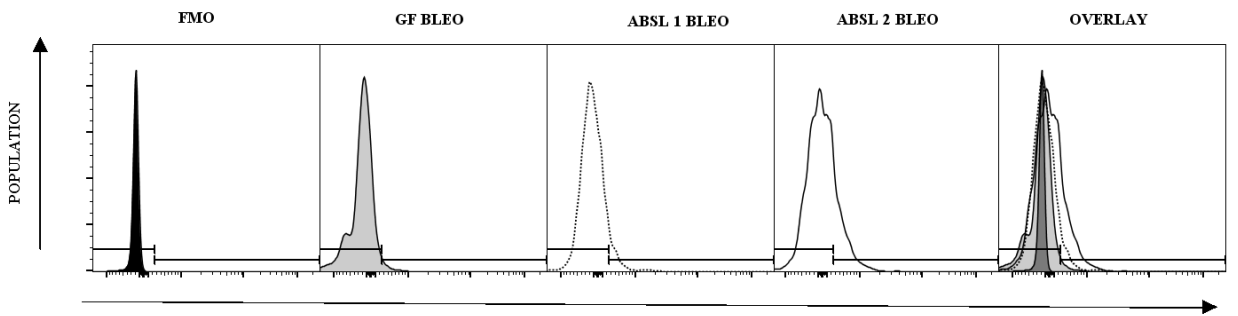
Supplementary Fig. 5a-d: Representative histogram depicting IL-6 expression on CD4+ T cells in GF, ABSL-1 and ABSL-2 bleomycin treated mice.

a. CD4⁺ IL-6⁺ T Cells (%)



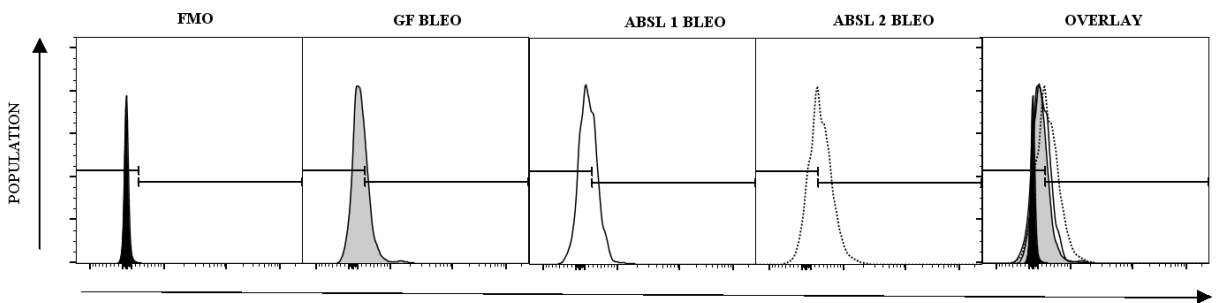
b. CD4⁺ GP130⁺ T Cells (%)

CD4⁺ IL6⁺ T CELLS



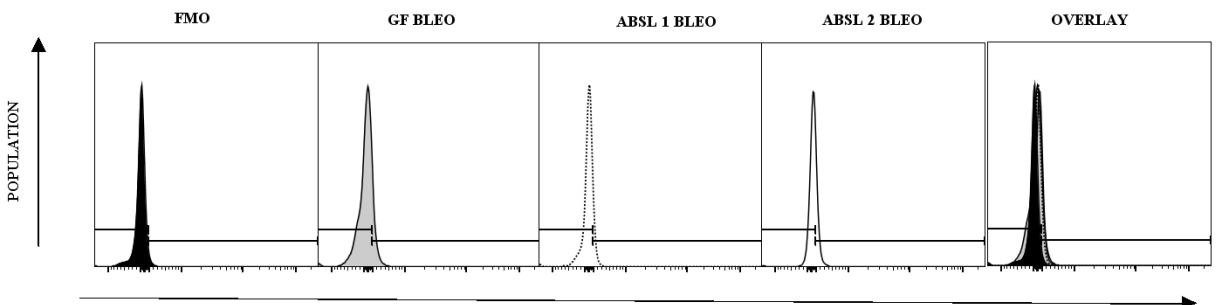
c. CD4⁺ PD-1⁺ IL-17A⁺ T Cells (%)

CD4⁺ GP130⁺ T CELLS



d. CD4⁺ pSTAT3⁺ T Cells (%)

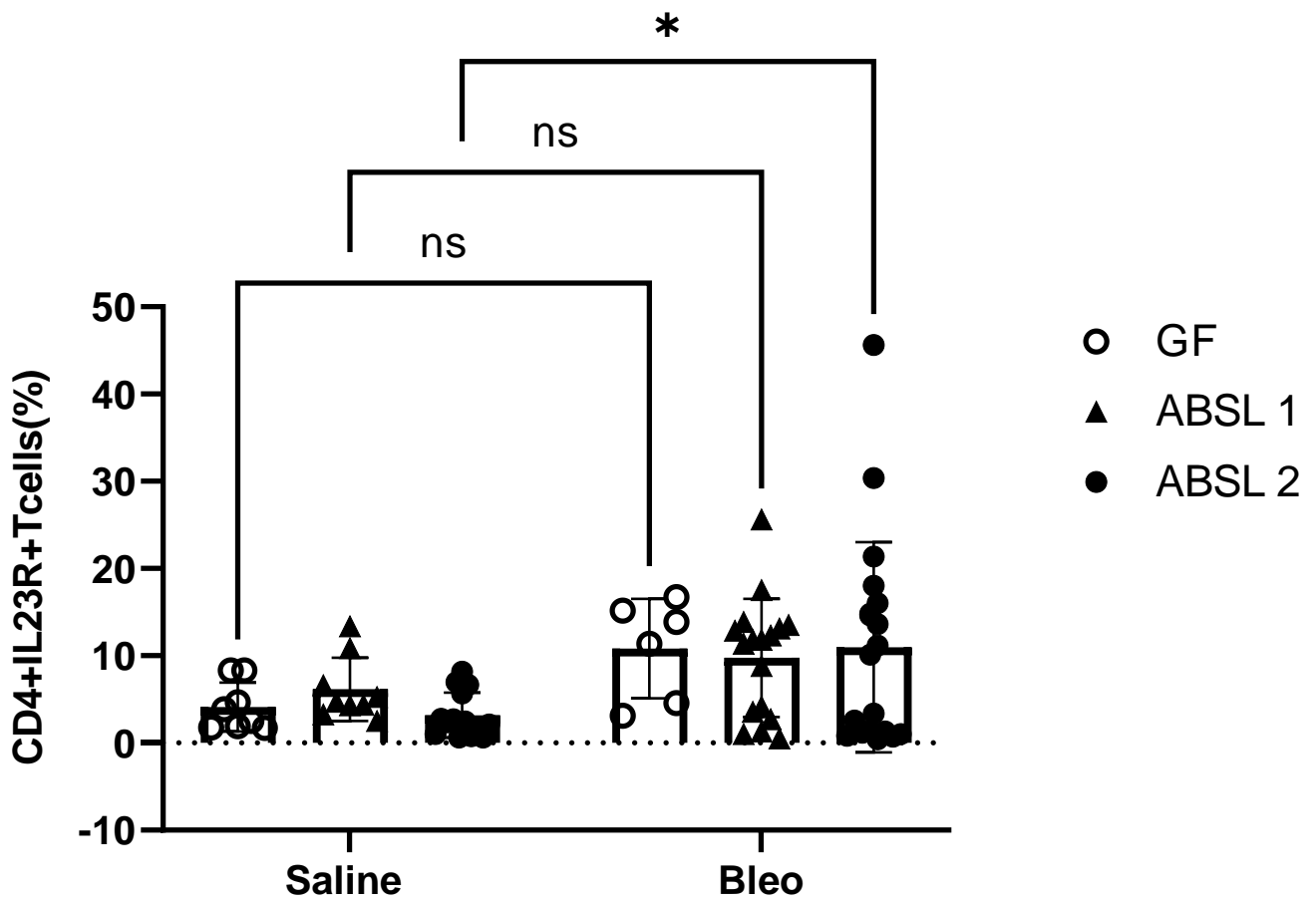
CD4⁺ IL17A⁺ PD1⁺ T CELLS



CD4⁺ pSTAT3⁺ T CELLS

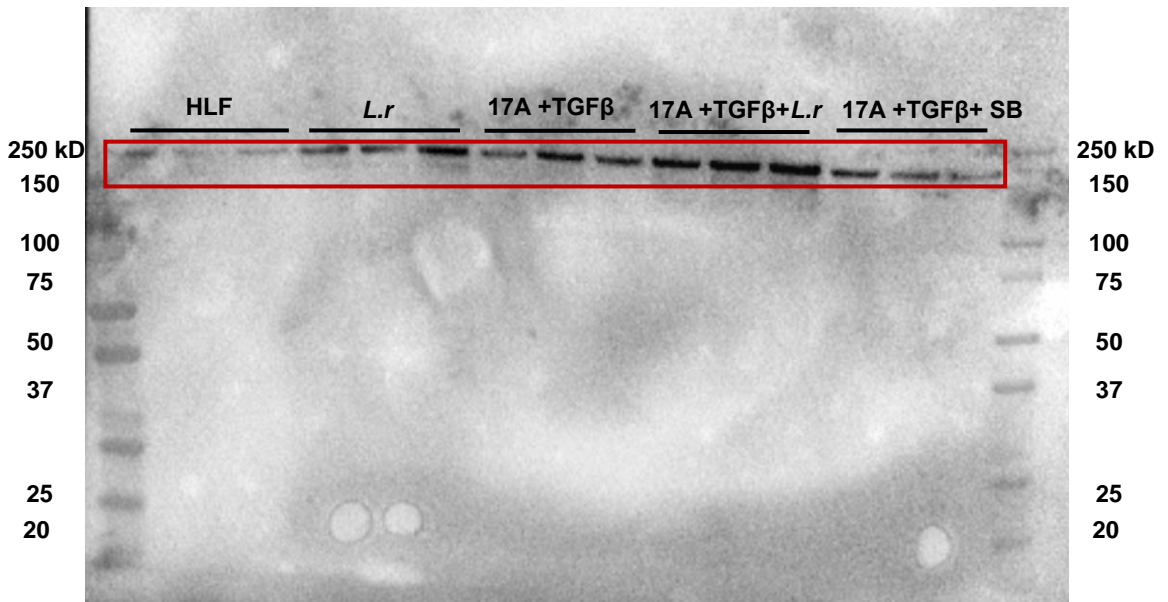
Representative histogram depicting (a) IL-6A+, (b) GP130, (c) PD-1+ IL-17A+, and (d) pSTAT3 expression on CD4+ T cells in GF mice, mice housed in ABSL 1 and ABSL2 Facilities. GF: Germ-free, ABSL-1: Animal Biosafety Level 1, ABSL-2: Animal Biosafety Level 2

Supplemental Fig. 6: Gut microbial diversity and taxa associate with inflammatory pulmonary lung responses.



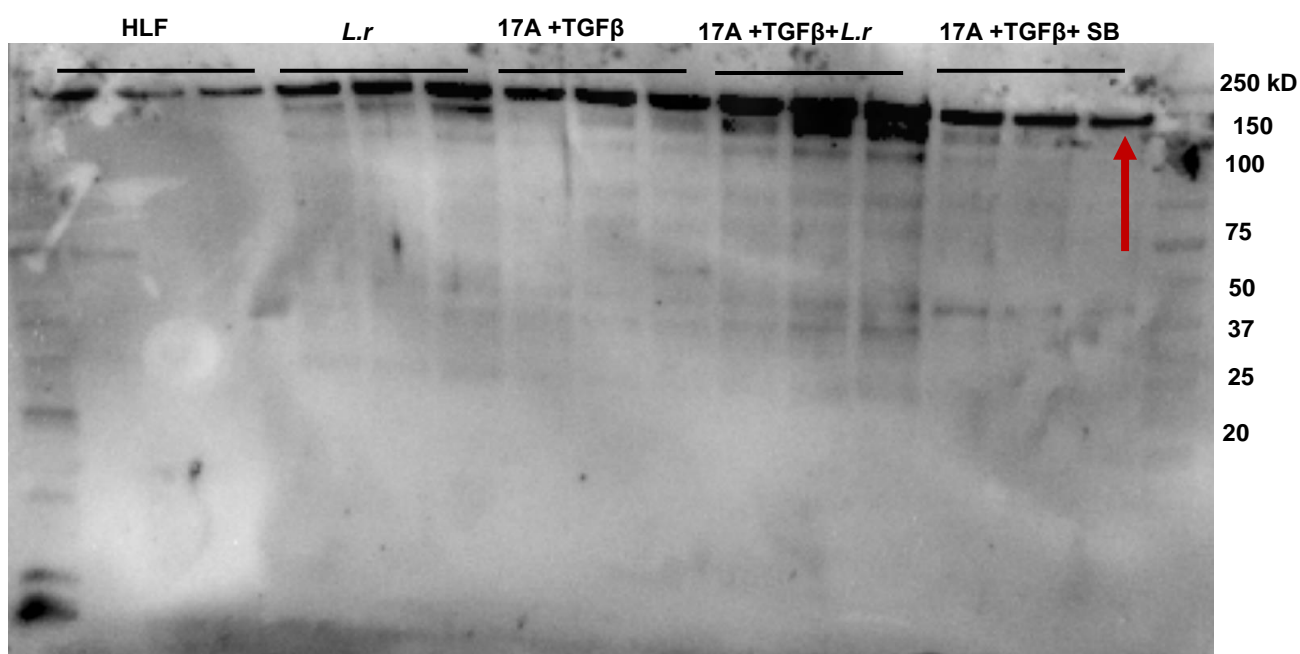
Single cell lung suspensions were obtained from mice housed in three different facilities (GF, ABSL-1 and ABSL-2). Flow cytometry analysis was conducted for IL-23R expression on CD4+T cells. Statistical significance was assessed using a two-way ANOVA with Tukey's multiple comparison's test. *P<0.05, **P < 0.01, ****P< 0.0001, ns = no significance n=6-19.

Supplemental Fig. 7: Western Blot image of Collagen-1 alpha (COL1A)



Immunoblot of Human Lung Fibroblast (HLF) assayed for collagen type 1A (col1a) production after HLF's are treated with rIL-17A and TGFβ1, supernatant from *Lactobacillus* grown to log phase and rIL-17A + TGFβ1 and sodium butyrate. Collagen-1 alpha (COL1A) shown at 210 kDA

Supplemental Fig. 8: Western Blot image of GAPDH



Immunoblot of Human Lung Fibroblast (HLF) assayed for GAPDH production after HLF's are treated with rIL-17A and TGFβ1, supernatant from *Lactobacillus rhamnosus* grown to log phase and rIL-17A + TGFβ1 and sodium butyrate. GAPDH Tetramer shown at 146 kDA.

Supplemental Table 1: Antibodies used for flow cytometric analysis of murine single cell suspensions.

Antibody	Vendor Catalog No	Host organism; antibody type	Clone	Conc (µg)/volume (µL) per test*
CD3 Alexa Fluor 700	Biolegend 100216	Rat Monoclonal	17A2	0.25 µg/µL
CD4 APC-Cy7	Biolegend 100414	Mouse Monoclonal	GK1.5	0.1 µg/µL
IL-6 APC	Biolegend 504808	Rat Monoclonal	MP5-20F3	0.04 µg/µL
GP-130 PE	Biolegend 149404	Rat Monoclonal	4H1B35	0.02 µg/µL
PD-1 PE-Cy7	Biolegend 135216	Rat Monoclonal	29F.1A12	0.1 µg/µL
IL-17A PE	Biolegend 506904	Rat Monoclonal	TCII-18H10.1	0.1 µg/µL
STAT3 (pY703) Pacific Blue	BD 560312	Mouse Monoclonal	4/P-STAT3	10 µg/µL
IL-23R APC	Biolegend 150906	Rat Monoclonal	12B2B64	0.02 µg/µL
*Test = 1 million cells in 50µl of staining buffer				

Supplemental Table 2: Microbiome comparison of parent and gavaged murine fecal pellets.

Group 1	Group 2	Bray-Curtis distance (mean±stddev)	Jaccard distance (mean±stddev)
Stool of ABSL1 mice	Stool of ABSL2 mice	0.158±0.116	0.301±0.088
Stool of germ-free mice gavaged with ABSL1 stool	Stool of ABSL1 mice	0.980±0.039	0.866±0.138
Stool of germ-free mice gavaged with ABSL2 stool	Stool of ABSL2 mice	0.942±0.101	0.739±0.144
Stool of germ-free mice gavaged with ABSL1 stool	Stool of germ-free mice gavaged with ABSL2 stool	0.890±0.136	0.772±0.239
Stool of germ-free mice gavaged with ABSL1 stool	Stool of ABSL2 mice	0.982±0.033	0.861±0.143
Stool of germ-free mice gavaged with ABSL2 stool	Stool of ABSL1 mice	0.965±0.056	0.755±0.139

Average Bray-Curtis dissimilarities and Jaccard distances between ABSL1 stool, stool from ABSL1-gavaged germ-free mice, ABSL2 stool, and stool from ABSL2-gavaged germ-free mice.