

## Gut microbiome and its products promote endometriosis in animal model

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Generation of microbiota-depleted mice using antibiotics. A Schematic of experimental timeline and procedures. B Quantification of relative abundances of Bacteroidetes, Firmicutes, and Gamma-proteobacteria in feces from vehicle and Microbiota-depleted (MD) mice. C-D Wet weights of C spleen and D caecum in indicated treatment groups at a sacrifice. E The number of Peyer's patches from indicated treatment groups. F Representative images of Hematoxylin and Eosin-stained uterine cross-sections from the indicated treatment groups. LE, Luminal Epithelium; GE, Glandular Epithelium; S, Stroma. Yellow arrows indicate the gland. G-H Mouse G body weight and H water consumption at indicated time points in vehicle and MD mice. I-J Relative



level of **I** 17 $\beta$ -Estradiol in serum and **J** IL-1 $\beta$  in peritoneal fluid of indicated treatment groups. Data are presented as mean  $\pm$  SE (n = 5 mice per group). \*\*P

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