



Weill Cornell Medicine

CRISPR-Mediated SATB2 Inhibition for the Treatment of Short Bowel Syndrome

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Background & Unmet Need

- Short bowel syndrome (SBS) results from surgical resection or congenital disease of the small intestine, and leads to an inability to absorb sufficient nutrients
- Over time, the remaining small bowel and colon undergo structural and functional changes to increase nutrient absorption, but ~50% of patients will require long-term total parenteral nutrition (TPN)
- Teduglutide (GLP-2 agonist) is the only FDA-approved therapy for SBS patients who are dependent on TPN
- While teduglutide led to a significant reduction in TPN frequency, only 11% of patients were completely weaned from TPN
- **Unmet Need:** Novel treatments for SBS patients that reduce the need for TPN and improve the ability of the remaining small bowel / colon to absorb nutrients

Technology Overview

- **The Technology:** Inhibition of SATB2 as a therapeutic strategy for the treatment of SBS, exemplified using CRISPR gene therapy
- **The Discovery:** Loss of the transcription factor SATB2 transforms colonic epithelium into ileum-like tissue in mice and human colonic organoids
- Intestinal deletion of SATB2 in mice led to significant remodeling of colonic tissue, with marked changes in tissue morphology, gene expression, and cell type composition
- **PoC Data:** Of note, SATB2 inhibition led to the generation of bona fide nutrient-absorbing enterocytes, with significantly enhanced nutrient absorption in *Satb2^{ckO}* mice colon compared to negative control
- CRISPR-mediated deletion of SATB2 using an optimized guide RNA (gRNA) replicated the findings observed in mice, suggesting a potential therapeutic strategy for SBS patients

Inventors:

Joe Zhou

Patents:

[US Application Filed](#)

[PCT Filed](#)

Publications:

[Gu et al. Cell Stem Cell. 2022.](#)

[Gu et al. Nat Commun. 2024.](#)

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Technology Applications

- SATB2-targeting CRISPR gene therapy using unique gRNA to restore small bowel function in SBS patients
- Cell therapy in which colonic stem cells are harvested from the patient, modified to disrupt the SATB2 gene, and then reimplanted into patients
- Gut-targeting siRNA therapy that reduces SATB2 expression

Technology Advantages

- Inhibition of SATB2 leads to durable gut remodeling that may reduce or eliminate the need for TPN
- SATB2 inhibition may be achieved through a variety of therapeutic approaches
- A single course of treatment may be sufficient to achieve lasting results

Supporting Data / Figures

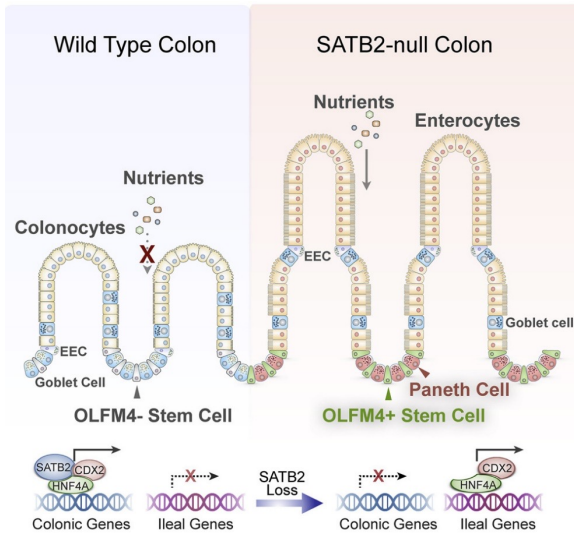


Figure 1: Loss of the transcription factor SATB2 leads to the downregulation of colonic genes and the upregulation of ileal genes, transforming colonic tissue into ileum-like tissue that absorbs key nutrients.

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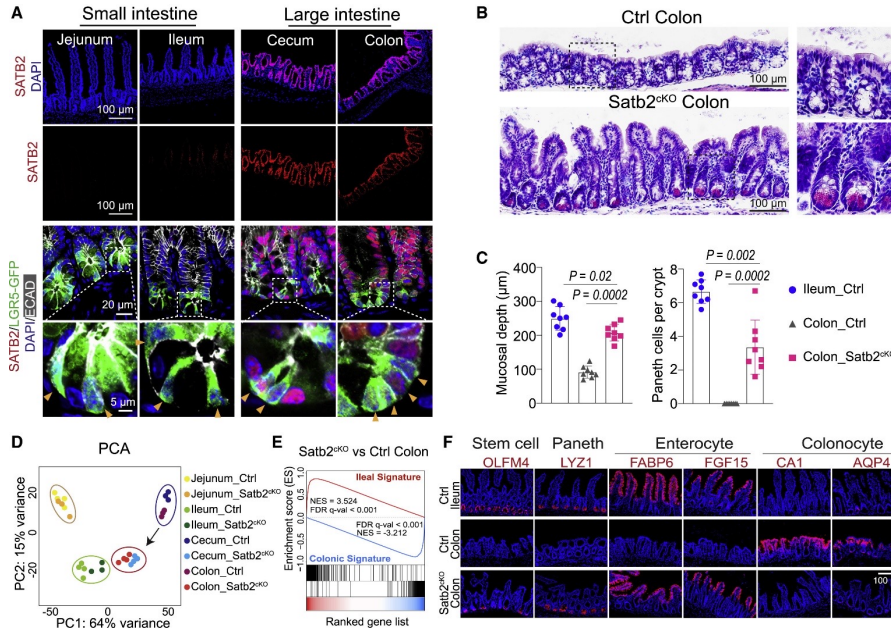


Figure 2: Conversion of large intestine mucosa to one that resembles ileal small intestine in Satb2^{CKO} mice. **A:** SATB2 is expressed in adult murine large intestinal epithelial cells but is absent in the small intestine. **B-C:** Intestinal deletion of *Satb2* in mice leads to morphological changes from colonic to ileum-like tissue. **D-E:** RNA-seq revealed a shift towards small intestine transcriptomes in colonic tissue of Satb2^{CKO} mice. **F:** Immunofluorescence demonstrated changes in cell types consistent with conversion of colonic tissue to ileum-like tissue.

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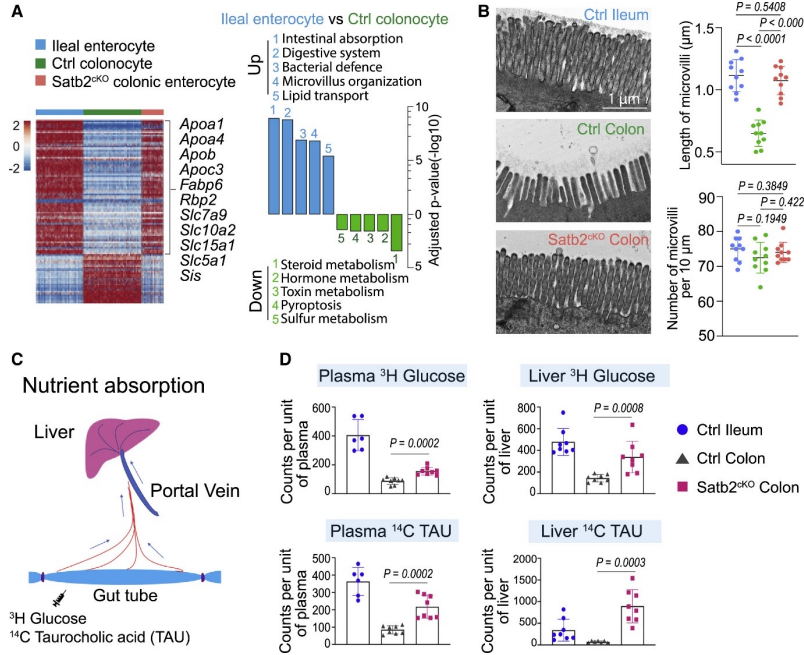


Figure 3: Generation of bona fide nutrient-absorbing enterocytes in *Satb2*^{CKO} colon. **A:** The scRNA profiles of *Satb2*^{CKO} colonic enterocytes closely resemble ileal enterocytes. **B:** The microvilli of *Satb2*^{CKO} enterocytes were significantly longer than those of control colon. **C:** Assay used to measure nutrient absorption. **D:** The amount of glucose and taurocholic acid absorbed is significantly higher in the *Satb2*^{CKO} colon compared to control colon and approaches that of a healthy ileum.

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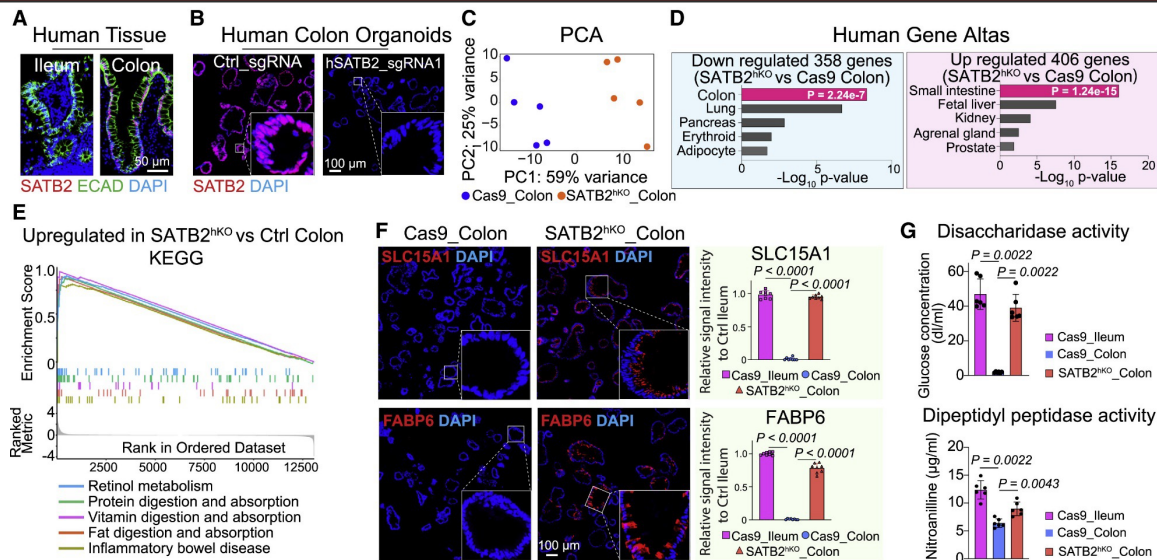


Figure 4: Colonic-to-ileal plasticity after CRISPR-mediated SATB2 deletion in human colonic organoids. **A-B:** SATB2 expression in normal tissues and human colon organoids. **C-E:** Transcriptomes of SATB2-deleted colonic organoids shifted towards the ileum. **F:** The ileal markers SLC15A1 and FABP6 were activated in colonic organoids after SATB2 loss. **G:** Significant activities of key enzymes were detected in SATB2-deleted colonic organoids.

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