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3 **Enterendoctrine cells couple nutrient sensing to nutrient absorption by regulating ion**  
4 **transport.**

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20 **Summary**

21 The ability to absorb ingested nutrients is an essential function of all metazoans and utilizes a  
22 wide array of nutrient transporters found on the absorptive enterocytes of the small intestine. A  
23 unique population of patients has previously been identified with severe congenital  
24 malabsorptive diarrhea upon ingestion of any enteral nutrition. The intestines of these patients  
25 are macroscopically normal, but lack enteroendocrine cells (EECs), suggesting an essential role  
26 for this rare population of nutrient-sensing cells in regulating macronutrient absorption. We used  
27 human and mouse models of EEC deficiency to identify a new role for the EEC hormone  
28 peptide YY in regulating ion-coupled absorption of glucose and dipeptides the small intestine.  
29 We found that peptide YY is required in to maintain normal electrophysiology in the presence of  
30 vasoactive intestinal polypeptide, a potent stimulator of ion secretion produced by enteric  
31 neurons. Administration of peptide YY to EEC-deficient mice restored normal electrophysiology,  
32 improved glucose and peptide absorption, diminished diarrhea and rescued postnatal survival.  
33 These data suggest that peptide YY is a key regulator of macronutrient absorption in the small  
34 intestine and may be a viable therapeutic option to treat patients with malabsorption.

36

37 **Keywords:** enteroendocrine, organoid, malabsorption, peptide YY, NEUROG3, vasoactive  
38 intestinal polypeptide, ion transport, diarrhea, nutrient absorption

39 **Introduction**

40 Enteroendocrine cells (EECs) are a rare population of cells found in the gastrointestinal  
41 epithelium that sense nutrients that are passing through the gut and in response secrete more  
42 than 20 distinct biologically active peptides. These peptides act in an endocrine or paracrine  
43 fashion to regulate all aspects of nutrient homeostasis including satiety, mechanical and  
44 chemical digestion, nutrient absorption, storage and utilization (Gribble and Reimann, 2019).  
45 Humans (Wang et al., 2006) and mice (Mellitzer et al., 2010) with genetic mutations that impact  
46 formation or function of EECs have intractable malabsorptive diarrhea, metabolic acidosis, and  
47 require parenteral nutrition or small-bowel transplant for survival. These findings were the first to  
48 link EECs to the absorption of macronutrients; however, the mechanism by which EECs  
49 contribute to this vital process is unknown. Poor absorption of macronutrients is a global health  
50 concern, with underlying etiology including short-gut syndrome, enteric pathogen infection, and  
51 malnutrition. Therefore, identification of factors regulating nutrient absorption has significant  
52 therapeutic potential.

53 Absorption of carbohydrate and protein requires coordinated activity of nutrient and ion  
54 transporters in the small intestine. Glucose is primarily absorbed via sodium-glucose  
55 cotransporter SGLT1, which uses a downhill  $\text{Na}^+$  gradient to transport one glucose or galactose  
56 molecule with two sodium ions from the lumen into the enterocyte (Wright et al., 2011). The  
57 majority of dietary protein absorption occurs via  $\text{Na}^+$ - and  $\text{H}^+$ -linked amino acid transporters and  
58 PEPT1, which imports di- and tri-peptides coupled with a hydrogen ion (Chen et al., 2010). The  
59 electrochemical gradients that drive nutrient absorption are maintained in part by ion  
60 transporters, including the cystic fibrosis transmembrane receptor (CFTR), which exports  
61 chloride (Wright et al., 1997), and sodium-hydrogen exchanger NHE3, which maintains  $\text{Na}^+$  and  
62  $\text{H}^+$  microclimates across the apical membrane (Thwaites et al., 2002). Activity of CFTR and

63 NHE3 are, in turn, regulated by levels of cyclic AMP (cAMP) (Burleigh and Banks, 2007; Yun et  
64 al., 1997).

65 Most secreted EEC peptides signal via G protein-coupled receptors that act via second  
66 messenger cascade effectors like cAMP. Given the requirement for EECs in regulating  
67 macronutrient absorption, we investigated the possibility that EEC-derived peptides coupled  
68 nutrient sensing to nutrient absorption by regulating electrogenic transport in neighboring  
69 enterocytes. Two well-studied peptides governing ion and water homeostasis in the colon are  
70 vasoactive intestinal peptide (VIP) and peptide YY (PYY). VIP, secreted from enteric neurons,  
71 signals via the  $G_{\alpha s}$ -coupled VIPR1 (VPAC1) on epithelial cells to raise levels of intracellular  
72 cAMP. In contrast, EEC-derived PYY acts in a paracrine fashion on colonocytes to lower cAMP  
73 via the epithelial  $G_{\alpha i}$  coupled receptor NPY1R (Cox et al., 2010; Hyland et al., 2003; Moodaley  
74 et al., 2017; Tough et al., 2011). We posited that the mechanism underlying malabsorptive  
75 diarrhea in patients lacking EECs might be due to loss of a similar EEC-ENS regulatory  
76 feedback in the small intestine, thus disrupting electrogenic nutrient absorption. Here, we found  
77 that PYY regulates normal ion transport and ion-coupled nutrient absorption in mouse and  
78 human small intestine, and that administration of exogenous PYY was sufficient to restore  
79 normal electrophysiology, nutrient absorption, and survival in EEC-deficient animals.

80

## 81 **Results**

### 82 **The PYY-VIP axis regulates ion and water transport in mouse and human small intestine.**

83 If EECs were required for regulating the normal electrophysiology of the small intestine,  
84 we would expect to see deranged ion transport in intestinal tissues lacking EECs. To investigate  
85 this, we used EEC-deficient mice (*VillinCre;Neurog3<sup>fl/fl</sup>*) (Mellitzer et al., 2010) and three  
86 different human small intestinal tissue models all derived from pluripotent stem cells (PSCs):  
87 human intestinal organoids (HIOs) derived *in vitro* (Spence et al., 2011), HIOs that were

88 matured to robust crypt-villus architecture *in vivo* (Watson et al., 2014), and epithelial organoids  
89 (enteroids) derived from crypts of matured HIO tissues (Watson et al., 2014). We generated  
90 EEC-deficient human small intestinal tissue by using PSC lines that had a null mutation in  
91 *NEUROG3* (McGrath et al., 2015), the basic helix-loop-helix transcription factor required for  
92 EEC formation in mice (Jenny et al., 2002) and humans (Wang et al., 2006). As previously  
93 reported (Zhang et al., 2019), *NEUROG3*<sup>-/-</sup> small intestinal organoids completely lacked EECs,  
94 but were otherwise normal in appearance (Figure S1).

95 Ion and water transport in the colon are regulated by EEC-derived PYY and ENS-  
96 derived VIP. To formally test whether the PYY-VIP axis operated in human and mouse small  
97 intestine, we performed experiments in EEC-deficient tissues without a functional ENS wherein  
98 we controlled PYY and VIP levels experimentally. We first determined the effects of the PYY-  
99 VIP axis on small intestine by measuring CFTR-mediated ion and water efflux (Dekkers et al.,  
100 2013) following exposure of human HIO-derived enteroids to the potent secretagogue VIP  
101 (Figure 1A). EEC-deficient enteroids swelled significantly more than did wild-type but blocking  
102 NPY1R in wild-type enteroids mimicked the EEC-deficient response (Figure 1A). Exogenous  
103 PYY blocked VIP-induced swelling in both wild-type and EEC-deficient enteroids in an NPY1R-  
104 dependent manner (Figure 1A), demonstrating that the PYY-VIP axis regulates ion and water  
105 secretion in human small intestine. We next tested the activity of NHE3 as a measure of Na<sup>+</sup>-  
106 dependent intracellular pH recovery after acidic challenge (Foulke-Abel et al., 2016) and found  
107 that EEC-deficient enteroids displayed impaired NHE3 function (Figure 1B). There was no  
108 difference in expression of *CFTR*, *SLC9A3* (encoding NHE3), *VIPR1* or *NPY1R* between wild-  
109 type and EEC-deficient human small intestinal organoids or enteroids (Figure 1C and Figure  
110 S2A-B). Together, these data suggest that PYY plays an important role in the regulation of ion  
111 transport in the small intestine, and that the abnormal response to VIP in EEC-deficient  
112 enteroids can be normalized by the addition of exogenous PYY.

113            If PYY were required to regulate electrochemical transport in the small intestine, we  
114    would expect that disruption of PYY signaling in wild-type small intestinal tissue would cause  
115    abnormal basal short-circuit current ( $I_{sc}$ ). To investigate this we isolated full thickness intestinal  
116    mucosa from *in vivo* matured human intestinal organoids and from the jejunum of wild-type mice  
117    and measured basal  $I_{sc}$  in a modified Ussing chamber (Clarke, 2009). Chemical inhibition of  
118    NPY1R in wild-type mouse jejunum and human intestinal organoids was sufficient to elevate the  
119    basal  $I_{sc}$  to EEC-deficient levels (Figure S2C). Conversely, treatment of EEC-deficient mouse  
120    and human tissues with exogenous PYY reduced the basal  $I_{sc}$  to wild-type levels in an NPY1R-  
121    dependent manner (Figure S2C). These data indicated that endogenous PYY signaling plays an  
122    essential role in maintaining normal electrophysiology in the small intestine.

123            We then investigated if PYY was required to modulate the stimulatory effects of VIP in  
124    mouse and human small intestine. We inhibited voltage-gated neuronal firing in mouse jejunum  
125    by including tetrodotoxin (Hyland et al., 2003) in all experiments so that we could precisely  
126    monitor epithelial response to exogenous VIP. Chemical inhibition of NPY1R in isolated wild-  
127    type tissues was sufficient to cause an elevated response to VIP (Figure 1D). This indicated that  
128    endogenous PYY signaling was required in the small intestine to modulate the stimulatory  
129    effects of VIP. Consistent with this, EEC-deficient mouse and human small intestinal tissue  
130    similarly displayed an exaggerated  $I_{sc}$  response to exogenous VIP compared to wild-type  
131    (Figure 1D). Addition of exogenous PYY to EEC-deficient small intestine was sufficient to  
132    restore the  $I_{sc}$  to normal (Figure 1D). These data suggested that PYY is required for maintaining  
133    a normal electrochemical response to VIP in the small intestine and that PYY can normalize this  
134    process in EEC-deficient small intestinal tissue. Furthermore, these data suggest that imbalance  
135    of this axis may be a mechanism underlying malabsorptive diarrhea suffered by patients without  
136    EECs.

137

138 **PYY restores normal glucose absorption in EEC-deficient human and mouse small  
139 intestine.**

140        While it is known that EECs sense nutrients, the mechanism linking sensing to the  
141 control of nutrient absorption is unclear. A hint came from the effects of enteral feeding of EEC-  
142 deficient patients, which resulted in a massive diarrheal response. This suggests that an inability  
143 to sense luminal nutrients uncoupled the ability to adequately absorb them. To explore this  
144 possibility we evaluated ion-coupled nutrient absorption in EEC-deficient small intestine. We  
145 observed an accelerated initial response to luminal glucose in the presence of VIP in EEC-  
146 deficient mouse and human intestinal tissues in the Ussing chamber (Figure 2C), as predicted if  
147 the normal electrochemical gradients were perturbed (Figure 2A-B). This recapitulated the  
148 exacerbated diarrhea observed in patients without EECs when they were fed with carbohydrate  
149 (Wang et al., 2006). Exogenous PYY restored a normal glucose response in EEC-deficient  
150 mouse and human tissue, and inhibition of NPY1R in wild-type caused an exaggerated initial  
151 response to glucose (Figure 2C). These data indicate that PYY is both necessary and sufficient  
152 to modulate glucose absorption in the small intestine. We found no defects in expression of  
153 SGLT1, GLUT2, (Figure 2D and S2A) or maximum absorptive competency of  $\text{Na}^+$ -coupled  
154 glucose transport (Figure 2E-G) in human epithelium without EECs. These data suggest that  
155 SGLT1 is competent to absorb glucose, but activity is dysregulated in the context of abnormal  
156 ion transport in the absence of EECs.

157

158  **$\text{H}^+$ -coupled dipeptide absorption is impaired in EEC-deficient small intestine.**

159        Approximately 80% of ingested amino acids were recovered in the stool of the index  
160 EEC-deficient patient (Wang et al., 2006), suggesting a critical role for EECs in regulating  
161 protein absorption. Consistent with this, we observed a striking loss of ion-coupled dipeptide  
162 absorption when human and mouse EEC-deficient small intestine were challenged with VIP

163 (Figure 3A), despite normal expression of PEPT1 (Figures 3B and S2A). VIP has an established  
164 role in inhibition of NHE3 and PEPT1-mediated dipeptide absorption (Anderson et al., 2003;  
165 Thwaites et al., 2002), but we were surprised to find that EEC-deficient intestine remained  
166 unable to respond to dipeptide when PYY was provided (Figure 3A). This suggested that  
167 dysregulated H<sup>+</sup> gradients may be a more stable phenotype in EEC-deficient intestine, and not  
168 easily reversed by PYY within minutes. To explore this possibility, we treated enteroids with or  
169 without PYY for one week *in vitro* in the presence of VIP. Wild-type enteroids were able to  
170 maintain their intracellular pH in the presence of VIP but EEC-deficient enteroids became  
171 significantly more acidic (Figure 3C). However, EEC-deficient enteroids were restored to normal  
172 intracellular pH levels and normal *SLC9A3* expression (encoding NHE3) in the presence of PYY  
173 (Figures 3C and S3). This suggested that long-term exposure to an imbalanced EEC-ENS axis  
174 dysregulates intestinal physiology, and that, over time, PYY may be sufficient to restore  
175 intracellular pH and dipeptide absorption in EEC-deficient small intestine.

176 We have demonstrated that inhibiting PYY signaling in isolated wild-type small intestinal  
177 tissues was sufficient to perturb normal electrophysiology in both human and mouse. This  
178 suggests that *in vivo* the mechanism of action of PYY could be paracrine rather than endocrine.  
179 PYY-expressing EECs are abundant in mouse and human small intestine (Egerod et al., 2012)  
180 (Figure S4). Moreover, PYY-expressing EECs extend long basal processes which underlie  
181 several neighboring epithelial cells (Bohorquez et al., 2014; Bohorquez et al., 2015), raising the  
182 possibility that they may exert paracrine effects on whole populations of nearby enterocytes. We  
183 therefore investigated whether the effects of PYY on ion transport in the small intestine occurred  
184 via paracrine mechanisms. To do this, we exploited the mosaicism of *VillinCre* mice to  
185 determine if regions of EEC-deficient epithelium had different transporter activities as compared  
186 to regions of epithelium that still had EECs. We observed in *VillinCre;Neurog3*<sup>flox/flox</sup> mice that  
187 4.38 $\pm$ 2.56% of jejunum escaped tdTomato labeling (Figure S5) and that in regions that had

188 EECs, neighboring enterocytes had a normal intracellular pH indicating normal ion transport. In  
189 contrast, enterocytes in EEC-deficient regions were significantly more acidic indicating  
190 perturbed H<sup>+</sup> transport (Figures 3D and S5). Together these data suggest that EECs control  
191 local H<sup>+</sup> transporter activity and dipeptide responsiveness in the small intestine via paracrine  
192 mechanisms.

193

194 **Exogenous PYY rescues EEC-deficient mice from malabsorptive diarrhea and death and  
195 restores normal glucose and dipeptide transport.**

196 As previously reported (Mellitzer et al., 2010), *VillinCre;Neurog3*<sup>flox/flox</sup> mice suffer from  
197 malabsorptive diarrhea and exhibit severely impaired postnatal survival, with only a small  
198 fraction of mice surviving weaning. Our data suggested that treatment with PYY might restore  
199 normal carbohydrate and protein absorption the intestines of EEC-deficient animals. We  
200 therefore used *VillinCre;Neurog3*<sup>flox/flox</sup> mice as a preclinical model to test if PYY could reverse  
201 malabsorptive diarrhea and improve postnatal survival (Figure 4A-B). We began daily treatment  
202 of mutant mice at postnatal day 10 with 10 µg PYY(1-36) by intraperitoneal injection. PYY can  
203 be converted to PYY(3-36) by the protease DPP4 (Mentlein et al., 1993), and this form of PYY  
204 has potent anorexic effects in the brain (Batterham et al.). We therefore co-injected PYY(1-36)  
205 and a DPP4 inhibitor to prevent PYY cleavage and to better target the epithelial NPY1R  
206 receptor that preferentially binds the 1-36 form (Hyland et al., 2003; Mentlein et al., 1993; Tough  
207 et al., 2011). Patients with EEC-deficiency die without total parenteral nutrition, and similarly  
208 very few EEC-deficient mice survive without treatment within the first few weeks. However, PYY  
209 injections dramatically improved mutant survival up to 88% (Figure 4A). Moreover, PYY  
210 treatment reduced diarrhea and improved fecal output of mutant mice to either be  
211 indistinguishable from wild-type or only slightly wet but well-defined pellets, which was  
212 independent of intestinal motility (Figures 4B and S6). Treatment of mutant mice with vehicle,

213 DPP4 inhibitor diluted in water, prolonged their survival but did not impact their fecal output or  
214 basal electrophysiology (Figure 4A-C), consistent with therapeutic administration of supportive  
215 fluids in diarrheal disease.

216 We investigated if the animals that survived in response to PYY injections had restored  
217 electrophysiology and improved nutrient absorption in the small intestine. We found that PYY-  
218 injections restored the basal  $I_{sc}$  of jejunum to normal (Figure 4C). Additionally, the response to  
219 VIP (Figure 4D) and the response to luminal glucose (Figure 4E) were both normalized  
220 indicating that PYY injections stably restored electrophysiology. Importantly, mice received their  
221 last injection of PYY approximately 16 hours prior to sacrifice, demonstrating sustained action of  
222 the peptide *in vivo*. The rescue of EEC-deficient intestinal tissue also extended to the human  
223 model, where EEC-deficient HIOs were grown and matured *in vivo* and then host animals were  
224 injected with exogenous PYY for 10 days prior to harvest. These EEC-deficient HIOs exposed  
225 to PYY demonstrated electrogenic response to glucose that was indistinguishable from wild-  
226 type (Figure 4E). Lastly, we investigated whether the PYY treated groups had improved amino  
227 acid absorption as measured by  $H^+$  export and response to the dipeptide Gly-Sar. By  
228 administering PYY to the mosaic EEC-deficient reporter mice, we found PYY injections restored  
229 intracellular pH in EEC-deficient intestinal cells to normal levels which would support PEPT1-  
230 mediated dipeptide absorption (Figure 4F). Consistent with this, PYY-injected mouse and  
231 human small intestine displayed a significantly improved electrogenic response to dipeptides  
232 (Figure 4G), indicating that dipeptide absorption was restored. These data demonstrated  
233 functional efficacy of PYY on improved ion and nutrient transport in EEC-deficient intestine.

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235

236

237 **Discussion**

238 In this study, we found that loss of all EECs resulted in profound imbalance of ion  
239 transport in the small intestine, with subsequent impairment of nutrient absorption. We  
240 demonstrated that PYY is an essential regulator of normal electrophysiology and absorptive  
241 function in the small intestine. Chemical inhibition of the epithelial NPY1R receptor in wild-type  
242 small intestine isolated from human intestinal organoids and mouse demonstrated the  
243 requirement of this pathway in the modulation of VIP-induced ion secretion. Administration of  
244 PYY to EEC-deficient animals resulted in improvements in survival, diarrheal symptoms,  
245 glucose absorption and protein absorption in the absence of all other EEC peptides.

246 Historically, mouse models have been exceedingly tolerant of loss of individual EEC  
247 populations, largely due to functional overlap between EEC-derived peptides (McCauley, 2019).  
248 This has rendered it difficult to assign roles of individual EEC peptides to physiologic functions.  
249 Here, we were able to exploit a model which lacks all EECs to functionally evaluate the role of  
250 one EEC peptide, PYY. However, other peptides like somatostatin have similar activities to PYY  
251 and likely play a similar regulatory role *in vivo*. Somatostatin has many systemic targets (Patel,  
252 1999) and the use of the somatostatin-analogue octreotide in the treatment of chylous effusion  
253 and hyperinsulinemia causes an increased risk of necrotizing enterocolitis in infants (Chandran  
254 et al., 2020). We therefore chose to use PYY in our preclinical model of malabsorptive diarrhea.

255 PYY has been classically defined as a satiety hormone that acts in an endocrine manner  
256 wherein the DPP4-cleaved PYY(3-36) signals to the brain to reduce food intake (Batterham et  
257 al.). However PYY(1-36) has been shown to act in a paracrine manner in the colon using  
258 combination of genetic and pharmacological approaches (Cox, 2008; Hyland et al., 2003; Tough  
259 et al., 2011). We and others (Egerod et al., 2012) observe abundant PYY+ cells in the small  
260 bowel, suggesting that these cells may reprise their paracrine role described in the colon in the  
261 regulation of ion and water transport in the small intestine, linking EECs to glucose and protein

262 absorption. These findings lend some clarity on how EECs integrate their nutrient sensing  
263 function with nutrient absorption, providing us with a new way to approach management of  
264 malabsorptive diseases and those in which EECs are commonly dysregulated.

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267

268 **Figure Legends**  
269

270 **Supplemental Figure 1. NEUROG3 is required for enteroendocrine cell development in**  
271 **human intestinal organoids.**

272 A. Human intestinal organoids (HIOs) derived from human pluripotent stem cells with a null  
273 mutation in *NEUROG3* lacked enteroendocrine cells (EECs) but otherwise had a normal  
274 morphology. The epithelial morphology was assessed using a PSC line expressing a CDH1-  
275 mRuby2 fusion protein (Ouchi et al., 2019) (red, bottom panels) and by co-staining with an  
276 anti-CDH1 antibody (red, top panels). Loss of NEUROG3 did not alter markers of intestinal  
277 identity (CDX2, purple). Only wild-type (top) and wild-type CDH1-mRuby2 (bottom) HIOs  
278 generated Chromogranin A (CHGA)- expressing EECs (green). Scale bars = 50  $\mu$ m.

279 B. After maturation *in vivo*, HIOs develop well-defined crypt-villus architecture. Transplantation  
280 of HIOs (~1 mm) into mice for 10-12 weeks results in growth (1-2 cm), morphogenesis and  
281 maturation (Watson et al., 2014). The epithelium is labeled by CDH1-mRuby2. Scale bar =  
282 500  $\mu$ m.

283 C. Transplanted HIOs with disrupted *NEUROG3* lacked EECs as marked by CHGA+, but were  
284 otherwise morphologically normal. Scale bars = 100  $\mu$ m. Enteroids derived from the crypts  
285 of transplanted HIOs produced EECs when differentiated, whereas those derived from EEC-  
286 deficient HIOs never did. DAPI and CDH1 mark nuclei and epithelium, respectively. Scale  
287 bars = 20  $\mu$ m.

288 D. Transplanted HIOs generated PYY+ and somatostatin (SST)+ EECs, which were never  
289 detected in NEUROG3-deficient transplanted HIOs. DAPI and CDH1 mark nuclei and  
290 epithelium, respectively. Scale bars = 100  $\mu$ m.

291 E. EEC-deficient transplanted HIOs (top) and EEC-deficient HIO-derived enteroids (bottom) did  
292 not express mRNA for EEC markers CHGA (\*\*\*\*P<0.001), PYY (\*\*P=0.001) or SST

293        (\*\*\*/P<0.0001). Neither wild-type nor EEC-deficient tissues expressed mRNA for *VIP* (n=9).

294        Error bars are  $\pm$  SEM.; statistics calculated by unpaired, two-tailed Student's *t*-test.

295        F. Regional patterning of transplanted HIOs was independent of NEUROG3. Transplanted  
296        HIOs, with and without EECs, coexpressed CDX2 and the proximal small intestinal marker  
297        PDX1. DAPI and CDH1 mark nuclei and epithelium, respectively. Scale bars = 100  $\mu$ m.

298        G. Regional identity of transplanted HIOs was maintained in enteroid culture. There was no  
299        difference in *CDX2* or *PDX1* mRNA expression between wild-type and EEC-deficient  
300        transplanted HIOs, or between wild-type and EEC-deficient HIO-derived enteroids. Error  
301        bars are  $\pm$  SEM.; statistics calculated by unpaired, two-tailed Student's *t*-test.

302

303

304 **Figure 1. The PYY-VIP axis regulates ion and water transport in mouse and human small**  
305 **intestine.**

306 A. PYY and VIP regulate ion and water transport in HIO-derived small intestinal enteroids.  
307 Addition of VIP to enteroids induced ion and water transport as measured by swelling. EEC-  
308 deficient enteroids had an elevated response to VIP compared to wild-type enteroids  
309 (\*\*P=0.004). Upon addition of PYY, there was no difference in swelling between wild-type  
310 and EEC-deficient enteroids, and inhibition of VIP-induced swelling. Chemical inhibition of  
311 the PYY receptor NPY1R with BIBO3304 resulted in swelling of wild-type enteroids to EEC-  
312 deficient levels, and abolished the inhibitory effects of PYY in both wild-type and EEC-  
313 deficient enteroids. VIP-induced enteroid swelling was CFTR dependent and blocked by the  
314 CFTR inhibitor CFTR-172. Scale bars = 500  $\mu$ m. Black bars represent wild-type and gray  
315 bars represent EEC-deficient enteroids. Error bars are  $\pm$  SEM. n=283 wild-type and n=351  
316 EEC-deficient enteroids over three biologically independent enteroid lines. Statistics  
317 calculated by two-way ANOVA with Sidak's multiple comparisons test.

318 B. EEC-deficient enteroids displayed impaired NHE3 activity. EEC-deficient enteroids exhibited  
319 reduced  $\text{Na}^+$ -dependent recovery of intracellular pH after an acid load using the ratiometric  
320 pH indicator SNARF-4F. Quantification is of initial rate of  $\text{Na}^+$ -dependent pH recovery (red  
321 line). n=16 wild-type, n=18 mutant enteroids; \*P=0.01. Error bars are  $\pm$  SEM; statistics  
322 calculated by unpaired, two-tailed Student's *t*-test.

323 C. The localization of the VIP receptor VIPR1 and PYY receptor NPY1R was comparable  
324 between wild-type and EEC-deficient human intestinal epithelium. PYY+ and CHGA+ cells  
325 were only found in wild-type HIOs. Scale bars = 100  $\mu$ m.

326 D. PYY modulates the stimulatory effects of VIP in mouse and human small intestine. In the  
327 Ussing chamber, EEC-deficient small intestine displayed a greater response ( $\Delta I_{sc}$ ) to 10 nM  
328 VIP than did wild-type (mouse, n=20 wild-type, 8 mutant, \*\*\*P<0.0001; human, n=15 wild-

329 type, 9 mutant, \*\*P=0.001). Inhibition of NPY1R in wild-type tissue with BIBO3304 resulted  
330 in an elevated response to VIP compared to untreated wild-type (mouse, n=24, \*P=0.01;  
331 human, n=7, \*P=0.04). Addition of exogenous PYY reduced the magnitude of EEC-deficient  
332 response to VIP (n=8 mutant mice, \*\*\*\*P<0.0001; n=7 mutant HIOs, \*\*P=.007) to wild-type  
333 levels. Electrogenic responses to VIP were blocked by the CFTR inhibitor CFTR-172m  
334 (dotted lines). One representative trace is shown (mouse), with baseline  $I_{sc}$  normalized to 0  
335  $\mu\text{A}/\text{cm}^2$ . Error bars are  $\pm$  SEM; statistics calculated by one-way ANOVA with Tukey's  
336 multiple comparisons test.

337

338 **Supplemental Figure 2. PYY is required to maintain normal electrophysiology in mouse**  
339 **and human small intestine.**

340 A. There was no difference in *NPY1R*, *VIPR1*, *SGLT1*, *GLUT2*, or *PEPT1* mRNA expression  
341 between transplanted HIOs with EECs and those without EECs. Error bars are  $\pm$  SEM.  
342 B. There was no difference in *NPY1R*, *VIPR1*, *CFTR* or *SLC9A3* mRNA expression between  
343 enteroids generated from wild-type or EEC-deficient HIOs. Error bars are  $\pm$  SEM.  
344 C. PYY modulates basal  $I_{sc}$  in human and mouse small intestine. EEC-deficient mouse and  
345 human small intestine had significantly higher basal  $I_{sc}$  than wild-type (mouse, n=36 wild-  
346 type, n=11 mutant, \*\*\*\*P<0.0001; HIO, n=7 wild-type, n=12 mutant, \*p=0.03) after  
347 equilibration in the Ussing chamber. Addition of 300 nM NPY1R inhibitor BIBO3304 to wild-  
348 type tissues reproducibly increased the basal  $I_{sc}$  (mouse, n= 26, human, n=10), whereas  
349 addition of 10 nM PYY lowered the basal  $I_{sc}$  in mutant mouse and human tissue (mouse,  
350 n=9, human, n=9). Blocking NPY1R with BIBO3304 abolished the effect of PYY in both wild-  
351 type and mutant tissues. Arrow indicates time of PYY or BIBO3304 application to the  
352 experiment. One representative trace is shown (mouse). Error bars are  $\pm$  SEM; statistics  
353 calculated by unpaired, two-tailed Student's *t*-test.

354

355 **Figure 2. PYY restores normal glucose absorption in EEC-deficient human and mouse**

356 **small intestine.**

357 A. Schematic depicting the PYY-VIP paracrine axis regulating ion and water homeostasis.

358 EEC-derived PYY and ENS-derived VIP both act via G-protein coupled receptors (NPY1R  
359 and VIPR1, respectively) on enterocytes. VIP signaling raises intracellular cAMP levels  
360 resulting in activation of CFTR and efflux of chloride ions while concurrently inhibiting the  
361 sodium-hydrogen exchanger NHE3. The downstream results are that water and sodium are  
362 drawn to the intestinal lumen via paracellular spaces to balance the secreted chloride. PYY  
363 is secreted in response to luminal nutrients and acts as a counterbalance to VIP by lowering  
364 intracellular cAMP levels. Transport of luminal nutrients into the enterocyte depends on  
365 these ion gradients; SGLT1 transports glucose with two  $\text{Na}^+$  ions and PEPT1 transports di-  
366 /tri-peptides with an  $\text{H}^+$  ion.

367 B. In the absence of EECs, ion and water homeostasis is deregulated due to loss of one arm of  
368 the PYY-VIP axis. In EEC-deficient small intestine, loss of PYY results in increased cAMP-  
369 signaling, increased chloride transport, and increased water and sodium accumulation in the  
370 intestinal lumen. Reduced NHE3 transport activity would cause accumulation of cytosolic  $\text{H}^+$   
371 and a decrease in pH. Subsequently, nutrient absorption would be dysregulated, with  
372 diminished di-/tri-peptide absorption due to increased intracellular proton accumulation and  
373 with increased uptake of glucose due to an exaggerated  $\text{Na}^+$  gradient across the apical  
374 membrane.

375 C.  $\text{Na}^+$ -coupled glucose transport is deranged in EEC-deficient human and mouse small  
376 intestine. Wild-type and EEC-deficient human and mouse intestinal tissues were treated with  
377 VIP, then 25 mM D-Glucose was added to the luminal chamber. EEC-deficient intestine had  
378 an elevated initial response to glucose (mouse,  $n=28$  wild-type,  $n=9$  mutant,  $**P=0.001$ ;  
379 HIO,  $n=6$  wild-type,  $n=4$  mutant,  $**P=0.002$ ) that was returned to wild-type levels by pre-  
380 treatment with 10 nM exogenous PYY (mouse,  $n=7$ ,  $*P=0.04$ ; HIO,  $n=3$ ). Inhibition of the

381 NPY1R in wild-type tissues using the antagonist BIBO3304 caused an abnormal initial  
382 response to glucose that mimicked EEC-deficient tissues (mouse, n=12, \*\*P=0.005; HIO,  
383 n=6). Bar graphs represent the slope of the curve depicted within the boxed area. Error bars  
384 are  $\pm$  SEM; statistics calculated by one-way ANOVA with Tukey's multiple comparisons test.

385 D. The subcellular distribution of glucose transporters SGLT1 and GLUT2 is normal in human  
386 intestinal tissue lacking EECs. Scale bars = 50  $\mu$ m.

387 E. SGLT1 is functional in EEC-deficient human small intestine. Human small intestinal tissue  
388 was isolated and transport of glucose in response to saturating amounts of NaCl were  
389 measured using the glucose analog 6-NBDG. EEC-deficient human small intestinal cells  
390 displayed similar total 6-NBDG uptake in the presence of NaCl (\*P=0.01) to wild-type human  
391 intestinal cells (\*P=0.01) and wild-type mouse jejunum cells (\*P=0.01), demonstrating  
392 functional SGLT1-mediated transport. Statistics calculated by one-way ANOVA with Tukey's  
393 multiple comparisons test.

394 F. The ability of SGLT1 to transport Na<sup>+</sup> is not altered in EEC-deficient enteroids. Enteroids  
395 were stained with the Na<sup>+</sup> fluorescent indicator NaGreen in the presence or absence of 25  
396 mM glucose. The Na<sup>+</sup> transport activity of SGLT1 in the presence of glucose is similar in  
397 both wild-type and EEC-deficient epithelium as measured by fluorescence intensity (MFI)  
398 (\*P=0.01). Data represents 4 independent experiments. Statistics calculated by one-way  
399 ANOVA with Tukey's multiple comparisons test.

400 G. Total glucose transport is similar in wild-type and EEC-deficient monolayer cultures. Wild-  
401 type and EEC-deficient enteroids were cultured as monolayers on transwell inserts and  
402 exposed to 25 mM D-glucose with 1 mM fluorescent glucose analog 2-NBDG on the apical  
403 surface. The fluorescence intensity of the basal chamber was quantified after 30 minutes  
404 (lower graph). The epithelium was then analyzed for 2-NBDG within CDH1-mRuby2-positive  
405 epithelium. Data represents 8 independent experiments. Statistics calculated by unpaired t-  
406 test.

407 **Figure 3. H<sup>+</sup>-coupled dipeptide absorption is impaired in EEC-deficient small intestine.**

408 A. EEC-deficient human and mouse small intestine did not respond to luminal Gly-Sar, a  
409 nonhydrolyzable dipeptide, in the Ussing chamber when exposed to 10nM VIP (mouse, n= 9  
410 wild-type, n=6 mutant, \*\*\*\*P<0.0001; human, n=11 wild-type, n=5 mutant, \*\*P=0.006). 10  
411 minutes pre-treatment of EEC-deficient tissue with 10 nM exogenous PYY (mouse, n=6), or  
412 of wild-type tissue with 300 nM NPY1R inhibitor BIBO3304 (mouse, n=9; human, n=6) did  
413 not alter the  $I_{sc}$  response to Gly-Sar. Error bars are  $\pm$  SEM; statistics calculated by one-way  
414 ANOVA with Tukey's multiple comparisons test.

415 B. Expression and localization of peptide transporter PEPT1 is unchanged in EEC-deficient  
416 human small intestine. Scale bars = 50  $\mu$ m.

417 C. The PYY-VIP axis regulates intracellular pH in human small intestinal cells. Wild-type and  
418 EEC-deficient enteroids were differentiated in the presence of 10 nM VIP for 5-7 days.  
419 EEC-deficient enteroids treated with VIP developed an H<sup>+</sup> imbalance with an acidic  
420 cytoplasm whereas wild-type enteroids were able to maintain their intracellular pH  
421 (\*\*P=0.004). Concurrent treatment with 10 nM PYY normalized the pH in EEC-deficient  
422 enteroids and was dependent on NPY1R. pHrodo mean fluorescence intensity (MFI) was  
423 normalized to vehicle-treated wild-type. n= 3 independent experiments. Error bars are  $\pm$   
424 SEM; statistics calculated by the Holm-Sidak method.

425 D. Small intestinal EECs regulate proton transport in a paracrine fashion. Using animals with  
426 mosaic loss of EECs we found that regions of epithelium that escaped recombination had  
427 normal pH and H<sup>+</sup> transport. Adjacent regions that expressed tdTomato, indicating Cre  
428 activity, had impaired elevated cytosolic H<sup>+</sup> as measured by flow cytometry using the  
429 fluorescent pH indicator dye pHrodo. There was no difference in pHrodo MFI between  
430 mosaic regions in wild-type jejunum (n=8), but a significant increase in pHrodo MFI,  
431 indicating relative acidic pH, in EEC-deficient jejunum compared to non-recombined

432 epithelial cells within the same segment of jejunum (n=4, \*\*\*P=0.0002). Error bars are  $\pm$

433 SEM; statistics calculated by two-way ANOVA with Sidak's multiple comparisons test.

434

435 **Supplemental Figure 3. VIP and PYY regulate NHE3 expression in human small intestinal  
436 epithelium.**

437 A. The PYY-VIP axis regulates *SLC9A3* expression. After 5-7 days of exposure to VIP,  
438 *SLC9A3* expression was reduced in wild-type and in EEC-deficient enteroids (\*P=0.04).  
439 Exposure to PYY concurrently with VIP in EEC-deficient enteroids restored *SLC9A3*  
440 expression to not significantly different from untreated. The effect of PYY was blocked with  
441 the NPY1R inhibitor BIBO3304 (\*P=0.02). While there was a trend for PYY treatment alone  
442 to increase *SLC9A3* expression, this did not reach significance. n=6 independent  
443 experiments. Error bars are  $\pm$  SEM; statistics calculated by one way ANOVA with Tukey's  
444 multiple comparisons test.

445

446 **Supplemental Figure 4. PYY is abundant in mouse and human small intestine.**

447 A. PYY+ EECs (arrows) are abundant in mouse and human small intestine. CDH1 labels  
448 epithelium in purple. Scale bars = 100  $\mu$ M.

449

450

451 **Supplemental Figure 5. *VillinCre; Neurog3*<sup>flox/flox</sup>; *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup> mice display**  
452 **incomplete recombination.**

453 A. The tdTomato reporter revealed regions of jejunal epithelium that escaped recombination by  
454 *VillinCre*. ChgA+ EECs were abundant in tdTomato+ and tdTomato negative regions of wild-  
455 type jejunum, but were only detected in tdTomato negative epithelium of *Neurog3*<sup>f/f</sup> animals  
456 (arrow). Scale bars = 20 $\mu$ m.

457 B. Representative dot plots and gating strategy from flow cytometric analysis of *VillinCre*;  
458 *Neurog3*<sup>+/+</sup>; *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup> and *VillinCre; Neurog3*<sup>flox/flox</sup>; *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup>  
459 jejunum.

460 C. Quantification of efficiency of recombination of *VillinCre*. Jejunum of *VillinCre; Neurog3*<sup>+/+</sup>;  
461 *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup>, *VillinCre; Neurog3*<sup>f/+</sup>; *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup> and *VillinCre*;  
462 *Neurog3*<sup>f/f</sup>; *Rosa26*<sup>Flox-STOP-flox-tdTomato</sup> were subjected to flow cytometry. After doublet  
463 discrimination, live, EpCam<sup>+</sup> cells were analyzed for tdTomato expression. Approximately  
464 5.675  $\pm$  1.98% of wild-type (n=8), 5.678  $\pm$  3.2% of heterozygous (n=9), and 4.38  $\pm$  2.56% of  
465 mutant jejunum (n=5) escaped labeling with the tdTomato reporter.

466

467

468

469 **Figure 4. Exogenous PYY rescues EEC-deficient mice from malabsorptive diarrhea and**  
470 **death and restores normal glucose and dipeptide transport.**

471 A. PYY treatment promotes survival of EEC-deficient mice. Survival curve of wild-type (n=100),  
472 EEC-deficient (n=34) and EEC-deficient mice treated once daily with 10  $\mu$ g PYY (n=25)  
473 beginning at postnatal day 10 (P10). Vehicle-treated mice received DPP4 inhibitor diluted in  
474 100 $\mu$ l water (n=18). Mice were weaned at P21. Statistics calculated by log-rank Mantel-Cox  
475 test.

476 B. Daily treatment of EEC-deficient mice with PYY reverses intractable diarrhea. As compared  
477 to control, EEC-deficient mice have intractable watery diarrhea from birth (given score of 3,  
478 gray bar; n=34; \*\*\*\*P<0.0001). Within 48 hours of PYY treatment, EEC-deficient animals  
479 had an average score of 1 with slightly soft yet well-defined fecal pellets (n=25,  
480 \*\*\*\*P<0.0001 from untreated mutant). Mutant mice treated with vehicle did not gain  
481 improvement in diarrhea score (n=18, \*\*\*\*P<0.0001 from PYY-treated mutant, and not  
482 significant from untreated mutant). Wild-type littermates produce well-defined fecal pellets  
483 (given score of 0, black bar; n=100). Error bars are  $\pm$  SEM; statistics calculated by one-way  
484 ANOVA with Tukey's multiple comparisons test.

485 C. PYY treatment of EEC-deficient animals restored a normal resting  $I_{sc}$  to small intestine.  
486 Jejunum from wild-type (black), *VillinCre; Neurog3*<sup>fl/fl</sup> (gray), *VillinCre; Neurog3*<sup>fl/fl</sup> +  
487 PYY injected (red) and *VillinCre; Neurog3*<sup>fl/fl</sup> + vehicle injected (purple) mice were  
488 mounted in the Ussing chamber. Mutant jejunum exhibited a significantly increased basal  $I_{sc}$   
489 compared to wild-type, which was significantly decreased after *in vivo* injections of PYY  
490 (n=6, \*\*\*\*P<0.0001). Treatment of mutant mice with vehicle did not result in improved basal  
491  $I_{sc}$  (n=6). Wild-type and untreated mutant data points are the same as Supplemental Figure  
492 2. Error bars are  $\pm$  SEM; statistics calculated by one-way ANOVA with Tukey's multiple  
493 comparisons test.

494 D. Electrogenic response to VIP was elevated in EEC-deficient animals but restored to wild-  
495 type levels in mutant mice treated with PYY (n=6, \*\*\*\*P<0.0001). Wild-type and untreated  
496 mutant data points are the same as Figure 1. Error bars are  $\pm$  SEM; statistics calculated by  
497 one-way ANOVA with Tukey's multiple comparisons test.

498 E. PYY treatment restores a normal glucose response in EEC-deficient mouse and human  
499 intestine. (mouse, n=6, \*\*P=.003; HIO, n=5, \*\*P=0.004). Wild-type and untreated mutant  
500 data points are the same as Figure 2. Error bars are  $\pm$  SEM; statistics calculated by one-way  
501 ANOVA with Tukey's multiple comparisons test.

502 F. Proton transport is normalized in EEC-deficient animals following PYY treatment. Mean  
503 fluorescent intensity (MFI) of pHrodo was normalized between EEC-deficient and EEC-rich  
504 regions of the mosaic jejunum (n=2). MFI was normalized to untreated wild-type. Wild-type  
505 and untreated mutant data points are the same as Figure 3. Error bars are  $\pm$  SEM; statistics  
506 calculated by two-way ANOVA with Sidak's multiple comparisons test.

507 G. PYY improves dipeptide transport in EEC-deficient mouse and human intestine. Long-term  
508 treatment of EEC-deficient animals and animals hosting transplanted HIOs with PYY  
509 resulted in improved  $I_{sc}$  response to luminal Gly-Sar compared to untreated mutant tissue  
510 (mouse, n=6, \*\*P=.009; HIO, n=5, \*\*\*P=0.0001). Wild-type and untreated mutant data points  
511 are the same as Figure 3. Error bars are  $\pm$  SEM; statistics calculated by one-way ANOVA  
512 with Tukey's multiple comparisons test.

513

514 **Supplemental Figure 6. PYY(1-36) does not slow intestinal motility in EEC-deficient mice.**

515 A. The mechanism of improved survival and diarrhea in PYY-treated mutant animals does  
516 not include slowing intestinal motility. Animals fed ad-lib were orally gavaged with dye-  
517 colored water then sacrificed 30 minutes later. The distance traveled by the dye-front  
518 was reported as percent of small intestinal length. n=15 wild-type mice, 6 wild-type +  
519 vehicle mice, 5 mutant mice (\*\*P=0.0002), 3 mutant + PYY mice, 7 mutant + vehicle  
520 mice. Error bars are  $\pm$ SEM. Statistics calculated by one-way ANOVA with Tukey's  
521 multiple comparisons test.

522 **Methods**

523 **Pluripotent stem cell culture and directed differentiation of HIOs**

524 Human embryonic stem cell (ESC) line WA01 (H1) was purchased from WiCell. We used H1  
525 cells with a CRISPR/Cas9 generated null mutation in *NEUROG3* as previously described  
526 (McGrath et al., 2015). Additionally, we inserted the CDH1-mRuby2 reporter construct (Ouchi et  
527 al., 2019) into *NEUROG3*-/- H1 hESCs. CDH1-mRuby2 and non-reporter hESCs were used  
528 interchangeably. hESCs were maintained in feeder-free culture. Cells were plated on hESC-  
529 qualified Matrigel (BD Biosciences, San Jose, CA) and maintained at 37 °C with 5% CO<sub>2</sub> with  
530 daily removal of differentiated cells and replacement of mTeSR1 media (STEMCELL  
531 Technologies, Vancouver, Canada). Cells were passaged routinely every 4 days using Dispase  
532 (STEMCELL Technologies). HIOs were generated according to protocols established in our lab  
533 (Múnера and Wells, 2017; Spence et al., 2011).

534 ***In vivo* transplant of HIOs**

535 28-35 days after spheroid generation, HIOs were removed from Matrigel and transplanted under  
536 the kidney capsule of immune deficient NOD.Cg-*Prkdc*<sup>scid</sup>/*Il2rg*<sup>tm1Wjl</sup>/SzJ (NSG) mice as  
537 previously described (Watson et al., 2014). NSG mice were maintained on Bactrim chow for a  
538 minimum of 2 weeks prior to transplantation and thereafter for the duration of the experiment (8-  
539 14 weeks).

540 **Generation and maintenance of HIO-derived enteroids**

541 After approximately 10 weeks of *in vivo* growth, crypts were isolated from transplanted HIOs  
542 and plated in 3D as previous described (Mahe et al., 2015). To promote growth, enteroids were  
543 maintained in Human IntestiCult components A+B (STEMCELL Technologies). To promote  
544 differentiation, HIOEs were cultured in gut media (Múnера and Wells, 2017) with 100 µg/ml EGF

545 for 5-7 days. Undifferentiated enteroids were passaged every 7-10 days into fresh Matrigel  
546 (Corning) using a 25G x1/2 needle.

547 **Immunofluorescence**

548 Tissue was fixed in 4% paraformaldehyde, cryopreserved in 30% sucrose, embedded in OCT,  
549 and frozen at -80 °C until cryosectioned. 8 µm cryosections were mounted on Superfrost Plus  
550 slides and permeabilized, blocked and stained according to standard protocol. Primary  
551 antibodies used are listed in the table below, and all secondary antibodies were conjugated to  
552 Alexa Fluor 488, 546/555/568 or 647 (Invitrogen). Images were acquired using a Nikon A1  
553 GaAsP LUNV inverted confocal microscope and NIS Elements software (Nikon).

<i>Primary antibody</i>	<i>Company</i>	<i>Host</i>	<i>Dilution</i>
CDX2	BioGenex	Mouse	1:300
CDX2	Cell Marquis	Rabbit	1:500
Chromogranin A	DSHB	Mouse	1:500
Chromogranin A	ImmunoStar	Rabbit	1:250
E-Cadherin (CDH1)	R&D	Goat	1:500
GLUT2	Santa Cruz	Goat	1:500
Muc2	Santa Cruz	Rabbit	1:250
NPY1R	Abcam	Rabbit	1:250
PDX1	Abcam	goat	1:5000
PEPT1	Santa Cruz	Rabbit	1:500

PYY	Abcam	Rabbit	1:1000
SGLT1	Santa Cruz	Rabbit	1:250
Somatostatin	Santa Cruz	Goat	1:200
VIPR1	Millipore	Mouse	1:200

554 **qPCR**

555 RNA was extracted using Nucleospin RNA extraction kit (Macharey-Nagel) and reverse  
556 transcribed into cDNA using Superscript VILO (Invitrogen) according to manufacturer's  
557 instruction. qPCR primers were designed using NCBI PrimerBlast. Primer sequences are listed  
558 in the table below. qPCR was performed using Quantitect SYBR® Green PCR kit (QIAGEN)  
559 and a QuantStudio 3 Flex Real-Time PCR System (Applied Biosystems). Relative expression  
560 was determined using the  $\Delta\Delta Ct$  method and normalizing to PPIA (cyclophilin A). Samples from  
561 at least three independent passages were used for quantification.

PPIA (CPHA) FWD	CCCACCGTGGTCTTCGACATT
PPIA (CPHA) REV	GGACCCGTATGCTTTAGGATGA
CHGA FWD	TGTGTCGGAGATGACCTCAA
CHGA REV	GTCCTGGCTTTCTGCTCTG
PYY FWD	CGAGACTAAATGTGGCGGGT
PYY REV	GAGCATGCAGTTCTGAGGGT
SST FWD	TGGGTTCAGACAGCAGCTC
SST REV	CCCAGACTCCGTCAGTTCT

VIP FWD	CCCTGTACCAGTCAAACGTCA
VIP REV	GAGTCTCCATGCAGGCTTCT
PDX1 FWD	CGTCCGCTTGTCTCCTC
PDX1 REV	CCTTCCCATGGATGAAGTC
CDX2 FWD	GGGCTCTCTGAGAGGCAGGT
CDX2 REV	GGTGACGGTGGGTTAGCA
NPY1R FWD	ATTCCTAGGCAATGCTTCCCC
NPY1R REV	ACGCCTCCTAAAGCCGAAC
VIPR1 FWD	GATAGGAGCCTGCTGGTCAC
VIPR1 REV	GGGAAACCAAGCCAATCCAA
CFTR FWD	GGCACCCAGAGTAGTAGGTC
CFTR REV	AGGCGCTGTCTGTATCCTTT
SLC9A3 (NHE3) FWD	GCTGGTCTTCATCTCCGTGT
SLC9A3 (NHE3) REV	CCAGAGGCTTGATGGTCAGG

562

563 **Swelling assay**

564 Enteroids were plated in 10  $\mu$ L Matrigel on an 8-chamber glass bottom slide (Ibidi) and  
565 maintained as described above. 3-5 days post-plating, the slide was mounted on an inverted  
566 confocal microscope (Nikon) fitted with an incubation chamber set to 37 °C and 5% CO<sub>2</sub>. Media  
567 was changed to include 10 nM VIP (Tocris). In some experiments, the media was changed 24

568 hours prior to imaging to include 300nM BIBO3304 trifluoroacetate (Tocris), 20  $\mu$ M CFTR(inh)-  
569 172 (Millipore Sigma) and/or 10 nM PYY (Phoenix Pharmaceuticals). Images were acquired  
570 every 5 minutes at 4X magnification. After 6 hours, some HIOEs swelled to the point of bursting;  
571 therefore, we used images acquired at time 0 and at 6 hours for quantification. The area of 10  
572 representative enteroids per well was quantified using NIS Elements software at both time  
573 points. The outline of individual enteroids was traced manually and the area calculated by NIS  
574 Elements. Fold change at 6 hours over baseline was reported. Data include a minimum of three  
575 independent experiments per condition on three wild-type and three EEC-deficient HIO-derived  
576 enteroid lines.

### 577 **NHE3 activity assay**

578 NHE3 activity was determined as previously described (Foulke-Abel et al., 2016) with minor  
579 modifications. Enteroids were plated in 5  $\mu$ L Matrigel on an 8-chamber glass bottom slide (Ibidi)  
580 and maintained as described above. 3-5 days post-plating, media was changed to  $\text{Na}^+$  media  
581 containing 5  $\mu$ M SNARF-4F 5-(and-6)- carboxylic acid, acetoxyethyl ester, acetate (Molecular  
582 Probes) and allowed to incubate for 30 minutes. The slide was then mounted on an inverted  
583 confocal microscope (Nikon), fitted with an incubation chamber set to 37 °C and 5%  $\text{CO}_2$ . Fresh  
584  $\text{Na}^+$  media was provided before image acquisition. Images were acquired every 2 minutes for 2  
585 hours at 10X magnification with excitation at 488 nm and emission at 561 nm and 640 nm.  
586 Media was changed to  $\text{NH}_4\text{Cl}$  to acid-load the epithelium, then to tetramethylammonium (TMA)  
587 media to withdraw  $\text{Na}^+$ .  $\text{Na}^+$  containing media was then added and NHE3 activity quantified as a  
588 measure of initial pH recovery. 1 mM probenecid and 5  $\mu$ M SNARF were present in all buffers,  
589 and all buffers were set to pH 7.4. Intracellular pH was calibrated using the Intracellular pH  
590 Calibration Buffer kit (Invitrogen) at pH 7.5, 6.5 and 5.5 in the presence of 10  $\mu$ M valinomycin  
591 and 10  $\mu$ M nigericin at the conclusion of each experiment. The ratio of 561/640 was determined  
592 using NIS Elements software by drawing a region of interest and quantifying the fluorescence

593 intensity of each wavelength over the period of the experiment. A minimum of 3 enteroids in 3  
594 wells over two independent passages were quantified. The ratio of 561/640 was converted to  
595 intracellular pH using the equation provided by the manufacturer.

596  $\text{Na}^+$  media: 130 mM NaCl, 5 mM KCl, 2 mM  $\text{CaCl}_2$ , 1 mM  $\text{MgSO}_4$ , 20 mM HEPES, 5 mM NaOH,  
597 1 mM (Na)PO<sub>4</sub>, 25 mM D-glucose

598 NH<sub>4</sub>Cl media: 25 mM NH<sub>4</sub>Cl, 105 mM NaCl, 2 mM  $\text{CaCl}_2$ , 1 mM  $\text{MgSO}_4$ , 20 mM HEPES, 8 mM  
599 NaOH, 5 mM KCl, 1 mM (Na)PO<sub>4</sub>, 25 mM D-glucose

600 TMA media: 130 mM TMA-Cl, 5 mM KCl, 2 mM  $\text{CaCl}_2$ , 1 mM  $\text{MgSO}_4$ , 20 mM HEPES, 8 mM  
601 TMA-OH, 1 mM (TMA)PO<sub>4</sub>, 25 mM D-glucose

## 602 **Electrophysiology**

603 Electrophysiological experiments were conducted as described (Clarke, 2009) with minor  
604 modifications. Mouse jejunum and transplanted HIOs were dissected and immediately placed in  
605 ice-cold Krebs-Ringer solution. Tissues were opened to create a flat epithelial surface. Because  
606 seromuscular stripping is associated with release of cyclooxygenases and prostaglandins  
607 (Clarke, 2009), and prostaglandins can stimulate L-cells to release GLP1, GLP2 and PYY  
608 (Briere et al., 2013), we performed the Ussing chamber experiments in intestinal tissue with an  
609 intact muscular layer. Tissues were mounted into sliders (0.031 cm<sup>2</sup> area slider, P2307,  
610 Physiological Instruments) and placed in an Ussing chamber with reservoirs containing 5 mL  
611 buffer (115 mM NaCl, 1.2 mM  $\text{CaCl}_2$ , 1.2 mM  $\text{MgCl}_2$ , 25 mM  $\text{NaHCO}_3$ , 2.4 mM  $\text{K}_2\text{HPO}_4$  and 0.4  
612 mM KH<sub>2</sub>PO<sub>4</sub>). The mucosal and serosal tissue surfaces were bathed in the same solution, with  
613 the exception of 10 mM glucose in the serosal buffer and 10 mM mannitol in the luminal buffer.  
614 Mucosal and serosal reservoir solutions were gassed with 95 % O<sub>2</sub> and 5 % CO<sub>2</sub> to pH 7.4 and  
615 maintained at 37 °C by a circulating water bath behind the reservoir  
616 chambers. Electrophysiology parameters were recorded as previously described (Matthis et al.,

617 2019). Tissue was allowed to equilibrate to a basal steady-state for a minimum of 30 minutes  
618 before the addition of chemicals or peptides. 10 nM tetrodotoxin (Tocris) was added to the  
619 serosal buffer bathing mouse intestine to inhibit voltage-gated neuronal firing, and allowed to  
620 incubate for a minimum of 10 minutes before basal  $I_{sc}$  recording. D-glucose and Gly-Sar were  
621 added to the luminal side of the chamber once the VIP-induced  $I_{sc}$  had stabilized at a maximum  
622 value.

Tetrodotoxin	Tocris	10 nM
BIBO3304 trifluoroacetate	Tocris	300 nM
VIP	Tocris	10 nM
PYY(1-36)	Phoenix Pharmaceuticals	10 nM
CFTR-172	Millipore Sigma	20 $\mu$ M
D-glucose	Sigma Aldrich	25 mM
Gly-Sar	Sigma Aldrich	20 mM

623

624 **Glucose uptake assays**

625 **6-NBDG**

626 Transplanted HIOs were removed from the murine kidney, bisected to expose the lumen, and  
627 incubated with 100 mM 6-(*N*-(7-Nitrobenz-2-oxa-1,3-diazol-4-yl)Amino)-2-Deoxyglucose (6-  
628 NBDG) (Life Technologies) in 10 nM Tris/HEPES buffer containing 150 mM KCl or 150 mM  
629 NaCl for 30 minutes at 37 °C. Tissues were washed with ice-cold 10 mM Tris/HEPES buffer,  
630 then dissociated to single-cell suspension in 5 mL Tryple Select (Gibco) + 10  $\mu$ M Y-27632  
631 (Tocris), filtered, and subjected to analysis by flow cytometry.

632 **Sodium Green**

633 HIOEs were differentiated for 5-7 days, then were removed from Matrigel and enzymatically  
634 dissociated into single-cell suspension using 0.25% Trypsin-EDTA. Each cell preparation was  
635 split into two samples: one incubated with 25 mM D-glucose and one incubated in the absence  
636 of glucose. Each sample was incubated in Live Cell Imaging Solution (Invitrogen) containing 5  
637  $\mu$ M final concentration of Sodium Green tetraacetate (Molecular Probes) for 30 minutes at 37  
638 °C, washed with ice-cold PBS and analyzed by flow cytometry.

639 **2-NBDG on Transwell filters**

640 Undifferentiated enteroids that were “ready to split” were dissociated and plated on transwell  
641 inserts (Corning) as previously described (Moon et al., 2014), with the exception of first coating  
642 the transwells with Collagen IV (Sigma-Aldrich). 300,000 cells were plated per 6.5 mm transwell  
643 insert. Differentiation was initiated at 24 hours post-plating and monolayers were analyzed after  
644 5-7 days. 1 mM fluorescent glucose analog 2-(*N*-(7-Nitrobenz-2-oxa-1,3-diazol-4-yl)Amino)-2-  
645 Deoxyglucose (2-NBDG, Life Technologies) was diluted in Live Cell Imaging Solution  
646 (Invitrogen) containing 25 mM D-glucose, added to the apical surface of HIOE monolayers and  
647 the fluorescence intensity of fresh Live Cell Imaging Solution in the basal chamber was  
648 quantified after 30 minutes at 37 °C. Intact barrier function was confirmed by co-incubation,  
649 quantification and exclusion of Cascade Blue conjugated 3000 MW dextran (Life Technologies)  
650 in every experiment.

651 **Intracellular pH assay**

652 Enteroids were differentiated for 5-7 days in the presence of vehicle (water or DMSO), 10 nM  
653 VIP (Tocris), 10 nM PYY (Phoenix Pharmaceuticals) and/or 300 nM BIBO3304. On the final  
654 day, enteroids were removed from Matrigel and enzymatically dissociated into single-cell  
655 suspension using 0.25% Trypsin-EDTA. Cell suspensions were counted and equal cell numbers

656 of dissociated HIOEs were incubated in pHrodo Green AM Intracellular pH indicator  
657 (ThermoFisher Scientific) according to manufacturer's directions for 30 minutes at 37C, washed  
658 with 1X PBS, and analyzed by flow cytometry.

659 **Flow cytometry**

660 After mechanical and enzymatic dissociation, tissues were filtered through a 40  $\mu$ m cell strainer  
661 to obtain a single-cell suspension. In all experiments, samples were labeled with either CDH1-  
662 mRuby2 or Anti-EpCam-APC (BD Biosciences) to distinguish epithelial cells and incubated with  
663 SYTOX Blue dead cell stain (Life Technologies) or 7-AAD (BD Pharmingen). Forward scatter  
664 and side scatter were used to discriminate doublets and cellular debris. A minimum of 50,000  
665 events per sample was recorded using an LSR Fortessa flow cytometer (BD Biosciences) and  
666 data was analyzed using FACSDiva software (BD Biosciences).

667 **Mice**

668 B6.Cg-*Tg(Vil1-cre)<sup>997Gum/J</sup>* (*VillinCre*) (JAX stock 004586), *Neurog3*<sup>flox/flox</sup> (Mellitzer et al., 2010)  
669 and B6.Cg-*Gt(ROSA)26Sor<sup>tm1(CAG-tdTomato)Hze</sup>*/J (tdTomato) (Madisen et al., 2010) mice were  
670 maintained on a C57BL/6 background and genotyped as previously described. Mice were  
671 housed in a specific pathogen free barrier facility in accordance with NIH Guidelines for the  
672 Care and Use of Laboratory Animals. All experiments were approved by the Cincinnati  
673 Children's Hospital Research Foundation Institutional Animal Care and Use Committee  
674 (IACUC2019-0006) and carried out using standard procedures. Mice were maintained on a 12-  
675 hour light/dark cycle and had *ad libitum* access to standard chow and water.

676 *VillinCre;Neurog3*<sup>flox/flox</sup> mice<sup>4</sup> and their littermates were weighed, genotyped and visually  
677 examined for liquid feces daily beginning at postnatal day 10. We established a diarrhea score,  
678 with 3 representing wet, yellow feces that smeared the perianal fur, and 0 representing normal,  
679 dry, brown, well-defined pellets. Mutant mice which suffered from diarrhea score 3 were

680 included in the rescue experiment. 10  $\mu$ g PYY (Phoenix Pharmaceuticals) was diluted in water  
681 and added to 20  $\mu$ l DPP4 inhibitor (Millipore) to a final volume of 100  $\mu$ l per mouse. Mice were  
682 injected intraperitoneally with this cocktail within 2 hours of the onset of the dark cycle (7pm)  
683 daily until analysis at postnatal day 28-35. Mice were given access to solid chow on the floor of  
684 the cage beginning at postnatal day 10 and weaned at postnatal day 21. Small intestinal transit  
685 was determined by oral gavage of food coloring diluted in 100 $\mu$ l to ad-lib fed mice, then sacrifice  
686 and measurement of the distance traveled by the dye-front 30 minutes post-gavage.  
687 NSG mice hosting HIOs were treated with 25  $\mu$ g PYY (Phoenix Pharmaceuticals) diluted in  
688 water to 100  $\mu$ L by intraperitoneal injection. Mice were treated daily for a minimum of 10 days  
689 after HIOs had been maturing for 8 weeks, then dissected and analyzed.

690 **Statistics**

691 Data is presented as the mean  $\pm$  SEM unless otherwise indicated. Significance was determined  
692 using appropriate tests in Graph Pad Prism, with  $P \geq 0.05$  not significant; \* $P < 0.05$ , \*\* $P < 0.01$ ,  
693 \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$ .

694

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706 Center (P30 DK078392).

707 **Author contributions**

708 HAM and JMW conceived and initiated the project, designed experiments, and wrote the  
709 manuscript, with conceptual input from MAH, MHM and EA. HAM performed all experiments in  
710 collaboration with: JRE, JGS and WJS on mouse transplantation; NS and MAH in generating  
711 HIO-derived enteroids; ALM, MHM and EA on electrophysiological studies. HAM, EA and JMW  
712 interpreted data. JMW supervised the project. All authors have edited and approved the  
713 manuscript.

714 **Competing interests**

715 The authors declare no competing interests.

716 **Materials and correspondence**

717 Requests for materials and correspondence should be directed to [james.wells@cchmc.org](mailto:james.wells@cchmc.org).

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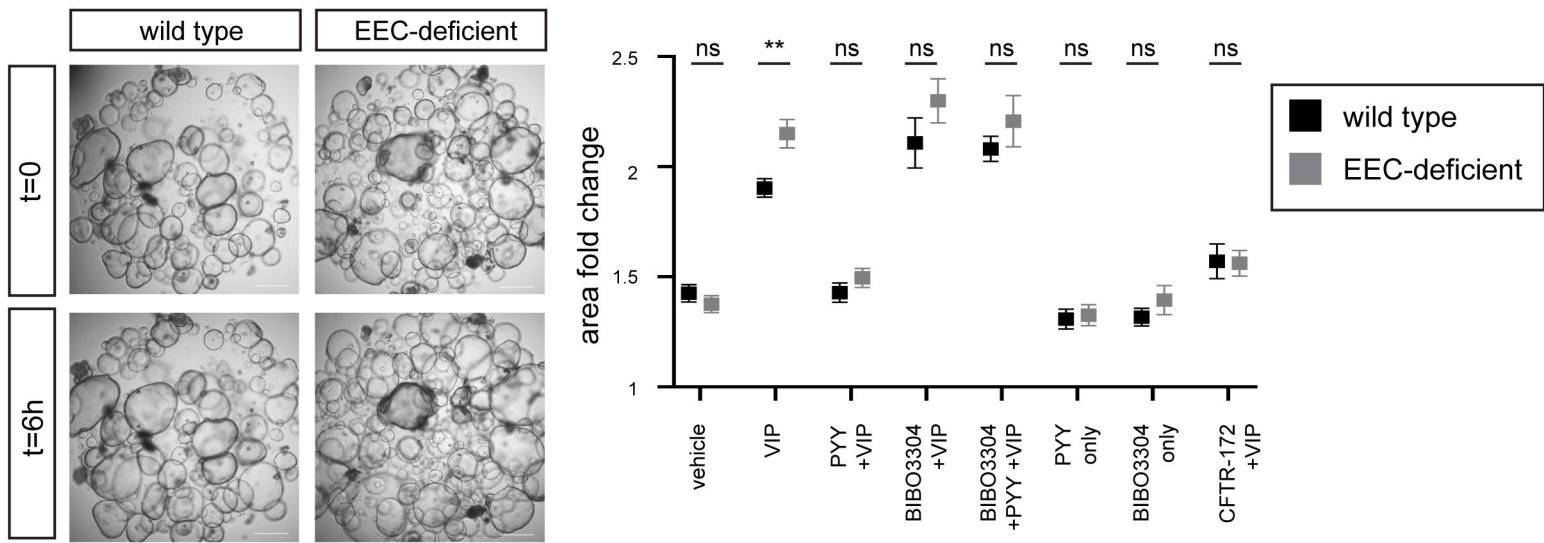
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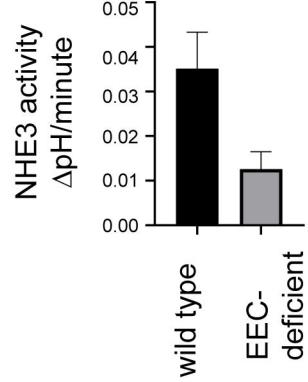
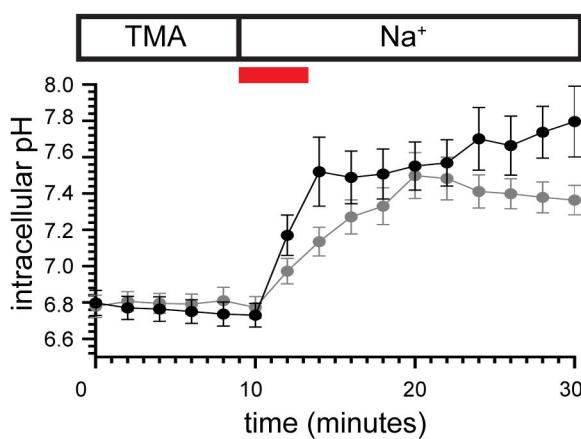
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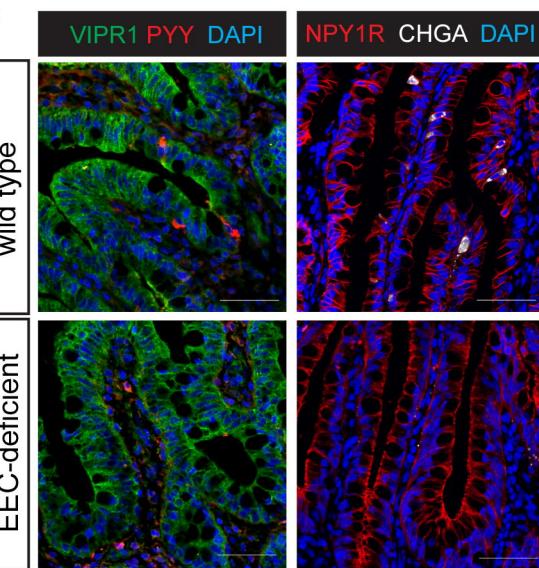
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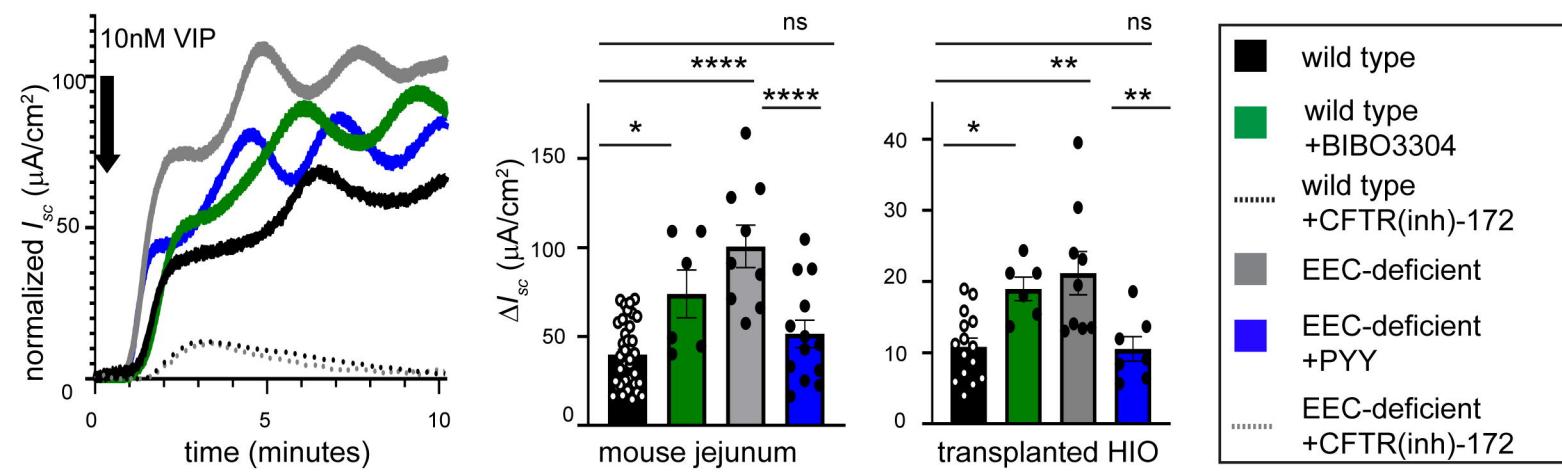
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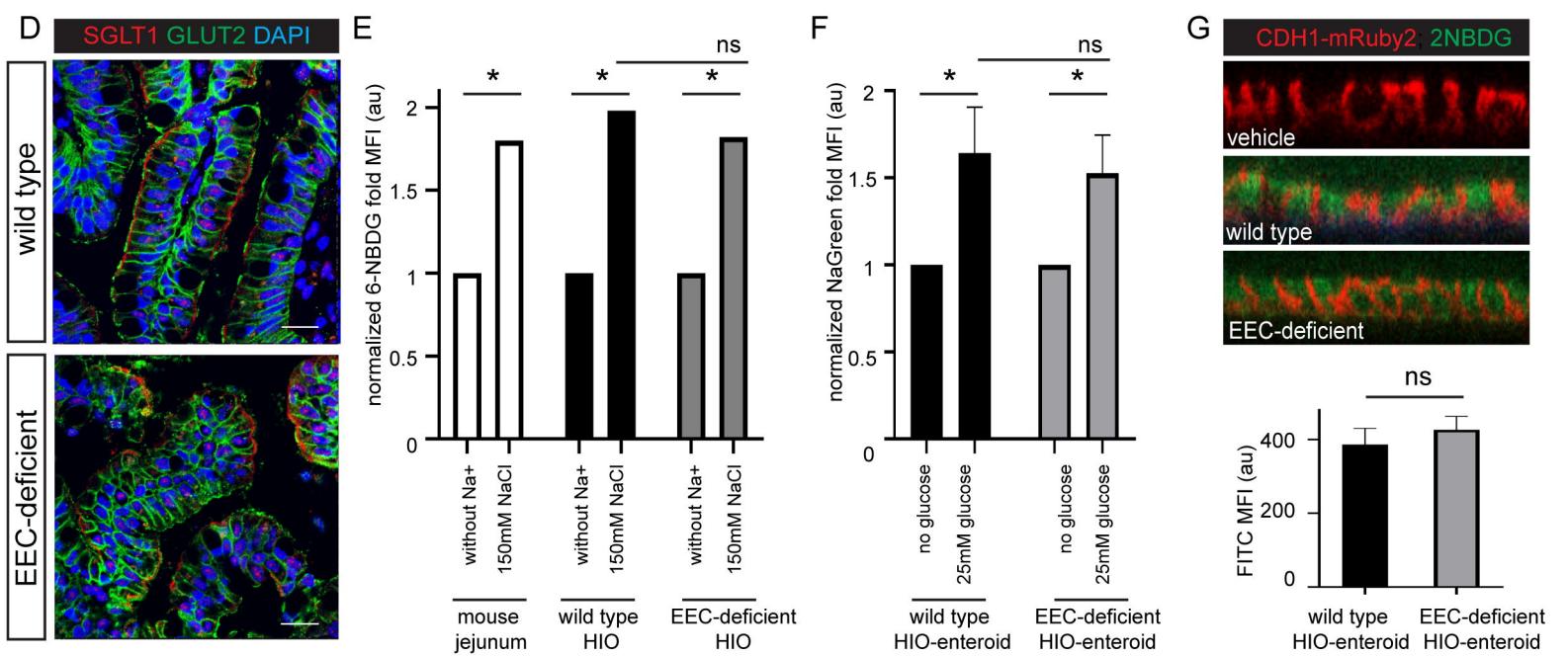
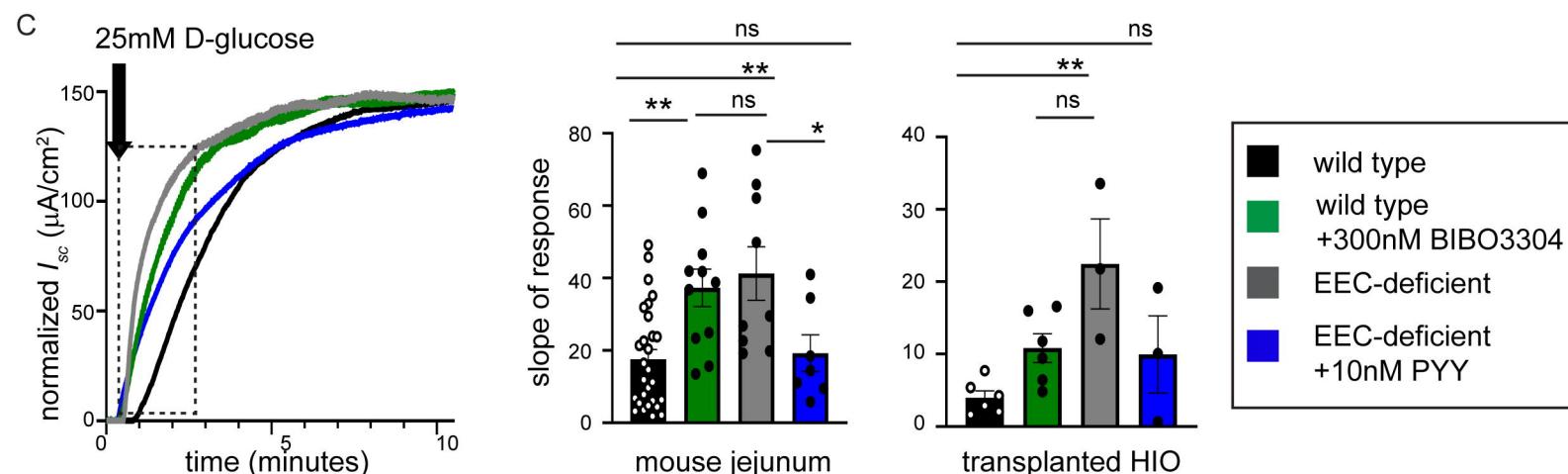
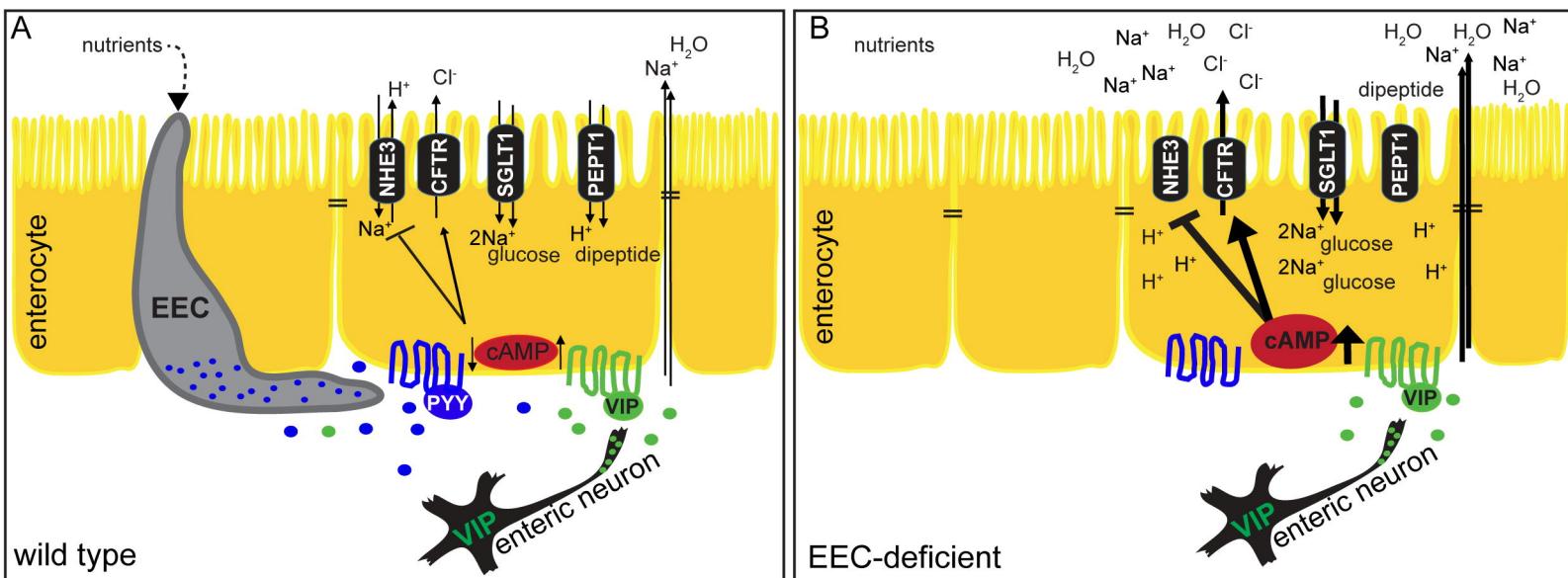


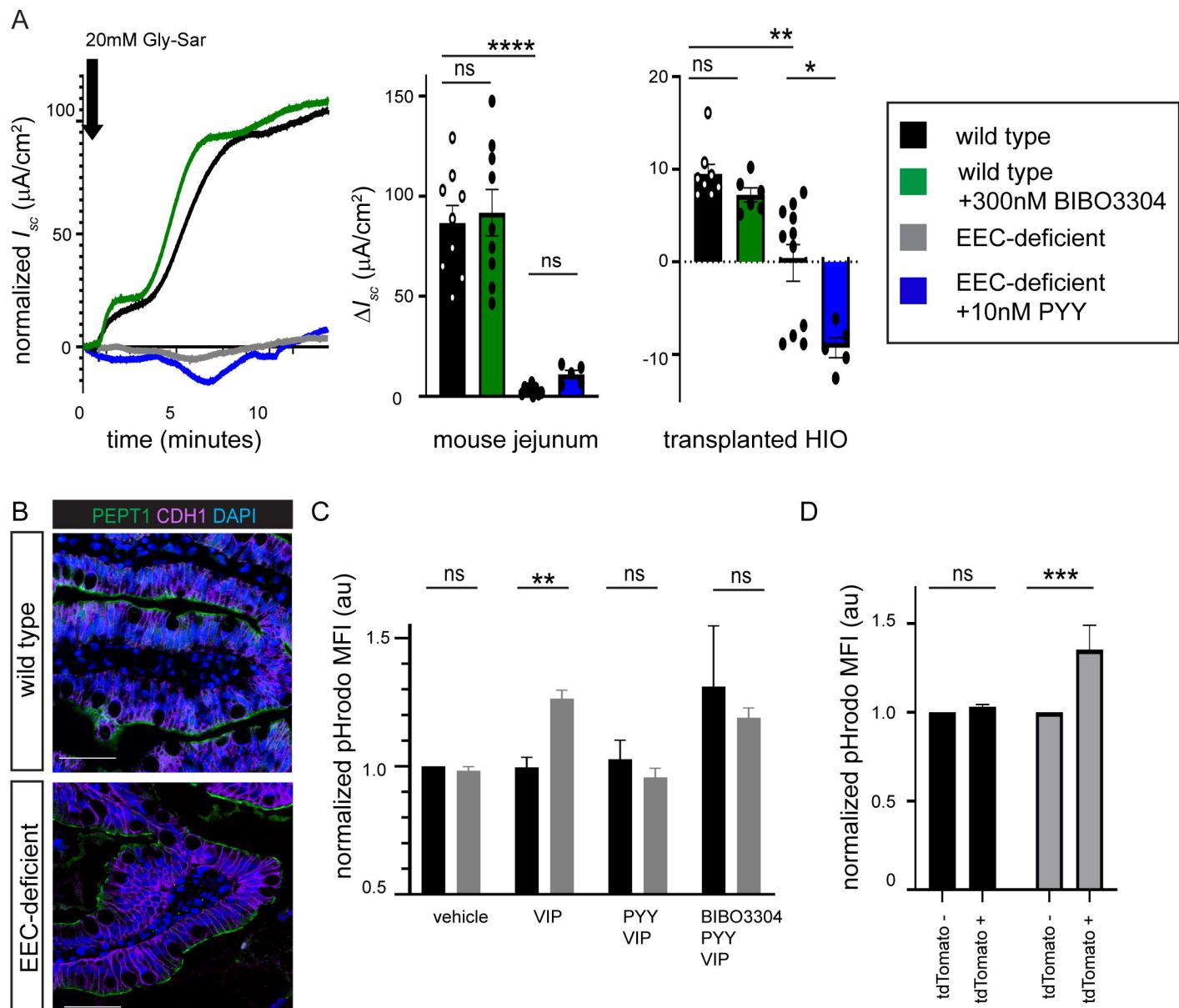
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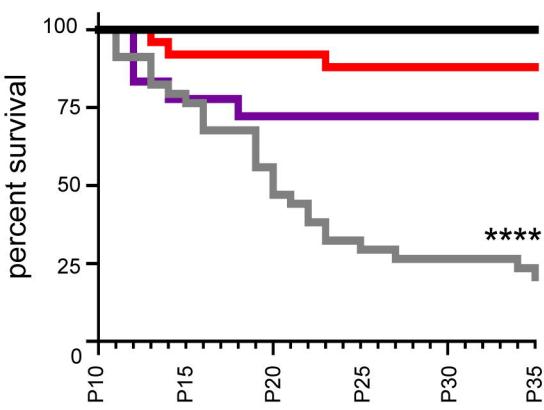
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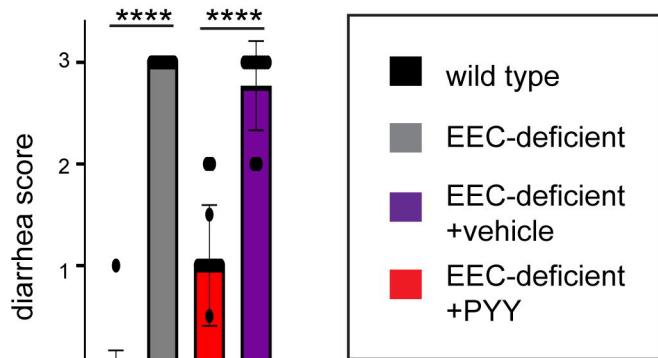




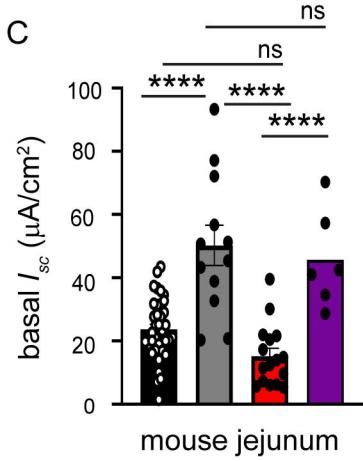
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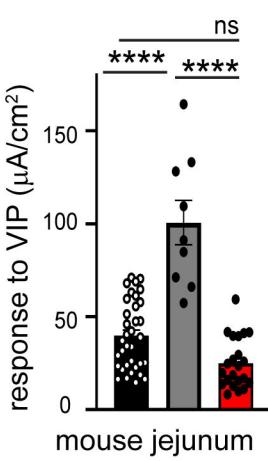
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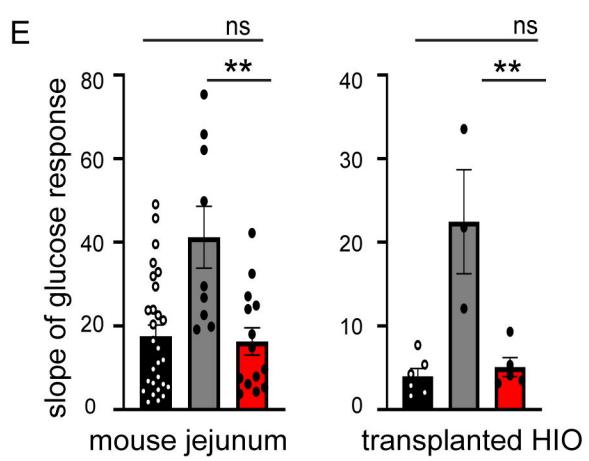
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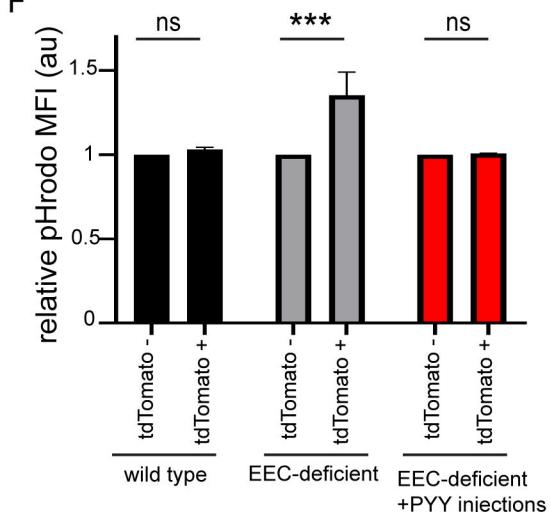
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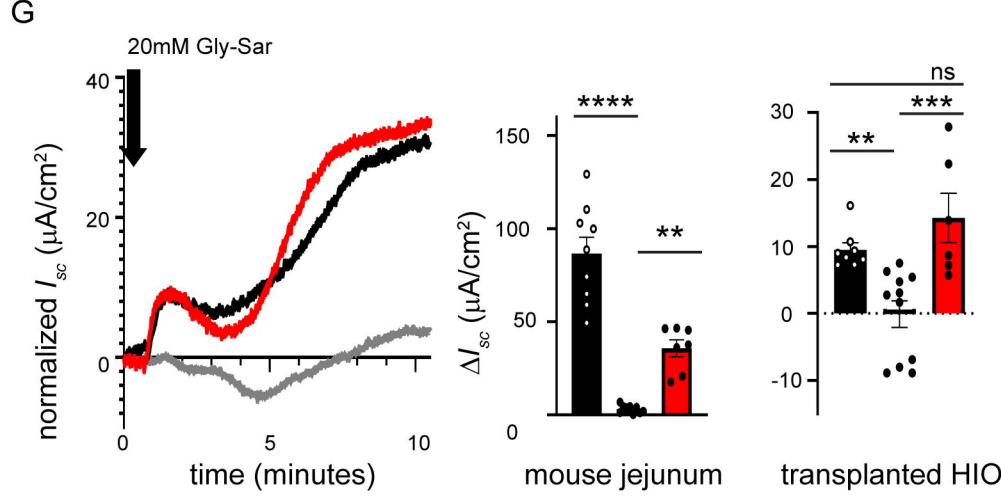
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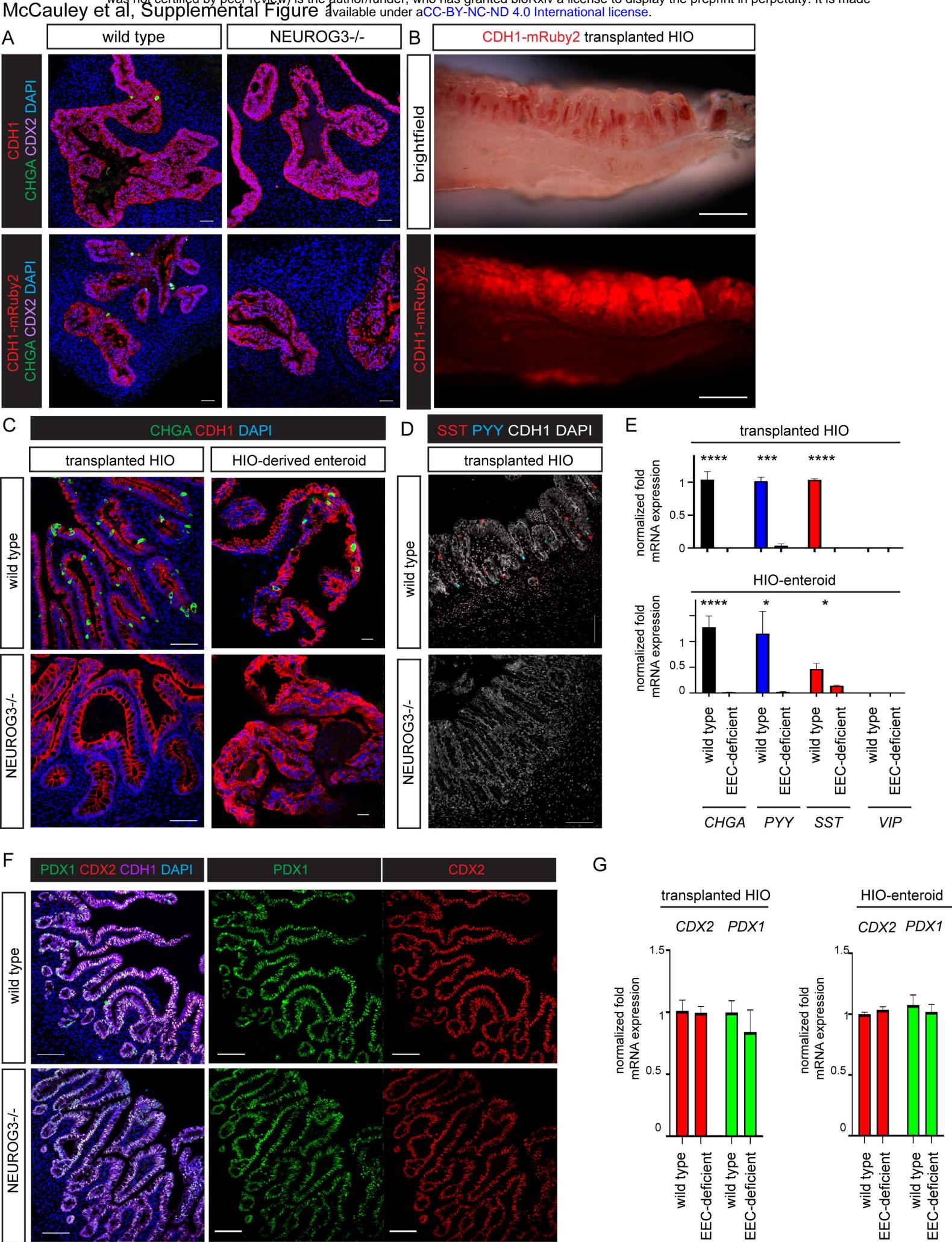


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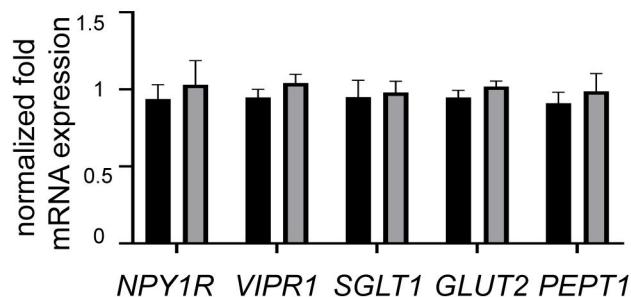
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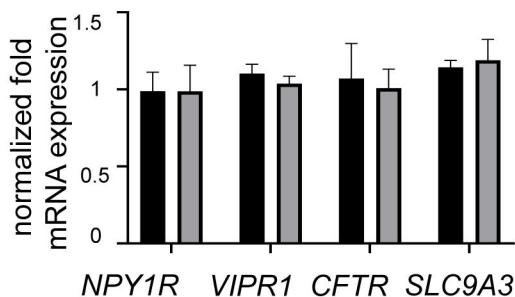
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transplanted HIO

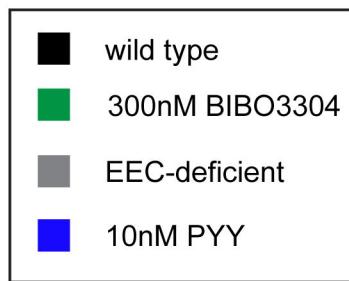
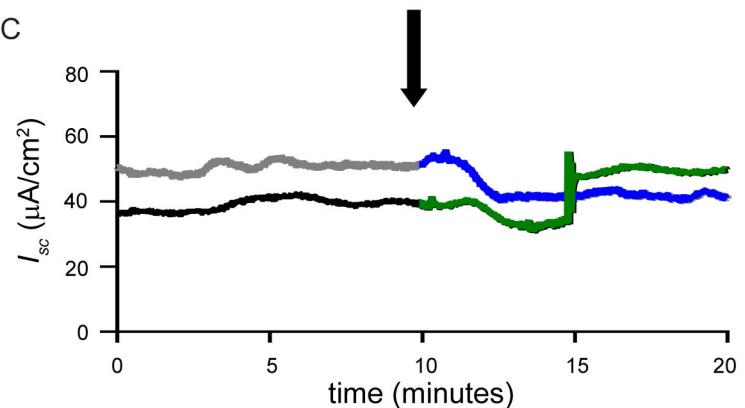


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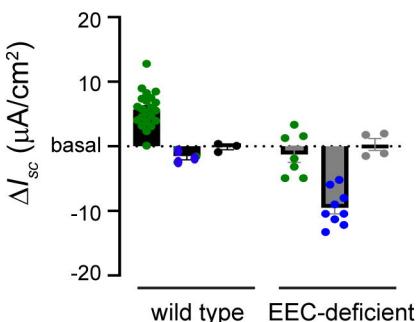
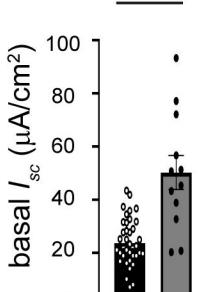
HIO-derived enteroids



C

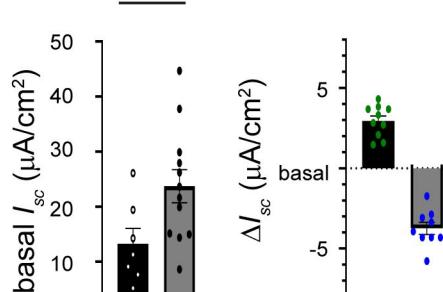


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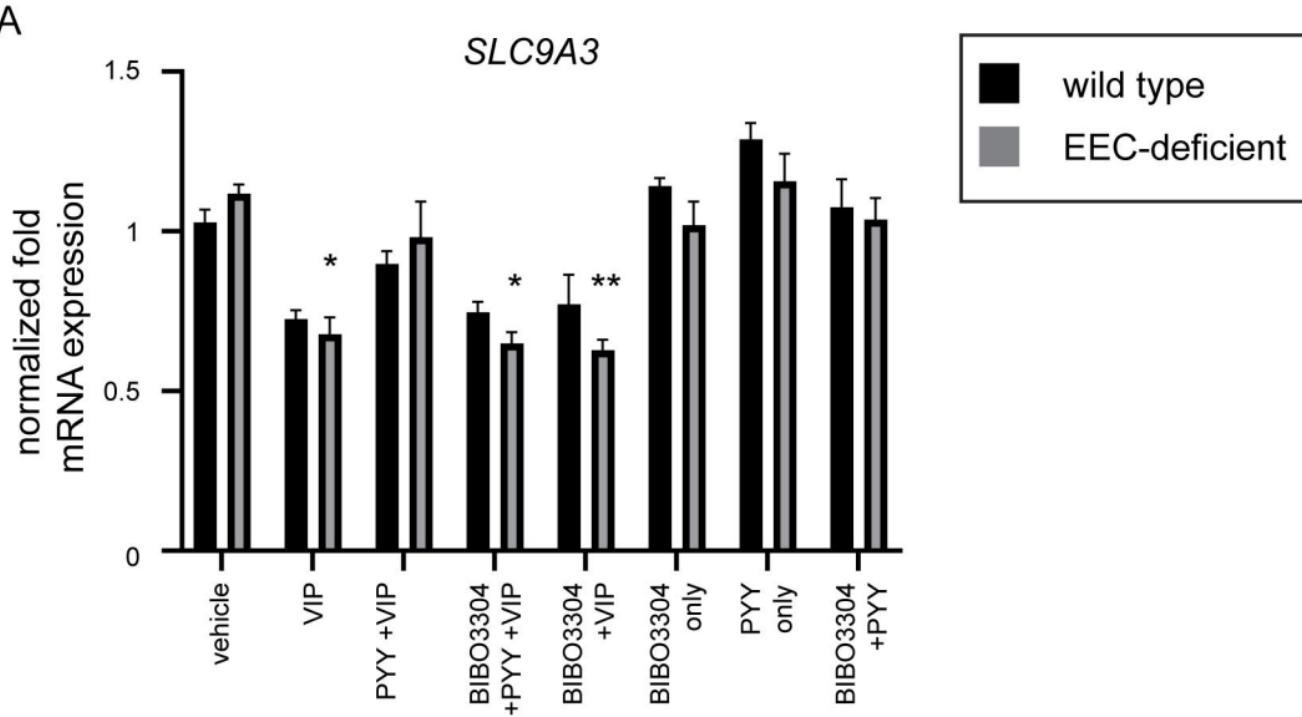
mouse jejunum

\*



transplanted HIO

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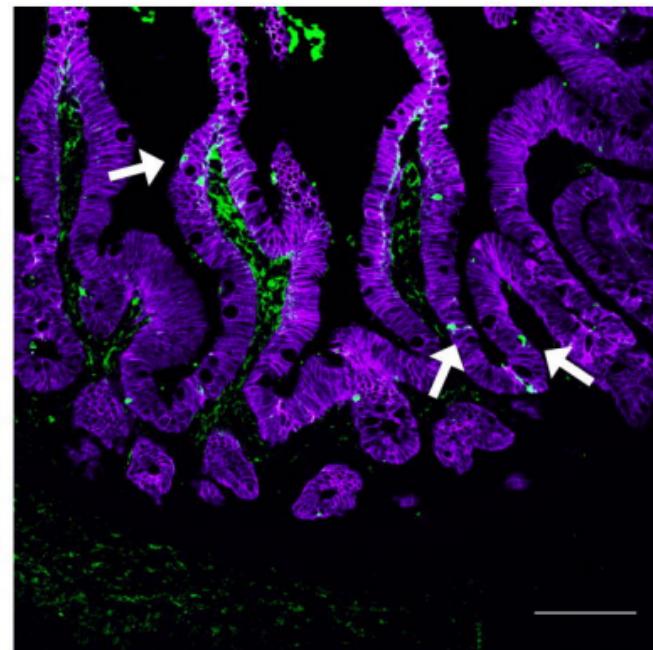
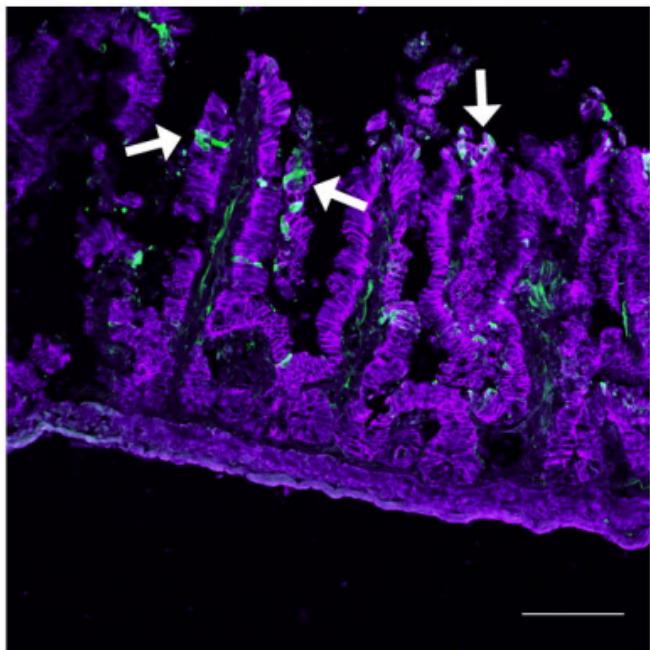


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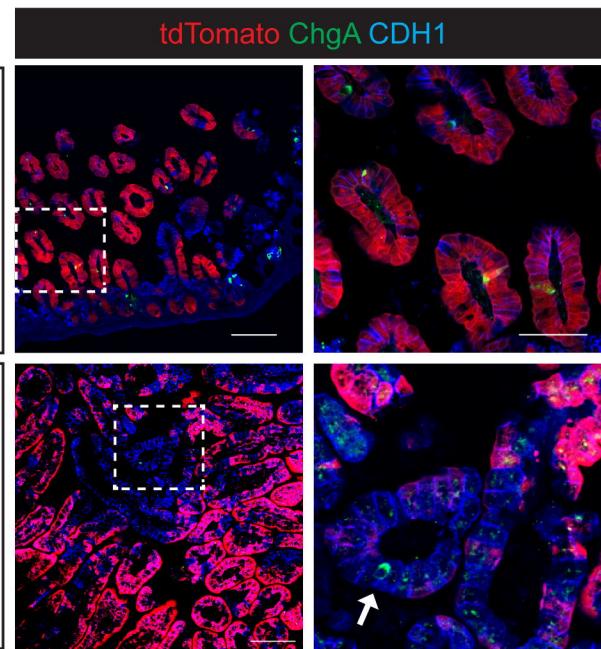
mouse jejunum

transplanted HIO

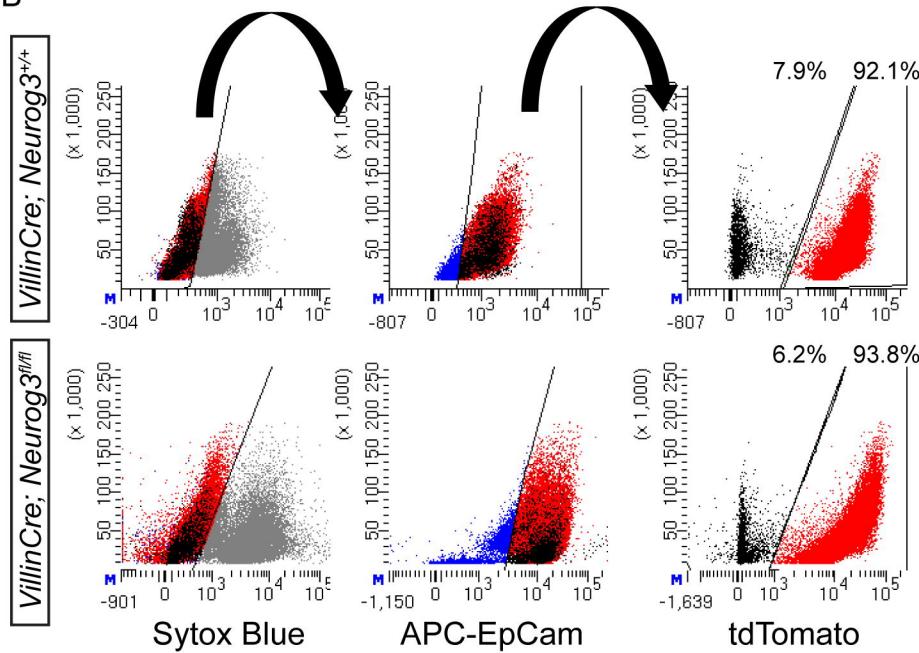
PYY CDH1



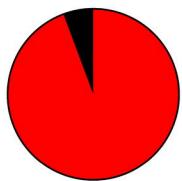
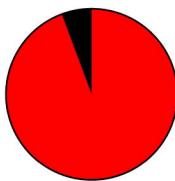
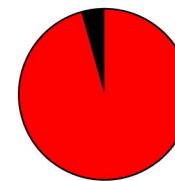
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C

VillinCre; Neurog3<sup>+/+</sup>VillinCre; Neurog3<sup>fl/+</sup>VillinCre; Neurog3<sup>fl/fl</sup>

A

