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Krüppel-Like Factor 5 Protects Against Dextran Sulfate Sodium-Induced Colonic Injury by Promoting Epithelial Repair in Mice

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Abstract

BACKGROUND & AIMS—Krüppel-like factor 5 (KLF5) is a transcription factor that promotes proliferation; is highly expressed in dividing crypt cells of the gastrointestinal epithelium and is induced by various stress stimuli. We sought to determine the role of KLF5 in colonic inflammation and recovery by studying mice with dextran sulfate sodium (DSS)-induced colitis.

METHODS—Wild-type (WT) and *Klf5*^{+/-} mice were given DSS in the drinking water to induce colitis. For recovery experiments, mice were given normal drinking water for 5 days after DSS administration. The extent of colitis was determined using established clinical and histological scoring systems. Immunohistochemical and immunoblotting analyses were used to examine proliferation, migration, and expression of the epidermal growth factor receptor (EGFR).

RESULTS—*Klf5* expression was increased in colonic tissues of WT mice given DSS; induction of *Klf5* was downstream of mitogen-activated protein kinase signaling. In DSS-induced colitis, *Klf5*^{+/-} mice exhibited greater sensitivity to DSS than WT mice, with significantly higher clinical and histological colitis scores. In recovery experiments, *Klf5*^{+/-} mice showed poor recovery, with continued weight loss and higher mortality than WT mice. *Klf5*^{+/-} mice from the recovery period had reduced epithelial proliferation and cell migration at sites of ulceration compared to WT mice; these reductions correlated with reduced expression of EGFR.

CONCLUSIONS—Epithelial repair is an important aspect of recovery from DSS-induced colitis.

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Authors' contributions

BBM designed and performed experiments, analyzed data and wrote the manuscript

SSK managed mouse strains and conducted mouse experiments and *in vivo* staining

ABB conducted *in vitro* wounding assays and Western blots

KY performed Western blots for time course experiments

SVS provided histological analysis and technical expertise and advised on study design

VWY advised on experimental design, analyzed data and edited the manuscript

All authors read and approved the final manuscript.

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The transcription factor KLF5 regulates mucosal healing through its effects on epithelial proliferation and migration.

Keywords

inflammatory bowel disease; animal model; inflammatory response; mouse

The mammalian gut epithelium forms a dynamic barrier between the organism and various external factors present within the lumen, including diet, microorganisms, and drugs. In the setting of rapid cell turnover and constant changes in luminal content, the intestinal mucosa maintains its integrity through stringent regulation of pathways involving proliferation, migration, differentiation, and cell death. Maintaining tissue homeostasis becomes critical in instances of inflammatory bowel disease (IBD), where dysregulation of these pathways can result in excessive tissue injury, inadequate regeneration and an increased risk of developing cancer. While normal maintenance of the intestinal epithelium by the Wnt signaling pathway has been well characterized^{1, 2}, little is known about the molecular events that regulate restoration of tissue homeostasis in response to external stress or injury.

Krüppel-like factor 5 (KLF5) is a zinc finger-containing transcription factor that belongs to a family of mammalian Krüppel-like transcription factors. These factors are often tissue-specific and serve diverse roles in processes such as cell proliferation, differentiation and embryogenesis^{3, 4}. KLF5 is highly expressed in the gut and is enriched in epithelial cells in the proliferative compartment of the intestinal tract⁵, indicating a role as a pro-proliferative factor. Indeed, several *in vitro* studies show that ectopic expression of KLF5 enhances proliferation of non-transformed cultured epithelial cells^{6–8}. KLF5 has also been shown to mediate the growth and transforming effects of oncogenic *KRAS* in intestinal epithelial cells⁹. Furthermore, mouse studies reveal that *Klf5* promotes proliferation in the settings of bacterial infection and intestinal tumor initiation^{10, 11}. The pro-proliferative effects of KLF5 are thought to be mediated through its transcriptional targets, which include the growth factor, platelet-derived growth factor alpha polypeptide (PDGF-A), and cell cycle proteins, cyclin D1, cyclin B1 and Cdc2^{12–14}.

In addition to its role in regulating proliferation, a number of studies indicate that KLF5 may act as a mediator of external stress responses. KLF5 expression is shown to be induced in vascular smooth muscle cells following balloon injury in rat aorta¹⁵. This induction is mediated through mitogen-activated protein kinase (MAPK) signaling by the early response gene, early growth response factor 1 (*Egr1*)^{15, 16}. Additional studies in heterozygous *Klf5* knockout (*Klf5*^{+/-}) mice reveal that KLF5 is a critical factor for tissue remodeling in injured vascular tissues by participating in proliferative and angiogenic responses¹⁴. In intestinal epithelial cells (IECs), KLF5 is activated by the bacterial component, lipopolysaccharide, and enhances inflammatory responses in these cells through its effects on NF-κB signaling¹⁷. Furthermore, *in vivo* studies show that KLF5 is induced by colonization of the mouse colon with the bacterial pathogen, *Citrobacter rodentium*, and that *Klf5* mediates colonic epithelial hyperplasia activated by this infection¹¹. Taken together, these reports indicate a critical role for KLF5 in cellular responses to cardiovascular injury and suggest a similar function for KLF5 in intestinal tissues.

In the present study, we utilize the dextran sulfate sodium (DSS)-induced model of colitis to examine the role of *Klf5* in promoting restoration of tissue homeostasis. WT and *Klf5*^{+/-} mice were treated with the chemical irritant, DSS, and examined for the response and the ability to recover from DSS-induced injury. Results revealed that reduced levels of *Klf5* heightened susceptibility to DSS-induced damage, indicating a protective role for *Klf5* in response to chemical-induced injury. Moreover, *Klf5*^{+/-} mice exhibited poor recovery from

DSS treatment, indicating that KLF5 is important for restoration of intestinal epithelial homeostasis.

Materials and Methods

Mice

C57BL/6 mice were purchased from The Jackson Laboratory (Bar Harbor, ME). *Klf5*^{+/-} mice on a C57BL/6 background were generated as described previously¹¹. Mouse strains were bred and housed in the Whitehead Animal Research Facility at Emory University. Animal care and procedures were conducted in compliance with Emory University Institutional Animal Care and Use Committee guidelines.

Cell Culture

Caco-2, HCT 116 and DLD-1 colon cancer cells were maintained at 37° C in 5% CO₂. For inhibitor studies, Caco-2 cells were pretreated for 10 min prior to addition of DSS with U0126 at 50 μM (Cell Signaling Technology; Beverly, MA) or Bay 11-7082 at 20μM (Calbiochem; San Diego, CA).

Induction of DSS Colitis

Colitis was induced in 7- to 8-week old gender-matched WT and *Klf5*^{+/-} mice by by addition of dextran sulfate sodium (DSS) (molecular weight, 35,000–50,000; MP Biomedicals, Solon, OH) to the drinking water at 3.5% (wt/vol) for 7 days. Controls received normal drinking water. For recovery experiments, WT and *Klf5*^{+/-} mice were administered 3.5% or 2.5% DSS, respectively, for 7 days, followed by 5 days of recovery with normal dH₂O.

Protein Isolation from Mouse Colon

Colon sections were flushed with phosphate-buffered saline (PBS), inverted and scraped to isolate mucosal protein. Scraped tissues were placed in Cell Extraction Buffer (Invitrogen, Carlsbad, CA) and homogenized with silicone beads using a Bullet Blender (Next Advance, Inc., Averill Park, NY) at 4° C for 5 min at Speed 5.

Antibodies

Antibodies are listed in Supplementary materials.

Histology

Colons were prepared as Swiss rolls for embedding in paraffin, and sections were stained with hemotoxylin and eosin (H&E). Images of tissue sections were captured using a Zeiss Axioskop2 plus microscope (Carl Zeiss MicroImaging, Inc., Thornwood, NY).

Immunohistochemistry

Immunohistochemistochemical staining was performed as described previously¹¹. Fluorescent images of *Klf5* staining were analyzed for nuclear intensities using Metamorph Imaging software (Molecular Devices, Sunnyvale, CA).

Clinical and Histological Assessment of Colitis

Clinical and histological scores for measuring the degree of colitis were determined by the method described by Cooper *et al.*¹⁸

MPO Activity

Neutrophil infiltration was quantified by measuring myeloperoxidase (MPO) activity. Colon tissue was homogenized in HTAB buffer (50 mM phosphate buffer, pH 6.0, 0.5% hexadecyltrimethyl ammonium bromide). Lysates (10 μ l) were added to 1 mg/ml of *o*-dianisidine hydrochloride and 5×10^{-4} % hydrogen peroxide, and the absorbance measured at 460 nm. The results were expressed as U activity per mg protein.

In Vitro Wounding Assays

HCT 116 cells or DLD-1 cells were plated in 6-well plates. Cells were transfected in triplicate wells with pCI-neo or pCI-neo-KLF5 expression vectors. At 24 hrs after transfection, confluent monolayers were wounded with a 10 μ l pipette tip. Images were taken at wounding and 18 hrs later. Wound widths were determined by averaging 6 measurements per image.

Statistical Analysis

The data are presented as mean \pm SEM. Equality of variances was tested by Levene's test. Multigroup data were analyzed by ANOVA with multiple pair-wise comparisons made among groups using Tukey's test. Data comparing 2 groups were analyzed using a 2-tailed *t* test. *P* values of < 0.05 were considered statistically significant.

Results

Induction of Klf5 Following DSS Treatment

The dextran sulfate sodium (DSS) mouse model of colitis exhibits many of the histological characteristics of human IBD, including disruption of crypt architecture, mucosal ulceration, and infiltration of neutrophils, monocytes and lymphocytes¹⁹ To investigate whether KLF5 expression is altered in the setting of colitis, C57BL/6 (WT) mice were treated with DSS and examined for levels and cellular localization of Klf5. Mice were administered 3.5% DSS in the drinking water for seven days to induce colitis, after which they were euthanized and colonic tissues collected for protein analysis and immunohistochemistry.

Immunohistochemical staining revealed that Klf5 was expressed at high levels in DSS-treated mice, particularly in epithelial cells contiguous to sites of ulceration. Intense KLF5 staining was seen in the crypts adjacent to ulcerated areas and at luminal surfaces covering the ulcers (Figure 1A; WT DSS). This staining pattern was in contrast to untreated WT mice in which staining of Klf5 was restricted to the proliferative zones in the lower two-thirds of the crypts, with no expression at the luminal surface (Figure 1A; WT H₂O). Increased levels of Klf5 were confirmed by immunofluorescence staining, which showed a trend towards higher KLF5 intensity per cell as well as a higher percentage of Klf5-positive cells per crypt (Figures 3A–3C; WT H₂O versus WT DSS). During a 5 day time course of DSS treatment, Western blot analysis indicated that Klf5 was induced in the colons of DSS-treated WT mice as early as 1 day after DSS treatment (Figure 1B). Increased levels of Klf5 began to decline at days 4 and 5, likely as epithelial cells were lost with progressive mucosal erosion.

To investigate the mechanism of Klf5 activation, two major signaling pathways were evaluated in response to DSS. MAPK signaling via activation of MAPK kinase (MEK) and extracellular-signal-regulated kinases 1 and 2 (ERK1/2) is known to induce Klf5 expression in several cell types^{13, 16, 20}. Phosphorylation of ERK1/2, which can occur in both epithelial and infiltrating immune cells, was increased at day 1 of DSS treatment and was induced further at days 4 and 5 (Figure 2A). NF- κ B, which is a key regulator of inflammatory responses, also appeared to be activated by DSS treatment, as indicated by a decrease in levels of the NF- κ B inhibitor I κ B beginning at day 3. As both of these pathways

are activated by DSS treatment *in vivo*, an *in vitro* system was used to examine the dependence of KLF5 induction on ERK1/2 and NF- κ B signaling.

In Caco-2 colon cancer cells treated with 3% DSS during an 8 hour period, KLF5 levels increased two hours after treatment with DSS and reached maximal induction at five hours (Figure 2B). Since DSS preparations can be contaminated with lipopolysaccharide (LPS) ²¹, cells were treated with DSS and polymyxin B (a scavenger for LPS) to verify that the effects of DSS on KLF5 were not skewed by the presence of LPS. At the 5 hour time point, KLF5 levels continued to be induced by DSS in the presence of polymyxin B, indicating no involvement of LPS (Supplementary Figure 1). ERK1/2 showed acute activation one hour after addition of DSS, and, NF- κ B was activated beginning at two hours after DSS (Figure 2B). To determine whether activation of these signaling pathways were required for induction of KLF5, Caco-2 cells were pretreated with inhibitors to MEK/ERK or NF- κ B prior to the addition of DSS for 5 hours. KLF5 was induced by DSS in the absence of inhibitors as well as in the presence of Bay 11-7082 (Figure 2C). However, KLF5 induction was blocked by U0126, indicating that KLF5 induction is dependent on MEK/ERK signaling.

Reduced Levels of Klf5 in Regenerating Tissues of *Klf5*^{+/-} Mice Treated with DSS

To examine the role of KLF5 in DSS-induced colitis, mice with heterozygous deletion of *Klf5* (*Klf5*^{+/-}) ¹¹ were compared to WT mice for responses to DSS treatment (homozygous deletion of *Klf5* is embryonic lethal ¹⁴). *Klf5*^{+/-} mice develop normally and are fertile. No histological differences have been noted in the colonic mucosal architecture of *Klf5*^{+/-} mice compared to WT mice, and epithelial barrier function is comparable between the two genotypes (Supplementary Figures 2A and 2B). Also, blood analysis revealed no abnormalities in the development of hematopoietic cells (Supplemental Figure 2C). WT and *Klf5*^{+/-} mice were treated for 7 days with DSS, and Klf5 levels were analyzed in colon tissues by immunofluorescence. In untreated mice, basal levels of Klf5 in the heterozygous mice were reduced in intensity compared to WT mice, confirming haploinsufficiency of the *Klf5* gene (Figures 3A and 3B). Upon treatment with DSS, Klf5 exhibited strong immunofluorescence staining in regenerative crypts and at the luminal surface of WT mice, with a trend towards increased Klf5 intensity per cell compared to untreated WT mice. In addition, WT mice treated with DSS had a higher percentage of Klf5-positive cells per crypt compared to untreated WT mice (Figure 3C). In *Klf5*^{+/-} mice treated with DSS, Klf5 levels were significantly lower than in WT treated mice, with reduced intensity of Klf5 staining per cell and a lower percentage of Klf5-positive cells per crypt (Figures 3A–3C).

Increased Sensitivity of *Klf5*^{+/-} Mice to DSS Treatment

The extent of colitis in both WT and *Klf5*^{+/-} mice was determined after 7 days of DSS treatment using clinical and histological characterization. Weights of WT and *Klf5*^{+/-} mice were recorded daily over the course of DSS treatment (Figure 4A). A significant difference was seen in the percentage weight loss between treated WT mice compared to treated *Klf5*^{+/-} mice, with an average weight loss of 5% in WT animals versus an average of 13% in *Klf5*^{+/-} mice (Figure 4A). H&E staining of colon from DSS-treated WT mice indicated moderate inflammation and ulceration, with some disruption of normal crypt architecture (Figure 4B). However, in *Klf5*^{+/-} mice, the damage was more severe, with extensive infiltration of neutrophils and a complete loss of crypt architecture in large, continuous sections. Extent of neutrophil infiltration was determined by myeloperoxidase (MPO) activity, a lysosomal enzyme abundant in neutrophils (Figure 4C). MPO activity was approximately 3-fold higher in the *Klf5*^{+/-} DSS-treated mice compared to WT-treated mice, correlating with an increased presence of granulocytes visualized by histology. As an indicator of colitis, colon lengths were significantly shorter in the *Klf5*^{+/-} mice compared to

WT mice, with lengths measuring $4.4 \text{ cm} \pm 0.2$ and $5.4 \text{ cm} \pm 0.2$, respectively ($n = 7$; $P = 0.01$). The extent of colitis was further quantified using a well-established method of clinical and histological scoring¹⁸. Clinical scores, based on the parameters of weight loss, stool consistency and presence of fecal blood, indicated more extensive colitis in *Klf5*^{+/-} mice (10.0 ± 0.6) compared to WT mice (5.9 ± 0.6 ; $n = 7$; $P < 0.001$) (Figure 4D). Similarly, histological scores, based on crypt destruction, mucosal damage, epithelial erosion and inflammatory infiltrate, mirrored the clinical findings, with *Klf5*^{+/-} mice scoring significantly higher than WT mice (10.0 ± 0.3 versus 6.9 ± 0.5 ; $n = 7$, $P < 0.001$) (Figure 4E). Thus, the data support a protective role for *Klf5* in responding to DSS-induced mucosal damage as indicated by increased sensitivity of *Klf5*^{+/-} mice to DSS treatment.

Poor Recovery of *Klf5*^{+/-} Mice Following DSS Treatment

To determine whether recovery from DSS treatment is impaired in *Klf5*^{+/-} mice, WT and *Klf5*^{+/-} mice were examined during a recovery phase, in which the mice were returned to normal drinking water for 5 days following 7 days of DSS treatment. To begin the recovery phase with similar degrees of colitis in WT and *Klf5*^{+/-} mice, different percentages of DSS were tested to identify treatment conditions that would yield similar effects in the two mouse strains. Consequently, *Klf5*^{+/-} mice were treated with 2.5% DSS for 7 days whereas WT mice were treated with 3.5% DSS. While WT and *Klf5*^{+/-} mice entered the 5 day recovery phase with similar degrees of colitis based on weights and colitis scores (Figures 5A, 5C and 5D, Day 7 DSS), *Klf5*^{+/-} mice exhibited poorer recovery, with shorter colon lengths at the end of the 5 day recovery period ($4.7 \pm 0.2 \text{ cm}$ versus $5.8 \pm 0.2 \text{ cm}$ in WT mice; $n = 14$, $P < 0.001$) and a continued decline in average body weight versus unchanged weight in the WT mice (Figure 5A). In addition, the *Klf5*^{+/-} mice exhibited significantly higher mortality than the WT mice, with 36% of the mice reaching IACUC endpoints during the recovery phase ($n = 14$, $P = 0.02$) (Figure 5B). Clinical and histological scores also showed an inability of *Klf5*^{+/-} mice to recover, as both measures indicated worsening of colitis during the recovery phase compared to stabilization or improvement in WT mice (Figures 5C and 5D). Therefore, *Klf5*^{+/-} mice not only exhibited increased sensitivity to DSS compared to WT mice, but also failed to stabilize or improve during the recovery period, suggesting an impaired ability to heal injured mucosal tissues.

Reduced Epithelial Proliferation and Migration in *Klf5*^{+/-} Mice Treated with DSS

Because KLF5 is highly expressed in areas of ulceration, we wanted to determine whether KLF5 contributes to injury repair. To explore this possibility, two aspects of wound repair, proliferation and cell migration, were examined in colons of recovery mice. To determine relative rates of proliferation, actively dividing cells in WT and *Klf5*^{+/-} recovery mice were pulse-labeled with the nucleoside, bromodeoxyuridine (BrdU), and numbers of proliferating cells per crypt were calculated in areas adjacent to ulcers (Figure 6A). Whereas 80% of cells in regenerative glands of WT mice stained positive for BrdU, this percentage was reduced to 50% in *Klf5*^{+/-} mice, indicating a slower rate of proliferation in regions of repair in *Klf5*^{+/-} mice. To address the question of epithelial migration, dividing cells in the mice were labeled for 24 hrs with BrdU to track cell migration. Colonic tissues from recovery mice were co-stained for Ki67 and BrdU to distinguish between cells that were actively proliferating at the time of sacrifice (green cells expressing Ki67) and cells that had ceased to proliferate and were migrating into ulcerated areas (red cells stained for BrdU) (Figure 6B). Numerous cells in the WT mice were migrating vertically within the crypts and across the luminal surface, whereas only minimal cell migration was visible in *Klf5*^{+/-} mice.

To examine further the effects of KLF5 on epithelial cell migration, scratch wound assays were used to measure differences in migration in colon cancer cells transfected with empty vector or with a vector for ectopic expression of KLF5. Confluent monolayers of DLD-1

cells were wounded with a pipette tip and examined for closure of the wound at 18 hrs post-wounding. Cells overexpressing KLF5 showed increased wound closure compared to cells transfected with empty vector (Figure 6C and 6D), indicating that KLF5 promotes cell migration into the wound site. In a reciprocal experiment, KLF5 expression was blocked by transfection of DLD-1 cells with two different siRNAs specific for KLF5. Cells transfected with KLF5-specific siRNAs exhibited reduced wound closure compared to cells transfected with a non-specific siRNA (Figure 6E). Similar results were obtained in HCT 116 colon cancer cells with overexpression and inhibition of KLF5 (Supplementary Figure 3). Thus, KLF5 promotes epithelial cell migration during wound repair.

Reduced EGFR Expression in *Klf5*^{+/-} Mice Treated with DSS

Activation of EGFR is a key signaling event that drives cells to proliferate, migrate and cover wounds in several types of epithelia²²⁻²⁴. EGFR is expressed at sites of ulcer healing in the gastrointestinal tract and is detected predominantly in the immature regenerative epithelium around ulcer margins²⁵. Previously, *EGFR* was reported to be a direct transcriptional target of KLF5 in primary esophageal keratinocytes by promoter analysis and chromatin immunoprecipitation assays²⁰. Given the participation of EGFR in wound healing, we examined whether EGFR was expressed differentially in WT and *Klf5*^{+/-} mice treated with DSS. After five days of DSS treatment, WT mice showed positive staining of EGFR in areas of ulceration, particularly in regenerative glands and regions of epithelial restitution at the luminal surface (Figure 7A, panel 1). In contrast, *Klf5*^{+/-} mice exhibited markedly reduced EGFR staining at comparable sites (Figure 7A, panel 2). Western blot analysis of lysates from DSS treated mice showed a pattern of induction of EGFR in WT mice that correlated with the induction of Klf5 (Figure 7B). In *Klf5*^{+/-} mice, EGFR levels were induced to a lesser extent than in WT mice at each time point analyzed, suggesting that EGFR levels are dependent on Klf5 expression. Likewise, in recovery experiments, EGFR was detected in epithelial cells at the luminal surface of ulcers of WT mice (Figure 7A, panels 3 and 5), and this expression was reduced in *Klf5* heterozygous animals (Figure 7A, panels 4 and 6). To examine directly whether KLF5 expression can regulate EGFR levels in colonic cells, we overexpressed or blocked expression of KLF5 in colon cancer cells. Suppression of KLF5 in DLD-1 cells with relatively high levels of endogenous KLF5 resulted in a marked reduction of EGFR protein in DLD-1 cells. Conversely, overexpression of KLF5 in HCT 116 cells with relatively low levels of endogenous KLF5 resulted in increased levels of EGFR. Taken together, these results indicate that EGFR levels are regulated by Klf5 expression in colonic tissues.

Discussion

The intestinal epithelium serves as a critical barrier against a broad spectrum of noxious and immunogenic substances within the intestinal lumen. When this barrier is disrupted, the integrity of the surface epithelium is quickly reestablished, even following severe damage, due to the remarkable regenerative capacity of the intestinal mucosa²⁶. However, in various intestinal disorders, including IBD, the mucosal barrier is repeatedly disrupted, allowing for increased exposure to infectious agents and toxins that can prolong the inflammatory condition²⁷. Indeed, barrier defects in IBD are thought to play a critical role in the pathobiology of disease, being intimately connected to chronic inflammatory responses²⁸. Thus, rapid repair of the mucosal barrier is critical for limiting inflammation and healing ulcerated lesions in chronic inflammatory diseases.

In this paper, we show that KLF5 serves a novel role in epithelial regeneration of DSS-damaged tissues by contributing both to proliferative and migratory processes essential for epithelial repair. The participation of KLF5 in proliferative responses to DSS damage is consistent with *in vitro* and *in vivo* studies which indicate a pro-proliferative role for KLF5

6–8, 11 However, the ability of KLF5 to promote proliferation can have pathogenic effects in the context of oncogenic mutations. Previously, we have reported that KLF5 mediates cell transformation during KRAS-induced intestinal tumorigenesis and that KLF5 is required for intestinal tumor initiation in the *Apc*^{Min/+} tumorigenesis model^{9–10}.

In addition to promoting proliferation in the setting of tissue repair, our studies provide the first evidence that KLF5 functions in epithelial cell migration in the colonic mucosa. Other KLF proteins have been implicated in regulating cell migration, including KLF2 in T cells and smooth muscle cells^{29–31}, KLF8 in immortalized human breast epithelial cells³² and KLF6 in endothelial cells³³. Also, KLF5 was recently shown to enhance migration of esophageal keratinocytes through activation of the transcriptional target, integrin-linked kinase (ILK)³⁴. In our experiments, however, we saw no change in ILK levels with induction of *Klf5* in the colons of mice treated with DSS (data not shown).

During ulcer healing, the mucosa at the ulcer margin forms a “healing zone”, in which the gastrointestinal glands become dilated, the cells lining the glands become de-differentiated and begin to express EGFR^{35–37}. This process is activated by growth factors that include epidermal growth factor (EGF), hepatocyte growth factor (HGF), platelet derived growth factor (PDGF) and basic fibroblast growth factor (bFGF), which stimulate proliferation and cell migration to restore mucosal surfaces and reconstitute mucosal glands³⁷. Several lines of evidence suggest that KLF5 may participate in intestinal wound healing through its effects on growth factor signaling. KLF5 has been shown to regulate expression of PDGF-A in vascular smooth muscle cells in response to vascular injury^{14, 38}. In addition, KLF5 has been reported to promote EGF signal amplification in primary esophageal cells through direct transcriptional activation of EGFR²⁰. Here, we show that EGFR levels correlate with changes in *Klf5* levels in the DSS mouse injury model, and that EGFR expression is reduced in healing zones of *Klf5*^{+/-} DSS-treated mice. In light of these results, future studies will explore the dependence of KLF5 on EGFR in promoting epithelial cell migration and wound healing.

In summary, these results demonstrate the importance of efficient epithelial repair in recovery from colonic injury. We show that KLF5 plays a critical role in epithelial repair following DSS treatment by promoting proliferation of colonic epithelial cells at sites of ulceration. In addition, we report a novel role for KLF5 in promoting epithelial cell migration as part of the restitution process.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

BrdU	bromodeoxyuridine
BSA	bovine serum albumin
DSS	dextran sulfate sodium
EGFR	epidermal growth factor receptor

FBS	fetal bovine serum
H&E	hematoxylin and eosin
IBD	inflammatory bowel disease
KLF5	Krüppel-like factor 5
MAPK	mitogen-activated protein kinase
MPO	myeloperoxidase
PBS	phosphate-buffered saline
PDGF-A	platelet-derived growth factor alpha polypeptide
WT	wild-type

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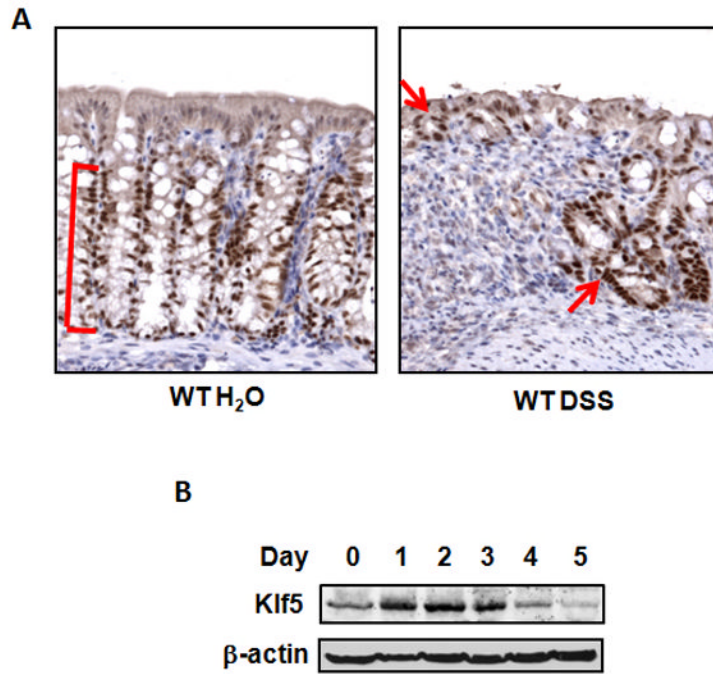


Figure 1. Klf5 is induced in wild-type (WT) mice treated with DSS. Eight week-old WT mice were given water or treated with 3.5% DSS (wt/vol). (A) Immunohistochemical staining of colon of Klf5 in WT mice untreated or treated with DSS for 7 days (brown, Klf5; blue; counterstain). Red bars and arrows indicate regions of Klf5 expression. (B) Western blots of lysates from WT mice treated with DSS over a 5 day time course. Lysates for each time point were pooled from 3 separate mice.

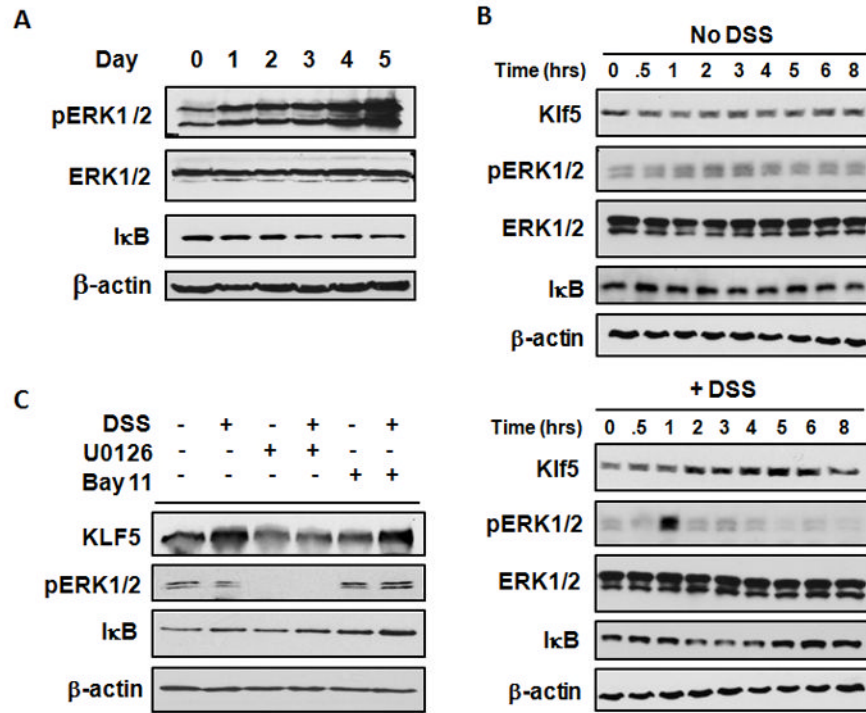


Figure 2. Induction of KLF5 with DSS treatment is dependent on ERK1/2 signaling. (A) Detection of ERK1/2 and NF-κB activation by Western blotting, using pooled lysates from WT mice treated with 3.5% DSS over a 5 day time course (n=3). (B) Caco-2 cells were treated with 3% DSS for up to 8 hrs, and lysates were examined by Western blotting. (C) Caco-2 cells were pretreated with inhibitors of MEK/ERK (U0126) or NF-κB (Bay 11-7082), and treated for 5 hrs with 3% DSS. Lysates were examined by Western blotting.

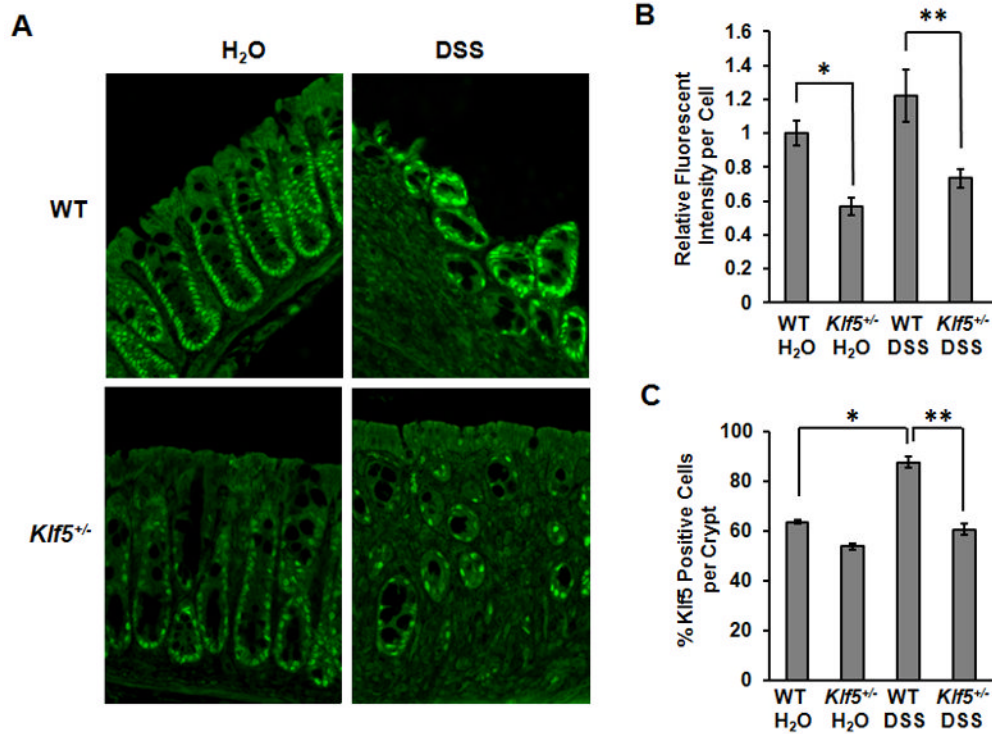


Figure 3. Klf5 expression is reduced at sites of ulceration in DSS-treated *Klf5*^{+/-} mice. (A) Immunofluorescence staining of Klf5 in WT and *Klf5*^{+/-} mice untreated or treated with DSS for 7 days. White arrows indicate Klf5 staining at luminal surface (B) Quantification of fluorescent intensities of Klf5 staining per cell. Data represent the mean ± SEM of 5 mice per group with at least 50 cells quantified per sample. * *P* = 0.01; ** *P* = 0.04. (C) Quantification of Klf5-positive cells per crypt in untreated and DSS-treated WT and *Klf5*^{+/-} mice. Data represent the mean ± SEM of 3 mice per group, counting 5 crypts per mouse. * *P* < 0.001. ** *P* = 0.001.

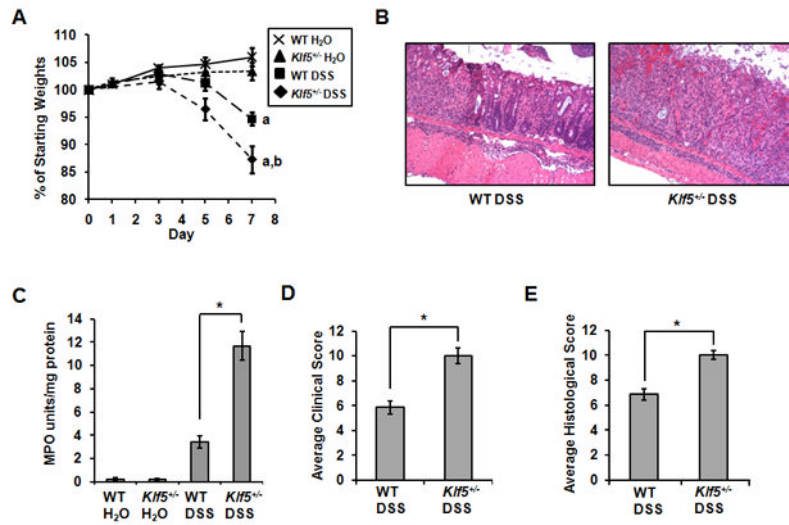


Figure 4. *Klf5*^{+/-} mice exhibit increased susceptibility to colitis with DSS treatment. Eight-week old WT and *Klf5*^{+/-} mice were treated with 3.5% DSS for 7 days and examined for clinical signs of colitis. Colon tissues were processed for enzymatic and histological analysis. (A) Weights of mice during 7 day treatment period; n = 9 mice per group, **a**: *P* < 0.001 versus control; **b**: *P* = 0.02 versus WT DSS. (B) Representative H&E staining of proximal colons from DSS-treated mice. (C) Myeloperoxidase (MPO) activity as a measure of neutrophil infiltration; n = 5, * *P* < 0.001. (D) Mean clinical scores of colitis with a maximum score of 12. n = 7, * *P* < 0.001. (E) Mean histological scores of colitis with a maximum score of 11. n = 7, * *P* < 0.001.

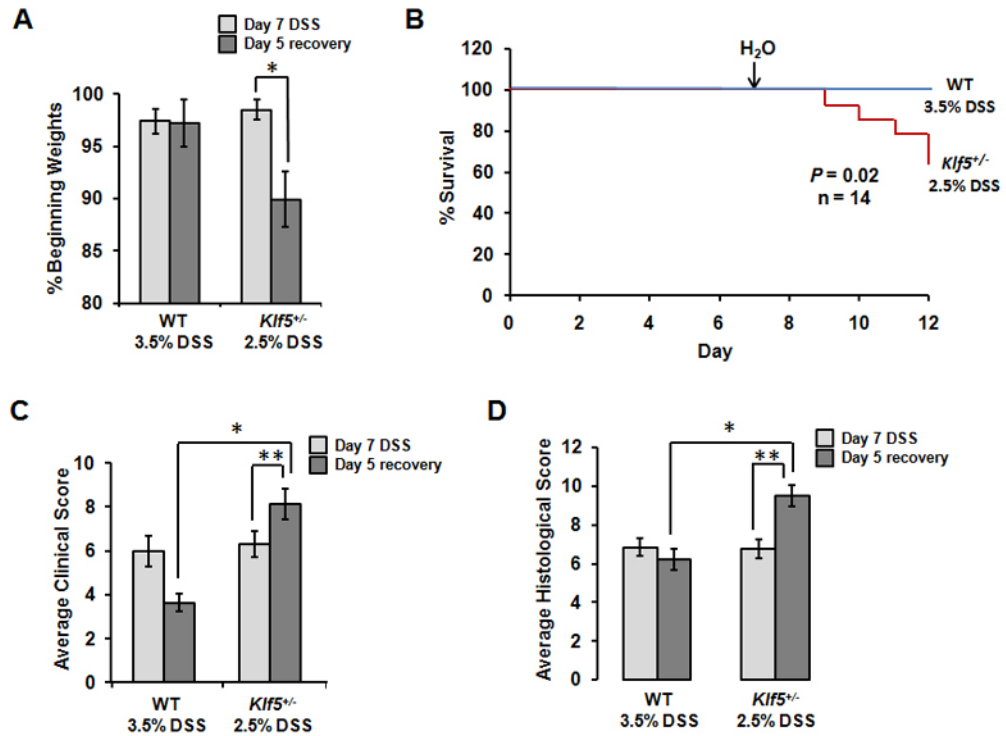


Figure 5. *Klf5*^{+/-} mice show impaired recovery during the DSS recovery phase. WT and *Klf5*^{+/-} mice were treated with DSS (3.5% and 2.5%, respectively) for 7 days to achieve similar degrees of colitis prior to 5 day recovery phase. (A) Weights of WT and *Klf5*^{+/-} mice after 7 days of DSS treatment and after 5 days of recovery (n = 14, * P = 0.02.) (B) Kaplan-Meier survival curve of WT and *Klf5*^{+/-} mice during DSS treatment and recovery phase (n = 14; P = 0.02). (C&D) Clinical and histological scores after 7 days of DSS treatment or after 5 days of recovery. (C) n = 7; * P = 0.003; ** P = 0.05 (D) n = 7; * P < 0.001; ** P = 0.006.

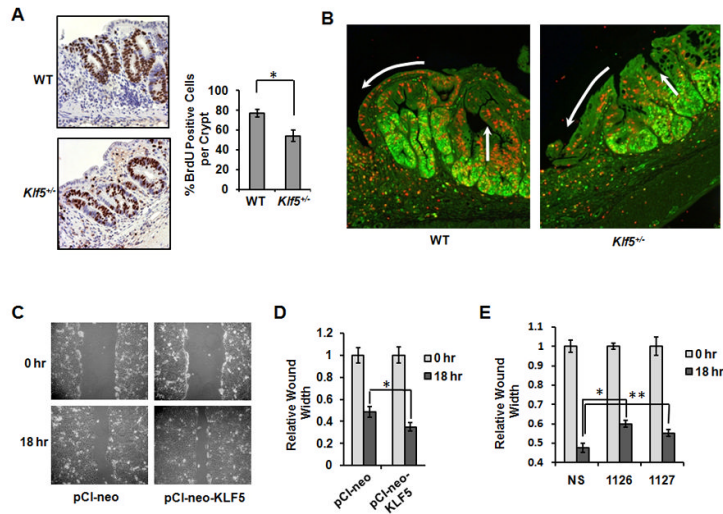


Figure 6. Epithelial proliferation and migration are reduced at sites of ulceration in *Klf5*^{+/-} mice. Colon tissues were isolated after 5 days of recovery and subjected to immunohistochemical staining. (A) BrdU staining of tissues from mice injected with BrdU 4 hrs prior to sacrifice. Images show crypts in regions adjacent to ulcers. Cell counts are from 6 crypts per mouse, (**P* = 0.009; n = 5). (B) Immunofluorescence staining of Ki67 (green) and BrdU (red) in colon tissues from mice injected with BrdU 24 hrs prior to sacrifice. White arrows indicate areas of migration. (C) Wounding assays in DLD-1 cells transfected with empty vector (pCI-neo) or an expression construct for KLF5 (pCI-neo-KLF5). (D) Graphical data of relative wound widths from (C). **P* = 0.01, n=3. (E) Wounding assays in DLD-1 cells transfected with non-specific siRNA (NS) or two different siRNAs specific for KLF5 (1126 or 1127). **P* < 0.001; n=3. ***P* = 0.03; n=3.

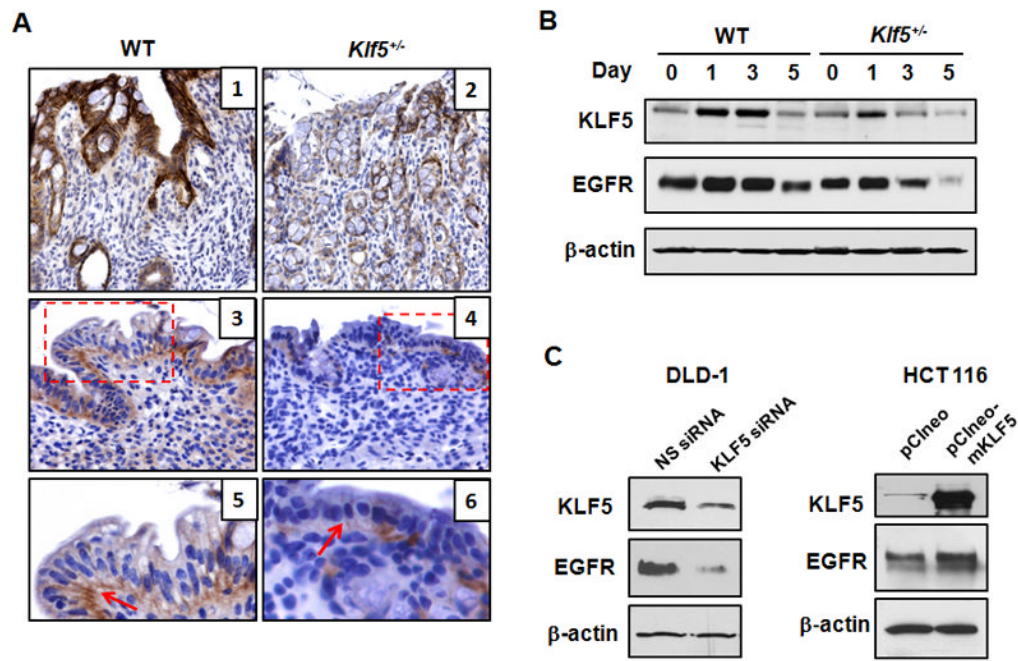


Figure 7. EGFR levels are reduced at sites of ulceration in *Klf5*^{+/-} mice treated with DSS. (A) EGFR staining in colons of mice after 5 days of DSS treatment (acute phase); panel 1, WT; panel 2, *Klf5*^{+/-}. EGFR staining after 5 days of recovery from DSS treatment. Panel 3, WT (enlargement, panel 5); panel 4, *Klf5*^{+/-} (enlargement, panel 6) Red arrows indicate luminal epithelial cells. (B) Western blots of colon lysates from WT and *Klf5*^{+/-} mice treated with 3.5% DSS, days 0–5. Lysates are pooled from 3 separate mice. (C) Effects of Klf5 expression on EGFR levels in colon cancer cells. Western blot analysis with inhibition of KLF5 by siRNA in DLD-1 cells or overexpression of KLF5 in HCT 116 cells.