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Exogenous Shh Protein Does Not Rescue Cultured Intestine from Atresia Formation

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Abstract

Introduction—The mechanism of intestinal atresia formation remains undefined. Atresia in *Fgfr2IIIb*^{-/-} mutant mouse embryos is preceded by endodermal apoptosis and involution of the surrounding mesoderm. We have observed that involution of the atretic segment is preceded by down regulation of Sonic hedgehog (SHH) in the endoderm which is a critical organizer of the intestinal mesoderm. We hypothesized that supplementation of *Fgfr2IIIb*^{-/-} intestinal tracts with exogenous SHH protein prior to atresia formation would prevent involution of the mesoderm and rescue normal intestinal development.

Methods—In situ hybridization were performed on control and *Fgfr2IIIb*^{-/-} intestinal tracts for *Shh* or *Foxf1* between embryonic (E) day 11.5 and E12.0. Control and *Fgfr2IIIb*^{-/-} intestinal tracts were harvested at E10.5 and cultured in media supplemented with FGF10 + SHH, or FGF10 with a SHH-coated bead. In situs were performed at E12.5 for *Foxf1*.

Results—*Shh* and *Foxf1* expression were down-regulated during intestinal atresia formation. Media containing exogenous FGF10 + SHH did not prevent colonic atresia formation (involution). A SHH protein point source bead did induce *Foxf1* expression in controls and mutants.

Discussion—*Shh* and *Foxf1* expression are disrupted in atresia formation of distal colon, thereby serving as potential markers of atretic events. Application of exogenous SHH (in media

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supplement or as a point source bead) is sufficient to induce *Foxf1* expression but insufficient to rescue development of distal colonic mesoderm in *Fgfr2IIIb*^{-/-} mutant embryos. *Shh* signal disruption is not the critical mechanism by which loss of *Fgfr2IIIb* function results in atresia formation.

Keywords

Shh; intestinal atresia; expression; *Foxf1*; *Fgfr2IIIb*^{-/-}; organ culture

Introduction

The mechanism of intestinal atresia remains undefined. Homozygous mutation of *Fibroblast growth factor receptor 2IIIb* (*Fgfr2IIIb*) early in intestinal development causes atresia formation in the colon of mice (1). Atresia formation is preceded by endodermal apoptosis in the areas where the atresia will form (1, 2). *Fgfr2IIIb* expression is limited to the intestinal endoderm at early stages of development (3) indicating that loss of the receptor in the endoderm causes endodermal apoptosis, loss of intestinal endoderm, and involution of the affected segment of intestine as a result of a loss of instructive signal to the surrounding mesoderm.

Shh is expressed in the early intestinal endoderm. It is a critical organizer of the intestine during development, instructing both radial and longitudinal growth (4, 5). Loss of *Shh* expression results in impaired organization of the intestinal mesoderm, although the intestinal tube remains continuous and does not form atresias. Tissue specific and spatial-temporal overlap is seen in the expression of *Shh* and *Fgfr2IIIb*, each performing a critical role in instructing intestinal development. Therefore, we hypothesized that following endodermal apoptosis, *Shh* signaling would be disrupted in the atretic precursor region (where the atresia will form).

We set out to investigate the role of *Shh* signaling pathway disruptions during atresia formation by examining expression patterns of *Shh* and its downstream mesodermal target, *Foxf1*, (6–8) by in situ hybridization prior to involution of the atretic segment. Further, we tested whether the addition of exogenous SHH protein to cultured embryonic intestines prior to atresia formation would rescue mesodermal development, prevent involution of the intestinal tube, and halt intestinal atresia formation.

Materials and Methods

Animals

IACUC approval for these studies was obtained from the University of Wisconsin School of Medicine and Public Health (P.F.N. protocol # M02258). All animals were maintained in a clean facility with *ad libitum* access to fresh food and water under a 12-hour alternating light/dark cycle.

Generation of mutant fetuses

Fgfr2IIIb^{-/-} mutant and *Fgfr2IIIb*^{+/-} littermate control embryos were generated using the *HprtCre* breeding strategy (9) as has been described previously (2).

Whole mount in situ hybridization

Fgfr2IIIb^{+/-} and *Fgfr2IIIb*^{-/-} embryos were harvested at Embryonic Day (E) 11.5 and E12.0 into cold PBS and fixed overnight in 4% PFA at 4°C. Fixed samples were dissected and dehydrated to 100% MeOH through a series of escalating Methanol/PBS-Tween steps

and stored at -20°C . The in situ hybridization protocol has been published elsewhere (10) and included incubation with antisense riboprobes, at 68°C for *Shh* and 70°C for *Foxf1* (constructs kindly provided by H. Hamada and Y. Saijoh). Photographs were taken using a dissecting light microscope.

Organ culture

For the media supplemented SHH protein experiments, *Fgfr2IIIb*^{+/-} and *Fgfr2IIIb*^{-/-} embryos were harvested at E10.5 and the developing intestinal tracts were isolated. Intestinal tracts were cultured in Matrigel (BD Biosciences, Bedford, MA) and allowed to polymerize at 37°C for 30 minutes within Millicell EZslide wells (Millipore, Billerica, MA). Matrigel embedded tracts were overlaid with a base media of DMEM/F-12 (HyClone, Logan, UT) containing L-Glutamine, penicillin/streptomycin, and fetal bovine serum. In this set of experiments, the base culture media was supplemented with FGF10 (PeproTech, Rocky Hill, NJ) and SHH (R&D Systems, Minneapolis, MN) to a final concentration of 500 ng/mL each. Media-supplemented organ culture were conducted 5 times following previously published protocols (10), with 12 normal (*Fgfr2IIIb*^{+/-}) littermate controls and 8 mutant (*Fgfr2IIIb*^{-/-}) tracts.

For the point source SHH protein experiments, E10.5 intestinal tracts were harvested as described above. These 3 additional culture experiments included 17 littermate controls and 9 mutant intestinal tracts. Intestinal explants were cultured in Matrigel with SHH-laden Affi-Gel beads (Bio-Rad, Hercules, CA, 153-7302) at 80 to 150 μm diameter. Beads were incubated with approximately 50 $\mu\text{g/mL}$ of SHH protein before being polymerized within Matrigel and overlaid with media containing 500 ng/mL of FGF10 within EZslide wells. Each well contained at least 1 control tract, 1 mutant tract and 1 SHH-laden bead. All organ cultures were maintained at 37°C and 5% CO_2 conditions. Media was refreshed after 24 hours. After 48 hours of culture, tissues were fixed with 4% PFA within culture wells and then processed for in situ hybridization with a *Foxf1* antisense riboprobe at 70°C . Photographs were taken at time of culture (E10.5 + 0 hours), E10.5 + 24 hours, and E10.5 + 48 hours, under a dissecting light microscope.

Results

At E11.5 on littermate *Shh* expression was down regulated in the colons of mutant *Fgfr2IIIb*^{-/-} embryos compared to controls but expressed equaling the small intestine of *Fgfr2IIIb*^{+/-} and *Fgfr2IIIb*^{-/-} embryos (Figure 1A and B).

To determine if the above colonic *Shh* expression loss was associated with a downstream signaling disruption, *Foxf1* expression was examined by whole mount in situ hybridization at E12.0. *Fgfr2IIIb*^{-/-} embryos exhibited a complete absence of *Foxf1* expression in the atretic precursor region (mid and distal colon) in contrast to controls (Figure 2A and B), indicating disruptions in the *Shh* signaling cascade in the atretic precursor region of the colon.

To determine whether exogenous SHH protein could rescue *Fgfr2IIIb*^{-/-} colons from an atretic fate, intestinal explants were cultured in the presence of SHH, either as a media supplement or as a point source on an Affi-Gel bead. In the presence of SHH protein, control *Fgfr2IIIb*^{+/-} intestines developed normally, exhibiting cecal bud formation and presence of an intact colon (Figure 3A-C). In contrast, *Fgfr2IIIb*^{-/-} colons failed to develop normally with the colons ending in a blind end (Figure 3). Exogenous SHH alone could not rescue colonic development from an atretic fate. When SHH was supplied on an Affi-Gel bead, *Foxf1* expression was induced at the boundary of the endoderm and mesoderm of the control intestines and along the surface of the mesoderm of the *Fgfr2IIIb*^{-/-}

– intestine. Control intestines maintained normal architecture with both the colon and the proximal intestine maintaining an open tube at either end. In contrast, *Fgfr2IIIb*^{-/-} colons ended blindly at the cecal region indicating the formation of an atresia (Figure 4).

Discussion

The atretic precursor region of the colon exhibits loss of *Shh* signaling prior to involution in *Fgfr2IIIb*^{-/-} mutants. The absence of *Shh* expression is nearly pan colonic at this stage (Figure 1). Further, *Foxf1* expression is absent in the mid and distal colon 12 hours after the loss of *Shh* expression (Figure 2). Interestingly, there is some preservation of *Foxf1* expression in the proximal colon of *Fgfr2IIIb*^{-/-} mutants. We suspect that the expression of *Foxf1* in the proximal colon may be a result of *Shh* expression arising from the distal small intestine (precursor of the terminal ileum) and acting longitudinally over a short distance.

Exogenous SHH protein, however, fails to rescue colon explants from an atretic fate in organ culture (Figures 3–4) regardless of method of supplementation: addition to culture media or via a SHH impregnated bead in proximity to the intestinal tract. The data suggest that the loss of intestinal endoderm through apoptosis leads to the absence of *Shh* expression and downstream *Foxf1* expression, which is critical for mesodermal development (4, 5). However, simply restoring SHH protein availability is insufficient to rescue development. The data also suggest that exogenous SHH protein does not prevent endoderm loss, directly or indirectly, via a feedback loop from the mesoderm to the endoderm.

In summary, atresia formation arising from homozygous mutation of *Fgfr2IIIb* is associated with endodermal apoptosis as reported previously (1,2) and with a loss of *Shh* expression and signaling. However, our data indicate that disruptions in *Shh* signaling are not the critical event that leads to the involution of the atretic precursor. In fact, supplementation with exogenous *Shh* protein can restore some level of *Shh* signaling in the mesoderm as evidenced by induction of *Foxf1* expression in the cultured intestinal tracts of *Fgfr2IIIb*^{-/-} mutants. Yet this is insufficient to rescue intestinal development. The findings indicate that disruptions in *Shh* signaling may serve as a marker, but do not support them as a critical event in atresia formation.

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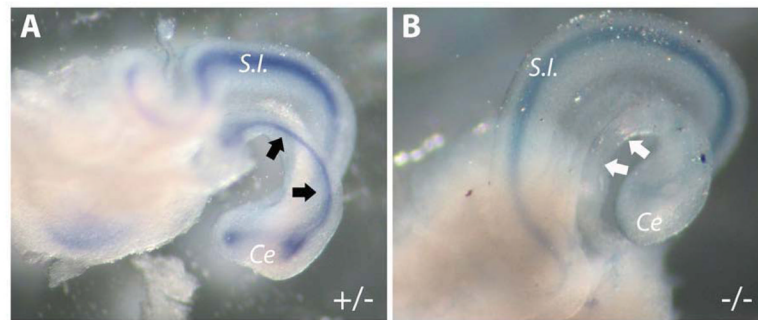


Figure 1. *Shh* Expression is absent in *Fgfr2IIIb*^{-/-} colon at E 11.5

Whole mount in situ hybridization for *Shh* in (A) control *Fgfr2IIIb*^{+/-} and (B) mutant *Fgfr2IIIb*^{-/-} colons. Presence of staining in control colon is indicated by black arrows, while absence of staining in mutant colon is indicated by white arrows. (Ce) cecum, (S.I.) small intestine.

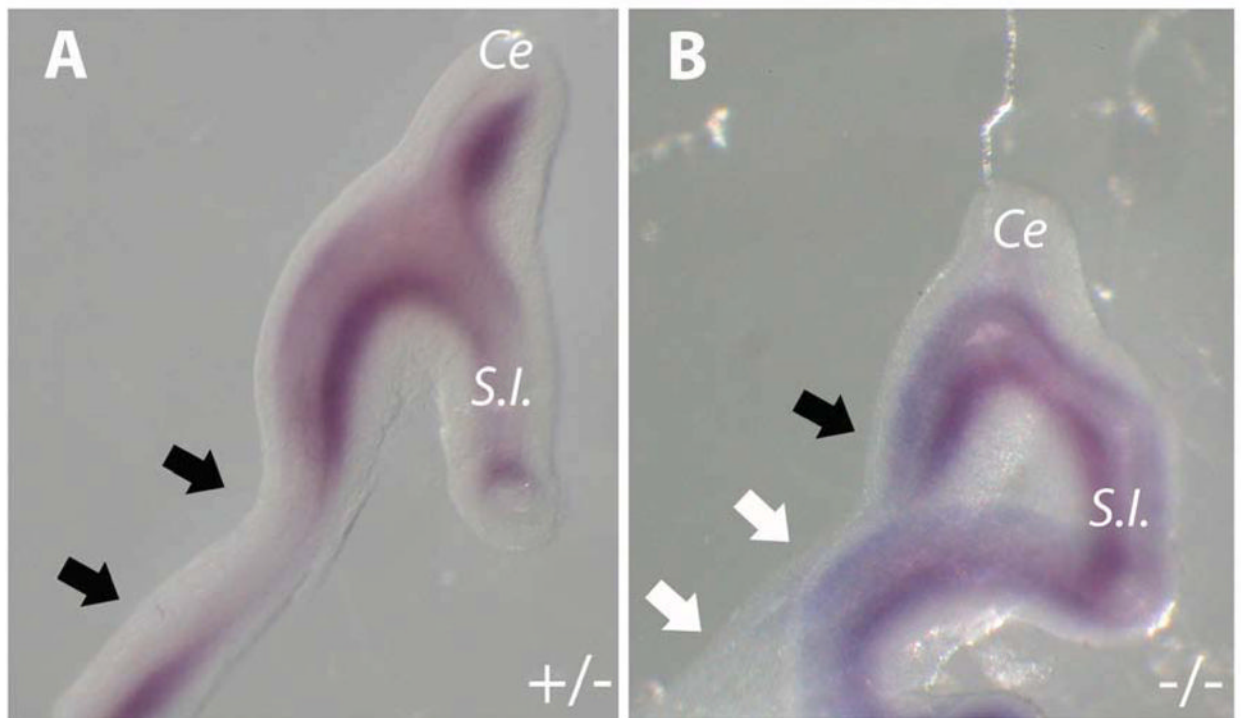


Figure 2. *Foxf1* Expression is absent in the mid and distal colon of *Fgfr2IIIb*^{-/-} embryos E 12.0
Whole mount in situ hybridization for *Foxf1* in (A) control *Fgfr2IIIb*^{+/-} and (B) mutant *Fgfr2IIIb*^{-/-} colons. Presence of staining in colon is indicated by black arrows, while absence of staining is indicated by white arrows. (Ce) cecum, (S.I.) small intestine.

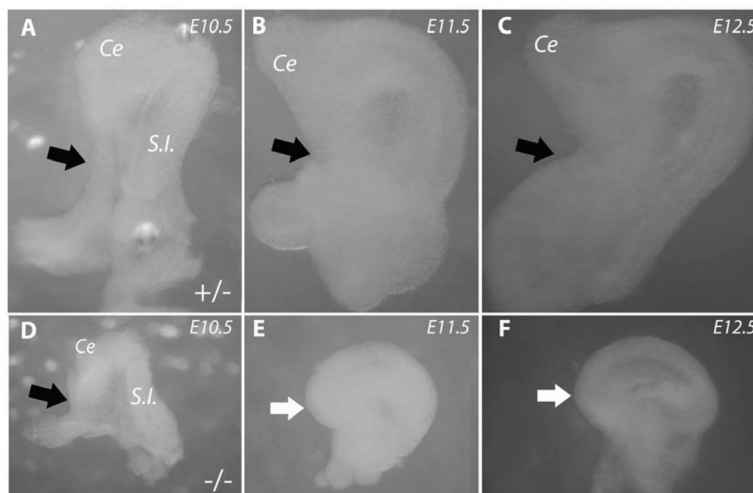


Figure 3. Media supplemented with exogenous SHH protein does not rescue cultured *Fgfr2IIIb*^{-/-} – colons from an atretic fate

Whole mount photographs of control (A–C) and mutant (D–F) colons at E10.5 (A and D), E10.5 + 24 hour culture (B and E) and E10.5 + 48 hours (C and F) in the presence of SHH protein. Colonic development (Black arrows) proceeds normally in *Fgfr2IIIb*^{+/-} controls. In *Fgfr2IIIb*^{-/-} mutants the colon ends blindly (White arrows).

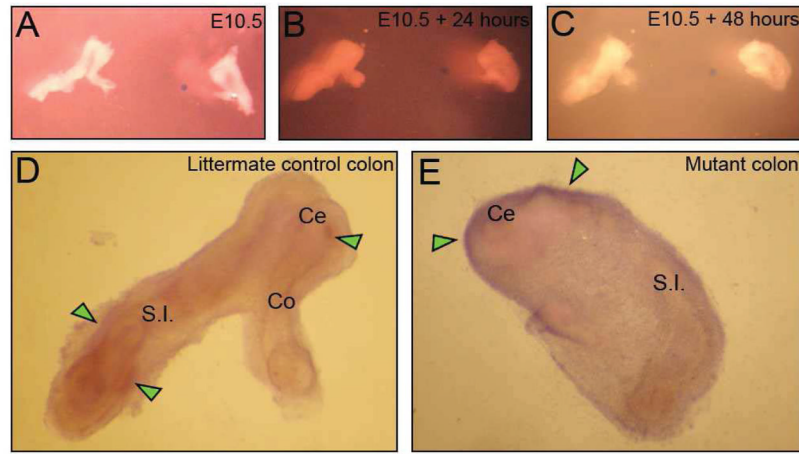


Figure 4. Exogenous SHH protein provided on an Affi-Gel bead does not rescue cultured *Fgfr2IIIb*^{-/-} colons from an atretic fate

Representative whole mount photographs of littermate control and mutant colon at E10.5 (A), E10.5 + 24 hours (B), E10.5 + 48 hours (C), in the presence of a SHH-laden bead (blue circle to the right of center), and *Foxf1* in situ after 48 hours of culture (D and E). Control intestine and colon display normal architecture and defined *Foxf1* expression in the mesoderm adjacent to the endoderm (Green arrows indicate *Foxf1* staining at the boundary between the endoderm and mesoderm) (D). *Fgfr2IIIb*^{-/-} mutant colon ends blindly and expresses *Foxf1* on the surface of the mesoderm (Green arrows) (E). (Ce) cecum, (Co) colon, (S.I.) small intestine.