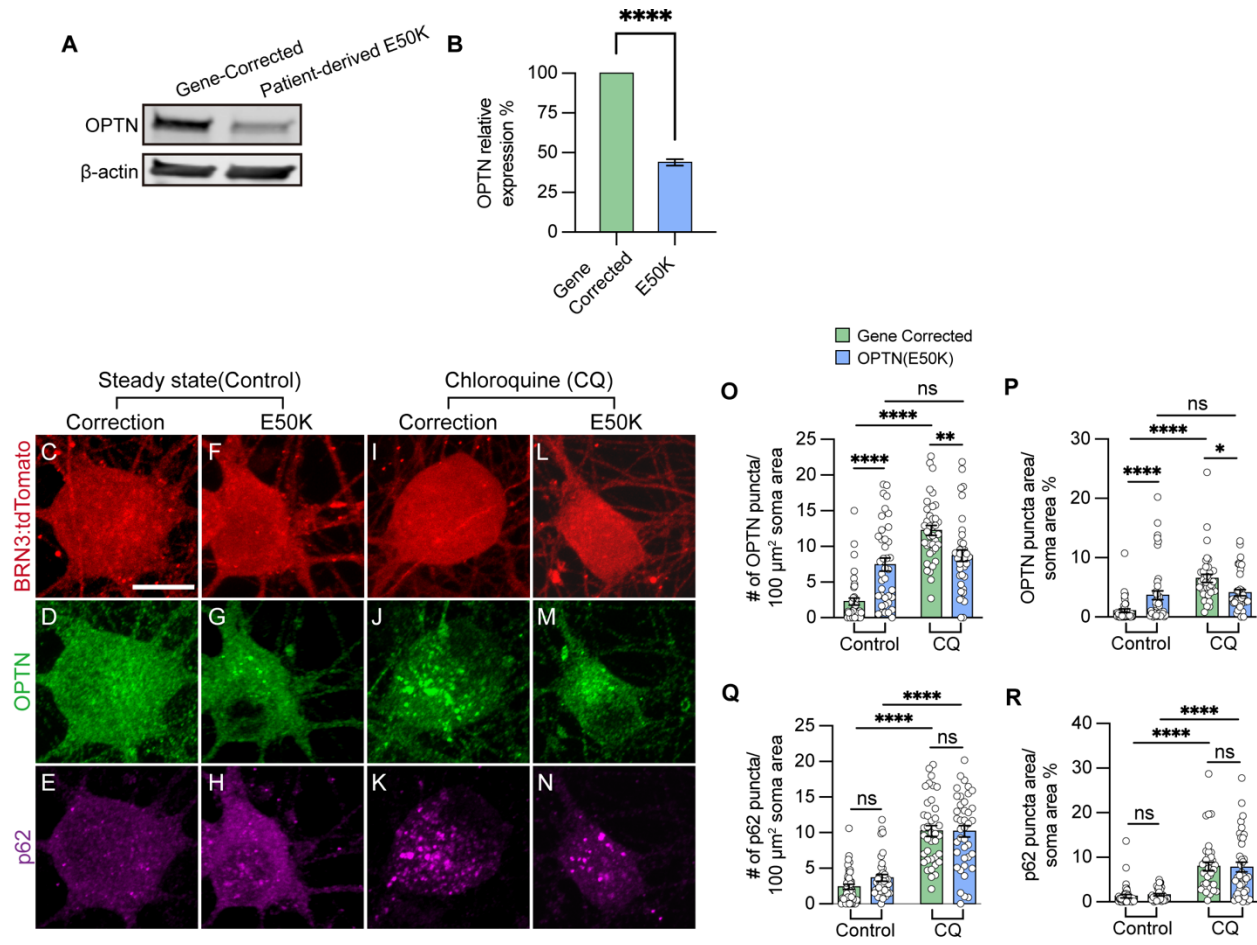


1194 **Supporting information**



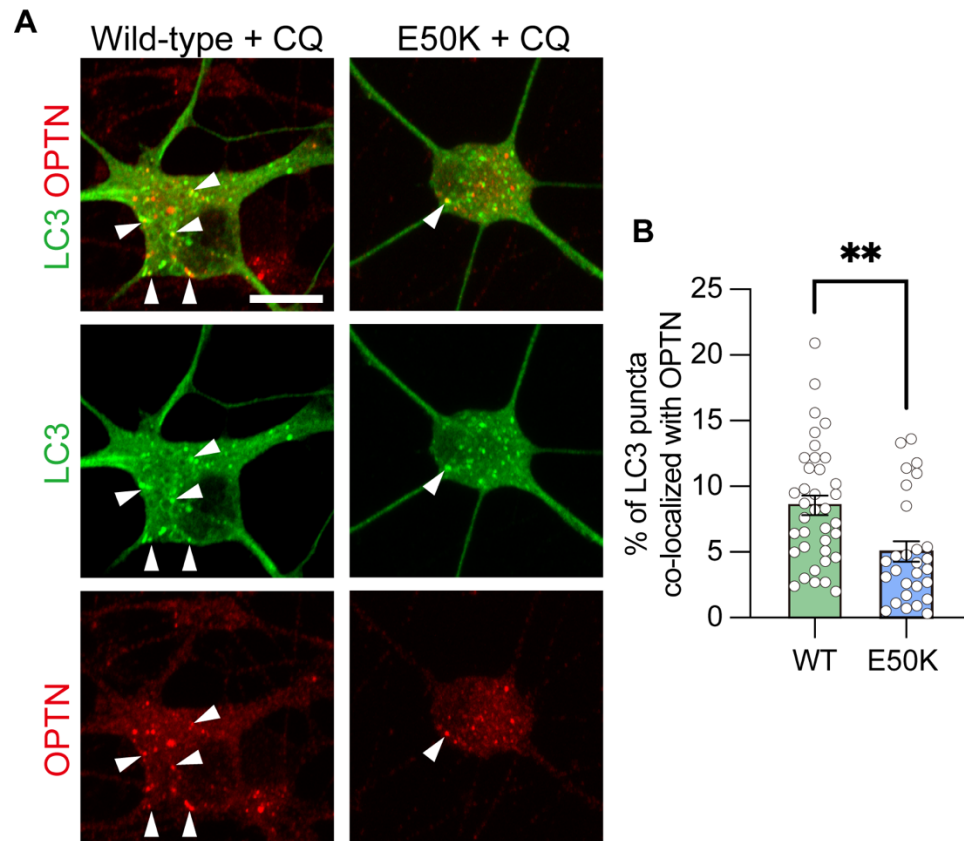
1195

1196 **Supplemental Figure 1. Confirmation of changes in OPTN levels in RGC-derived from**  
 1197 **patient-specific iPSCs with OPTN(E50K) mutation compared to corresponding isogenic**  
 1198 **control line.** (A-B) Western blot of OPTN relative to  $\beta$ -actin in patient-derived OPTN(E50K) and  
 1199 gene-corrected isogenic control iPSC-RGCs (n=5 for each WT and E50K; t-test, \*\*\*\*p<0.0001).  
 1200 (C-N) Immunostaining revealed the aggregation of OPTN and p62 puncta in BRN3:tdTomato  
 1201 iPSC-RGCs in patient-derived OPTN(E50K) mutant and gene-corrected isogenic control lines  
 1202 after steady state (control) (C-H) and chloroquine (CQ) treatment (I-N). Scale bar: 10  $\mu$ m. (O-R)  
 1203 Quantification of OPTN puncta (O and P) or p62 puncta (Q and R) in iPSC-RGC (n=3 biological  
 1204 replicates using Ctrl-WT n=42, Ctrl-E50K n=39, CQ-WT n=38 and CQ-E50K n=40 technical

1205 replicates; One-way ANOVA, Tukey post hoc test. \*\*\*\*p<0.0001, \*\*p<0.01, \*p<0.05, ns= not  
1206 significant). Data are all represented as mean values  $\pm$  S.E.M.

1207 **Supplemental figure 1-source data 1. The alternation of OPTN protein in gene corrected**  
1208 **and patient derived-OPTN(E50K) mutation hPSC-RGCs.**

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1211 **Supplemental Figure 2. Confirmation of reduced recruitment of LC3 by OPTN in patient-**

1212 **derived OPTN(E50K) iPSC-RGCs after chloroquine treatment.** (A) Representative images of

1213 OPTN and LC3 localization in patient-derived iPSC-RGCs from wild-type and OPTN(E50K) cell

1214 lines after chloroquine treatment. White arrows identify puncta colocalized with OPTN and LC3.

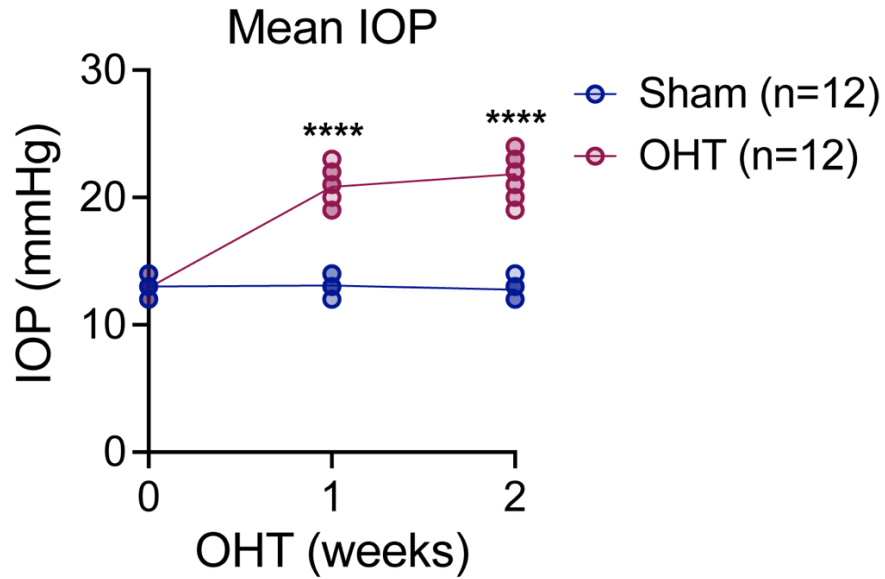
1215 (B) Quantification of colocalization between OPTN and LC3 in patient-derived iPSC-RGCs (n=3

1216 biological replicates using WT n=37 and E50K n=28 technical replicates; t-test, \*\*p>0.005).

1217 Scale bar: 10  $\mu$ m. Data are all represented as mean values  $\pm$  S.E.M.

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1221 **Supplemental Figure 3. Elevation of intraocular pressure in an ocular hypertension**

1222 **glaucoma model.** Maintained elevation of intraocular pressure in mouse eyes following the

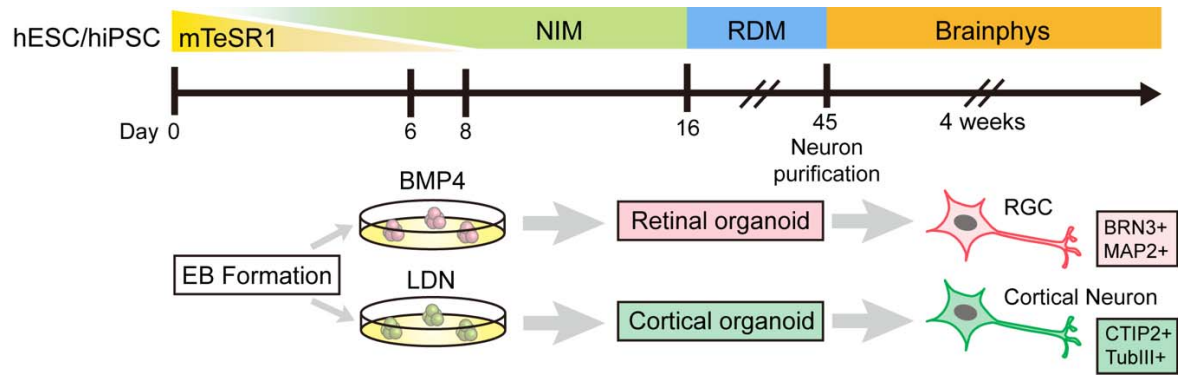
1223 injection of magnetic microbeads into the mouse anterior chamber, compared to sham-injected

1224 controls. Two-way ANOVA with Tukey's multiple comparison pos hoc test, \*\*\*\*p < 0.0001,

1225 n = 12 mice/group.

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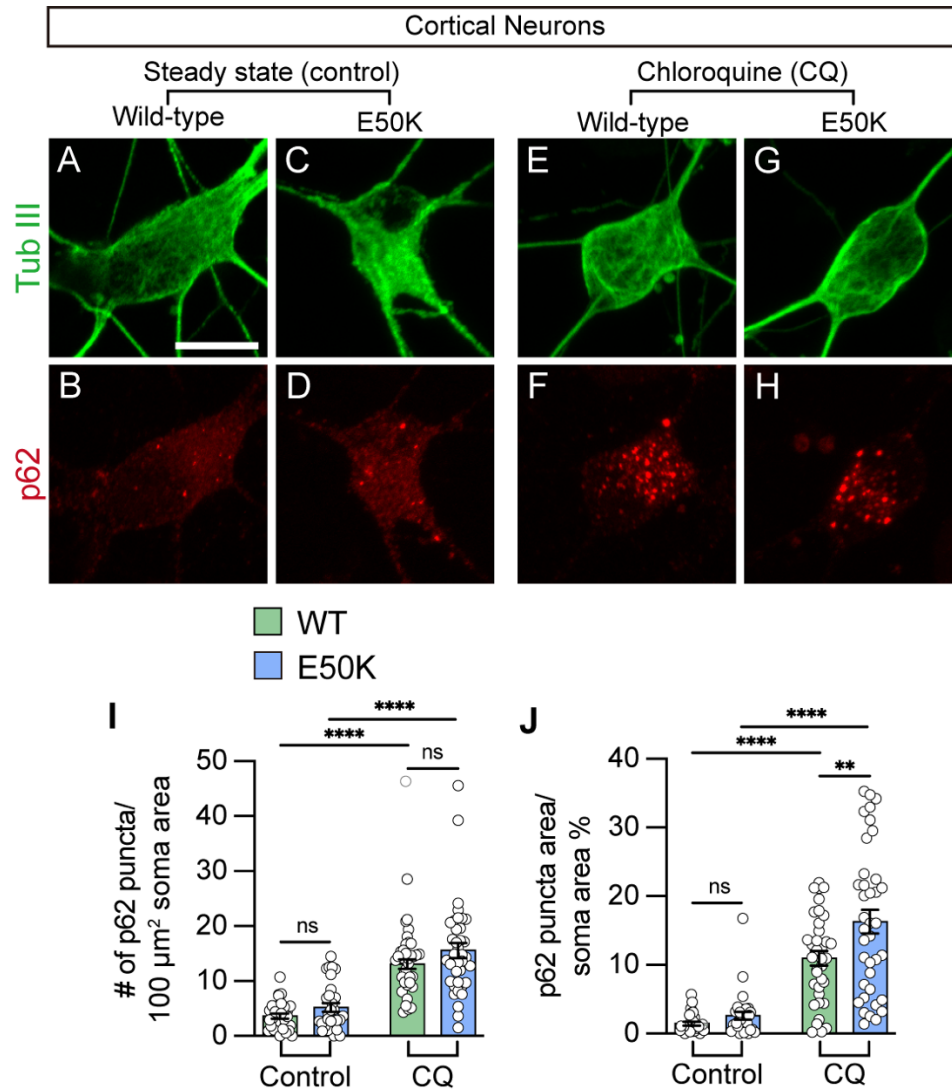
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1229 **Supplemental Figure 4. Schematic diagram outlining the differentiation of RGCs and**

1230 **cortical neurons from hPSCs.**

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1234 **Supplemental Figure 5. Characterization of p62 expression in hPSC-cortical neurons**

1235 **from wild-type and OPTN(E50K) cell lines. (A-H) Representative images of p62 puncta in**

1236 **iPSC-derived cortical neurons from WT and OPTN(E50K) cell lines under steady state (control)**

1237 **(A-D) and chloroquine (CQ) treatment (E-H). Scale bar: 10  $\mu\text{m}$ . (I-J) Quantification of p62**

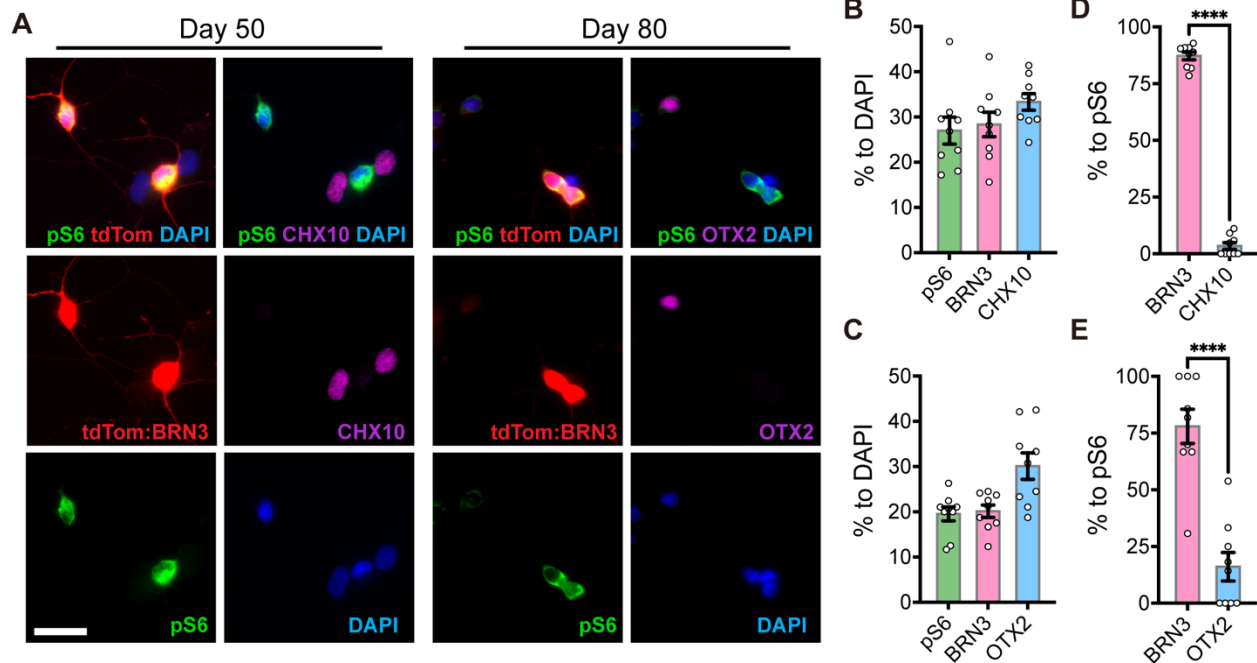
1238 **puncta in iPSC-derived cortical neurons (n=3 biological replicates using Ctrl-WT n=30, Ctrl-**

1239 **E50K n=30, CQ-WT n=36 and CQ-E50K n=38 technical replicates; One-way ANOVA, Tukey**

1240 **post hoc test. \*\*\*\*p<0.0001, \*\*p<0.01, ns= not significant, p>0.05). Data are all represented as**

1241 **mean values  $\pm$  S.E.M.**

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1244 **Supplemental Figure 6. mTORC1 activity is preferentially observed within RGCs among**

1245 **hPSC-derived retinal cells.** (A) Retinal organoids were dissociated and plated onto laminin-

1246 coated coverslips at either day 50 or day 80 of differentiation to acquire the majority of major

1247 retinal cell types, including RGCs (BRN3:tdTomato), retinal progenitor cells (CHX10), and

1248 photoreceptors (OTX2), following by analysis of mTORC1 activity based upon co-staining with

1249 pS6Ser240/244. Scale bar: 25  $\mu$ m. (B-C) Quantification of results showing the percentage of

1250 retinal cell types observed at day 50 or day 80 (n=9 images from three technical replicates). (D-

1251 E) Quantification of pS6Ser240/244 expression colocalized with either BRN3B:tdTomato,

1252 CHX10 or OTX2, suggesting that mTORC1 signaling is highly expressed in RGCs, with little

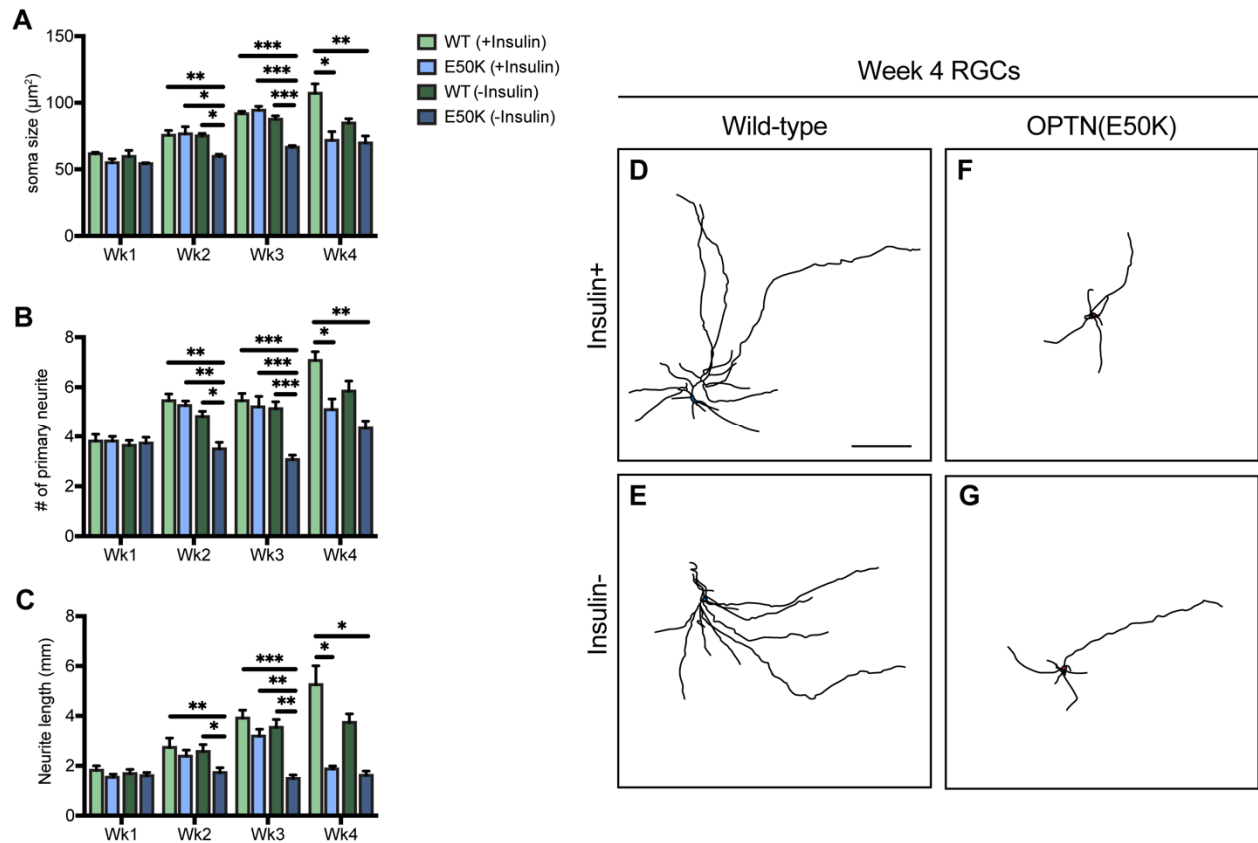
1253 expression in retinal progenitor cells or photoreceptors (n=9 images from three technical

1254 replicates; t-test, BRN3 vs CHX10:  $p < 0.0001$ , BRN3 vs OTX2:  $p < 0.0001$ ). Data are all

1255 represented as mean values  $\pm$  S.E.M.

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**Supplemental Figure 7. Insulin deprivation expedites the onset of degenerative**

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**phenotypes in hPSC-RGCs with the OPTN(E50K) mutation.** (A-C) Quantitative analysis of

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neurite measurements over the course of 4 weeks of differentiation in wild-type and

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OPTN(E50K) hPSC-RGCs grown with or without insulin, as measured by soma size ( $n \geq 30$  each

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condition with 4 biological replicates; One-way ANOVA, Tukey post hoc test.  $***p < 0.001$ ,

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$**p < 0.01$ ,  $*p < 0.05$ ) (A), number of primary neurites ( $n \geq 10$  each condition with 4 biological

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replicates; One-way ANOVA, Tukey post hoc test.  $***p < 0.001$ ,  $**p < 0.01$ ,  $*p < 0.05$ ) (B), total

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neurite length ( $n \geq 10$  each condition with 4 biological replicates; One-way ANOVA, Tukey post

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hoc test.  $***p < 0.001$ ,  $**p < 0.01$ ,  $*p < 0.05$ ) (C). Data are all represented as mean values  $\pm$  S.E.M.

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(D-G) Representative neurite tracings of WT and OPTN(E50K) hPSC-RGCs after 4 weeks of

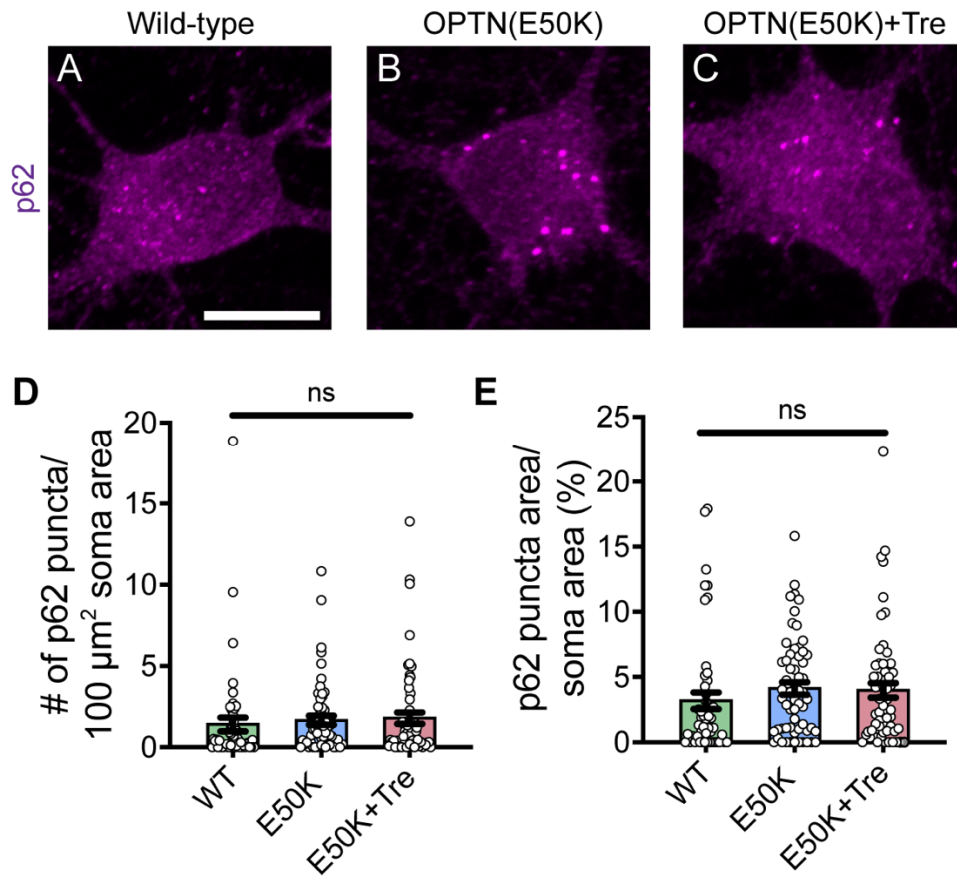
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growth either with or without insulin. Scale bar: 200  $\mu$ m.

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1273 **Supplemental Figure 8. p62 expression remains unchanged in hPSC-RGCs comparing**

1274 **wild-type, OPTN(E50K) and OPTN(E50K) plus trehalose conditions. (A-C) Representative**

1275 images of p62 puncta in hPSC-RGCs. Scale bar: 10  $\mu\text{m}$ . (D-E) Quantification of p62 puncta in

1276 hPSC-RGCs (n=3 biological replicates using WT n=51, E50K n=60, and E50K-trehalose n=61

1277 technical replicates; One-way ANOVA, Tukey post hoc test. ns= not significant,  $p>0.05$ ). Data

1278 are all represented as mean values  $\pm$  S.E.M.

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1281 **Supplemental Table. List of antibodies.**

Antibody	Type	Source	Catalog	RRID	WB dilution	IF dilution
AKT	Mouse monoclonal	Cell Signaling Technology	2920	AB_1147620	1:2000	
AMPK	Rabbit polyclonal	Cell Signaling Technology	2532	AB_330331	1:1000	
$\beta$ -actin	Mouse monoclonal	Sigma	A5441	AB_476744	1:10000	
BRN3	Goat polyclonal	Santa Cruz	SC-6026	AB_673441		1:200
CHX10	Goat polyclonal	Santa Cruz	SC-21690	AB_2216006		1:200
CTIP2	Rat monoclonal	Abcam	Ab18465	AB_2064130		1:500
LAMP1	Rabbit monoclonal	Cell signaling Technology	9091	AB_2687579	1:1000	
LAMP1	Rat monoclonal	DSHB	1D4B	AB_2134500	1:200	1:20
LC3 A/B	Rabbit monoclonal	Cell Signaling Technology	12741	AB_2617131	1:1000	
MAP1LC3A	Rabbit monoclonal	Abcam	ab185036	AB_881226		1:200
MAP2	Mouse monoclonal	Synaptic Systems	188011	AB_11042001		1:200
OPTN	Rabbit polyclonal	Novus	NBP1-84682	AB_11032496	1:1000	1:200
OTX2	Goat polyclonal	R&D Systems	AF1979	AB_2157172		1:2000
p62	Mouse monoclonal	Abcam	ab56416	AB_945626	1:2000	1:50
pAKT	Rabbit monoclonal	Cell Signaling Technology	4060	AB_2315049	1:2000	
pAMPK	Rabbit monoclonal	Cell Signaling Technology	2535	AB_331250	1:1000	
pS6	Rabbit polyclonal	Cell Signaling Technology	2215	AB_331682	1:1000	1:200
RBPMS	Guinea pig polyclonal	PhosphoSolutions	1832-RBPMS	AB_2492226		1:500
RFP	Rabbit polyclonal	Rockland	600-401-379	AB_2209751		1:200
RFP	Goat polyclonal	Origene	AB1140-100	AB_2877097		1:200
RFP	Mouse monoclonal	Rockland	200-301-379	AB_2611063		1:200
S6	Rabbit monoclonal	Cell Signaling Technology	2217	AB_331355	1:1000	

Tubulin, $\beta$ -III	Rabbit polyclonal	Biologend	802001	AB_2564645		1:500
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