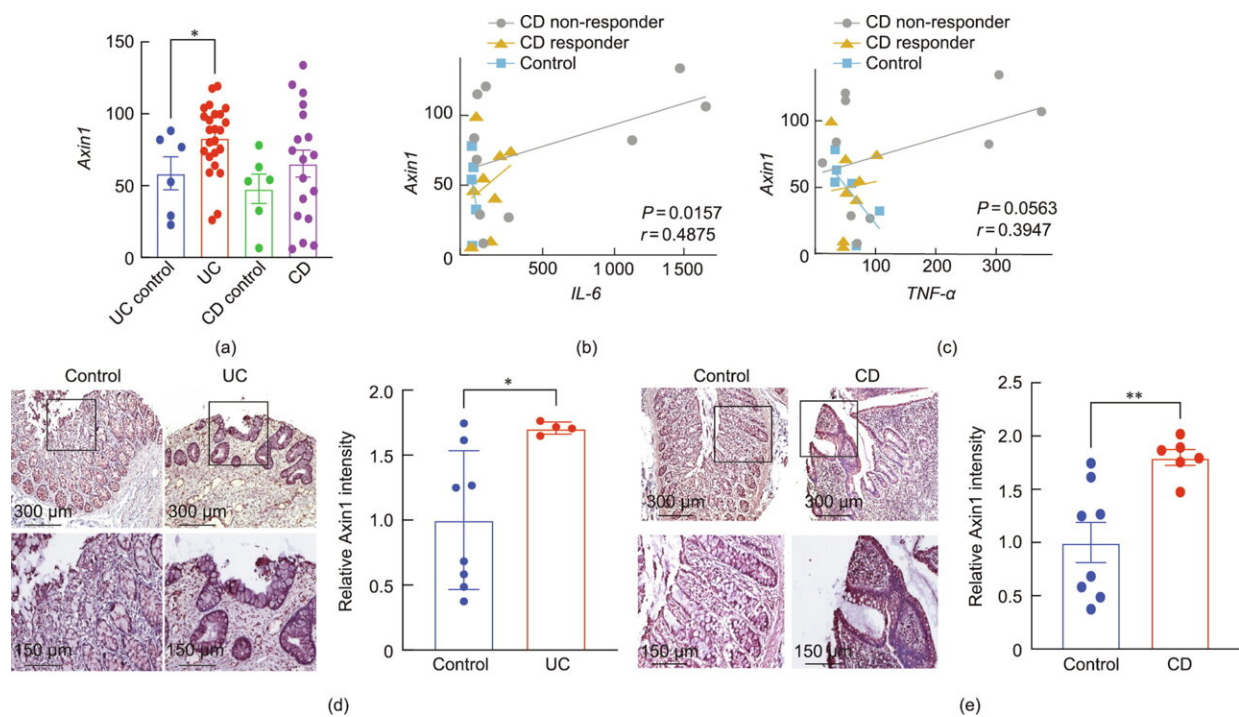


New research reveals gut microbiota link to colitis

September 22 2023



Expression of Axin1 is upregulated in human IBD. (a) Axin1 mRNA expression in patients with UC and CD. Values for healthy control, CD, and UC patients were obtained from GEO database GSE 16879. Data are expressed as mean \pm SEM; UC controls (n = 6), UC (n = 24), CD controls (n = 6), CD (n = 18); one-way ANOVA; *P Engineering (2023). DOI: 10.1016/j.eng.2023.06.007

A study conducted by Jun Sun's research team at the University of Illinois Chicago has revealed a new and critical role of Axin1 in

regulating intestinal epithelial development and microbial homeostasis. The research, titled "Intestinal Epithelial Axin1 Deficiency Protects Against Colitis via Altered Gut Microbiota," and published in the journal *Engineering*, highlights the potential therapeutic strategies for human inflammatory bowel disease (IBD).

IBD, a chronic inflammatory disorder affecting the [gastrointestinal tract](#), has been a significant health concern worldwide. The study focused on understanding the role of Axin1, a negative regulator of Wnt/ β -catenin signaling, in maintaining gut homeostasis and host response to inflammation.

The research team analyzed Axin1 expression in human inflammatory bowel disease datasets and found increased Axin1 expression in the colonic epithelium of IBD patients. To further investigate the effects and mechanism of intestinal Axin1 in regulating intestinal homeostasis and colitis, the team generated new mouse models with Axin1 conditional knockout in intestinal epithelial cells (Axin1 ^{Δ IEC}) and Paneth cells (Axin1 ^{Δ PC}).

The results showed that Axin1 ^{Δ IEC} mice exhibited altered goblet cell [spatial distribution](#), Paneth cell morphology, reduced lysozyme expression, and an enriched presence of *Akkermansia muciniphila* (*A. muciniphila*) in the gut microbiota. Importantly, the absence of intestinal epithelial and Paneth cell Axin1 led to decreased susceptibility to dextran sulfate sodium-induced colitis in vivo.

Furthermore, when Axin1 ^{Δ IEC} and Axin1 ^{Δ PC} mice were cohoused with control mice, they became more susceptible to dextran sulfate sodium (DSS)-colitis, suggesting the protective role of Axin1 in the presence of a healthy gut microbiota. Treatment with *A. muciniphila* further reduced the severity of DSS-colitis, highlighting its potential as a therapeutic target.

Interestingly, antibiotic treatment did not change the proliferation of [intestinal epithelial cells](#) in the control mice. However, in Axin1^{ΔIEC} mice with antibiotic treatment, the intestinal proliferative cells were significantly reduced, indicating the non-colitogenic effects driven by the gut microbiome.

These findings demonstrate the novel role of Axin1 in mediating intestinal homeostasis and the microbiota. The loss of intestinal Axin1 protects against colitis, likely through the regulation of epithelial Axin1 and Axin1-associated *A. muciniphila*. Further [mechanistic studies](#) using specific Axin1 mutations will be crucial in elucidating how Axin1 modulates the microbiome and host [inflammatory response](#), paving the way for new therapeutic strategies for human IBD.

Jiaming Wu, editor of the subject of medicine and health of engineering, commented, "This study provides valuable insights into the development of inflammatory bowel disease and offers potential therapeutic strategies for its treatment. By understanding the intricate interactions between Axin1, the [gut microbiota](#), and host immunity, researchers can develop targeted interventions to restore intestinal homeostasis and alleviate the symptoms of IBD."

The research team's findings have significant implications for the field of gastroenterology and hold promise for the development of novel treatments for IBD. As further studies are conducted, the [scientific community](#) eagerly awaits the potential therapeutic breakthroughs that may arise from this research.

More information: Shari Garrett et al, Intestinal Epithelial Axin1 Deficiency Protects Against Colitis via Altered Gut Microbiota, *Engineering* (2023). [DOI: 10.1016/j.eng.2023.06.007](https://doi.org/10.1016/j.eng.2023.06.007)

Provided by Engineering

Citation: New research reveals gut microbiota link to colitis (2023, September 22) retrieved 28 April 2024 from

<https://medicalxpress.com/news/2023-09-reveals-gut-microbiota-link-colitis.html>

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