

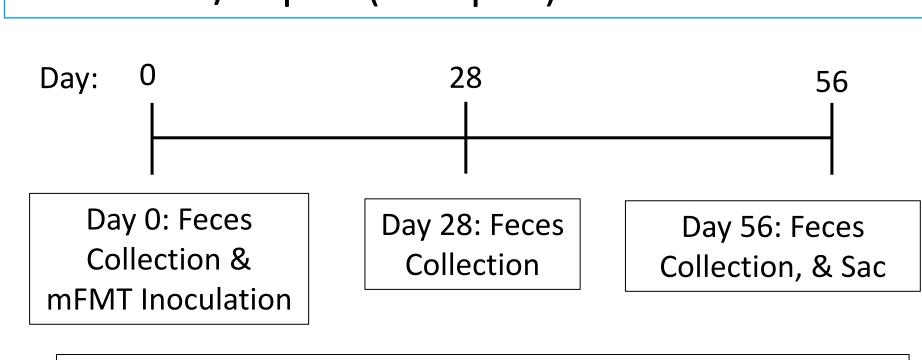
Validation of a Microbiome Dependent Colitis Model via Fecal Microbial Transplant in Germ-Free IL-10 Deficient Mice

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ABSTRACT

Background: Inflammatory Bowel Diseases (IBD) are immune-mediated intestinal tract diseases. Though specific etiologies remain undefined, proposed pathogenic mechanisms include abnormal inflammatory response to the constitutive intestinal microbiome. Mice deficient in IL-10, a critical cytokine for mucosal immune homeostasis, develop enterocolitis; spontaneously phenotypes are microbiome sensitive.

dependent colitis model with improved construct validity and translational clinical relevance, we assessed colitis development in female Germ-Free IL-10 knockout mice (GF IL-10 KO; Taconic Biosciences) following inoculation with fecal microbial transplant (FMT) from wild-type C57BL/6NTac mice at the Murine Pathogen Free™ (MPF™) health standard and assessed clinical responses to anti-IL-12/23p40 (anti-p40).

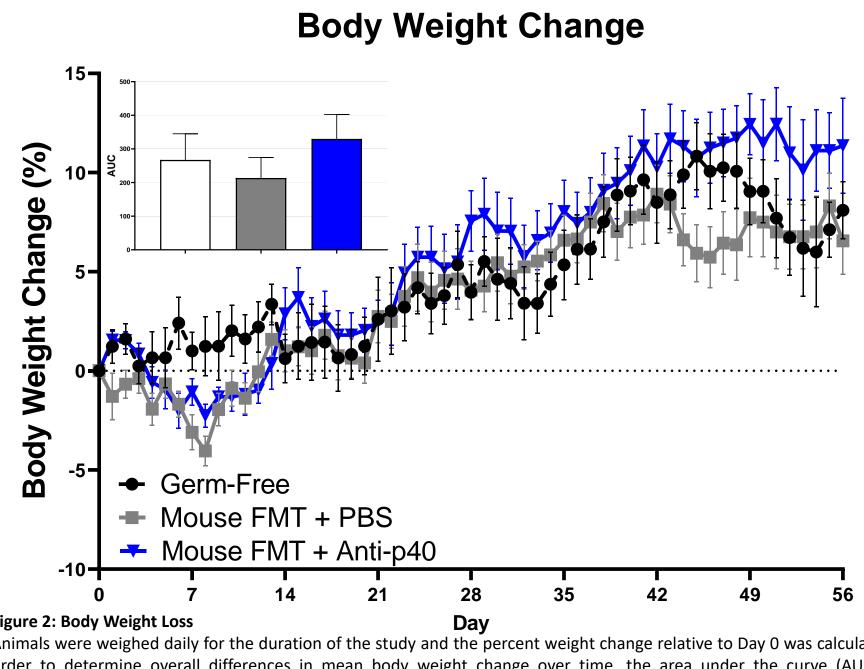


Q3D, Days 0-56: Vehicle or Anti-p40 IP

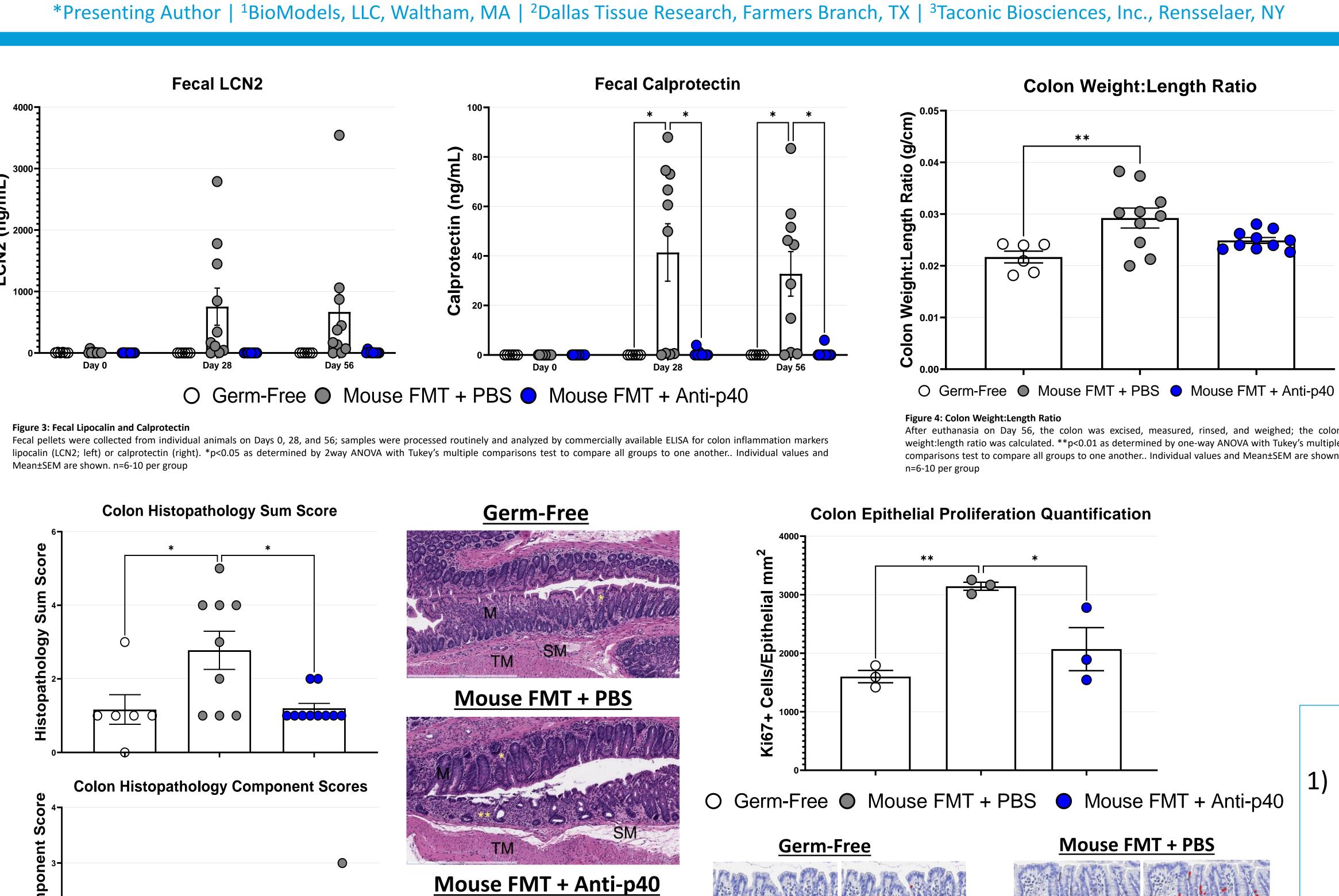


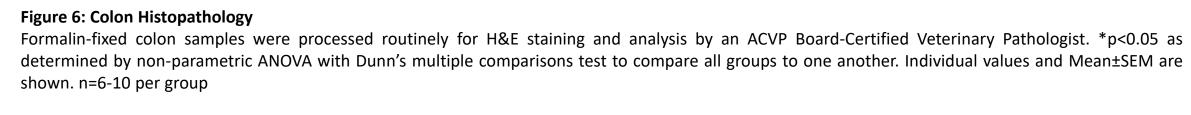
Taconic model #16006, II10 Knockout on C57BL/6 strain background at Germ Free health standard

Figure 1: Study Design Germ-free IL-10 Knockout Mice were inoculated with FMT from wild type C57BL/6NTac mice at the MPF™ health standard on Day 0; a group of mice were maintained germ-free for study duration as naïve controls. FMT-inoculated animals were treated every three days with vehicle or reference compound via intraperitoneal (IP) administration. Animals were assessed for survival and weight loss, as well as presence/severity of diarrhea and bloody stool, on a daily basis. Feces were collected from all animals on Days 28 and 56. At euthanasia on Day 56, colon was excised, measured, rinsed, weighed, and stored for downstream analysis. Feces was processed for ELISA analysis of colitis markers lipocalin 2 (LCN2) and calprotectin. Formalin-fixed colon samples were processed routinely and analyzed for inflammation, gland loss, erosion, and hyperplasia (H&E). Samples were also analyzed for T cells and epithelial proliferation via IHC. Flash-frozen colon samples were processed for multiplex cytokine analysis (Luminex).



calculated using the trapezoidal rule transformation and is shown in the figure inset. Mean±SEM is shown. n=6-10 per group

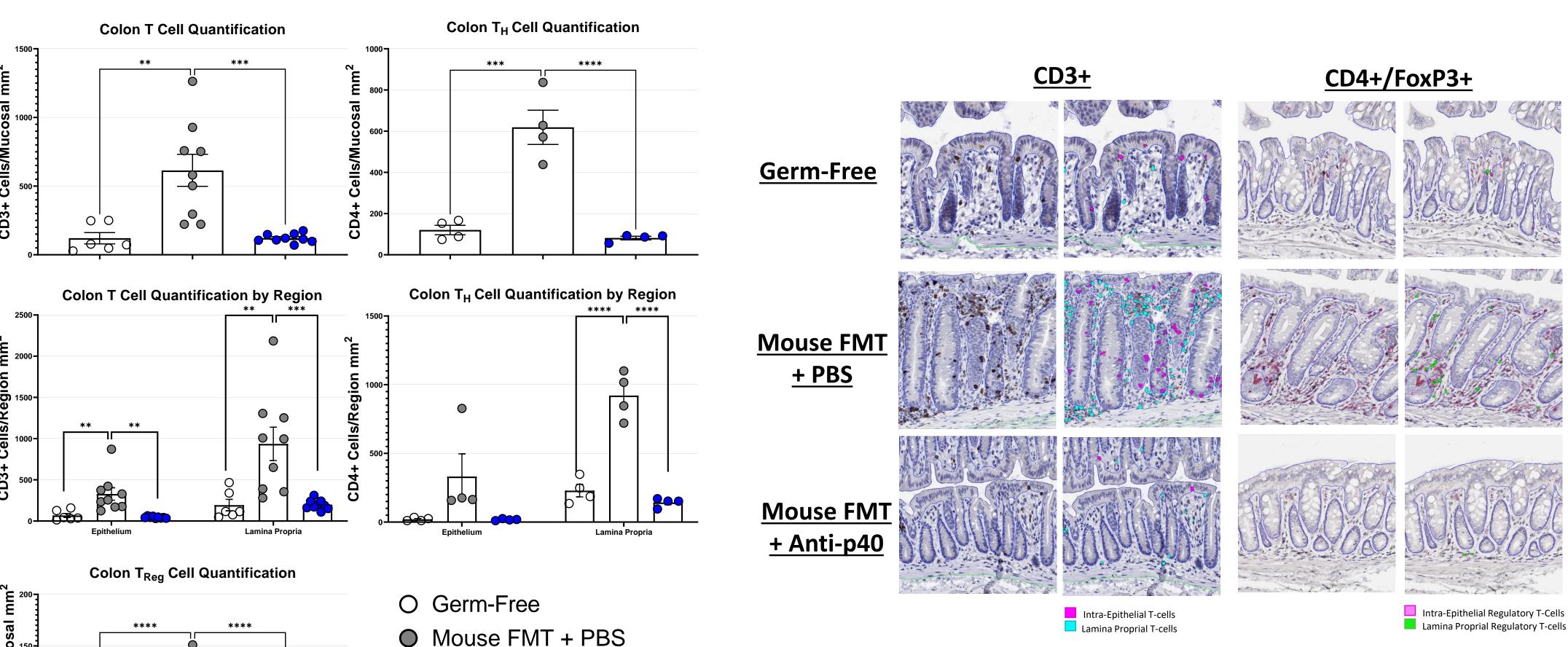




Mouse FMT + Anti-p40

groups to one another. Individual values and Mean±SEM are shown. n=6-10 per group

O Germ-Free Mouse FMT + PBS Mouse FMT + Anti-p40



Formalin-fixed colon samples were processed routinely for CD3 (upper left), CD4 (upper right) and CD4+FoxP3 (lower left) IHC and

quantification by image analysis. Middle panel shows intraepithelial and lamina propria quantification. Far right panels show representative images. **p<0.01; ***p<0.005; ****p<0.001 as determined by one-way ANOVA with Tukey's multiple comparisons test to compare all

Mouse FMT + Anti-p40

67 IHC and image analysis. *p<0.05; **p<0.01 as determined

TISSUE

Mean±SEM are shown. n=6-10 per group

Ki-67+ epithelial cells

-Mouse FMT + PBS Mouse FMT + Anti-p40 Flash-Frozen colon samples were processed routinely and analyzed by multiplex (Luminex) for panel of cytokines

RESULTS

including IL-1β, IL-17, IP-10, MIP-1α, RANTES, and TNFα. *p<0.05; **p<0.01 as determined by one-way ANOVA with Tukey's multiple comparisons test to compare all groups to one another.. Individual values and Mean±SEM

- Compared to control GF IL-10 KO mice, FMTinoculated mice demonstrated reduced overall weight gain, elevated levels of fecal lipocalin 2 calprotectin, and increased colon weight:length ratio at 8 weeks following FMT.
- 2) Colon protein levels of IL-1β, IL-17, IP-10, MIP- 1α , RANTES, and TNF α were elevated in diseased animals.
- Histopathology and IHC analysis showed that FMT-inoculated mice demonstrated elevated histopathology sum scores, increased colonic CD3+ T cells, CD4+ T_{H} cells, and increased FoxP3+ T_{regulatory} cells, and epithelial hyperplasia (increased Ki-67 immunolabeling density).
- Phenotypes were responsive to treatment with anti-p40.

CONCLUSIONS

These data provide a validated colitis model with relevant mechanisms for assessing the role of the microbiome and response to therapeutics in IBD.

Studies were performed at BioModels' facility in Waltham, MA, under BioModels' IACUC Protocol 21-1214-1.

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