

RESEARCH HIGHLIGHT

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# Dedifferentiation: the return road to repair the intestinal epithelium



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## Abstract

In the March 5 issue of *Cell Stem Cell*, (Murata K et al. *Cell Stem Cell*. 26(377–390):e376 2020) reported that intestinal stem cell recovery after injury is principally through *Ascl2*-dependent dedifferentiation of absorptive and secretory precursors in mice. This study provides evidence for robust regenerative capability of the intestinal epithelium via dedifferentiation of absorptive and secretory progenitors in the crypt.

## Main Text

Many tissues in adults contain a population of tissue-specific stem cells, which are required for tissue homeostasis and repair (Blanpain and Fuchs, 2014; Li and Clevers, 2010). Most cells in the intestinal epithelium renew every 5 days. This quick renewal process is driven by the crypt base-resident *Lgr5*-marked intestinal stem cells (ISCs) (Barker, 2014). However, the intestinal epithelium harbors an extraordinary regenerative capacity to deal with repeatedly challenges, like various injuries or inflammation, to regenerate all cell-types and restore its normal architecture even without *Lgr5*<sup>+</sup> ISCs.

Two models have been proposed to explain regeneration of the intestinal epithelium after damage (Barker et al., 2012). In the first model, known as “reserve stem cell” model, the quiescent reserve stem cells (also called +4 cells), marked by *Bmi1*, *Hopx*, *mTert* and *Lrig1*, are regarded to be the regenerative cell type that can survive upon damage and repopulate *Lgr5*<sup>+</sup> ISCs. The major challenge in this model is that expression of these markers traditionally seen as the quiescent stem cells is often not restricted to a given cell type, and whether the quiescent +4 stem cells exist is still under debate

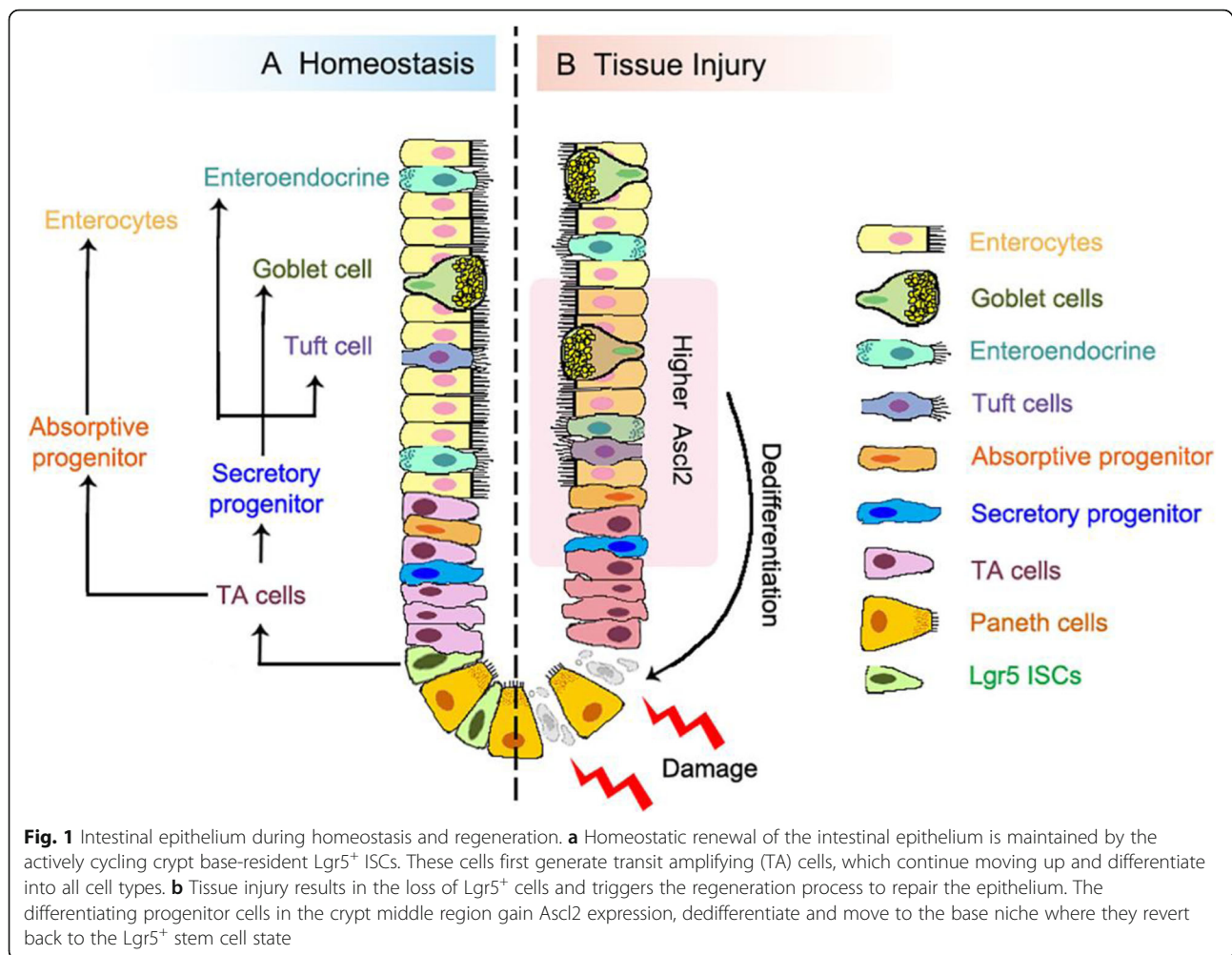
(Barker et al., 2012). Recently, the other model has gained wide attention. In the “dedifferentiation” model, various cell types may serve as a source to dedifferentiate back to intestinal stem cells after epithelium damage (Bankaitis et al., 2018), which is accordance with the high plasticity of the intestinal epithelium (de Sousa and de Sauvage, 2019).

The transcription factor *Ascl2* has been shown to play an important role in *Lgr5*<sup>+</sup> ISC maintenance, and its expression is restricted to these stem cells (van der Flier et al., 2009). However, in the recent work, Murata et al. found that *Ascl2* is largely dispensable for normal intestinal homeostasis, while *Ascl2*-deficiency impairs the refill of the *Lgr5*<sup>+</sup> ISCs pool in both small intestinal and colonic regions after irradiation or ablation of *Lgr5*<sup>+</sup> ISCs (Murata et al., 2020). These *Ascl2*-null mice display much shorter lifespan upon the lethal irradiation treatment. Upon the Diphtheria toxin-mediated *Lgr5*<sup>+</sup> ISC depletion, the mCherry-marked *Ascl2*<sup>+</sup> stem cells, which normally locate in the bottom of the crypt, disappear in the first few days and then re-appear in the middle region of colonic crypts at day 8. These cells move down into the bottom of the crypt at day 10, where they eventually express *Lgr5* and regain stemness (Fig. 1). These results suggest that ISCs ablation-activated *Ascl2* expression in the cells of the middle crypt region drives these cells to de-differentiate and repopulate the stem cell pool in the bottom region of the

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crypt. Supporting this, RNA-seq analysis revealed that regenerating *Ascl2*<sup>+</sup> cells have a gene expression signature associated with absorptive cells or secretory cells and exhibit ISC-oriented dedifferentiation, and pseudotime analysis indicates a transition from the absorptive or secretory state to dedifferentiating state, and eventually reach the endpoint that acquire ISC characteristics. The authors further exploited the *Ascl2* targets that mediate dedifferentiation. Among the upregulated genes in the middle region *Ascl2*<sup>+</sup> cells, the interleukin-11 (IL-11) receptor *Il11ra1* is a functional target of *Ascl2* during regeneration. Consistently, recombinant IL-11 augments the organoid formation of the middle region *Ascl2*<sup>+</sup> cells but not resting ISCs.

Their work prompts additional questions. Firstly, the important question is how epithelium damage or ISCs ablation triggers the dedifferentiation process. In another word, how do precursor cells sense the injury and initiate dedifferentiation? Extracellular matrix remodeling and YAP/TAZ signaling activation have been shown to

be involved in the colonic regeneration in the dextran sulfate sodium (DSS) colitis mouse model (Yui et al., 2018), while no YAP/TAZ signaling signature is enriched in regenerating cells in the Murata et al.'s study. Secondly, single-cell RNA sequencing analysis revealed that the upper regenerative *Ascl2*<sup>+</sup> cells comprise both absorptive and goblet progenitors. Do these two types of cells have an equal ability to revert to stem cells? Thirdly, there are two subpopulations with different proliferative activity in the upper *Ascl2*<sup>+</sup> cells. Does there exist relationship between the actively cycling and regeneration? Fourthly, it remains unclear how IL-11 induces the dedifferentiation process. Fifthly, the complexity and redundancy of signals supporting ISC regeneration indicate the possibility that other target genes or signals may also contribute to the *Ascl2*-dependent dedifferentiation. Sixthly, this study demonstrated that *Ascl2* re-expression in progenitors is critical for ISC repopulation after stem cell ablation or irradiation. However, it is unclear if this mechanism also applies to colitis-related epithelium repair. Moreover, after specific

ablation of tumor Lgr5<sup>+</sup> cells that have long-term self-renewal and differentiation capacities, differentiated KRT20<sup>+</sup> cells can dedifferentiate into Lgr5<sup>+</sup> cells (Shimokawa et al., 2017). Therefore, it will be interesting to compare signaling differences between physiological and tumor conditions. Addressing these questions will surely advance our understanding of tissue regeneration and provide solutions for regenerative medicine.

#### Authors' contributions

YL, XX and YGC wrote the paper. The authors read and approved the final manuscript.

#### Competing interests

The authors declare that they have no competing interest.

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