Supplementary Information

A nanobody-based fluorescent reporter reveals human α-synuclein in the cell cytosol

Christoph Gerdes¹, Natalia Waal², Thomas Offner^{3,4}, Eugenio F. Fornasiero¹, Nora Wender¹, Hannes Verbarg¹, Ivan Manzini^{3,4}, Claudia Trenkwalder^{5,6}, Brit Mollenhauer^{6,7}, Timo Strohäker⁸, Markus Zweckstetter^{8,9}, Stefan Becker⁹, Silvio O. Rizzoli¹, Fitnat Buket Basmanav^{1,10,§}, Felipe Opazo^{1,2,§,*}

¹Department of Neuro- and Sensory Physiology, University Medical Center

Göttingen, D-37073 Göttingen, Germany

²Center for Biostructural Imaging of Neurodegeneration (BIN), University Medical

Center Göttingen, D-37073 Göttingen, Germany

³Institute of Animal Physiology, Department of Animal Physiology and Molecular

Biomedicine, Justus-Liebig University Giessen, 35390 Giessen, Germany.

⁴Institute of Neurophysiology and Cellular Biophysics, University of Göttingen,

Göttingen, Germany

⁵Department of Neurosurgery, University Medical Center Göttingen, D-37075

Göttingen, Germany

⁶Paracelsus-Elena-Klinik, Klinikstraße 16, 34128 Kassel, Germany

⁷Department of Neurology, University Medical Center Göttingen, D-37075 Göttingen, Germany

⁸German Center for Neurodegenerative Diseases (DZNE), Von-Siebold-Str. 3a,

37075 Göttingen, Germany

⁹Department for NMR-based Structural Biology, Max Planck Institute for Biophysical

Chemistry, Am Faßberg 11, 37077 Göttingen, Germany

¹⁰Campus Laboratory for Advanced Imaging, Microscopy and Spectroscopy,

University of Göttingen, D-37073 Göttingen, Germany

§ These authors contributed equally

*Correspondence should be addressed to:

Felipe Opazo (fopazo@gwdg.de)



Supplementary Figure 1 | NbSyn87 binding to h α Syn and hRpn10 and the biochemical characterization of Reporter-cells. (a, b) Increasing amounts of h α Syn (a) or hRpn10 (b) were spotted on a nitrocellulose membrane and detected using NbSyn87 directly coupled to Alexa 647. Bovine serum albumin (BSA) was spotted as control protein and its signal was used as background for normalization. Error bars represent the SEM from 3 independent experiments. (c) Induction response curve of the Reporter-cells to determine the optimal duration of doxycycline administration at a concentration of 0.5 μ g/ml. Error bars represent the SEM from 4 independent experiments. (d) Full Western blots membranes from Fig. 1f.



Supplementary Figure 2 | Determination of the optimal concentration for MG132 administration. (a) Fully induced Reporter-cells were exposed to different concentrations of MG132 for 16h. The mCherry signal (red) indicate that the cells are producing FluoReSyn, however, only starting from 500 nM of MG132, the FluoReSyn signal begins to appear (green). (b) Quantification of the FluoReSyn signal (green) in arbitrary units (a.u.) or normalized to the signal of mCherry. Error bars represent the SEM from 3 independent experiments with several hundreds of cells analyzed per experiment.



Supplementary Figure 3 | Specificity of FluoReSyn for hαSyn. (a) Quantification of the signal obtained after immunostaining synucleins with a pan-Synuclein antibody. Lines represent the mean and the error bars the SEM from n= 8. (b) Pearson's correlation analysis between the signal from FluoReSyn and immunostaining of α Syn. Signals from α Syn immunofluorescence (IF) and FluoReSyn were obtained by analyzing regions of interest (ROI) determined automatically. The ROIs were obtained by identifying all cell nuclei in the DAPI channel, followed by a morphological dilation of the nucleus ROIs, to account for the rest of the cell surfaces. (b) Raw values, in arbitrary units. (c) Double logarithmic graph of the same values. The Pearson's correlation coefficient (r) was 0.8438 for (b) with a two-tailed p < 0.0001 (****), and r = 0.8131 for (c) with a two-tailed p < 0.0001(****). (d) Amino acid sequence alignment of the putative epitope recognized by NbSyn87 for h α Syn (letters in light purple), mouse and rat aSyn or human BSyn. (accession numbers for each sequence are on the figure). Red letters show a match amino acid to the putative epitope, black letters represent mismatches. The core difference between the sequences is highlighted by a background box.



Supplementary Figure 4 | TetON induction of stably-transfected HEK293 cells expressing untagged haSyn. (left plot) Fluorescence intensity of cells immunoassayed for synuclein after being induced or not with 0.5 μ g/ml of doxycycline for 16h. (right plot) Percentage of cells displaying a positive immunofluorescense (IF) signal for synuclein. Error bars represent the SEM from 3 independent experiments (n= 3).



Supplementary Figure 5 | Co-culture of Reporter cells with TetON-haSyn cells. (a) Reporter-cells co-cultured with either a doxycycline inducible haSyn expressing cell line (TetON-haSyn) or the wildtype HEK293 cell line without endogenous haSyn expression. Induced Reporter-cells can be identified by their mCherry signal while induced TetON-haSyn cells can be identified by the haSyn immunofluorescence (IF, yellow). (b) Quantitative analysis of FluoReSyn signal intensity of cells with mCherry above 300 arbitrary units (a.u.) after 2 or 5 days of co-culturing. Scatter plots show 3 independent experiments (n= 3) \pm SEM for all condition. Significance was assessed by One-Way ANOVA and Tukey's Post-hoc test. ns, non-significant. Per replication and condition more than 550 cells were analyzed.



Supplementary Figure 6 | Reporter cell line signals plotted according to the clinical category of the patients and to their haSyn content in CSF. (a) The same data as plotted in Fig 7b, displayed according to the patient clinical description. Note that some categories have only one patient. Each patient is represented as a data point (n), the mean values are depicted with a horizontal lines and error bars represent the SEM for each category. (b) The percentage of positive Reporter cells, plotted against (left) the mass percentage of h α Syn (within all CSF proteins) or (right) the concentration of haSyn present in the CSF (µg/ml). Note that the concentrations of haSyn or the total protein in CSF were not available for some cases, and are therefore missing in the graphs. CBD, corticobasal dementia; MPFS, muscular pain-fasciculation syndrome; FTD, frontotemporal dementia; NPH, normal pressure hydrocephalus; PNP, peripheral neuropathy; PSP, progressive supranuclear palsy; RLS, restless legs syndrome; SCA, spinocerebellar ataxia; SRE, steroid responsive encephalopathy. Pearson's correlation (r) was 0.6778 when h α Syn concentrations were normalized by total protein in the CSF (two-tailed p = 0.5260, not significant), and r = -0.3424 when using the direct h α Syn concentrations (two-tailed p =0.7775, not-significant).

Supplementary Table 1. Demographic and clinical characteristics of the individuals in the CSF study

ID	Gender	Ag	Diagnosis
		е	
1	f	65	Muscular pain-fasciculation syndrome (MPFS)
2	f	73	RLS
3	m	77	FTD
4	f	60	RLS
5	m	73	CBD
6	m	74	Cerebellar ataxia
7	m	68	PSP
8	f	82	PSP
9	f	71	PSP
10	f	64	FTD
11	m	87	Vascular parkinsonism
12	m	88	Essential tremor
13	m	76	Vascular parkinsonism
14	f	81	PSP
15	m	70	PSP
16	f	81	NPH without evidence for other neurodegenerative disorders
17	f	73	CBD
18	m	76	RLS
19	m	73	NPH without evidence for other neurodegenerative disorders
20	f	69	NPH without evidence for other neurodegenerative disorders
21	f	77	NPH without evidence for other neurodegenerative disorders
22	m	75	PSP
23	f	73	CBD
24	f	64	Cerebellar ataxia
25	m	60	PSP
26	f	75	SCA
27	f	77	Steroid responsive encephalopathy associated with athyroiditis
28	f	71	Essential tremor
29	m	69	PSP
30	f	75	Dystonic tremor
31	m	84	PNP
32	f	35	Paraesthesia of skin
33	f	65	Neuroleptic-induced dyskinesia
34	m	48	RLS
35	f	59	PSP
36	m	76	PSP
37	m	65	NPH without evidence for other neurodegenerative disorders
38	m	60	Cerebellar ataxia
39	f	71	CBD
40	f	80	Benign paroxysmal positional vertigo
41	f	73	RLS
42	m	67	PSP

m, male; f, female; CBD, corticobasal dementia; FTD, frontotemporal dementia; NPH, normal pressure hydrocephalus; PNP, peripheral neuropathy; PSP, progressive supranuclear palsy; RLS, restless legs syndrome; SCA, spinocerebellar ataxia.