Supplementary Material

Modeling Parkinson's disease in midbrain-like organoids

Lisa M. Smits^{1,§}, Lydia Reinhardt^{2,3,§}, Peter Reinhardt²⁻⁴, Michael Glatza^{2,3}, Anna S. Monzel¹, Nancy Stanslowsky⁵, Marcelo D. Rosato-Siri⁶, Alessandra Zanon⁶, Paul M Antony¹, Jessica Bellmann², Sarah M. Nicklas¹, Kathrin Hemmer¹, Xiaobing Qing¹, Emanuel Berger¹, Norman Kalmbach⁵, Marc Ehrlich³, Silvia Bolognin¹, Andrew A. Hicks⁶, Florian Wegner⁵, Jared L. Sterneckert^{2,3*} and Jens C. Schwamborn^{1*}

¹Luxembourg Centre for Systems Biomedicine (LCSB), Developmental and Cellular Biology, University of Luxembourg, Belvaux Luxembourg

²DFG-Center for Regenerative Therapies, Technische Universität Dresden, Dresden, Germany

³Department of Cell and Developmental Biology, Max Planck Institute for Molecular Biomedicine, Münster, Germany

⁴Currently located at AbbVie Deutschland GmbH $\&$ Co KG, Neuroscience Discovery–Biology Department, Ludwigshafen, Germany

⁵Department of Neurology, Hannover Medical School, Hannover, Germany

⁶Institute for Biomedicine, Eurac Research, Affiliated Institute of the University of Lübeck, Bolzano, Italy

§These authors contributed equally and thus share first authorship.

***Corresponding authors**

Prof. Dr. Jens C. Schwamborn jens.schwamborn@uni.lu Dr. Jared L. Sterneckert Jared.Sterneckert@tu-dresden.de

Supplementary Figures and Tables

Supplementary Figure 1 Derivation of mfNPCs.

Supplementary Figure 1: Derivation of mfNPCs. (a) Schematic illustration of mfNPC derivation and expansion. $SB = SB-431542$, $LDN = LDN-193189$, $SAG = \text{sonic hedgehog}$ agonist. (b) Microarray profiling displayed as a heatmap forselected markers of regional cell fate identity in small molecule neural precursor cells (smNPCs) and midbrain floor plate neural precursor cells (mfNPCs). (c) Immunostaining shows that mfNPCs lack PAX3 and PAX6, which are not expressed in the midbrain floor plate. Scale bar is 100 μm. (d) Immunofluorescence confirms that mfNPCs co-express FOXA2 and EN1, which are markers of the midbrain floor plate. Scale bar is 100 μm. Experiments were conducted with mfNPC lines H1, P1-GC, P2-GC, here representative images are shown for line P1-GC, which looked similar to the other cell lines.

Supplementary Figure 2 Differentiation potential of mfNPCs for mDANs.

3

 100 ms

Supplementary Figure 2: Differentiation potential of mfNPCs for mDANs. Where indicated, analyses were performed with multiple cell lines (details see below) and a representative image looking similar to the images obtained from all used lines, is displayed. (a) Schematic illustration of the conditions used to differentiate mfNPCs into mDANs under 2D conditions. BD = BDNF, GD = GDNF. (b) mfNPCs were differentiated for four days and immunostained for CORIN and OTX2, which are markers of mDAN progenitors. (c) Immunostaining for the midbrain dopaminergic markers EN1, FOXA2, and TH after 14 days of differentiation. Scale bars are 100 µM. Stainings were performed with mfNPC lines H1, P1-GC, P2-GC, here representative images are shown for line P1-GC. (d) Patch-clamp recordings of voltage-gated ion currents, action potentials and synaptic activity of mfNPC-derived neurons after four weeks of differentiation indicate the development of essential functional properties during differentiation. (i) Representative voltage-gated sodium inward and potassium outward currents of a mfNPCderived neuron recorded in whole-cell voltage-clamp mode by stepwise depolarizations in 10 mV increments from a holding potential of -70 to 40 mV. (ii-iii) Ion currents normalized for cell sizes based on the capacitance of the cell membrane (pA/pF) after differentiation for two weeks (ii) and four weeks (iii). Data are presented as means \pm SEM. (iv) Example of neuron firing repetitive action potentials upon depolarization in the current-clamp mode. (v) Some neurons were also able to spike single action potentials spontaneously (same cell as in i after four weeks of differentiation). (vi) Spontaneous synaptic activity of neurons with miniature postsynaptic currents recorded in voltage-clamp mode. Recordings were performed with neurons derived from mfNPC lines P1-GC and P2-GC, four cells were recorded after two weeks of differentiation, 22 cells were recorded after four weeks of differentiation.

Supplementary Figure 3 Characterization of midbrain-specific organoids.

Supplementary Figure 3: Characterization of midbrain-specific organoids. Where indicated, analyses were performed with multiple cell lines (details see below) and a representative image looking similar to the images obtained from all used lines, is displayed. (a) (i) Immunohistological staining of apoptosis marker cleaved-caspase 3 (CC3), cell proliferation marker KI67, and stem cell marker SOX2 in an organoid quadrant. Scale bar is 50 μm. (ii) SOX2 and CC3 positive pixels expressed as a percentage of the total Hoechst signal (mfNPC lines H1- 3, three passages each, n=9, here representative images are shown for line H1). (b) qRT-PCR analysis for mDAN markers FOXA2, LMX1A, EN1, and TH. Data presented as mean \pm SEM, * p-value < 0.05 , ** p-value < 0.01 (mfNPC lines H1-4, n=4). (c) Representative traces of voltage-gated sodium inward and potassium outward currents (i) evoked by stepwise depolarizations and sustained firing patterns of evoked action potentials (ii) upon a current-step application (i.e. 50 pA). These recordings were obtained from the same cell displayed in Figure 1 from a dissociated 3D culture of 75 days from line H4. Black plain line in (ii) indicates 0 mV. Experiment conducted with mfNPC line H4, P2-GC and P3-GC, in total 26 cells were recorded. (d) Immunostaining for indicated ventral midbrain markers of 35 old organoid middle (i)-(iv) and edge sections (v). Scale bars are $20 \mu m$. hMOs derived from mfNPC lines H1-4, here representative images are shown for line H1 (i and iii) and line H2 (ii, iv, v).

cell lines	Derivation conditions	Gender	Age at sampling	Genotype	Comment	hiPSC ID	Figure
H1	2D	Female	81	LRRK2 WT	Reinhardt et al., 2013	2.0.0.10.1.0	$S1C$, D, S2B, C
	3D	Female	81	LRRK2 WT	Reinhardt et al., 2013	2.0.0.10.1.0	1B, C, E, F, G, S3A, B, D
H ₂	3D	Male	n.a	LRRK2 WT	Alstem (iPS15)	2.0.0.33.0.0	1B, C, E, G, S3A, B, D
H ₃	3D	Female	n.a.	LRRK2 WT	Bill Skarnes, WTSI	2.0.0.19.0.0	1B, C, E, G, H, 2A, B, C, S3A, B, D
H3-G2019S (isogenic to H3)	3D	Female	n.a.	LRRK2 G2019S	Qing et al., 2017	2.0.8.19.0.7	2B, C
H ₄	3D	Female	cord blood	LRRK2 WT	Gibco (A13777)	2.0.0.15.0.0	1D, F, 2A, B, C, S3B, C, D
H4-G2019S (isogenic to H4)	3D	Female	cord blood	LRRK2 G2019S	Qing et al., 2017	2.0.8.15.0.7	2B, C
P1-GC	2D	Female	81	LRRK2 WT	Reinhardt et al., 2013	2.1.2.11.2.0	S1B, C, D, S2B, C, D
P ₂	3D	Female	54	LRRK2 G2019S	Reinhardt et al 2013	2.1.1.46.0.0	2A, B, C
P ₂ -GC (isogenic to P2)	2D	Female	54	LRRK2 WT	Reinhardt et al., 2013	2.1.2.46.0.0	S1B, C, D, S2B, C, D
	3D	Female	54	LRRK2 WT	Reinhardt et al., 2013	2.1.2.46.0.0	1D, S3C, 2B, C
P ₃	3D	Female	81	LRRK2 G2019S	Reinhardt et al 2013	2.1.1.11.3.0	2A, B, C
P3-GC (isogenic to P3)	3D	Female	81	LRRK2 WT	Reinhardt et al 2013	2.1.2.11.3.0	1D, S3C 2B, C

Supplemetary Table 1

Supplementary Table 1 related to experimental procedures: Cell lines used in this study to generate 2D or 3D cultures. Human mfNPCs were derived from iPSCs that have been previously published ^{1,2}. Our data set includes iPSCs of healthy or diseased origin (H=healthy, P=PD patient with LRRK2-G2019S mutation). Additionally, cells were derived from isogenic controls, with either inserted LRRK2 mutation (H-G2019S) or gene corrected LRRK2 mutation (P-GC). The last column shows the contribution of each cell line to the data in the respective figure.

Supplementary Table 2

Supplementary Table 2 related to experimental procedures: Antibodies used in this study.

Supplementary Table 3

Supplementary Table 3 related to Figure 1 and 2 and experimental procedures: Features from image analysis (adapted from Bolognin et al., 2018³).

Supplementary Table 4

Supplementary Table 4 related to Figure 2: Statistical evaluation of the image analysis for PD phenotyping. Several passages of PD patient-derived and healthy hMOs, as well as their isogenic controls (mfNPC H3, H3-G2019S, H4, H4-G2019S, P3, P3-GC, P4, and P4-GC, see Supplementary Table S1) were used. A 2way ANOVA, Tukey's multiple comparisons test was performed, asterisks and adjusted p values indicate significant differences between compared groups, * p-value < 0.05 , ** p-value < 0.01 , *** p-value < 0.001 , *** p-value $<$ 0.0001, ns= not significant.

Supplementary References

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