

# BRAIN COMMUNICATIONS

## Analysis of $\alpha$ -synuclein species enriched from cerebral cortex of humans with sporadic dementia with Lewy bodies

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Since researchers identified  $\alpha$ -synuclein as the principal component of Lewy bodies and Lewy neurites, studies have suggested that it plays a causative role in the pathogenesis of dementia with Lewy bodies and other 'synucleinopathies'. While  $\alpha$ -synuclein dyshomeostasis likely contributes to the neurodegeneration associated with the synucleinopathies, few direct biochemical analyses of  $\alpha$ -synuclein from diseased human brain tissue currently exist. In this study, we analysed sequential protein extracts from a substantial number of patients with neuropathological diagnoses of dementia with Lewy bodies and corresponding controls, detecting a shift of cytosolic and membrane-bound physiological  $\alpha$ -synuclein to highly aggregated forms. We then fractionated aqueous extracts (cytosol) from cerebral cortex using non-denaturing methods to search for soluble, disease-associated high molecular weight species potentially associated with toxicity. We applied these fractions and corresponding insoluble fractions containing Lewy-type aggregates to several reporter assays to determine their bioactivity and cytotoxicity. Ultimately, high molecular weight cytosolic fractions enhances phospholipid membrane permeability, while insoluble, Lewy-associated fractions induced morphological changes in the neurites of human stem cell-derived neurons. While the concentrations of soluble, high molecular weight  $\alpha$ -synuclein were only slightly elevated in brains of dementia with Lewy bodies patients compared to healthy, age-matched controls, these observations suggest that a small subset of soluble  $\alpha$ -synuclein aggregates in the brain may drive early pathogenic effects, while Lewy body-associated  $\alpha$ -synuclein can drive neurotoxicity.

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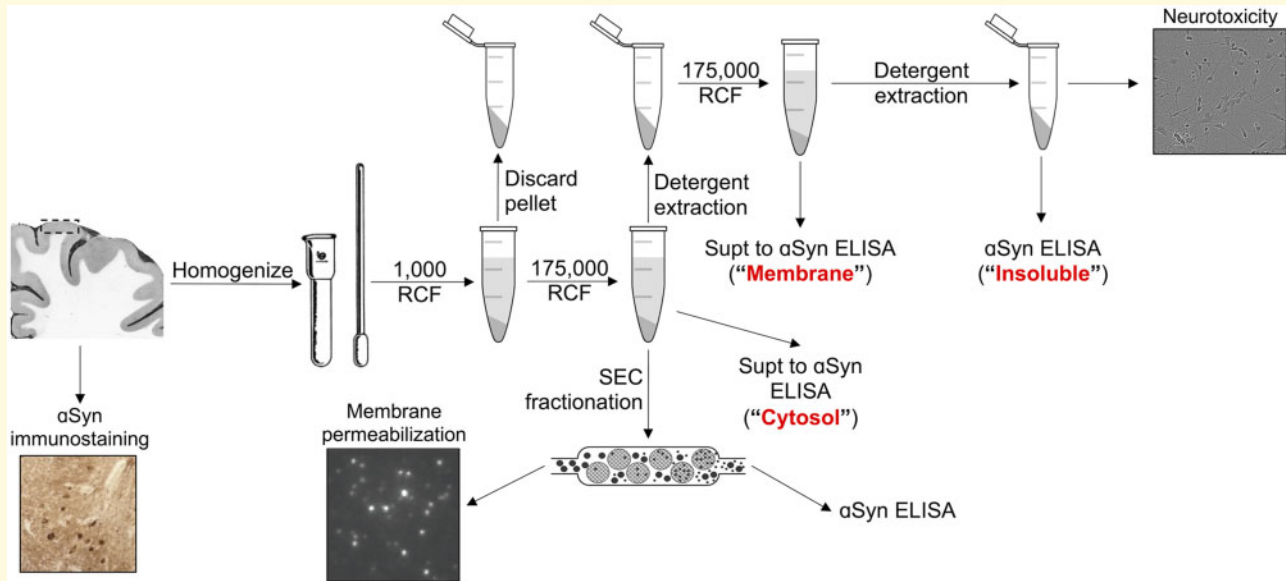
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**Abbreviations:**  $\alpha$ Syn =  $\alpha$ -synuclein;  $A\beta$  = beta-amyloid; CI = confidence interval; DLB = dementia with Lewy Bodies; ELISA = enzyme-linked immunosorbent assay; HMW = high molecular weight; HS = high salt buffer; IHC = immunohistochemistry; iPSC = induced pluripotent stem cell; LB = Lewy body; OG-RIPA = radioimmunoprecipitation assay; PBS = phosphate-buffered saline; PD = Parkinson's disease; PI = protease inhibitor; SDS = sodium dodecyl sulphate; SEC = size-exclusion chromatography; SEM = standard error of the mean.

## Graphical Abstract



## Introduction

Dementia with Lewy bodies (DLB) is an aggressive neurodegenerative disease associated with progressive cognitive dysfunction and hallucinations. Neuropathologically, the presence of intraneuronal aggregates of  $\alpha$ -synuclein ( $\alpha$ Syn), similar to those of Parkinson's disease (PD), is diagnostic (McKeith *et al.*, 1996; Spillantini *et al.*, 1997; McKeith *et al.*, 1999; McKeith *et al.*, 2005; McKeith *et al.*, 2017). In both PD and DLB,  $\alpha$ Syn forms hallmark perikaryal and neuritic aggregates called Lewy bodies and Lewy neurites, in addition to more recently described small synaptic aggregates (Kramer and Schulz-Schaeffer, 2007; Roberts *et al.*, 2015).

$\alpha$ Syn is a 140-amino-acid intracellular protein with a physiological role in regulating the trafficking of synaptic and other small vesicles (Cabin *et al.*, 2002; Burré *et al.*, 2010; Sharma *et al.*, 2011; Burré *et al.*, 2014; Logan *et al.*, 2017). To date, six different point mutations within the SNCA gene have been linked to early-onset PD and/or DLB, indicating that  $\alpha$ Syn abnormalities can be causative in the pathogenesis of these diseases (Polymeropoulos *et al.*, 1997; Krüger *et al.*, 1998; Singleton *et al.*, 2003; Chartier-Harlin *et al.*, 2004; Zarranz *et al.*, 2004; Appel-Cresswell *et al.*, 2013, Lesage *et al.*, 2013; Pasanen *et al.*, 2014).

Despite the genetic and neuropathological evidence suggesting a critical role for  $\alpha$ Syn in DLB and PD

pathogenesis, the mechanism by which  $\alpha$ Syn leads to widespread neurodegeneration remains poorly understood. The evidence in favour of the intrinsic toxicity of Lewy bodies is conflicting (Calne and Mizuno, 2004; Tanaka *et al.*, 2004; Greffard *et al.*, 2010; Osterberg *et al.*, 2015; Power *et al.*, 2017). This discordance suggests that the highly insoluble, presumably filamentous  $\alpha$ Syn in Lewy bodies may not be the principal neurotoxic species underlying clinical symptoms. A large body of work, predominantly assessing recombinant  $\alpha$ Syn, indicates that soluble oligomers show neurotoxic properties, including a propensity to alter cellular membranes (Volles *et al.*, 2001; Putcha *et al.*, 2010; Fusco *et al.*, 2016). Validation of these *in vitro* results in human tissue is incomplete, however. Regional staging of  $\alpha$ Syn aggregates in synucleinopathy patients suggests a spatio-temporal progression of pathology (Marui *et al.*, 2002; Braak *et al.*, 2004; Deramecourt *et al.*, 2006; Del Tredici and Braak, 2015), potentially mediated by similar diffusible  $\alpha$ Syn species. In fact,  $\alpha$ Syn-containing insoluble extracts from human synucleinopathy brain tissue can propagate the aggregation of endogenous  $\alpha$ Syn in cellular models (Prusiner *et al.*, 2015; Woerman *et al.*, 2015; Cavaliere *et al.*, 2017). In addition, previous *in vivo* and *in vitro* studies suggest the existence of a soluble, diffusible  $\alpha$ Syn species that might cause the spreading of Lewy pathology and neurotoxicity

(Volpicelli-Daley *et al.*, 2011; Luk *et al.*, 2012; Kovacs *et al.*, 2014; Volpicelli-Daley *et al.*, 2014; Xin *et al.*, 2015; Osterberg *et al.*, 2015, Mason *et al.*, 2016; Rey *et al.*, 2016; Sacino *et al.*, 2016; Blumenstock *et al.*, 2017; Peng *et al.*, 2018). Clinically, 'post-mortem' analyses of PD patients who had received dopaminergic neuron grafts showed  $\alpha$ Syn pathology (Kordower *et al.*, 2008; Li *et al.*, 2008), although this phenomenon was not observed in all neural graft recipients (Mendez *et al.*, 2008), nor does it explain the selective vulnerability of the respective neuronal populations observed in synucleinopathies (Walsh and Selkoe, 2016).

Some work does begin to characterize these potentially toxic  $\alpha$ Syn species. For example, application of oligomer-selective antibodies (Paleologou *et al.*, 2009; Kovacs *et al.*, 2012) or single-chain antibody fragments (Emadi *et al.*, 2007; Emadi *et al.*, 2009; Xin *et al.*, 2015) to soluble fractions of brain homogenates demonstrates the likely presence of misfolded oligomeric  $\alpha$ Syn species that are distinct from the physiological multimeric  $\alpha$ Syn that exist in complex equilibrium with monomeric  $\alpha$ Syn (Mor *et al.*, 2016; Bartels *et al.*, 2011; Wang *et al.*, 2011; Burré *et al.*, 2014; Wang *et al.*, 2014; Luth *et al.*, 2015). Similarly, the use of a proximity ligation assay on fixed brain tissue sections of synucleinopathy patients revealed abundant oligomeric  $\alpha$ Syn species previously undetectable by traditional immunostaining methods (Kramer and Schulz-Schaeffer, 2007; Roberts *et al.*, 2015). Other work has shown vesicle-associated, soluble  $\alpha$ Syn oligomers associated with DLB (Sharon *et al.*, 2003).

To enrich and further characterize oligomeric  $\alpha$ Syn from DLB patients, we obtained tissue from numerous *post mortem* brains with a neuropathological diagnosis of DLB, histochemically confirmed the presence  $\alpha$ Syn Lewy-type  $\alpha$ Syn neuronal inclusions in this tissue, and observed substantial levels of detergent-insoluble  $\alpha$ Syn, as has previously been reported (Campbell *et al.*, 2000; Klucken *et al.*, 2006). Fractionation of the brain tissue extracts by ultracentrifugation and size-exclusion chromatography (SEC) yielded buffer-soluble fractions separated by molecular radius. We quantified buffer-soluble, high molecular weight (HMW)  $\alpha$ Syn and then searched for specific neurotoxicity of these fractions compared with simultaneously prepared cytosolic fractions of non-neurodegenerative control brains. Given previous *in vitro* studies showing the ability of  $\alpha$ Syn to permeabilize membranes (Stefanovic *et al.*, 2015; Di Scala *et al.*, 2016; Ludtmann *et al.*, 2018), we then tested the ability of these enriched species to permeabilize lipid vesicles in a recently developed *in vitro* assay (Flagmeier *et al.*, 2017). In addition to testing these soluble species, we analysed detergent-insoluble material from the same brains for neurotoxicity to elucidate the potential mode of bioactivity of Lewy body (LB)-associated  $\alpha$ Syn. This study provides evidence from human samples to support previous *in vitro* and animal-based work on the role of  $\alpha$ Syn in synucleinopathy pathogenesis.

## Materials and methods

### Human tissue

Human brain tissue was provided by Brigham and Women's Hospital (Boston, MA, USA), Mayo Clinic (Jacksonville, FL, USA), Massachusetts General Hospital/Massachusetts Alzheimer's Disease Research Center (Boston, MA, USA), Newcastle Brain Tissue Resource (Newcastle upon Tyne, UK) and Queen Square Brain Bank for Neurological Disorders (London, UK). Information about the brain samples used in this study is summarized in Table 1. Consent was obtained from patients prior to death at each brain collection centre. All five brain banks approved of the proposal for the use of human tissue in this study, and the IRB at the first and last authors' institution deemed the planned use of this tissue to be appropriate and ethical.

### Sequential extraction of human tissue

Frontal cortex tissue pieces weighing between 250 and 600 mg were Dounce homogenized with 20 strokes at 2500 rpm using an overhead stirrer (Wheaton, Millville, NJ, USA) in four volumes (weight:volume) tris-buffered saline (TBS)/protease inhibitor (PI) (20 mM Tris-HCl, 500 mM NaCl, pH 7.5 with complete PI tablet; Sigma-Aldrich, St. Louis, MO, USA). Homogenates were centrifuged for 5 min at  $1000 \times g$  at  $4^{\circ}\text{C}$  to remove highly insoluble structures and debris from the tissue. The resulting supernatants were centrifuged for 30 min at  $175\,000 \times g$ . The high-speed supernatant was collected as 'cytosol', and a small aliquot was saved for enzyme-linked immunosorbent assay (ELISA)  $\alpha$ Syn quantification. The remaining supernatant was flash-frozen in liquid nitrogen and stored at  $-80^{\circ}\text{C}$  until SEC fractionation. The resulting pellet was resuspended in either 1% Triton X-100 (TX) in TBS or in modified radioimmunoprecipitation assay (OG-RIPA) buffer [0.5% Nonidet P-40 substitute, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulphate (SDS), 10 mM calcium chloride with 2% *n*-octyl- $\beta$ -D-glucoside; Abcam, Cambridge, MA, USA] (Kan *et al.*, 2013) and sonicated to homogeneity using a Dismembrator 300 Microtip Sonicator (Fisher Scientific, Hampton, NH, USA) set to 40% output. This homogenate was centrifuged for 30 min at  $175\,000 \times g$  at  $4^{\circ}\text{C}$ . The resulting OG-RIPA supernatants were collected, and the pellets were resuspended in 8 M urea with 5% SDS in TBS and boiled for 10 min at  $100^{\circ}\text{C}$ . The resulting TX supernatants were collected, and the remaining pellets were resuspended in 5% SDS in TBS, sonicated to homogeneity and centrifuged for 30 min at  $175\,000 \times g$  at  $4^{\circ}\text{C}$ . The resulting SDS pellets were resuspended in 8 M urea with 5% SDS in TBS and boiled for 10 min at  $100^{\circ}\text{C}$ .

**Table 1** Tissue samples were collected from five separate sources: Brigham and Women's Hospital (Boston, MA, USA); Mayo Clinic (Jacksonville, FL, USA); Massachusetts General Hospital/Massachusetts Alzheimer's Disease Research Center (Boston, MA, USA); Newcastle Brain Tissue Resource (Newcastle upon Tyne, UK); and Queen Square Brain Bank for Neurological Disorders (London, UK)

Case #	Source	Diagnosis	Age	Sex	Duration of disease (years)	PMI (h)
20060025	Newcastle	Probable DLB	76	M	8	13
20050022	Newcastle	PD with DLB	68	F	9	69
20070105	Newcastle	Probable DLB	71	M	7	8
20040085	Newcastle	Probable DLB	77	M	2	29
20100353	Newcastle	Cog. normal control	74	F	NA	53
20100150	Newcastle	Cog. normal control	77	M	NA	83
A01-064	Brigham and Women's	Interstitial pneumonitis	49	F	NA	25.5
A01-111	Brigham and Women's	Cog. normal control	43	F	NA	23.5
A01-213	Brigham and Women's	DLB	83	M	8	54
BN14D00074	Brigham and Women's	DLB	82	M	5	<24
P80/10	Queen Square	DLB	67	M	7	40
P11/11	Queen Square	DLB	60	M	8	24
P20/14	Queen Square	DLB	75	M	16	57
P48/07	Queen Square	Cog. normal control		F	NA	39
I687	MGH	DLB	73	M		24
I650	MGH	DLB	76	M		10
I594	MGH	DLB	81	F		8
I590	MGH	DLB	62	M		<24
I504	MGH	DLB	79	F		12
I751	MGH	DLB	87	F		6
I901	MGH	Control	54	M	NA	6
I887	MGH	Control	60	M	NA	14
CON 1	Mayo Clinic	PA/CAA	81	F	NA	
CON 2	Mayo Clinic	PA	75	F	NA	
CON 3	Mayo Clinic	PA	81	F	NA	22
CON 4	Mayo Clinic	PA/lacune (pseudo-dementia)	75	F	NA	
CON 5	Mayo Clinic	PA	81	M	NA	
CON 6	Mayo Clinic	PA	78	M	NA	13
CON 7	Mayo Clinic	SC/VaD	79	F	NA	8
CON 8	Mayo Clinic	SC	70	M	NA	20
CON 9	Mayo Clinic	Normal (pseudo-dementia)	63	M	NA	3
CON 10	Mayo Clinic	PA	87	F	NA	21
CON 11	Mayo Clinic	PA	88	F	NA	6
Syn1	Mayo Clinic	DLB	81	F		
Syn2	Mayo Clinic	DLB	67	F		
Syn3	Mayo Clinic	DLB	68	F	11	
Syn4	Mayo Clinic	DLB	66	F		16
Syn5	Mayo Clinic	DLB	79	M		
Syn6	Mayo Clinic	DLB	54	M		5
Syn7	Mayo Clinic	DLB	62	F		7

Frontal cortex grey matter was analysed from DLB patients and corresponding controls.

CAA: cerebral amyloid angiopathy; F: female; M: male; MGH: Massachusetts General Hospital; NA: not applicable; PA: pathological aging; PMI: post-mortem interval; SC: senile changes; VaD: vascular dementia.

## Comparison of membrane-associated $\alpha$ -synuclein from different extraction methods

To analyse membrane-associated  $\alpha$ Syn levels, the 1% Triton X-100 and 2% SDS extracts from our original extraction protocol were pooled and compared with the OG-RIPA extracts generated from our final quantitative extraction method (Supplementary Fig. 1). Both methods extract similar levels of  $\alpha$ Syn by the percentage of total, although extraction by OG-RIPA instead of TX and SDS yielded higher levels of membrane-associated  $\alpha$ Syn at the

expense of Lewy-associated  $\alpha$ Syn in terms of  $\alpha$ Syn amount normalized to total protein.

## Antibodies

2F12 and SOY1 human  $\alpha$ Syn-specific mouse monoclonal antibodies were used for ELISA (antibodies were developed in-house, commercially available as catalogue no. MABN1817 for 2F12, MABN1818 for SOY1, Sigma-Aldrich). Site-directed mutagenesis epitope mapping showed that 2F12 binds most strongly to a C-terminal epitope corresponding to amino acids 125–135, while SOY1



binds most strongly to a mid-region epitope corresponding to amino acids 91–110. For comparison, sections from the same blocks were stained with both 2F12 and LB509, a commonly used antibody in neuropathological diagnosis of synucleinopathies (Supplementary Fig. 2).

## Immunohistochemical staining and analysis of tissue

During frozen tissue dissection, a slice of  $\sim$ 4-mm thick was taken from the area adjacent to the section analysed for  $\alpha$ Syn content and stained as described (Sanderson, 2019). In short, the slice was submerged in 4% paraformaldehyde in phosphate-buffered saline (PBS) for between 72 and 96 h for fixation. Tissue was dehydrated and embedded in paraffin. Six-micrometre sections were cut and affixed to glass slides and dried overnight at 37°C. To remove the paraffin, slides were incubated for 3 min twice each in 100% ethanol, 95% ethanol and ultrapure water. Slides were then heated for 15 min by microwave in 10 mM sodium citrate and allowed to cool to ambient temperature before a 1-h ambient temperature incubation in 2% vol/vol fat-free milk in PBS-T. Staining was performed using 2F12 mouse monoclonal antibody to  $\alpha$ Syn at a concentration of 65 ng/ml for 1 h at ambient temperature. Following incubation in primary antibody, slides were rinsed three times in PBS-T and then incubated for 2 h at ambient temperature in biotinylated anti-mouse IgG antibody (Southern Biotech, Birmingham, AL, USA) prepared at 2.5  $\mu$ g/ml in 2% vol/vol fat-free milk in PBS-T. Following incubation, slides were rinsed twice in PBS-T and once in PBS, incubated in ABC avidin–biotin peroxidase complex (Vector Laboratories, Burlingame, CA, USA) for 1 h at ambient temperature, and then rinsed twice in PBS-T and once in PBS. Immunostaining was visualized using 3,3-diaminobenzidine horseradish peroxidase substrate (Vector Laboratories, Burlingame, CA, USA). Slides were rinsed three times in ultrapure water, counterstained with haematoxylin, differentiated in 0.3% acid alcohol and dehydrated by submerging in two baths of each of the following: 95% ethanol, 100% ethanol and 100% xylene. Glass coverslips were mounted in Permount (Thermo Fisher Scientific, Waltham, MA, USA).

## Denaturing $\alpha$ -synuclein sandwich enzyme-linked immunosorbent assay

All reagents were purchased from Meso Scale Discovery (Rockville, MD, USA). Prior to ELISA  $\alpha$ Syn quantification, aliquots of all previously non-denatured samples were boiled for 10 min at 100°C in 2% SDS. Single-spot, standard-bind plates were coated with 200 ng of 2F12 prepared in PBS and incubated overnight at 4°C. Before sample loading, excess capture antibody was rinsed out and

the wells were treated with 5% MSD Blocker A in TBS-T for 1 h at ambient temperature with shaking at 500 rpm. Blocking buffer was removed, and the plate was rinsed three times with TBS-T. Prior to loading, samples were diluted in 1% Blocker A with 0.5% Nonidet P-40 substitute in TBS-T to a final SDS concentration of no greater than 0.4%. Samples were loaded in duplicate, including amino acid sequenced recombinant  $\alpha$ Syn standards diluted in the appropriate detergent mixture. Plates were shaken at 500 rpm for 1 h at ambient temperature and then incubated overnight at 4°C. Sample liquid was removed, the plate was rinsed three times with TBS-T and 200 ng of sulfo-labelled SOY1 diluted in 1% Blocker A was added per well. The plate was light-shielded and shaken for 1 h at ambient temperature at 500  $\times$  g. The detection antibody solution was then tapped out, and the plate was washed three times in TBS-T. About 150  $\mu$ l of 2 $\times$  Read Buffer S (diluted in ultrapure water) was added per well, and the plate was immediately read using a SECTOR Imager 2400.

## Size-exclusion chromatography fractionation of cytosolic proteins

Between 1 and 2.5 mg of total protein (measured via Pierce BCA kit; Thermo Scientific, Waltham, MA, USA) in 400–600  $\mu$ l was injected into a 2-ml sample loop. The sample was passed over a Superose 6 Increase 10/300GL size-exclusion column (GE Healthcare, Pittsburgh, PA, USA) mounted on an ÄKTA chromatography system (GE Healthcare). The column was equilibrated with 50 mM ammonium acetate (pH 7.40). Using a flow rate of 1.5 ml/min, 1 ml of fractions were collected. Twenty microlitres from each fraction was mixed with 5  $\mu$ l of 10% (v/v) SDS (final SDS concentration: 2%) and boiled for 10 min at 100°C for ELISA quantification. The remaining portion of the fractions were flash-frozen in liquid nitrogen and either stored at  $-80^{\circ}\text{C}$  or lyophilized for future analysis. For the estimation of MW, a gel filtration molecular marker kit with standards ranging from 29 to 700 kDa in size (catalogue no. MWGF1000; Sigma-Aldrich) was used to calibrate the column prior to the analysis of brain material. These standards were used to generate a linear model relating elution volume to molecular radius.

## Permeabilization assays

The membrane permeabilization assay was performed as previously described (Flagmeier *et al.*, 2017). Briefly, 200 mM 1-palmitoyl-2-oleoyl-sn-glycero-3-phosphocholine (POPC) (16:0–18:1 PC: 18:1–12:0 Biotin PC—100:1) vesicles were prepared by extrusion and five freeze-thaw cycles. The lipid mixture was then hydrated in 100  $\mu$ M Cal-520 dye dissolved in HEPES buffer (50 mM, pH 6.5). Vesicles were tethered to PLL-g-PEG coated cover slides via biotin–neutravidin linkage. The HEPES buffer was

replaced with 50  $\mu\text{l}$   $\text{Ca}^{2+}$ -containing buffer solution and blank images ( $F_{\text{blank}}$ ) were recorded. Each field was selected using a homemade stage-control programme to avoid user bias. Once these fields were selected, 50  $\mu\text{l}$  of the soluble HMW fractions were added to the coverslips and incubated for 20 min before the same fields were re-imaged ( $F_{\text{sample}}$ ). Ionomycin was added as a positive control, and the same area containing vesicles are reimaged ( $F_{\text{ionomycin}}$ ). The relative influx of  $\text{Ca}^{2+}$  into an individual vesicle due to aggregates of  $\alpha\text{Syn}$  was then determined using the following equation

$$\text{Ca}^{2+} \text{ influx} = \frac{F_{\text{sample}} - F_{\text{blank}}}{F_{\text{ionomycin}} - F_{\text{blank}}}$$

For antibody experiments, the samples were incubated with 300 nM of antibody for 20 min before addition to the coverslips. Imaging were performed using a home-built total internal reflection fluorescence microscope equipped with 60 $\times$ , 1.49 NA oil immersion objective lens. A 488-nm laser was used to excite the Cal-520 dye, and the emitted fluorescence was collected by electron multiplying charge-coupled device (CCD) camera.

## Cell lines

All materials purchased from Thermo Fisher Scientific unless otherwise noted. Human embryonic kidney cells (HEK-293; ATCC number CRL-1573) were cultured at 37°C and 5%  $\text{CO}_2$  in Dulbecco's modified Eagle's medium supplemented with 10% foetal bovine serum, 100 U/ml penicillin, 100  $\mu\text{g}/\text{ml}$  streptomycin, 2 mM L-glutamine, 1 mM sodium pyruvate and 1 $\times$  minimum essential medium (MEM) non-essential amino acids solution. Monoclonal stable cell lines were generated by transfecting with pcDNA4-A53T-hu $\alpha\text{Syn}$ -yellow fluorescent protein using Lipofectamine 2000 according to the manufacturer's instructions. The transfected cells were cultured under zeocin selection, and individual clones were selected. N27 rat dopaminergic neural cell line was purchased from EMD Millipore (catalogue no. SCC048, Billerica, MA, USA) and cultured at 37°C with 5%  $\text{CO}_2$  in the Roswell Park Memorial Institute (RPMI) 1640 medium supplemented with 10% foetal bovine serum, 100 U/ml penicillin, 100  $\mu\text{g}/\text{ml}$  streptomycin and 2 mM L-glutamine. Monoclonal stable cell lines were generated by transfecting with pcDNA4-hu  $\alpha\text{Syn}$  using Lipofectamine 2000 according to the manufacturer's instructions. The transfected cells were cultured under zeocin selection, and individual clones were selected. The clone with the highest  $\alpha\text{Syn}$  expression was selected for our cell viability analysis.

## Cell viability analysis

Cells were seeded at a density of 6500 cells per well on a black poly-D-lysine (PDL) coated 96-well plate with clear bottom (catalogue no. 354363; Corning, Corning,

NY, USA). After 24 h of culturing, cells were treated with brain extract material for 72–96 h prior to viability measurement. Twenty-four hours prior to measurement, 1  $\mu\text{M}$  staurosporine was added to three wells as a positive control for cytotoxicity. Two hours prior to measurement, culture medium was replaced with 120  $\mu\text{l}$  of culture medium with 16% (v/v) CellTiter Blue reagent (Promega, Madison, WI, USA). Cells were cultured for 2 h, and then fluorescence was quantified using a BioTek H1 plate reader (BioTek, Winooski, VT, USA) with an excitation wavelength of 560 nm and an emission wavelength of 590 nm. The average fluorescence value of three non-cell containing wells was considered background and subtracted from all other values during analysis.

## A53T- $\alpha$ -synuclein-yellow fluorescent protein inclusion formation assay

This protocol has been adapted from [Woerman \*et al.\* \(2015\)](#). All reagents were purchased from Thermo Fisher Scientific unless otherwise noted. HEK293 cells stably expressing pcDNA4-A53T-hu $\alpha\text{Syn}$ -yellow fluorescent protein were seeded at a density of 5000 cells per well on a black PDL-coated 96-well, clear-bottom plate (catalogue no. 354363; Corning). Protein samples (either recombinant monomer or pre-formed fibrils) were prepared in Opti-MEM with 3% (volume:volume) Lipofectamine 2000 and incubated at room temperature for 2 h. Prior to cell treatment, protein-Opti-MEM-Lipofectamine mixtures were mixed 1:1 with HEK culture medium. Culture medium on cells was removed and replaced by 100  $\mu\text{l}$  per well of these proteofection mixtures. Cells were cultured for 4 h with the proteofection mixture, after which the mixture was removed and replaced with 250  $\mu\text{l}$  of culture medium. The cells were cultured for 96 h, and fluorescent images were captured every 2 h using the InCuCyte Zoom live cell imaging platform (Essen Biosciences, Ann Arbor, MI, USA). Inclusion formation assessment was performed using the InCuCyte analysis software.

## Enrichment of Lewy body-associated brain material for bioactivity analysis

LB-rich sequential extractions were conducted according to the protocol published by [Peng \*et al.\* \(2018\)](#). Brain pieces of 200–400 mg were Dounce homogenized in four volumes (weight:volume) of high salt buffer with PIs (HS/PI) (50 mM Tris-HCl, 750 mM NaCl, 5 mM ethylenediaminetetraacetic acid (EDTA), pH 7.4 with complete PI tablet; Sigma-Aldrich). Homogenates were centrifuged at 100 000  $\times g$  for 30 min at 4°C. Supernatants were collected, and bicinchoninic acid assay for total protein concentration was performed to assess total protein concentration. Pellets were then re-extracted with HS/PI

buffer, followed by sequential extractions with 1% Triton X-100-containing HS buffer, 1% Triton X-100-containing HS buffer with 30% (weight:volume) sucrose and finally 1% sodium lauroyl sarcosinate (sarkosyl) (weight:volume)-containing HS buffer. The HS + 1% sarkosyl homogenate was incubated with nutation overnight at 4°C. Following this incubation, the homogenate was centrifuged at  $100\,000 \times g$  for 30 min and the supernatant was collected. The pellet was resuspended for washing in a volume of PBS equal to the homogenate volume initially centrifuged and then centrifuged at  $100\,000 \times g$  for 30 min. The pellet was then resuspended and homogenized in 0.2 volumes of the initial homogenate volume of PBS using a probe sonicator (catalogue no. 431MPX, QSonica, Newtown, CT, USA) for 30 s (5 s ON, 5 s OFF) at amplitude 20. After sonication, these samples were centrifuged at  $100\,000 \times g$  for 20 min at 4°C. The supernatant was collected and filtered through a syringe filter (catalogue no. SLGV004SL; Millipore Sigma) in a cell culture hood. The sterile supernatant was then aliquoted in such that each aliquot contained 150 ng of total protein based on the bicinchoninic acid protein quantification. The aliquots were snap-frozen in liquid nitrogen in 0.5-ml Eppendorf Protein LoBind Tubes (catalogue no. 022431081, Eppendorf) and stored at  $-80^{\circ}\text{C}$ .

## Harvest, culture and treatment of hippocampal primary rat neurons

The hippocampi of E18 SD rats were dissected in Hank's Balanced Salt Solution buffered with HEPES and dissociated with 0.125% trypsin (Invitrogen, Carlsbad, CA, USA) for 15 min at 37°C followed by trituration. Dissociated cells were plated in 96-well plates pre-coated with poly-D-lysine (100  $\mu\text{g}/\text{ml}$ ) at a density of 15 000 cells/ $\text{cm}^2$ . After culturing for 4 days in the Neurobasal medium with B-27 supplement and Glutamax (Invitrogen), FUDR (Sigma-Aldrich) was added to block astrocyte growth to a final concentration of 50  $\mu\text{g}/\text{ml}$ . Half of the growth medium was exchanged every 4 days. Primary rat hippocampal neurons were cultured for 14 days *in vitro* (DIV14) as described previously (Jin *et al.*, 2011). At DIV14, pooled and lyophilized HMW SEC fractions (2–8) were resuspended in culture medium and applied to the cells in triplicate. Neurite length was monitored over 72 h using the IncuCyte live cell imaging platform (Essen Biosciences).

## Culture and treatment of human induced pluripotent stem cell-derived neurons

The induced pluripotent stem cell (iPSC) line used in this study (YZ1) was originally generated from the IMR-90 cell line (ATCC) and characterized as described

previously (Zeng *et al.*, 2010). Due to a karyotype abnormality in a small subset of cells, monoclonal isolates were obtained and confirmed to be karyotypically normal and pathogen-free prior to this study (Srikanth *et al.*, 2018). The iPSC line used was confirmed to be of the correct identity using short tandem repeat profiling (Genetica Cell Line Testing). iPSCs were maintained in media containing 400 ml of Dulbecco's modified Eagle's medium/F12 (Invitrogen), 100 ml of KnockOut Serum Replacement, 5 ml of penicillin/streptomycin/glutamine, 5 ml of MEM non-essential amino acids and 500  $\mu\text{l}$  of 2-mercaptoethanol (all from Invitrogen) with fresh addition of 10  $\mu\text{g}/\text{ml}$  fibroblast growth factor-basic (bFGF) (Millipore). The differentiation was achieved through neurogenin expression under a doxycycline promoter as previously reported with minor modifications (Zhang *et al.*, 2013). Induced neurons, which are described as a homogenous population of neurons expressing VGLUT2, a transporter for the excitatory neurotransmitter glutamate, were plated at DIV4 on 96-well plates (Greiner Bio-One, Monroe, NC, USA) pre-coated with Matrigel (Corning) at a density of 5000 cells per well and maintained in media consisting of 1% Glutamax (volume:volume), 0.3% dextrose (weight:volume) and 0.5% MEM non-essential amino acids (volume:volume) in the Neurobasal medium. Two percent B27 supplement (v/v), brain-derived neurotrophic factor (BDNF), ciliary neurotrophic factor (CNTF), glial cell line-derived neurotrophic factor (GDNF) (10 ng/ml each), puromycin (1  $\mu\text{g}/\text{ml}$ ) and doxycycline (2  $\mu\text{g}/\text{ml}$ ) were added just prior to feeding. Differentiated cells were treated between DIV21 and DIV28 with insoluble extracts prepared as described above containing 50 ng of total protein per well and brought to a final volume of 200  $\mu\text{l}$ . Cells were treated in triplicate wells with randomized locations. During the course of the treatment, the cells underwent a single medium replacement of 50% of the volume in each well.

## Neurite morphology analysis

The IncuCyte automated live cell imaging platform captured phase contrast images every 2 h over a total of 88–96 h. Neurite and cell body masks were applied to the images (Supplementary Fig. 3). Average neurite length and cell body cluster area were measured using the IncuCyte image capture and analysis software in Neurotrack mode. To compare neurite degeneration between wells, average neurite lengths at 88 h were divided by the normalized average neurite lengths in the same field at 0 h. This was compared to the same ratio in untreated wells.

## Statistical analysis

Prism (GraphPad, San Diego, CA, USA) was used for all statistical analyses and graph generation. Details of each analysis are provided in the relevant figure legends.

Means  $\pm$  standard error of the means (SEMs) are shown in cohort-wide analyses, while means  $\pm$  standard deviations are shown in single-patient analyses.

## Data availability

The data that support the findings of this study are available from the corresponding author upon request.

## Results

### $\alpha$ -Synuclein is differentially distributed in subcellular compartments in dementia with Lewy Bodies cortical tissue

Using sequential extraction with solvents of increasing denaturing capability, we quantitatively analysed the relative solubility of  $\alpha$ Syn in frontal cortical extracts from 22 DLB patients and 18 non-synucleinopathy control subjects (Table 1). Analysis of demographic data showed no difference in ages or post-mortem intervals between the two populations ( $P=0.929$  and  $P=0.980$ , respectively, unpaired  $T$ -tests). No post-mortem interval data were provided for 6 of the 22 DLB brains and 4 of the 18 control brains, and no ages were given for one of the 18 control brains. In addition, chi-square analysis showed no significant deviation from the expected gender ratios ( $P=0.262$ ) (Table 2). To minimize the significant variation we observed in both the levels of 'insoluble'  $\alpha$ Syn (insoluble in OG-RIPA but soluble in 5% SDS/8 M urea-soluble) and the degree of Lewy cytopathology by  $\alpha$ Syn immunohistochemistry (IHC) (Fig. 1), we homogenized between three and six different tissue pieces dissected from frozen frontal cortex of each DLB brain that showed Lewy pathology by  $\alpha$ Syn IHC in sections prepared from a piece of the adjacent frozen cortex. When thawed, each of these pieces weighed  $\sim 50$  mg. Small aliquots of each extraction fraction were denatured by boiling in 2% SDS to measure total  $\alpha$ Syn amount by sandwich ELISA. To account for possible shifts in cell populations in diseased tissue, such as the presumed loss of  $\alpha$ Syn-expressing neurons and the local invasion of non-expressing glia, we expressed the data in  $\alpha$ Syn levels as a percentage of the total  $\alpha$ Syn

recovered from all three fractions (Fig. 2). We also expressed the absolute amount of  $\alpha$ Syn in each sequential extract normalized to total protein (per a bicinchoninic acid total protein assay) in Supplementary Fig. 4.

While this extraction was designed primarily to identify small brain pieces with high pathology burden based on the established finding that detergent-insoluble  $\alpha$ Syn is associated with high histopathological LB density (Campbell et al., 2000; Klucken et al., 2006), we also included a membrane-associated fraction to examine the subcellular localization of  $\alpha$ Syn in our relatively large, multi-centre collection of human samples. Expressing the three extract fractions (cytosol, membrane extract and urea-solubilized extract) as a percentage of total recovered  $\alpha$ Syn showed differences in the solubility distributions of  $\alpha$ Syn extracts from control compared with synucleinopathy tissue. DLB cytosols contained a higher proportion of extracted  $\alpha$ Syn than those of controls, whereas membrane-associated fractions showed the opposite: DLB samples had a lower proportion of total extracted  $\alpha$ Syn (Fig. 2A). This observation suggests a partial redistribution of membrane-bound  $\alpha$ Syn to the cytosolic compartment in the affected DLB cortex. Although correlation analyses of related percentages, by definition, show strong associations, we interrogated whether the relative strengths of these associations on a sample-to-sample basis suggested a predictable redistribution of  $\alpha$ Syn from one compartment to another. Stratification of DLB brain extracts into low  $\alpha$ Syn burden (insoluble  $\alpha$ Syn  $< 1\%$ ) and high  $\alpha$ Syn burden (insoluble  $\alpha$ Syn  $> 1\%$ ), and analysis of membrane:cytosol  $\alpha$ Syn ratio showed a relative shift to the cytosolic compartment in high-insoluble  $\alpha$ Syn brains, suggesting that insoluble  $\alpha$ Syn may come from the membrane compartment (Fig. 2B). This result supports recent research showing an abundance of vesicle-bound  $\alpha$ Syn in Lewy bodies (Shahmoradian et al., 2019) and reinforces the hypothesis that vesicle binding increases  $\alpha$ Syn aggregation (Galvagnion et al., 2015). As expected, in control subject brain samples, there was nearly perfect linear correlation between the cytosolic and membrane-associated  $\alpha$ Syn, as physiological  $\alpha$ Syn likely exists in a rapid equilibrium between its soluble and membrane-bound forms [Spearman  $r = -1.000$ , 95% confidence interval (CI):  $-1.000$  to  $-1.000$ ,  $P < 0.0001$ ] (Fig. 2C). Because the  $\alpha$ Syn extracted from DLB patient brains was spread across all three fractions, with variable elevations in the final urea-solubilized extract (Fig. 2A, right), this correlation was not as strong in DLB (Spearman  $r = -0.609$ , 95% CI:  $-0.721$  to  $-0.462$ ,  $P < 0.0001$ ) (Fig. 2C). Looking exclusively at DLB patient brains, there were strong negative associations between the insoluble fractions and the cytosolic and membrane fractions (insoluble versus cytosol: Spearman  $r = -0.298$ , 95% CI:  $-0.469$  to  $-0.105$ ,  $P = 0.0023$ ; insoluble versus membrane: Spearman  $r = -0.315$ , 95% CI:  $-0.484$  to  $-0.124$ ,  $P = 0.0012$ ) (Fig. 2D and E). In analysis of the brains obtained from the Mayo Clinic brain bank, which

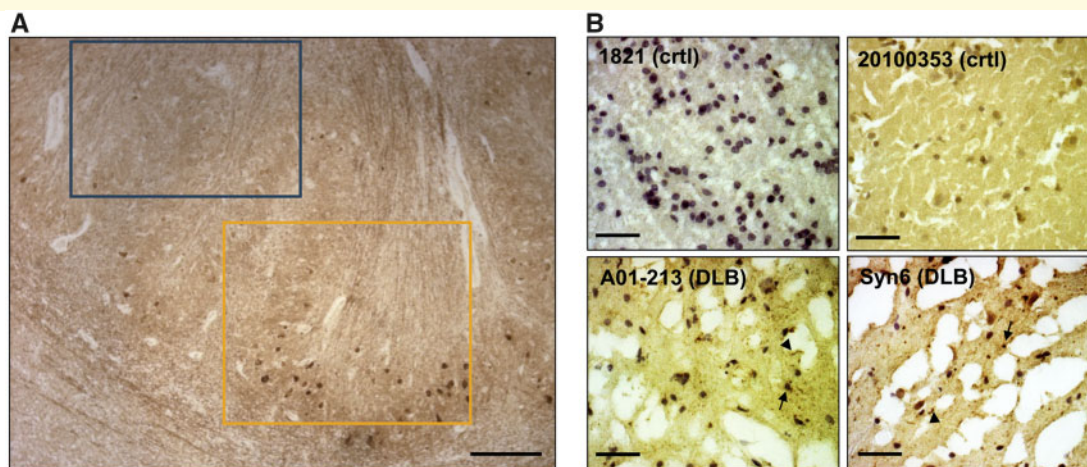
**Table 2 Unpaired  $T$ -tests showed no significant difference between the mean ages or PMIs of the control and DLB patient samples**

	Control (N = 18)	DLB (N = 22)	P-value
Age	72.8 $\pm$ 13.5	72.5 $\pm$ 8.6	0.93
PMI	24.1 $\pm$ 20.9	23.9 $\pm$ 20.5	0.98
Gender (F:M)	11:7	9:13	0.26

A chi-square test also showed no significant difference in the F:M reported gender ratios between the groups.

F: female; M: male; PMI: post-mortem interval.





**Figure 1 IHC confirmation of synucleinopathy in DLB frontal cortical samples.** (A) IHC of the formalin-fixed striatum of MGH DLB brain #1594 with  $\alpha$ Syn monoclonal antibody 2F12 demonstrates the striking local heterogeneity of cytopathology. Yellow box: area of high density of Lewy cytopathology; blue box: immediately adjacent area devoid of IHC-detectable  $\alpha$ Syn aggregates. Scale bars: 300  $\mu$ m. (B) Small frozen pieces of tissue were excised from locations adjacent (within  $\sim$ 5 mm) to the frozen pieces we used for our biochemical extractions. The resultant cryostat sections were stained with  $\alpha$ Syn monoclonal antibody 2F12 and nuclei counterstained with haematoxylin. Representative images from two control and two DLB brains show ice crystals resulting from variable freezing of the unfixed tissue. Arrows exemplify LBs; arrowheads exemplify Lewy neurites. Scale bars: 80  $\mu$ m.

had exceptionally high levels of insoluble  $\alpha$ Syn (Fig. 2F), both of these negative correlations were even stronger. This suggests that in patients with a high burden of LB pathology,  $\alpha$ Syn from both the cytosolic and membrane-associated physiological pools contributes to the formation of the insoluble aggregates (insoluble versus cytosol: Spearman  $r = -0.900$ , 95% CI:  $-0.954$  to  $-0.89$ ,  $P < 0.0001$ ; insoluble versus membrane: Spearman  $r = -0.415$ , 95% CI:  $-0.689$  to  $-0.386$ ,  $P = 0.0028$ ) (Fig. 2G and H).

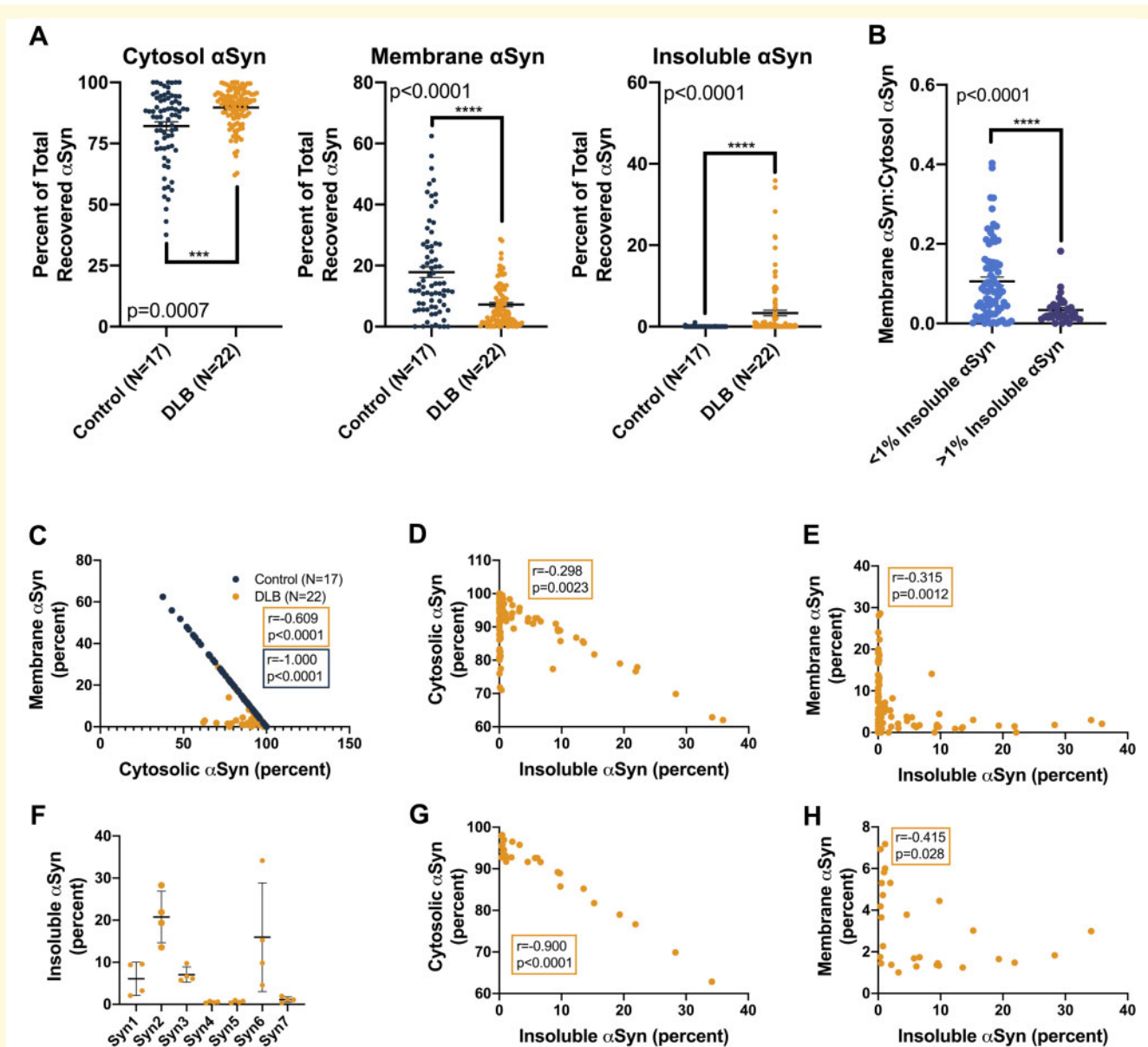
Looking at the absolute  $\alpha$ Syn concentrations in each fraction, we observed significantly lower levels of membrane-associated  $\alpha$ Syn in DLB than control cortex when normalized to total protein ( $P < 0.0001$ ) (Supplementary Fig. 4A) and robust regression and outlier removal outlier removal did not alter this finding.  $\alpha$ Syn content in the urea-soluble extracts confirmed that DLB cortex had greater levels of highly insoluble  $\alpha$ Syn than did control cortex ( $P < 0.0001$ ) (Supplementary Fig. 4A). As expected, only negligible amounts of  $\alpha$ Syn were detected in the urea-solubilized fraction of control brains, indicating that the  $\alpha$ Syn detected in this fraction derives from the hallmark aggregated  $\alpha$ Syn lesions of DLB.

Analyses of the of log-normalized absolute  $\alpha$ Syn levels (Supplementary Fig. 4B–D) in the three different extracts revealed associations of  $\alpha$ Syn levels between fractions. Cytosolic absolute  $\alpha$ Syn levels from frontal cortex (whether DLB or control) correlated strongly with membrane-associated absolute  $\alpha$ Syn levels (control: Spearman  $r = 0.656$ ,  $P < 0.0001$ , 95% CI:  $0.494$ – $0.770$ , DLB: Spearman  $r = 0.761$ ,

$P < 0.0001$ , 95% CI:  $0.662$ – $0.834$ ) (Supplementary Fig. 3B). Detergent-insoluble/urea-solubilized (Lewy-associated)  $\alpha$ Syn levels showed no significant relationships with membrane or cytosolic  $\alpha$ Syn levels in DLB extracts (insoluble versus cytosol: Spearman  $r = -0.078$ , 95% CI:  $-0.118$  to  $-0.267$ ,  $P = 0.436$ ; insoluble versus membrane: Spearman  $r = -0.026$ , 95% CI:  $-0.174$  to  $-0.224$ ,  $P = 0.793$ ) (Supplementary Fig. 4C and D), except for a subset of DLB samples with the most severe LB/Lewy neurite pathology burden by high levels of insoluble  $\alpha$ Syn in the corresponding urea extracts. Specifically, DLB frontal cortices from the Mayo Clinic brain bank that had very high levels of insoluble, Lewy-associated  $\alpha$ Syn (Supplementary Fig. 4E–G), showed strong negative correlations between those levels and both cytosolic and membrane-associated  $\alpha$ Syn (insoluble versus cytosol: Spearman  $r = -0.719$ , 95% CI:  $-0.864$  to  $-0.464$ ,  $P < 0.0001$ ; insoluble versus membrane: Spearman  $r = -0.563$ , 95% CI:  $-0.778$  to  $-0.230$ ,  $P = 0.0018$ ).

Analysis of this protein-normalized  $\alpha$ Syn content in each fraction revealed a significant decrease in the mean level of total extracted  $\alpha$ Syn in DLB frontal homogenates versus controls by  $\sim$ 25% ( $P = 0.0016$ , Mann–Whitney comparison of ranks) (Supplementary Fig. 4A). Analysis of cytosolic fractions yielded a similar result: on average, there was significantly less  $\alpha$ Syn in DLB extracts than in control ( $P = 0.010$ ) (Supplementary Fig. 4A).

Notably, analysis comparing the subcellular localization of  $\alpha$ Syn in extracts of two brains of patients with familial DLB (one A53T  $\alpha$ Syn mutant and one SNCA duplication carrier) did not show striking



**Figure 2**  $\alpha$ Syn solubility distribution in DLB versus control brains based on the percentage of total analysis. **(A)** For each tissue piece (represented by a single point), the ELISA-measured  $\alpha$ Syn concentration was multiplied by the original volume of the extract, yielding the total  $\alpha$ Syn in each fraction. This was then expressed as a percentage of the total  $\alpha$ Syn recovered in all three fractions. Graphs: percentage of total  $\alpha$ Syn in the cytosolic, membrane (1% Triton-soluble) and insoluble (SDS/8 M urea-soluble) extracts. Between three and eight tissue pieces were processed and analysed per brain, and each piece was measured in triplicate. Bars: means with SEMs. Control and DLB extracts were compared using the Mann–Whitney comparison of ranks. **(B)** DLB brain extracts were stratified based on insoluble  $\alpha$ Syn quantities (blue: <1% of total extracted  $\alpha$ Syn; purple: >1% of total extracted  $\alpha$ Syn), and the ratio of membranous to cytosolic  $\alpha$ Syn was calculated for each group. Proportionally, this analysis revealed higher membranous  $\alpha$ Syn in low-pathology DLB extracts ( $P < 0.0001$ , Mann–Whitney comparison of ranks). **(C)** Correlation analyses of the percentages of recovered  $\alpha$ Syn in the cytosolic and membrane cortical fractions, both from control subject and DLB patient brains, showed strong negative associations, an unsurprising finding given that these fractions contain most of the total extracted  $\alpha$ Syn (control: Spearman  $r = -1.000$ , 95% CI:  $-1.000$  to  $-1.000$ , 2-tailed  $P < 0.0001$ ; DLB: Spearman  $r = -0.609$ , 95% CI:  $-0.721$  to  $-0.462$ ,  $P < 0.0001$ ). **(D and E)** Strong negative associations also exist between the percentage of total extracted  $\alpha$ Syn in the insoluble fraction and the percentage of total extracted  $\alpha$ Syn in both the cytosolic and membrane-associated fractions across all cortical extracts from DLB patient brains (insoluble versus cytosol: Spearman  $r = -0.298$ , 95% CI:  $-0.469$  to  $-0.105$ ,  $P = 0.0023$ ; insoluble versus membrane: Spearman  $r = -0.315$ , 95% CI:  $-0.484$  to  $-0.124$ ,  $P = 0.0012$ ). **(F)** In terms of insoluble  $\alpha$ Syn as a percentage of total extracted  $\alpha$ Syn, DLB patient brains from the Mayo Clinic brain bank showed the highest pathological burden. Means with standard deviations are shown. **(G and H)** Correlation analyses comparing the percentage of total extracted  $\alpha$ Syn in insoluble fractions to cytosolic and membrane-associated fractions from these high-burden Mayo Clinic brains showed even stronger negative associations than the analyses that included DLB patient brains from all sources (insoluble versus cytosol: Spearman  $r = -0.900$ , 95% CI:  $-0.954$  to  $-0.89$ ,  $P < 0.0001$ ; insoluble versus membrane: Spearman  $r = -0.415$ , 95% CI:  $-0.689$  to  $-0.386$ ,  $P = 0.0028$ ).

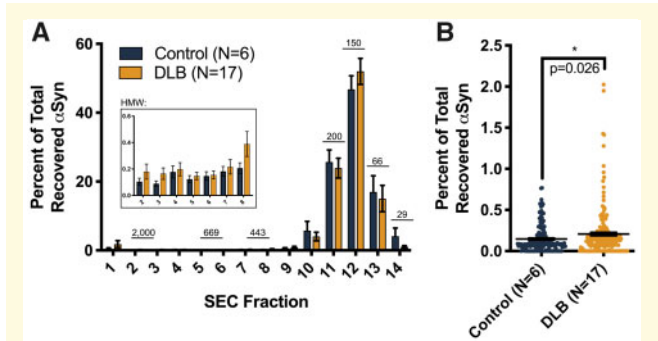
differences compared with the pooled data from all control extracts (Supplementary Fig. 5A). Statistical analyses were not performed due to the disparate sample sizes.

### Soluble, high molecular weight $\alpha$ -synuclein species are more abundant in dementia with Lewy Bodies than control cortical extracts

The cytosolic fractions of brain pieces with high levels of  $\alpha$ Syn in the urea extracts were fractionated using non-denaturing SEC on a column with a 3-MDa exclusion size to probe for relative abundance of a range of HMW forms of soluble  $\alpha$ Syn. After SEC fractionation, small aliquots of each fraction were boiled in 2% SDS for 10 min to denature the samples and maximally expose epitopes prior to  $\alpha$ Syn ELISA quantification. Across all samples, an average of 91.2% of soluble  $\alpha$ Syn eluted in SEC fractions 11–13, which correspond to the expected elution volume of physiological  $\alpha$ Syn (MW of  $\sim$ 80 kDa based on the elution of recombinant protein standards; Supplementary Fig. 6). Comparison of  $\alpha$ Syn levels in individual SEC fractions of control and DLB cytosols revealed no statistically significant differences (Fig. 3A). However, pooling the individual levels across the HMW SEC fractions (i.e. fractions 2–8, corresponding to  $\sim$ 2 MDa down to  $\sim$ 440 kDa, Supplementary Fig. 6) revealed a significant increase in relative levels of soluble, HMW  $\alpha$ Syn in cytosols of DLB versus control cortex ( $P=0.0256$ , Mann–Whitney comparison of ranks) (Fig. 3B). On average, the  $\alpha$ Syn content of these HMW SEC fractions comprised only 1.0% (in control cortex) and 1.4% (in DLB cortex) of the total eluted cytosolic  $\alpha$ Syn, indicating that in the diseased brain, HMW-soluble  $\alpha$ Syn species is relatively elevated but still not abundant, similar to previous observations of tau protein species present in Alzheimer's disease brain (Takeda *et al.*, 2015).

### Soluble $\alpha$ -synuclein species increase the permeability of lipid membranes *in vitro*

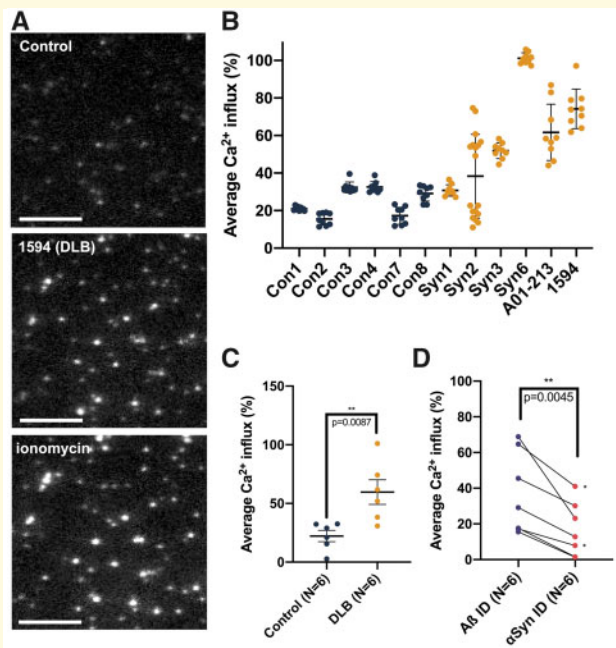
We then tested whether the soluble, HMW SEC fractions of brain cytosols were capable of phospholipid membrane permeabilization. We used a previously described membrane permeabilization assay that utilizes nanosized lipid vesicles tethered to a coated glass surface via biotin–neutravidin linkage and filled with a  $\text{Ca}^{2+}$ -sensitive dye (Flagmeier *et al.*, 2017). Influx of  $\text{Ca}^{2+}$  into these vesicles from the surrounding buffer was measured using total internal reflection fluorescence microscopy (Fig. 4A). We applied the pooled HMW SEC fractions (fractions 2–8, corresponding to a molecular



**Figure 3** Size distribution of soluble  $\alpha$ Syn from DLB and control frontal cortex. TBS-soluble, post-175 000  $\times$  g supernatants from cortical pieces with high levels of insoluble (Lewy-associated)  $\alpha$ Syn and corresponding controls were fractionated by SEC (3 MDa exclusion size; see Materials and methods). To correct for the amount of  $\alpha$ Syn injected into the SEC column for each case,  $\alpha$ Syn levels for each fraction were expressed as a percentage of total  $\alpha$ Syn recovered across all the SEC fractions. (A) Percentage of total  $\alpha$ Syn in each SEC fraction (means with SEMs are shown); elution of marker proteins indicated in kDa. Inset:  $\alpha$ Syn recovered in HMW fractions 2–8, corresponding to MWs from  $\sim$ 2 MDa to  $\sim$ 440 kDa. (B) Pooling the HMW fractions revealed significantly higher mean  $\alpha$ Syn levels in DLB cytosolic extracts than controls ( $P=0.0256$ , Mann–Whitney comparison of ranks). DLB:  $n=1-4$  cytosolic fractions per subject; controls:  $n=1-8$ .

weight of 2 MDa–440 kDa) from cytosolic extracts of six DLB and six control brains to this assay. Extracts from both subject groups caused influx of  $\text{Ca}^{2+}$ , indicating the destabilization of lipid membranes (Fig. 4B). On average, application of pooled HMW fractions cytosolic extracts from DLB brains destabilized membranes significantly more than corresponding extracts from control subject brains (Mann–Whitney comparison of ranks,  $P=0.0087$ ; Fig. 4C). Interestingly, samples from brains Syn3 and Syn6 caused complete permeabilization of lipid vesicles. Importantly, immunoneutralization of these samples with an  $\alpha$ Syn-specific monoclonal antibody (2F12), but not a beta-amyloid ( $\text{A}\beta$ )-specific monoclonal antibody (4G8), markedly decreased the  $\text{Ca}^{2+}$  influx (paired  $T$ -test, two-tailed  $P=0.0045$ ). This indicates that the active species in these DLB cortical fractions that potentially enhance membrane permeabilization are composed of  $\alpha$ Syn (Fig. 4D). This finding provides validating human support of similar results obtained in both *in vitro* and mouse models with  $\alpha$ Syn fibrils (Fusco *et al.*, 2016; Mor *et al.*, 2017). Interestingly, despite unremarkable shifts in subcellular localization of  $\alpha$ Syn compared with control extracts (Supplementary Fig. 5A), the two familial DLB brain extracts significantly increased the permeabilization of these vesicles (Supplementary Fig. 5B, Mann–Whitney comparison of ranks,  $P<0.0001$  for both brains). This increase in permeabilization appeared dependent on the presence of exposed  $\alpha$ Syn epitopes (Supplementary Fig. 5C),





**Figure 4** Quantitative measurement of membrane permeabilization upon applying HMW fractions of soluble brain extracts via fluorescence imaging. **(A)** Images obtained through TIRF microscopy show an untreated vesicle sample (*top*, ‘Control’), a vesicle sample treated with soluble, HMW brain material from DLB patient 1594 (*middle*) and a sample treated with ionomycin as a positive control (*bottom*). Scale bars are 3  $\mu\text{m}$ . **(B)** Brain homogenates display variable vesicle permeabilization capability between different cases but low variability within replicates of the same samples. Means with standard deviations are shown. **(C)** Mann–Whitney comparison of ranks analysing average  $\text{Ca}^{2+}$  influx measured from six DLB and six control brains reveals significantly elevated permeabilization of lipid membranes by soluble extracts from DLB patients ( $P = 0.0087$ ). For each patient sample, nine technical replicates were used to generate the data point. Means with SEMs are shown. **(D)** Immunoneutralization of DLB cytosolic extracts with 2F12 anti- $\alpha\text{Syn}$  antibody (2F12, right) reduces the DLB extract-induced permeabilization of the vesicles compared to immunoneutralization anti- $\text{A}\beta$  antibody (4G8, left) (paired  $T$ -test,  $P = 0.0045$ ). For each patient sample, nine technical replicates were used to generate the data point. Each pair of points connected by a line represents a single cytosolic HMW extract. The two pairs of points denoted by a ‘\*’ came from a single DLB brain (Syn2).

although this dependence did not meet the level of statistical significance (paired  $T$ -test,  $P = 0.206$ ).

## Insoluble $\alpha$ -synuclein from dementia with Lewy Bodies brain extracts induces morphological changes in human induced pluripotent stem cell neurons

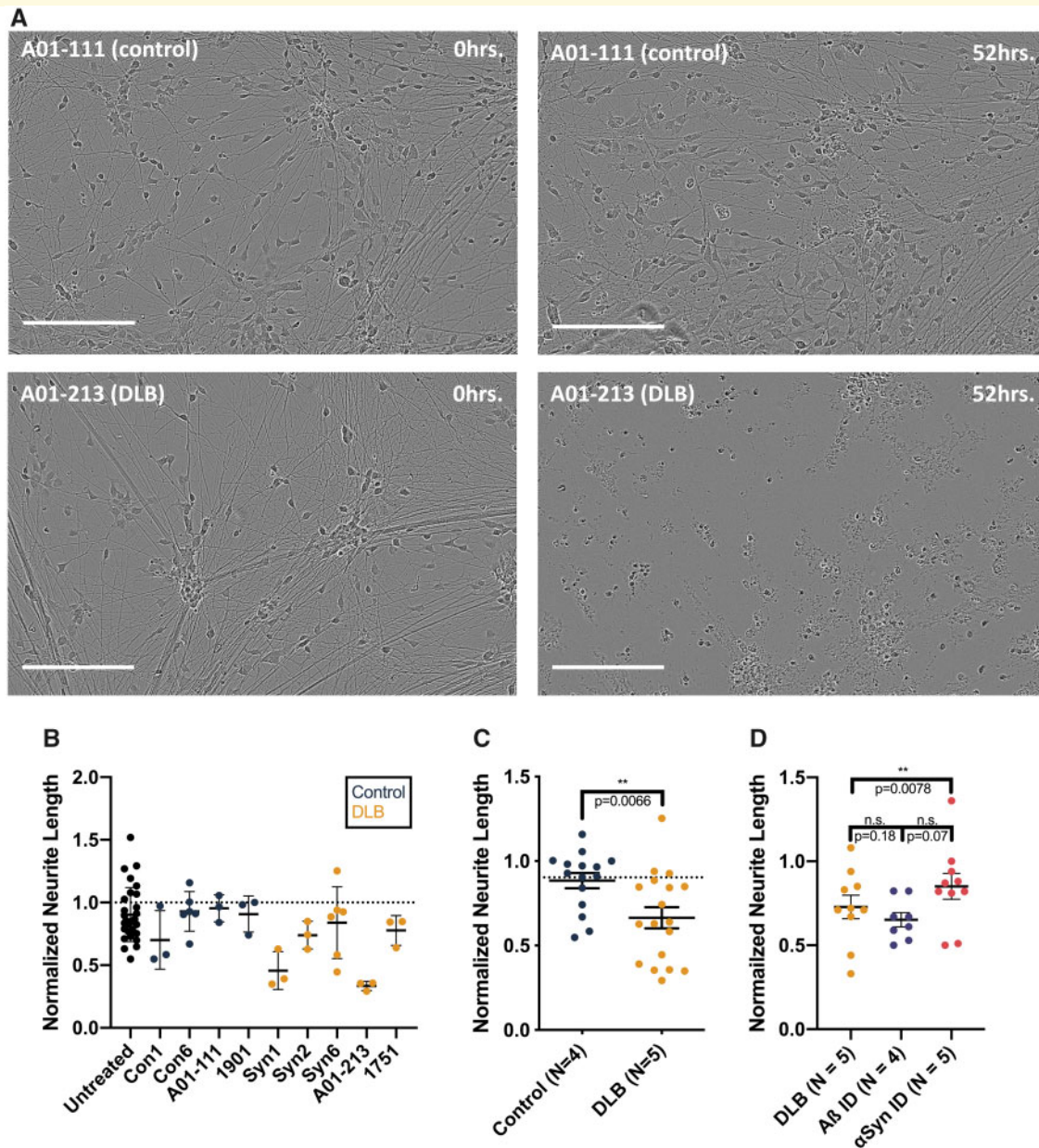
To assess the potential neurotoxicity of insoluble, Lewy-associated  $\alpha\text{Syn}$ , we generated a sonically

dispersed preparation of detergent-insoluble cortical protein using density gradient centrifugation and partial detergent treatment (Blumenstock *et al.*, 2017). These sonicated brain extracts were added to a homogeneous population of iPSC-derived, neurogenin-2-induced human excitatory neurons (Zhang *et al.*, 2013), and neurite morphology was monitored over 88 h (Fig. 5A and Supplementary Fig. 3). Previous works have shown that alterations in the morphology of neurite processes are sensitive indicators of neurotoxicity (Stephens *et al.*, 2005; Scott *et al.*, 2010; Domínguez-Álvaro *et al.*, 2018). Significantly, the particular commercial live-imaging platform used to assess morphological changes in neurons relies not on the expression of non-physiological, but rather on the light microscopy coupled with a computer algorithm for detecting and applying masks to either neurites or cell bodies. For this assay, we applied extracts from five DLB brains with the highest levels of detergent-insoluble/8M urea-solubilized  $\alpha\text{Syn}$  and four corresponding control brains. The effects of these extracts on neurite length showed significant variability across both control and DLB brain samples (Fig. 5B), but when data were aggregated based on neuropathological diagnosis, the DLB extracts led to significantly lower mean neurite length than did the control extracts ( $P = 0.0066$ , Mann–Whitney comparison of ranks) (Fig. 5C). Comparison of cells treated with  $\text{A}\beta$ -immunoneutralized extracts (4G8 antibody) with those treated with  $\alpha\text{Syn}$ -immunoneutralized extracts (2F12 antibody) did not show a significant difference. However, there was a trend towards shorter neurite length in the  $\text{A}\beta$ -immunoneutralized extracts ( $P = 0.0703$ , Wilcoxon paired comparison of ranks), suggesting that the toxicity of these extracts is driven primarily by pathological  $\alpha\text{Syn}$ , not  $\text{A}\beta$  (Fig. 5D).

## Discussion

Using non-denaturing biochemical fractionation of human frontal cortex to avoid potential denaturation of endogenous  $\alpha\text{Syn}$  species (Luth *et al.*, 2015), we have characterized the oligomeric landscape of a variety of DLB and control brain tissue samples and validated the existence of neurotoxic soluble  $\alpha\text{Syn}$  species in human brain. We report and describe a number of biochemical findings, tightly correlating with neuropathology, which could help elucidate the role of  $\alpha\text{Syn}$  in the inception and progression of synucleinopathies. In addition, we identified pieces of tissue from our collection with both biochemical and neuropathological evidence of significant  $\alpha\text{Syn}$  disease burden, establishing a method to identify pathology-rich tissue pieces for further analysis of corresponding soluble species. Notably, we observed a high degree of ‘microheterogeneity’ within single sections of tissue, both by IHC (Fig. 1) and biochemistry (Fig. 2). This observation drove





**Figure 5 Treating iPSC-derived human neurons with Lewy-associated brain extracts.** (A) Neurogenin-induced human neurons were treated with protein-normalized volumes of sonicated Lewy-associated brain extracts. Neurite morphology was monitored for 88–96 h. Images were analysed by applying a neurite mask (Supplementary Fig. 5) and automatically quantifying mean neurite length. In these representative images, neurons treated with  $\alpha$ Syn-enriched control brain extract (top) show largely intact neurites at 52 h, whereas neurons treated with A01-213 DLB  $\alpha$ Syn-enriched extract display neurite retraction (bottom). Images shown from each brain sample are taken from identical fields at different time points. Bars are 200  $\mu$ m. (B) Mean neurite length calculated across four images per well at 88 h was normalized to the mean neurite length at  $t = 0$ . While the amount of  $\alpha$ Syn levels applied to each well was not normalized, these levels did not correlate with the extent of neurite retraction (not shown). Means with standard deviations are shown. (C) Pooling of data collected from four non-synucleinopathy control extracts and five DLB patient extracts across three separate experiments shows a significantly greater neurite retraction in neurons treated with DLB extracts ( $P = 0.0066$ , Mann–Whitney comparison of ranks). Means with SEMs are shown. (D) Human iPSCs were treated with two extracts from five DLB brains, each split into three aliquots: one was applied directly; one was pre-incubated with 2F12, a monoclonal antibody to  $\alpha$ Syn; and a third was pre-incubated with 4G8, a monoclonal antibody to A $\beta$ . Immunoneutralization of  $\alpha$ Syn in these extracts restored the length of neurons treated compared with neurons treated with non-immunoneutralized aliquots of the same DLB extracts ( $P = 0.0078$ , Wilcoxon paired comparison of ranks). Each point represents the mean of technical triplicates from a single experiment. Means with SEMs are shown.

our decision to look at several small tissue pieces per brain, as we feared that unburdened areas of larger sections of tissue might dilute any pathology-specific effects and we recommend future studies to employ a similar approach.

Through the quantification of  $\alpha$ Syn levels in fractions generated through our extraction process, we observed differences in the subcellular distribution of  $\alpha$ Syn species in DLB brains compared with controls. Analysis of our most pathology-rich tissue indicated that the  $\alpha$ Syn levels in the cytosolic and membrane fractions of a single tissue piece were associated with the degree of Lewy neuropathology as determined by its level of insoluble  $\alpha$ Syn. The strong inverse relationship between the relative levels of pathological, Lewy-associated  $\alpha$ Syn and cytosolic or membrane-associated  $\alpha$ Syn implies that the accumulation of insoluble  $\alpha$ Syn is accompanied by a relative decrease in its physiological pools. Our observation that strong  $\alpha$ Syn pathology is directly associated with a decrease in physiological  $\alpha$ Syn in the same small samples of DLB cortex supports the possibility of a loss-of-function of physiological  $\alpha$ Syn in the disease (Gorbatyuk *et al.*, 2010a, b; Benskey *et al.*, 2016).

Our analysis of the absolute concentrations of  $\alpha$ Syn in different fractions points towards this possibility: in patients with DLB, there were lower total levels of  $\alpha$ Syn, an expected consequence of late-stage disease with accompanying neuronal loss. At the same time, according to our analyses, the remaining  $\alpha$ Syn redistributes to the cytosolic and urea-solubilized fractions. We found the relative biochemical increase in urea-solubilized  $\alpha$ Syn to be associated with the presence of insoluble  $\alpha$ Syn aggregates, which are the histopathological hallmarks of DLB and the other synucleinopathies. However, the relative increase in cytosolic  $\alpha$ Syn in DLB cortex is not as readily explained. Our native SEC-ELISA analyses do show a significant increase in soluble, HMW species that could contribute to this redistribution; however, these species are of very low abundance, comprising  $\sim 0.1$ – $1\%$  of the total extractable  $\alpha$ Syn in our assays. Alternatively, the shift in relative abundance away from the membrane-associated fraction to the cytosolic fraction may be related to a loss of physiological function. Previous work has shown that transient interactions with phospholipid membranes induce proper folding and assembly of aggregation-resistant multimeric  $\alpha$ Syn (Rovere *et al.*, 2018), and it appears that  $\alpha$ Syn is likely involved in the physiological recycling of exocytotic (including synaptic) vesicles (Logan *et al.*, 2017). In addition, recent studies have shown that altering the lipid composition of vesicles can mitigate  $\alpha$ Syn-related toxicity (Fanning *et al.*, 2019; Vincent *et al.*, 2018). Thus, it is possible that in the pathological state, more  $\alpha$ Syn is in a disordered, cytosolic state that is more prone to aggregation than the multimeric state (Bartels *et al.*, 2011).

In further analysis of the cytosolic fractions, SEC-ELISA fractionation and analysis showed that, proportionally, HMW  $\alpha$ Syn species are more abundant than in simultaneously and identically prepared extracts of age-matched control cortex. These soluble HMW species span a broad distribution of apparent sizes (440 kDa–3 MDa). Despite this finding, the low absolute levels of soluble HMW  $\alpha$ Syn in both DLB and control cortex posed a challenge for further analyses and characterizations. In fact, the total amount of  $\alpha$ Syn in these fractions, even when pooled, was close to the lower limit of quantification of our ELISA (0.5–1 ng/ml) than to the minimum concentration of synthetic  $\alpha$ Syn pre-formed fibrils required to see an effect in our assays (500 ng/ml). While we observed no frank neurotoxicity of these fractions in a neurite outgrowth assay, our vesicle permeabilization assay indicated that these fractions can decrease phospholipid membrane integrity in an  $\alpha$ Syn-dependent manner, an effect that likely contributes to neurotoxicity over time. The disparate sensitivities of our assays and the low concentrations of  $\alpha$ Syn in these soluble extracts might explain the apparent lack of neurotoxicity. However, our findings in a vesicle permeabilization assay indicate that low amounts of HMW  $\alpha$ Syn present in DLB brain cortex are qualitatively different than HMW  $\alpha$ Syn present in the control cortex of aged brains and suggest a possible mechanism of toxicity.

We also examined the toxicity of the Lewy-associated  $\alpha$ Syn species that are enriched in our detergent-insoluble, urea-solubilized fraction. These insoluble species alone demonstrated the ability to alter neurite morphology in human iPSC-derived neuronal cultures (Fig. 5), providing evidence that highly aggregated  $\alpha$ Syn species found in LBs and Lewy neurites may contribute to the cytotoxicity that leads to the neurodegeneration and clinical manifestation of synucleinopathies. Given that the levels of cytosolic  $\alpha$ Syn (including the soluble HMW  $\alpha$ Syn species) correlate inversely with the levels of insoluble  $\alpha$ Syn in our collection of DLB frontal cortices (Fig. 2F), we speculate that the soluble aggregates are an intermediate species that both directly contribute to neurotoxicity and deplete pools of physiological  $\alpha$ Syn, analogous to the proposed role of amyloid plaques in Alzheimer's disease (Martins *et al.*, 2008; Shankar *et al.*, 2008).

In conclusion, this study characterizes  $\alpha$ Syn-specific biochemical differences in a large sample of DLB and control brain tissue for the first time. We also report on challenges associated with working with multiple brain banks. We hope that reporting on these challenges may standardize future tissue collection and classification protocols. In addition, bioassays of  $\alpha$ Syn-enriched DLB samples provide more clues that should guide future research into the toxic mechanism of pathological  $\alpha$ Syn.

## Supplementary material

Supplementary material is available at *Brain Communications* online.

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## Competing interests

D.J.S. is a director and consultant to Prothena Biosciences. B.T.H. serves on the scientific advisory boards of Biogen and Dewpoint and owns stock in Dewpoint. A family member of B.T.H. works for and owns stock in Novartis. All other authors report no competing interests.

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