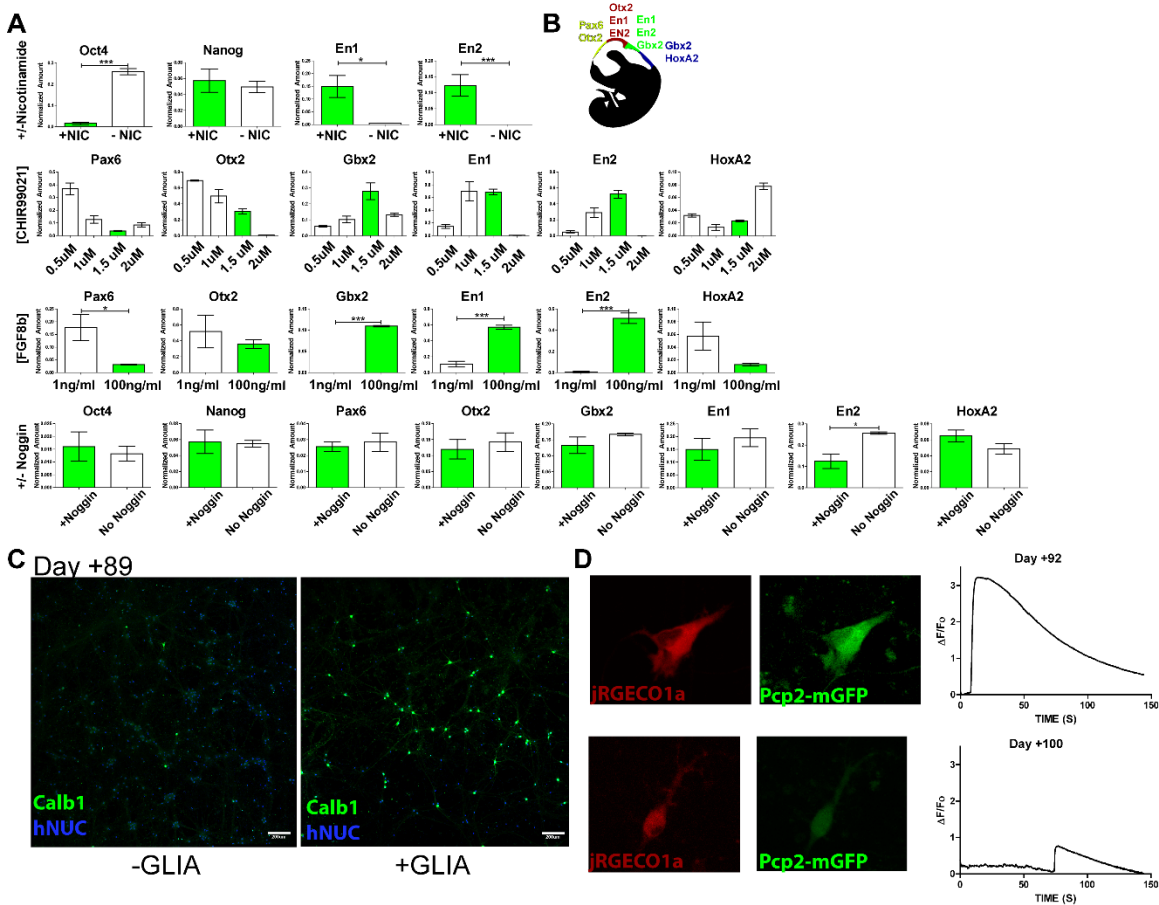
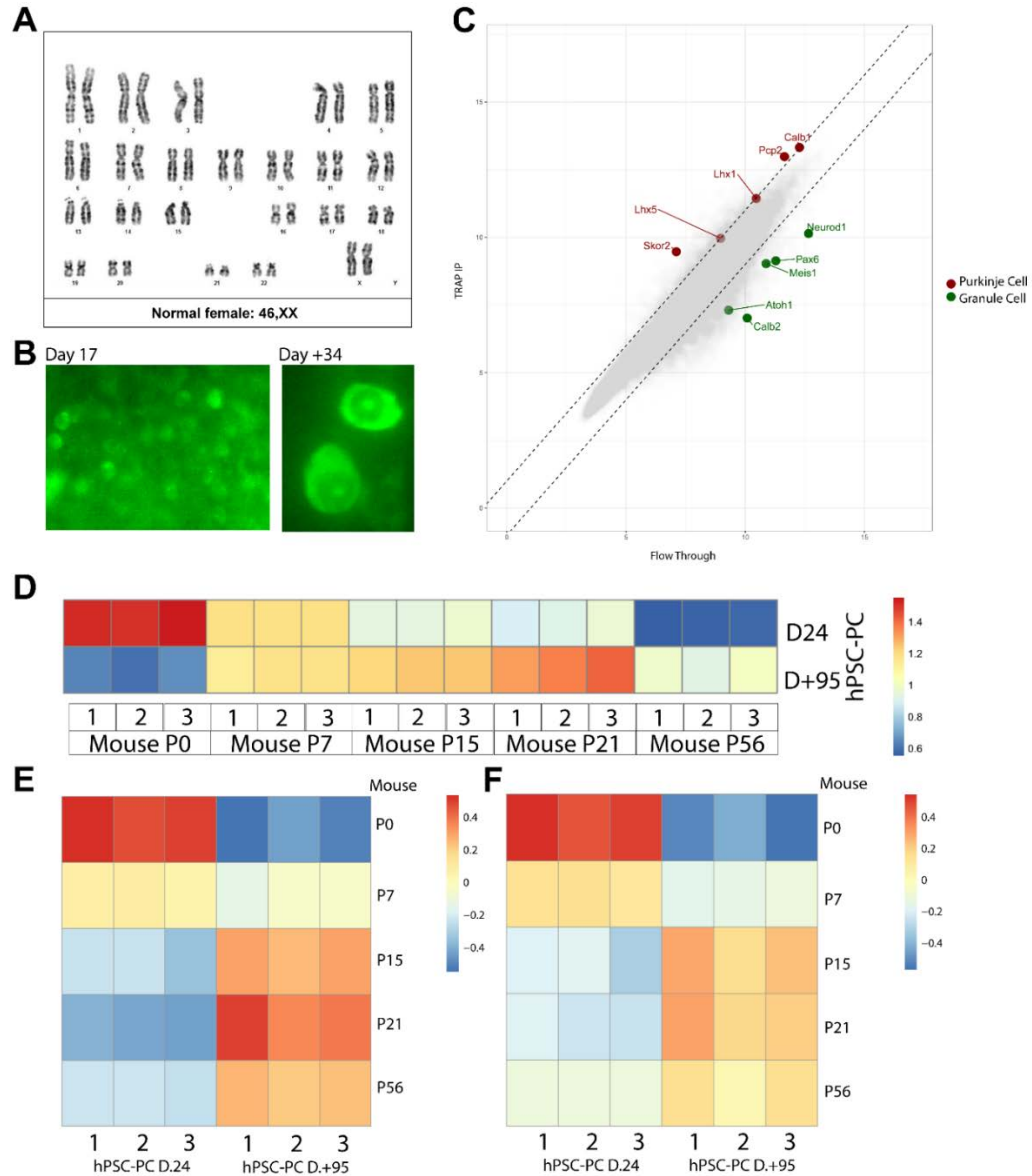
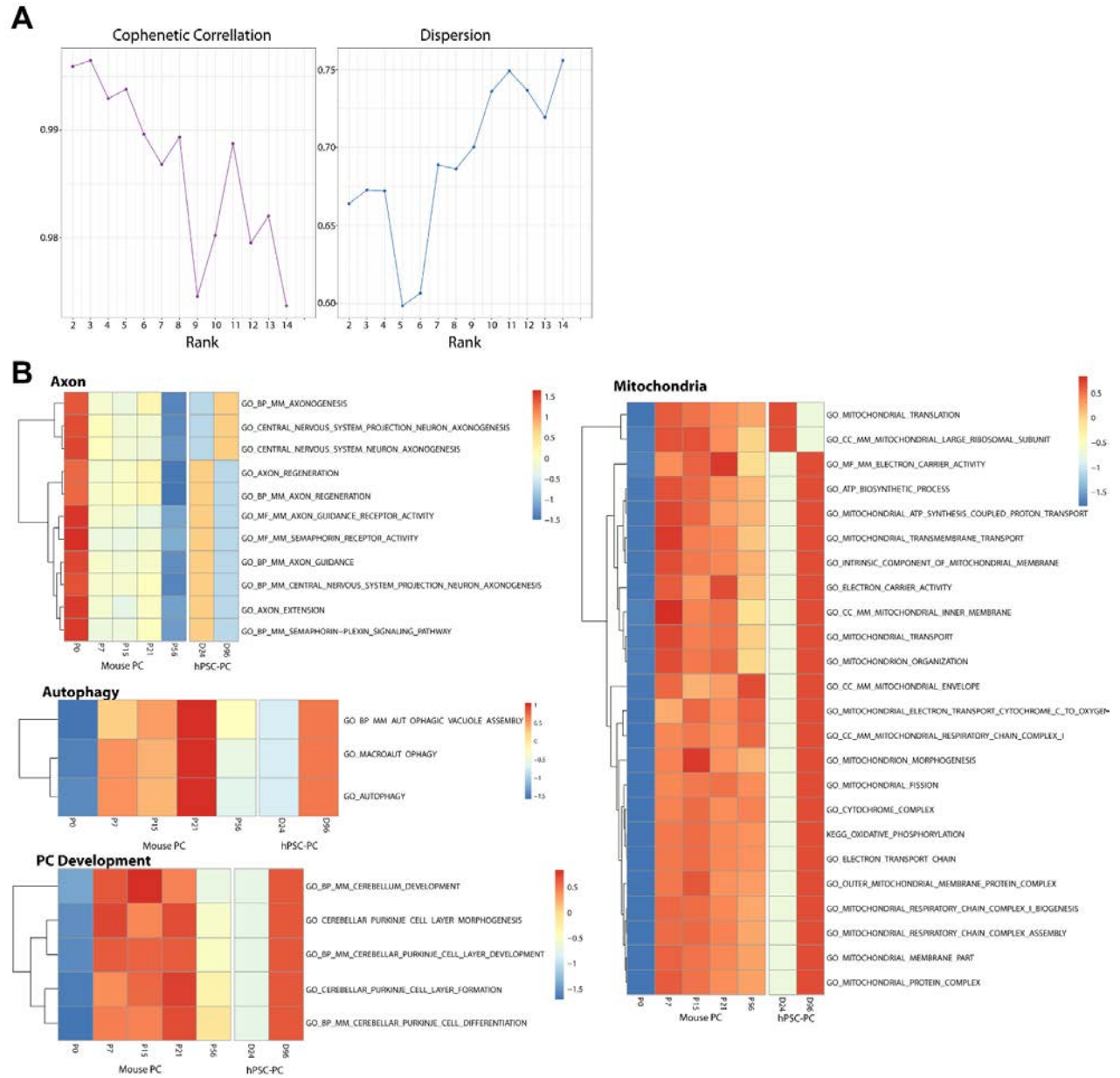


## Supplemental Figures



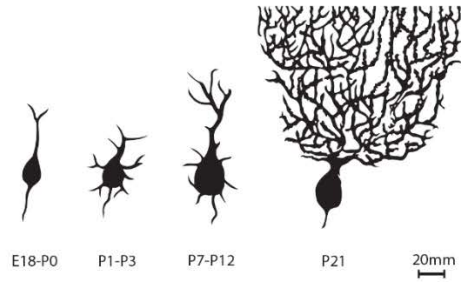


**Supplemental Figure 2. Translational Profiling in hPSC-PCs and Mouse PCs.** **A.** Karyotype analysis of RUES2-PCP2-L10a-EGFP stem cell line. **B.** Live imaging of the L10a-EGFP TRAP reporter after 17 days of differentiation and an additional +34 days of differentiation following isolation and co-culture. **C.** Test for enrichment of PC markers and depletion of GC markers from TRAP isolation of RNA from a P7 *Tg(Pcp2-L10a-Egfp)* TRAP mouse, dotted lines represent Log<sub>2</sub> fold change. **D.** Heat map depicting expression levels in mouse PCs over postnatal development of gene sets defined as log<sub>2</sub> four-fold change between Day 24 and Day +95 in hPSC-PCs following subtraction of mouse genes from hPSC-PC data using an in silico human-mouse reference genome (34). Day 24 hPSC-PCs are most similar to P0 mouse PCs ( $p=5.26 \times 10^{-17}$ ). Day +95 PCs are most similar to P21 mouse PCs ( $p=3.73 \times 10^{-5}$ ). **E.** Heat map depicting expression levels in hPSC-PCs of mouse gene sets defined as the 100 most expressed genes per time point. **F.** Heat map depicting expression level in hPSC-PCs of mouse gene sets defined as the 100 most expressed genes per time point following subtraction of mouse genes from hPSC-PC data using an in silico human-mouse reference genome (34).

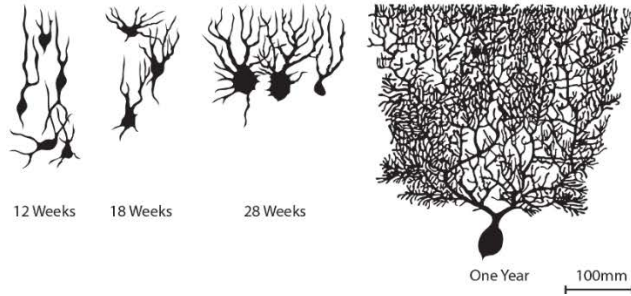


**Supplemental Figure 3. NMF metagene analysis of mouse PC TRAP Data. A.** Cophenetic correlation and dispersion by rank (metagene number). Rank 5 showed high cophenetic correlation with low dispersion. **B.** Heat maps depicting all gene ontology terms for the gene ontology signatures (Axon, Autophagy, PC Development, Mitochondria) in Figure 3.

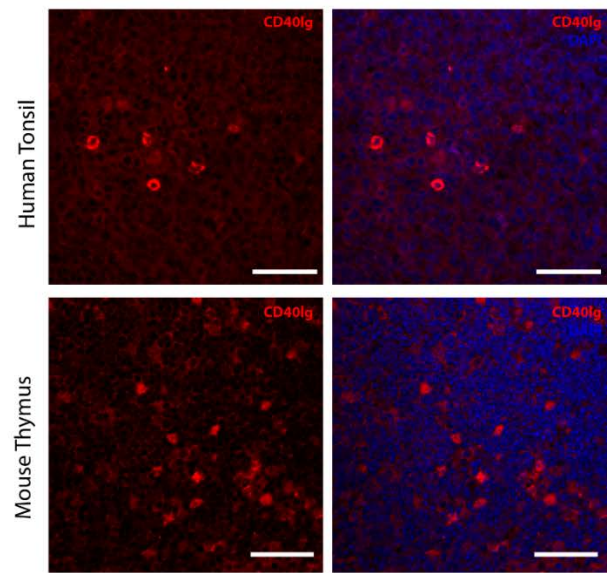
**A - Mouse PC Development**



**B - Human PC Development**



**C**



**Supplemental Figure 4. Species differences.** **A.** Schematic of mouse PC development. (Adapted from (76) and (77)). **B.** Schematic of human PC development (Adapted from (27)). **C.** Positive controls for anti-human CD40LG antibody and anti-mouse CD40LG antibody. Scale bars = 50µm.

Gene	Forward	Reverse	Source
Oct4	CAGTGCCCGAAACCCACAC	GGAGACCCAGCAGCCTCAA	(78)
Nanog	CAGAAGGCCTCAGCACCTA C	ATTGTTCCAGGTCTGGTTGC	(78)
Pax6	TCACCATGGCAAATAACCTG	CAGCATGCAGGAGTATGAGG	Designed on Primer3
Otx2	ACAAGTGGCCAATTCCTCC	GAGGTGGACAAGGGATCTGA	Designed on Primer3
Gbx2	GTTCCCGCCGTCGCTGATG AT	GCCGGTGTAGACGAAATGGCC G	Designed on Primer3
En1	GAGCGCAGGGCACCAAATA	CGAGTCAGTTTTGACCACGG	Primerbank (126090908c 1)
En2	GGCGTGGGTCTACTGTACG	TACCTGTTGGTCTGGAACCTCG	Designed on Primer3
HoxA2	CGTCGCTCGCTGAGTGCCT G	TGTCGAGTGTGAAAGCGTCGA GG	(24)
GPI3'	GGACCACGAGCCCTTAGC	AACACTTCAGCCAATTCTAACA C	(65)
GPI5'	CGTCATCAACATTGGCATTG G	GGGACCTCCTGAAGAGTATGG	(65)
HMBS	TGCTATCTGGGGAGTGATTA CC	GGCTGTTGCTTGACTTCTC	(65)
GAPD	AGCAAGAGCACAAAGAGGAA GAG	GAGCACAGGGTACTTTATTGAT GG	(65)

**Supplementary Table 1. qPCR Primer Sequences.**

<b>Antibody</b>	<b>% Triton X-100</b>	<b>Dilution</b>	<b>Source</b>
En1	0.3	1:10,000	Gift of T. Jessell
Otx2	0.3	1:2000	Millipore Cat# AB9566, RRID:AB_2157186
Kirrel2	0.3	1:250	R&D Systems
Corl2(Skor2)	0.3	1:200	Atlas Antibodies Cat# HPA046206, RRID:AB_2679588
Pcp2(L7)	0-0.1	1:500	Takara Cat#M202
Calb1	0.3	1:250	Swant Cat# 300, RRID:AB_10000347
Calb1	0.3	1:500	Swant Cat# CB38, RRID:AB_2721225
GD3	0	1:200	BioLegend Cat# 917701, RRID:AB_2565200
NCAM(CD56)	0	1:200	BioLegend Cat# 304601, RRID:AB_314443
hNUC	0.3	1:200	Millipore Cat# MAB1281, RRID:AB_94090
Syn1	0.3	1:500	Sigma-Aldrich Cat# S193, RRID:AB_261457
GFP	0.3	1:4000	Aves Labs Cat# GFP- 1020, RRID:AB_10000240
Human CD40lg	0.05 cell culture, 0.1 paraffin	1:50	R and D Systems Cat# MAB617, RRID:AB_2291414
Mouse CD40lg	0.05	1:50	R and D Systems Cat# AF1163, RRID:AB_35463

**Supplemental Table 2. Antibody List.**