

Analysis of Enteric Neural Crest Cell Migration Using Heterotopic Grafts of Embryonic Guts

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[Abstract] Hirschsprung disease (HSCR), also named aganglionic megacolon, is a severe congenital malformation characterized by a lack of enteric nervous system (ENS) in the terminal regions of the bowel (Bergeron *et al.*, 2013). As the ENS notably regulates motility in the whole gastrointestinal track, the segment without neurons remains tonically contracted, resulting in functional intestinal obstruction and accumulation of fecal material (megacolon). HSCR occurs when enteric neural progenitors of vagal neural crest origin fail to fully colonize the developing intestines. These “enteric” neural crest cells (ENCCs) have to migrate in a rostro-caudal direction during a fixed temporal window, which is between embryonic day (e) 9.5 and e14.5 in the mouse (Obermayr *et al.*, 2013). Recently, our group generated a new HSCR mouse model called Holstein in which migration of ENCCs is impaired because of increased collagen VI levels in their microenvironment (Soret *et al.*, 2015). Here, we describe the method that allowed us to demonstrate the cell-autonomous nature of this migration defect. In this system adapted from a previously described heterotopic grafting approach (Breau *et al.*, 2006), the donor tissue is a fully colonized segment of e12.5 midgut while the host tissue is an aneural segment of e12.5 hindgut. Extent of ENCC migration in host tissue is assessed after 24 h of culture and is greatly facilitated when donor tissue has a transgenic background such as the Gata4-RFP (Pilon *et al.*, 2008) that allows endogenous labeling of ENCCs with fluorescence. Depending of the genetic background of donor and host tissues, this approach can allow evaluating both cell-autonomous and non-cell-autonomous defects of ENCC migration.

Materials and Reagents

1. Petri dishes (Corning, catalog number: 70165-102)
2. 24-well plate
3. Nitrocellulose filter (Merck Millipore, catalog number: GSWP01300)
4. 8-chamber slides (ibiTreat μ -slide) (ibidi GmbH, catalog number: 80826)
5. Mature mice (\geq 2-month old)
6. Isoflurane for inhaled anesthesia (Henry Schein Animal Health, catalog number: 050031)
7. 70% ethanol
8. 1x phosphate-buffered saline (PBS)
9. DMEM/Ham's F-12 (WISSENT, catalog number: 319-085-CL)
10. Fetal Bovine Serum (WISSENT, catalog number: 920-040)

11. Penicillin/streptomycin (WISENT, catalog number: 450-201-EL)

Equipment

1. Dumont #5 dissection forceps (Fine Science Tools, catalog number: 11251-20)
2. Dumont #7 dissection forceps (Fine Science Tools, catalog number: 11274-20)
3. Dissection scissors (Moria Spring Scissors) (Fine Science Tools, catalog number: 15396-01)
4. Dissecting stereomicroscope (Leica Microsystems, model: M125)
5. CO₂ cell culture incubator (Sanyo Scientific, model: MCO-18AIC)
Note: This product has been discontinued.
6. Infinity-2 camera (Lumenera Corporation) mounted on a fluorescent stereomicroscope (Leica Microsystems, model: M205FA)

Software

1. ImageJ software

Procedure

1. Mate mature mice (\geq 2-month old) overnight and check for the presence of a vaginal plug the next morning. Noon of the day a vaginal plug is observed is considered embryonic day (e) 0.5. A typical experiment requires one wildtype couple and one couple bearing the mutation to study.
2. Twelve days later, euthanize isoflurane-anesthetized pregnant female(s) via CO₂ inhalation.
3. Use 70% ethanol to spray the mouse abdomen and open it with dissecting scissors to access the uterus.
4. Remove the uterus into a glass petri dish containing 15 ml of sterile ice-cold PBS and cut it between individual embryos.
5. Working on each embryo separately in another glass petri dish filled with 15 ml of ice-cold PBS, use fine forceps to remove the uterine muscle layers under a dissecting microscope (Figure 1 – step 1 and Video 1).
6. Open the extra-embryonic membranes to access the embryo by taking care not to sever the developing intestines that are intertwined with the blood vessels connecting the embryo to the placenta/extra-embryonic membranes (Figure 1 - step 2 and Video 1).
7. Cut embryo's head which can be used for PCR-based genotyping if necessary and open the abdominal cavity (Figure 1 - steps 3-4 and Video 1).
8. After removing the liver, cut the esophagus and pull the whole gastrointestinal tract out of the abdominal cavity while being careful not to damage the colon still attached to the anus (Figure 1 - step 5 and Video 1).

- Cut at the anus to free the gastrointestinal tract. The stomach and cecum will allow keeping track of sample's orientation (Figure 1 - step 6 and Video 1).

Video 1. Whole dissection procedure



- Transfer each sample into a well of a 24-well plate containing 1 ml of sterile PBS and kept on ice. If PCR genotyping is required, be careful to match intestine and head numbers for each sample.
- Assemble each graft by depositing and juxtaposing donor (wildtype or mutant) and host (wildtype or mutant) tissues onto a small 13 mm nitrocellulose membrane in a glass petri dish containing 15 ml of sterile ice-cold PBS. Prepare each half of a graft in a sequential manner, so that the first half has already adhered to the filter before processing the second half. Take care to respect the normal rostro-caudal orientation (esophagus to anus) of both the donor and host tissues in the assembled graft. Host tissue from a selected sample is prepared by cutting a ~0.5 cm section of the most caudal hindgut whereas donor tissue is prepared from another sample by cutting a ~0.5 cm section of the midgut just upstream of the cecum.
- Transfer each graft-bearing filter in a chamber of an 8-chamber slide containing 250 μ l of DMEM/F12 supplemented with 10% FBS and penicillin/streptomycin (Figure 2A).
- After 24 h of culture (37 °C, 5% CO₂), colonization of aneural hindgut tissues by midgut-derived ENCCs is quantified by measuring the distance separating the most distal ENCC from the midgut-hindgut graft junction (Figure 2B).

Representative data

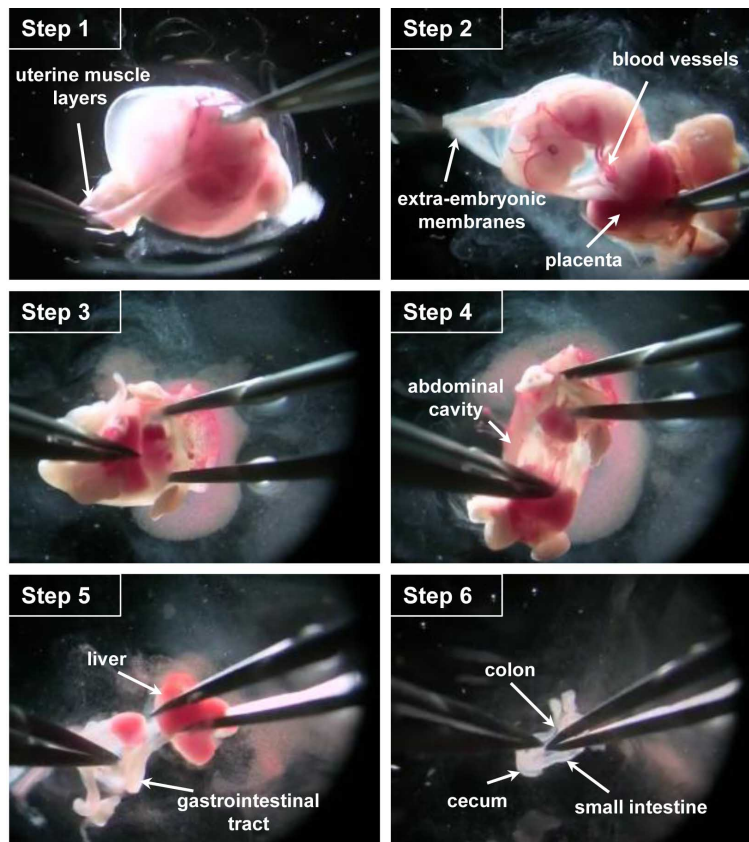


Figure 1. Key steps of the dissection procedure. Step 1. With fine forceps, remove the uterine muscle layers. Step 2. Open the extra-embryonic membrane to access the embryo, cut the blood vessels connecting the embryo to the placenta/extra-embryonic membranes. Step 3. Cut the embryo head and open the abdominal cavity. Step 4. Pull the whole gastrointestinal tract out of the abdominal cavity. Step 5. Remove the liver, cut the esophagus and release the digestive tract of the connective tissue. Step 6. Free the gastrointestinal tract by cutting it at the anus.

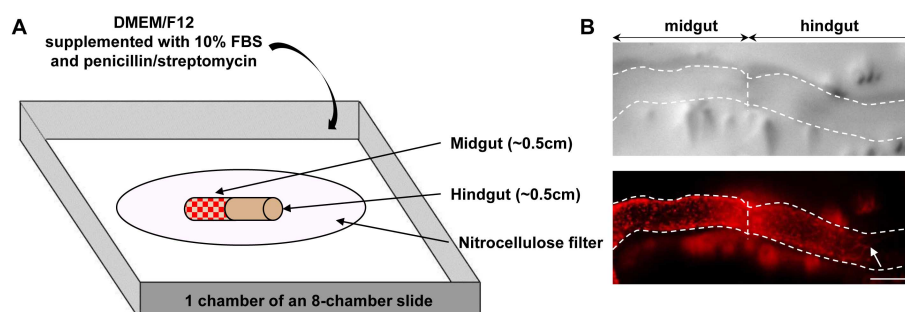


Figure 2. Overview of graft assembly and representative result. A. Schematic representation of a heterotopic e12.5 midgut-hindgut graft onto a nitrocellulose filter in a chamber of an 8-chamber slide. Red dots in midgut tissue represent ENCCs labeled by RFP

(DsRed2) fluorescence provided by the G4-RFP transgene; B. Representative images of a graft (delineated by dotted lines) after 24 h of culture, showing the colonization of a previously aneural hindgut host tissue by fluorescently labeled ENCCs from midgut donor tissue. Pictures were taken using an Infinity-2 camera mounted on a Leica M205FA fluorescent stereomicroscope and images were analyzed using the ImageJ software. The white arrow points to the location of the migration front at the end of the culture period. Scale bar: 150 μ m.

Notes

It is noteworthy that this method is greatly simplified when donor tissues are taken from embryos bearing a transgene such as the G4-RFP transgene (Pilon *et al.*, 2008) that labels ENCCs with fluorescence. At step 10, this can allow the identification of mutant tissues (*i.e.*, displaying delayed migration) by simple fluorescent microscopy instead of having to wait for genotyping results after graft assembly. Moreover, at step 13, such an intrinsic fluorescent labeling greatly facilitates the analysis of ENCC migration which otherwise requires immunofluorescence labeling using an antibody against a marker of undifferentiated enteric neural progenitors such as Sox10.

Acknowledgments

The Pilon laboratory is funded by grants from the Canadian Institute of Health Research (CIHR), the Natural Science and Engineering Research Council of Canada (NSERC) and the Fondation du grand défi Pierre Lavoie. RS holds a fellowship from the Fonds de la recherche du Québec Santé (FRQS) whereas NP is an FRQS Junior 2 Research Scholar as well as the recipient of a UQAM Research Chair on Rare Genetic Diseases. The authors thank the innovative work performed by Breau *et al.* (2006), on which this protocol was based.

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