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**Supplemental Information**

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*in vivo* and eliminate the risk of teratoma formation**

**Yasaman Aghazadeh, Farida Sarangi, Frankie Poon, Blessing Nkennor, Emily C. McGaugh, Sara S. Nunes, and M. Cristina Nostro**

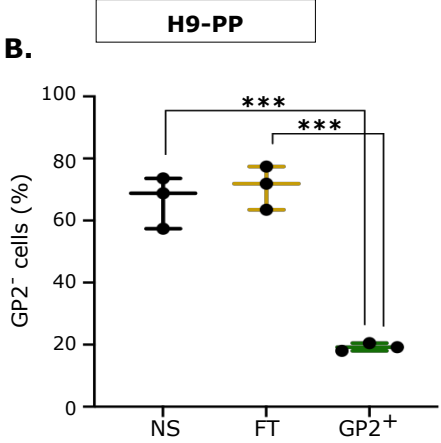
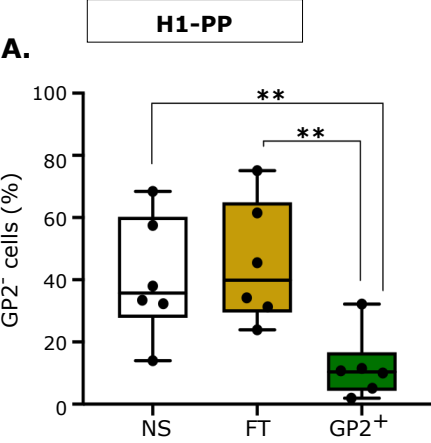
## **Supplemental Information**

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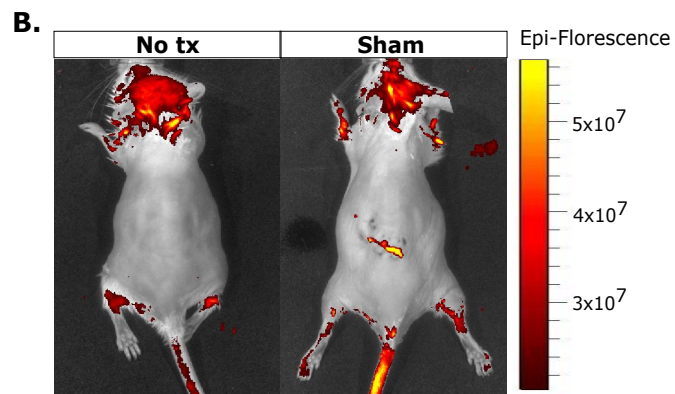
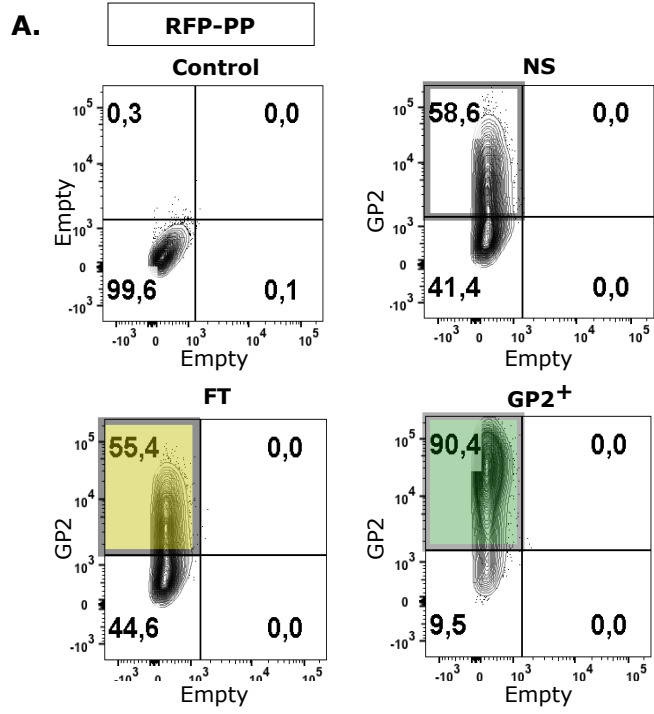
Sara S. Nunes, M. Cristina Nostro.

Supplementary Figure 1

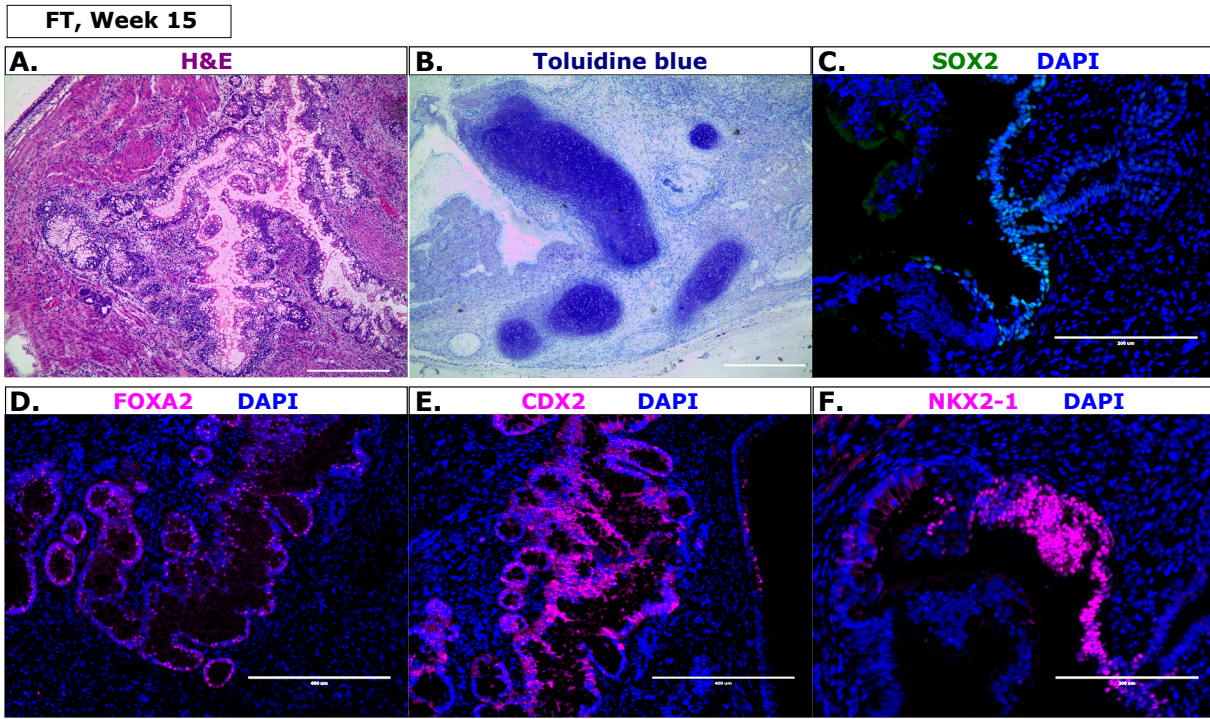


***Supplementary Figure 1. Related to Figure 1.*** (A-B) Frequency of GP2 negative (GP2-) cells in NS, FT and GP2<sup>+</sup> cell populations after MACS in H1 (A) and H9 (B)- derived cells. (A, n=6 independent experiments; B, n=3 independent experiments; One-way ANOVA analysis with Tukey's multiple comparisons; \*\* P < 0.01, \*\*\* P < 0.0001, error bars represent S.E.M.).

## Supplementary Figure 2

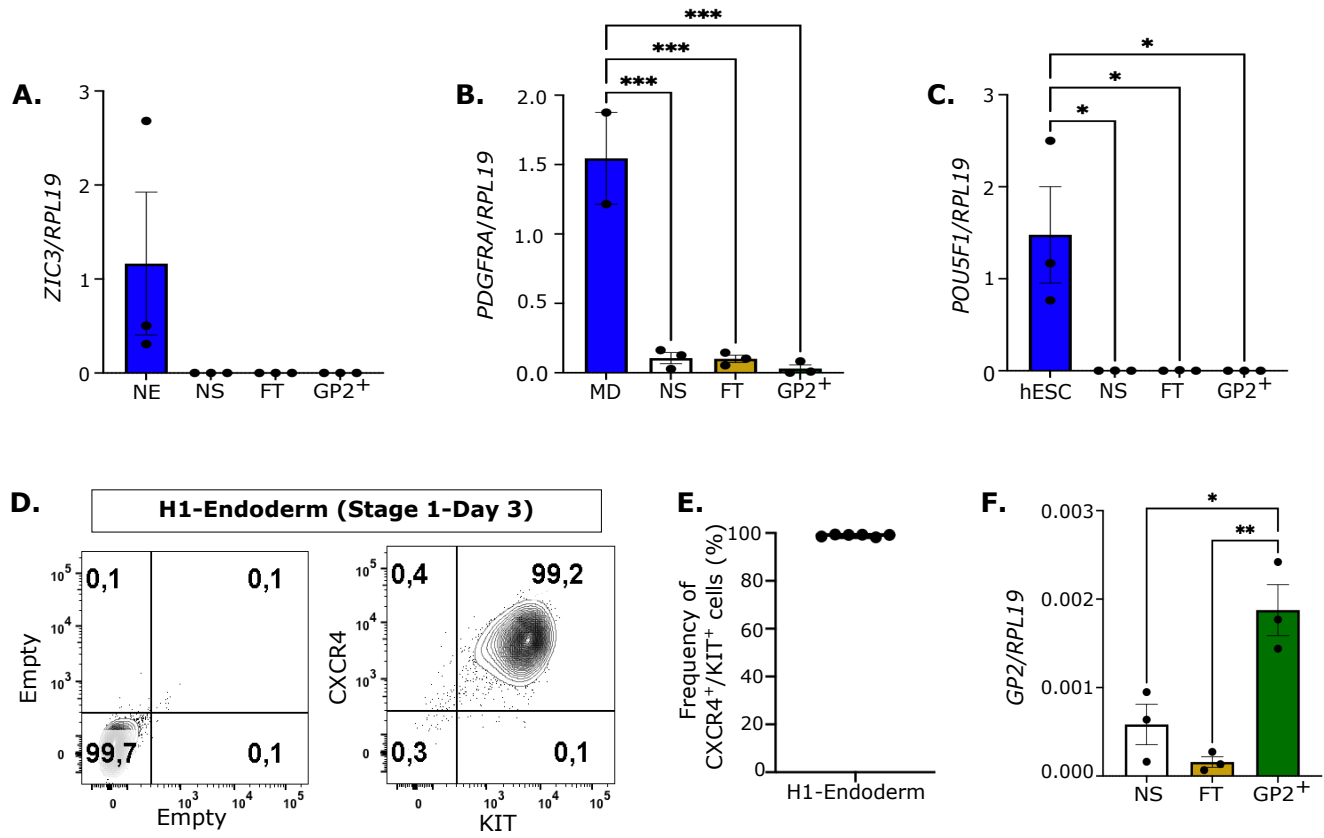


***Supplemental Figure 2. Related to Figure 2.*** (A) Flow cytometry plot quantifying GP2 expression in RFP-H1-derived PPs at stage 4. (B) Live imaging of mice without transplantation (No Tx), or transplantation of an empty collagen hydrogel (sham) at week 1 post-transplantation.



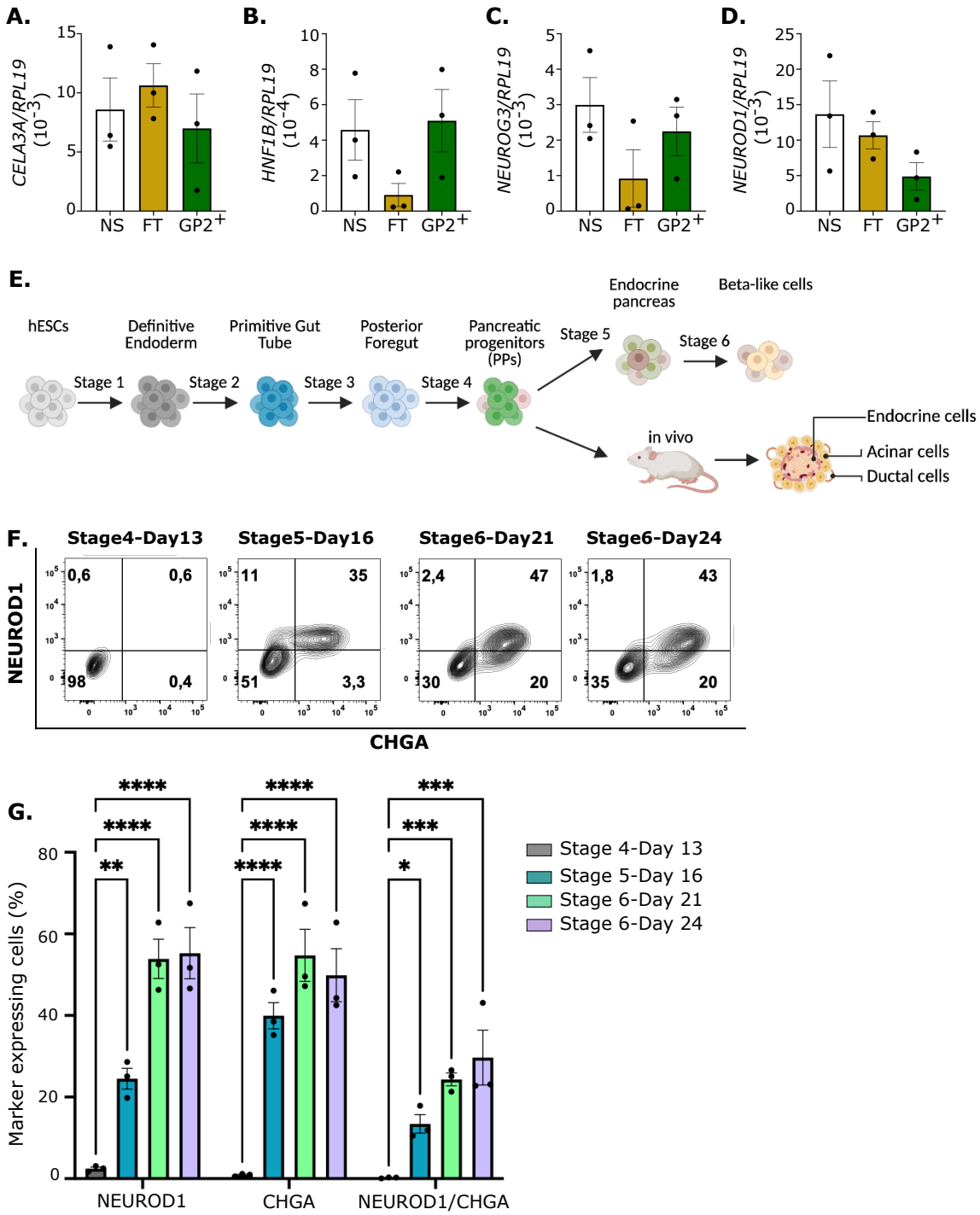
***Supplemental Figure 3. Related to Figure 2.*** (A-F) Representative images of FT-derived outgrowths stained with H&E (A), toluidine blue (B), and immunostained using SOX2 (C), FOXA2 (D), CDX2 (E), and NKX2-1 (F) antibodies at week 15 post-transplantation. DAPI marks cell nuclei. (A, scale bar is 100 $\mu$ m; C & F, scale bar is 200 $\mu$ m; B, D, E, scale bar is 400 $\mu$ m).





**Supplementary Figure 4. Related to Figure 3.** (A-C) RT-PCR quantification of *ZIC3* (neuroectoderm marker), *PDGFRA* (mesoderm marker) and *POU5F1* (pluripotency marker) levels in NS, FT and GP2<sup>+</sup> cell fractions after MACS normalized to *RPL19* housekeeping gene. HESC-derived NE (neuroectoderm), MD (mesoderm), and undifferentiated hESCs are used as a positive control (NE, NS, FT, GP2<sup>+</sup>, hESC, n=3 independent experiments; MD, n=2 independent experiments; One-way ANOVA analysis with Tukey's multiple comparisons; \* P< 0.05, \*\*\* P<0.001). (D, E) Representative flow cytometry plot and quantification of CXCR4 and KIT expression in hESC-derived endodermal cells, indicating robust differentiation efficacy (~ 100%). (F) RT-PCR quantification of *GP2* levels normalized to *RPL19* housekeeping gene in NS, FT, and GP2<sup>+</sup> fractions after MACS confirming enrichment of GP2<sup>+</sup> cells (n=3 independent experiments; One-way ANOVA analysis with Tukey's multiple comparisons; \* P< 0.05, \*\* P<0.01).

Supplementary Figure 5



**Supplementary Figure 5. Related to Figure 3.** (A-D) RT-PCR quantification of *CELA3A* (acinar marker), *HNFBI* (ductal marker), *NEUROG3* and *NEUROD1* (endocrine markers) normalized to *RPL19* housekeeping gene in NS, FT and GP2<sup>+</sup> fractions after MACS. (E) Schematic of hESC differentiation to PP (Stage 4, Days 8-13), which can be transplanted *in vivo* to generate endocrine pancreas, acinar and ductal cells or differentiated *in vitro* to endocrine pancreatic cells (Stage 5, Days 13-16), and Beta-like cells (Stage 6, Days 16-24). (F) Representative flow cytometry plots of NEUROD1 and Chromogranin (CHGA) expression, which indicates commitment to the endocrine pancreas. (G) Quantification of the frequency of endocrine-committed cells at various differentiation stages. (n=3 independent experiments; One-way ANOVA with Tukey's multiple comparisons; \* P<0.05, \*\* P<0.01, \*\*\* <0.001, \*\*\*\* P<0.0001).

**Supplementary Tables:**

**Supplementary Table 1.** List of antibodies used for immunostaining (I) and/or flow cytometry (F). Related to Figures 1-5.

Antigen	Species	Vendor	Dilution
CDX2	Rabbit	Abcam	1:300 (I), 1:2000 (F)
Chromogranin A (CHGA)	Rabbit	Abcam	1:100 (I), 1:1000 (F)
CD117 (KIT)-PE conjugated	Mouse	Invitrogen	1:100 (F)
CD184 (CXCR4)-APC conjugated	Mouse	BD Biosciences	1:200 (F)
Cytokeratin 19 (KRT19)	Rabbit	Abcam	1:800 (I)
C-peptide	Rat	Developmental Studies Hybridoma Bank	1:1000 (I)
FOXA2	Rabbit	Abcam	1:500 (I)
Glucagon (GCG)	Mouse	Sigma-Aldrich	1:500 (I)
GP2	Human	MBL	1:800 (F)
KI67	Rabbit	Abcam	1:1000 (I, F)
KU80	Rabbit	Cell Signaling	1:450 (I)
NKX2-1	Rabbit	Abcam	1: 250 (I)
NKX6-1	Mouse	Developmental Studies Hybridoma Bank	1:1000 (F)
NEUROD1-AF657 conjugated	Mouse	BD Bioscience	1:3 (F)
PDX1	Goat	Abcam	1:10000 (I)
PDX1	Goat	R&D Systems	1:100 (F)
SLC18A1	Rabbit	Sigma	1:1000 (I)
Somatostatin (SST)	Rabbit	Abcam	1:500 (I)
SOX2	Rabbit	Cell signaling	1:400 (I, F)
Trypsin (PRSS1)	Sheep	R&D Systems	1:300 (I)
IgG-AF488	Goat	Jackson Immuno Research Labs	1:400 (F)
IgG-AF488	Mouse	Jackson Immuno Research Labs	1:800 (I)
IgG-AF488	Rabbit	Jackson Immuno Research Labs	1:800 (F)

IgG-Cy5	Mouse	Jackson Immuno Research Labs	1:800 (I)
IgG-Cy5	Rabbit	Jackson Immuno Research Labs	1:800 (I)
IgG-Cy5	Rat	Jackson Immuno Research Labs	1:800 (I)
IgG-AF647	Mouse	Jackson Immuno Research Labs	1:400 (F), 1:800 (I)
IgG-AF549	Mouse	Jackson Immuno Research Labs	1:800 (I)
IgG-AF549	Rabbit	Jackson Immuno Research Labs	1:800 (I)
IgG-R-Phycoerythrin-conjugated	Mouse	Jackson Immuno Research Labs	1:800 (F)

**Supplementary Table 2.** List of QPCR primers. *Related to Figures 3 and 4.*

<i>Gene</i>	<i>Forward primer (5' to 3')</i>	<i>Reverse primer (3' to 5')</i>
<i>CDX2</i>	<i>AGC CAA CCT GGA CTT CCT GTC ATT</i>	<i>ACA CAG ACC AAC AAC CCA AAC AGC</i>
<i>CELA3A</i>	<i>ATG ACA TGC CCC TCA TCA AGC TCT</i>	<i>ATG TAG CAG GGT GTC TTG TTG GGA</i>
<i>GP2</i>	<i>AAC CCT TCC GAA GCA CAG AG</i>	<i>GGA CAC AGG TCT CCG ACA TC</i>
<i>HNF1B</i>	<i>AGA GTA ACA TGC CAG CTT CCT CCT GTG</i>	<i>TAT CAA ACA GCC AGT TTC CCT CCT GCC</i>
<i>MKI67</i>	<i>AAT TTG CTT CTG GCC TTC CC</i>	<i>GAC CCC GCT CCT TTT GAT AGT</i>
<i>NEUROD1</i>	<i>TCC CAT GTC TTC CAC GTT AAG CCT</i>	<i>CAT CAA AGG AAG GGC TGG TGC AAT</i>
<i>NEUROG3</i>	<i>GCG CAA TCG AAT GCA CAA CCT CAA</i>	<i>TTC GAG TCA GCG CCA AGA TGT AGT T</i>
<i>NKX2-1</i>	<i>GCC AAA CTG CTG GAC GTC TTT CTT</i>	<i>CCT TGA GAT TGG ATG CGC TTG GTT</i>
<i>POU5F1</i>	<i>ATG CAT TCA AAC TGA GGT GCC TGC</i>	<i>CCA CCC TTT GTG TTC CCA ATT CCT</i>
<i>PDGFRA</i>	<i>GGCAGTACCCCATGTCTGAAG</i>	<i>CGTCACAAAAAGGCCGCTG</i>
<i>RPL19</i>	<i>CTCGATGCCGGAAAAACACC</i>	<i>TTCTCTGGCATTCTGGGCATT</i>
<i>SOX2</i>	<i>GGA TAA GTA CAC GCT GCC CG</i>	<i>ATG TGC GCG TAA CTG TCC AT</i>
<i>TOP2A</i>	<i>GCT GCC CCA AAA GGA ACT AA</i>	<i>GGC GAT TCT TGG TTT TGG CA</i>
<i>ZIC3</i>	<i>TAT CAG TCT CGC GCT CAC</i>	<i>TGT CTT TTG CGG TTT ATC TTC CTG</i>