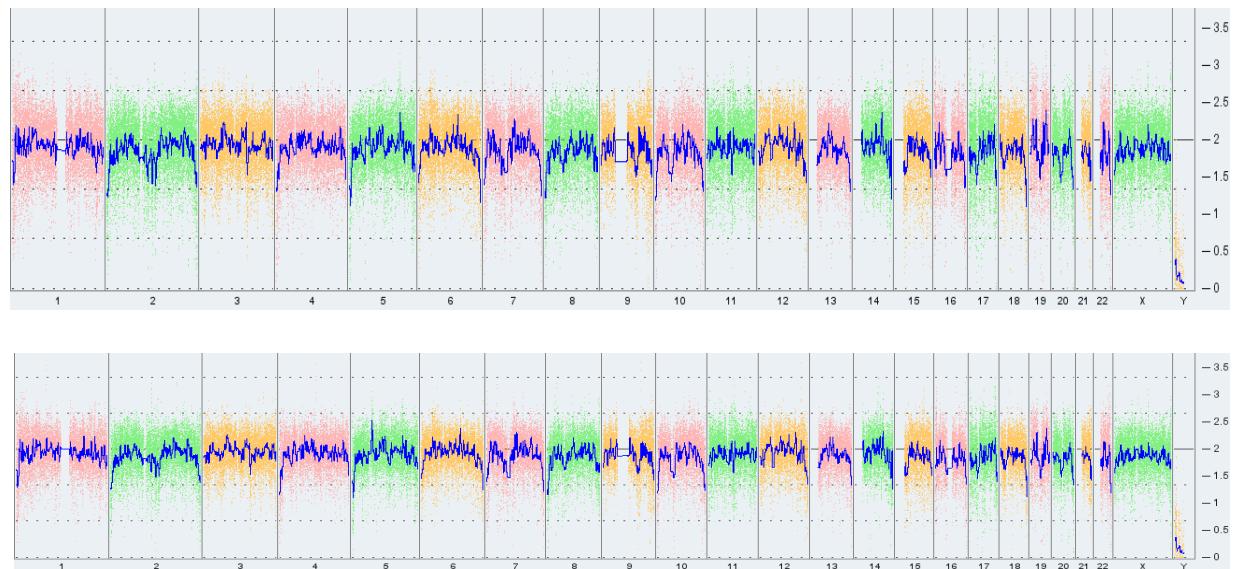


**Supplemental information**

**Microglia-containing cerebral organoids derived from induced pluripotent stem cells for the study of neurological diseases**

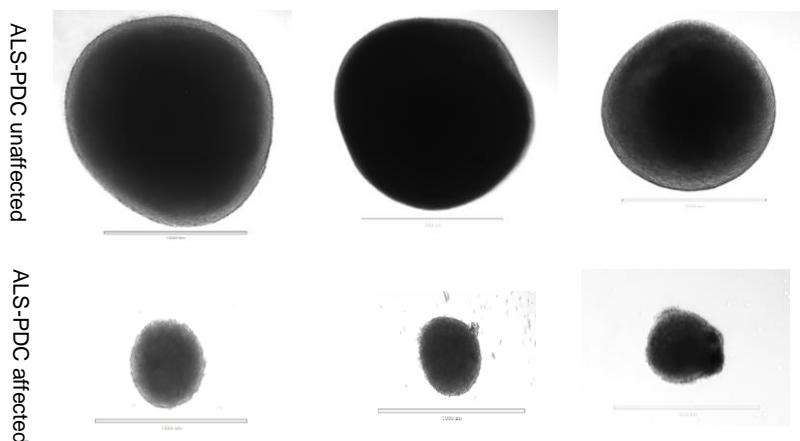
**Yiling Hong, Xu Dong, Lawrence Chang, Chen Xie, Mariann Chang, Jose S. Aguilar, Jimmy Lin, Juncheng Lin, and Qingshun Q. Li**

## Supplementary Figures

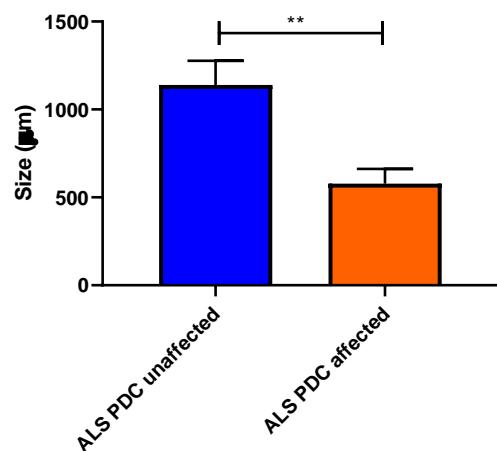


**Figure S1, related to figure 2. Karyotyping in whole genome views.** KaryoStat-based analysis revealed that the sample originated from a female. No chromosomal aberrations were found either in ALS-PDC-affected or ALS-PDC-unaffected iPSCs. The whole-genome view displays all somatic and sex chromosomes in one frame with high-level copy number. The smooth signal plot (right y-axis) is the smoothing of the log<sub>2</sub> ratios that depict the signal intensities of probes in the microarray. A value of 2 represents a normal copy number state (CN=2). A value of 3 represents chromosomal gain (CN=3). A value of 1 represents chromosomal loss (CN=1). The pink, green, and yellow colors indicate the raw signal for each chromosome probe; the blue signal represents the normalized probe signal used to identify copy number and aberrations (if any). The top panel refers to ALS-PDC-unaffected iPSCs and the bottom panel to ALS-PDC-affected iPSCs.

**A.**



**B.**



**Figure S2, related to figure 2. Organoid size comparisons.** (A) Phase-contrast images of selected ALS-PDC-affected and -unaffected organoids. (B) Quantification of organoid size. All data were obtained from three separate images and compared to the ALS-PDC-unaffected conditions: \*\* $p<0.01$ , error bars=SEM.

**Table S1, related to figure 6. Primers used for the qPCR analyses**

Gene	Forward	Reverse
PNPO	AGGCTGTTCAAGTGTCCCTGAC	AACATGCGAGCAGAGGGTT
INF- $\alpha$	GACTCCATCTGGCTGTGA	TGATTCTGCTCTGACAACCT
INF- $\beta$	CAACTTGCTTGGATT CCTACAAAG	TATTCAAGCCTCCCATTCAATTG
INF- $\rho$	TGGCTTTCAAGCTCTGCATC	CCGCTACATCTGAATGACCTG
IFNAR1	GTGATACACATCTCTCCTGG	GTATAATCCCATTAAAGAACATAG
IFNAR2	GAGTAAACCAGAAGATTGAAG	CGTGTGGAATTAACTTGTC
MDA5/ IFIH1	AGAGTGGCTGTTACATTGCC	GCTGTTCAACGTAGCAGTACCTT
IFITM1	ACTCCGTGAAGTCTAGGGACA	TGTCACAGAGCCGAATACCAG
IFITM2	ATTGTGCAAACCTCTCTCCTG	ACCCCCAGCATAGCCACTTCCT
TGF $\beta$ 1	CAATT CCTGGCGATACCTCAG	GCACA ACTCCGGTGACATCAA
TGF $\beta$ 2	CAGCACACTCGATATGGACCA	CCTCGGGCTCAGGATAGTCT
IL34	AAGGTGGAATCCGTGTTGTCC	AGCTTGTTACAGCAGGAGC
Col9A2	TCCCTGGTGAGATTGGAATCC	TTGGTTGGACACAGGAAATCC
IL1 $\beta$	ATGATGGCTTATTACAGTGGCAA	GTCGGAGATT CGTAGCTGGA
TNF $\alpha$	CCTCTCTCTAATCAGCCCTCTG	GAGGACCTGGAGTAGATGAG
iNOS	GCAGAATGTGACC CATCATGG	ACAA CCTTGGTGTGAAGGC
ROS	TTGTTATGCACCCATCCAAA	CCTGCTGCTCGGATATGAAT
Glast	TGTGGAGCAGCACTGATTTC	AGCCA AACCTCCTCTTTGGT
Iba1	ATGAGCCAAACCAGGGATTAC	GGGATCGTCTAGGAATTGCTTGT