SENP2-based N-terminal truncation of α-synuclein in Lewy pathology propagation

Short title:

α-Synuclein truncation by SENP2 in Lewy pathology propagation

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List of abbreviations

αSyn: α-Synuclein

BSA: bovine serum albumin

CM: culture medium

CPI: carboxypeptidase inhibitor derived from potato tuber

cRAP: common Repository of Adventitious Proteins

DDT: dithiothreitol

DLB: Dementia with Lewy bodies

DMSO: dimethyl sulfoxide

FDR: false discovery rate

HE: Hematoxylin and Eosin

HPLC: high-performance liquid chromatography

HRP: horseradish peroxidase

KO: knockout

LBs: Lewy bodies

LBD: Lewy body disease

LNs: Lewy neurites

NM: neuromelanin

PB: phosphate buffer

PBS: phosphate-buffered saline

PBST: phosphate-buffered saline containing Triton X-100

PD: Parkinson's disease

PFA: paraformaldehyde

PFFs: preformed fibrils

PTMs: posttranslational modifications

PVDF: polyvinylidene difluoride

SEC: size-exclusion chromatography

SEM: standard error of the mean

SN: substantia nigra

Str: striatum

WT: wildtype

Abstract

α-Synuclein (αSyn) is a major component of Lewy bodies (LBs) and Lewy neurites (LNs) which are pathological features of Parkinson's disease (PD) and Dementia with Lewy bodies. In the PD brain, with disease progression, LB/LN formation is propagated from the lower brainstem to the cerebral cortex. Prion-like cell-to-cell seed-transmission has been implicated as an underlying mechanism for Lewy-pathology propagation. However, the biochemical properties and production mechanism of those pathogenic seeds are unelucidated. In this study, we ascertained that the seeds released from pathological neurons that harbour LB/LN-like aggregates have the N-terminally truncated form of αSyn. This N-terminal truncation is directly catalysed by SENP2, which is a well-known deSUMOylation enzyme. After SENP2 processing of recombinant αSyn, the SDS-resistant high-molecular oligomer formation was promoted *in vitro*. Inhibition of SENP2 activity suppressed aggregate formation and propagation in cultured neurons and mouse brains. Thus, SENP2 might be a novel therapeutic target in LB diseases.

Introduction

α-Synuclein (αSyn) is one of the major constituents of Lewy bodies (LBs) and Lewy neurites (LNs) that constitute the well-known pathological hallmarks of synucleinopathies, including Parkinson's disease (PD) and dementia with Lewy bodies (DLB)[1-3]. In the sporadic PD brain, the propagation of LB/LN formation indicates a caudorostral process that is associated with disease progression from the lower brainstem, through the basal midbrain and forebrain, into the cerebral cortex[4, 5]. Interestingly, LBs/LNs were observed even in the grafted dopaminergic neurons derived from foetal mesencephalic tissue that has been transplanted into the striatum of a patient with PD[6]. This suggests the presence of prion-like propagation as the underlying mechanism in neurodegeneration[7]. In several experimental models, inoculation of preformed fibrils (PFFs) prepared from recombinant αSyn or PD brain-derived tissue extracts into the mouse striatum can induce LB/LN-like aggregate formation and propagation through the anatomical connections in the synaptic contacts within the brain[8-10].

Abnormal intracellular aggregates, such as LBs and LNs, are formed by the recruitment of intrinsic soluble αSyn into the insoluble aggregate core (the so-called "seeds")[7, 11]. Therefore, the endogenous expression of αSyn as well as seeds, which accelerate the polymerization of soluble αSyn, are required for the formation of aggregates. Next, the

cell-to-cell transmission of seeds is a crucial step for the propagation of Lewy pathology.

However, the biochemical properties and mechanism of production of these pathogenic seeds

that are released from pathological neurons, which harbour LBs/LNs, are unelucidated.

The evidence from recent studies implicates posttranslational modifications (PTMs) of

αSyn, such as phosphorylation, glycosylation, acetylation, ubiquitination, SUMOylation, and

truncation, in the pathophysiological propensities of αSyn[12, 13]. By using histological

methods and mass spectrometric analysis, the presence of various PTMs of αSyn has been

detected in the PD brain. Phosphorylation at the Ser129 residue is a well-known PTM of

αSyn-comprising LBs/LNs and is used as a marker for the detection of these abnormal

aggregates[14]. Furthermore, ubiquitination and SUMOylation play important roles in the

regulation of αSyn's stability and aggregation ability[15, 16]. The truncation of αSyn has

significant effects on the aggregation ability, cross-seeding kinetics, and fibril stability of the

resultant species[12, 17, 18]. Interestingly, there are some characteristic differences between

the C-terminal- and N-terminal-truncated forms of the αSyn[17, 19], wherein the former

shows higher aggregation ability and the latter shows lower aggregation ability and fibril

stability with higher cross-seeding activity in cultured cells and the mouse brain. The

injection of PFFs prepared from the N-terminal 10-residue- or 30-residue-truncated αSyn,

rather than PFFs prepared from full-length αSyn, into the mouse striatum induces more

abundant aggregate formation[17], which, given the consensus on the inverse correlation of

seeding activity with fibril stability in the field of prions, possibly contributes to the lower

stability of N-terminal-truncated αSyn-derived PFFs[20]. Among the various

αSyn-processing enzymes that have been identified thus far, caspase-1 and asparagine

endopeptidase are C-terminal cleavage enzymes, whereas 20S proteasome, calpain-1,

cathepsins, and matrix metalloproteases can truncate both C-terminal and N-terminal regions,

including the NAC region of αSyn[12]. Furthermore, PFFs are processed by lysosomal

cathepsin B, and this partial cleavage enhances the seeding ability of PFFs[21].

Studies using mass spectrometric analysis have identified many cleavage sites within the

tissue extracts obtained from the PD brain[12]. For example, calpain-mediated C-terminal

truncation at 122/123 residues of αSyn has been detected in LBs/LNs[22, 23]; however, the

relationship between αSyn truncation and the biochemical properties of the seeds remains

elusive.

In this study, we explored this relationship through the characterization of

seed-constituting αSyn released from PD-model neurons containing pathological αSyn

aggregates and identified novel cleavage sites in the N-terminal region.

Results

Isolation and characterization of cell-to-cell transmissible seeds

To isolate the cell-to-cell transmissible seeds released from pathological neurons that harbour

LB/LN-like abnormal aggregates, the oligomeric species of αSyn within the conditioned

medium of those pathological neurons were screened using sucrose density-gradient

ultracentrifugation and Western blotting analysis (S1 Fig. A–C and Methods). Accordingly,

140-kDa αSyn oligomers, separated by native-PAGE, were recovered from the interface

fraction between the concentrated conditioned medium and the 0.8 M sucrose-containing

PBS layer. Moreover, we confirmed that the LB/LN-like aggregate formation in the primary

cultured neurons was induced by the fraction-containing fresh medium (S1 Fig. D).

To more precisely isolate the seeds and obviate PFF contamination, isolation was achieved

with microfluidic devices (Fig. 1A, B; C1 and C3 chambers), wherein seeds released from

pathological neurons were recovered from the PFF-free culture medium. Furthermore, using

this microfluidic method, 140-kDa αSyn oligomers that were separated by native-PAGE were

recovered from the interface fraction as well as the results obtained by the batch method (Fig.

1C–E). Atomic force microscopy showed that this fraction contained a uniform population of

granular molecules with a diameter of 36 ± 1.2 nm (Fig. 1F, G). Next, to further confirm that

these granular molecules are composed of αSyn, these molecules were detected by several

antibodies against αSyn. For αSyn detection by immunoelectron microscopy, the reactivity of

secondary antibodies was optimized using PFFs as the positive control (S1 Fig. G). These

granular molecules (diameter 35 ± 2.3 nm) detected by electron microscopy were composed

of αSyn and thus labelled by certain monoclonal antibody 5G4, but not by the O2 antibody

(Fig. 1H–J). 5G4 recognizes protofibrils or large oligomeric species of αSyn[28]. To

determine the molecular weight of the intact seed structure, size-exclusion chromatography

(SEC)-high-performance liquid chromatography (HPLC) analysis was performed (S1 Fig. E,

F). The seed structure detected with immunoelectron microscopy was recovered in the SEC

Fraction No. 8 (corresponding to approximately 1000 kDa), as determined using a

high-molecular-weight standard kit. These results indicated that seeds of approximately

1000-kDa size were produced in the pathological neurons and subsequently released into the

culture medium.

This is the title and legend for Fig. 1

Fig. 1 Isolation and characterization of pathological seeds. **(A)** Schematic of a

microfluidic device for pathological seed preparation. The primary neurons were cultured in

separate chambers (C1, C2, and C3) connected through microgrooves. Preformed fibrils

(PFFs) were supplemented in C2 culture medium, with height kept constantly lower than that

of C1 and C3 media during cell culture. **(B)** Propagation of phosphorylated-αSyn

(pS-αSyn)-positive intracellular aggregates from C2 (upper panels) to C1 and C3 (lower

panels) areas. **(C)** Fractionation of concentrated culture medium (CM) recovered from C1

and C3 using sucrose density-gradient ultracentrifugation. **(D)** SDS-PAGE and

immunoblotting assay using anti-αSyn antibody (numbers indicate recovered fractions in **(C)**;

His-tagged and endogenous αSyn (arrows a and b, respectively)). **(E)** Native-PAGE and

immunoblotting assay of Fraction 4 **(C, D)**. A major band at approximately 140-kDa (arrow)

was detected using anti-αSyn antibody. **(F, G)** Atomic force microscopic imaging of the

molecules recovered in Fraction 4, containing uniform granular molecules (diameter 36 ±

1.2-nm; n=16). White square in **(F)** is magnified in **(G)**. Arrowheads in **(G)** indicate the

examples of those molecules. **(H–J)** Immunoelectron microscopic imaging of molecules in

Fraction 4; αSyn-positive granular molecules detected using anti-pan-αSyn antibody (black

arrowheads: 10-nm gold particles) in **(H–J)**. Granular molecules were recognized by

anti-aggregated αSyn antibody 5G4 (**J**; red arrowheads, 6-nm gold particles), but not by the

anti-oligomeric αSyn antibody O2 (**I**; negative for 6-nm gold particles). αSyn-positive

granular molecules (diameter 35 ± 2.3 nm; n=6). Data represent mean ± SEM. Scale bars: 20

μm and 1 μm in **B** and **F**, respectively; 100 nm in **G**, and 50 nm in **H–J**.

Seeds contain truncated forms of αSyn

To investigate whether seeds contained the truncated αSyn or not, the cleavage sites within

αSyn were screened using mass spectrometric analysis. As a large amount of seed protein

was required for identification of the cleavage sites, a large microfluidic device was

developed (S2 Fig.). Using immunoprecipitation, samples for mass spectrometric analysis

were prepared from seed fraction (Fig. 2A). PFF-constituting αSyn, which was derived from

the human αSyn sequence, was prepared as a control sample for mass spectrometric analysis

because results of the artificially cleaved sites detected in PFF-constituting αSyn were

subtracted from those of the cleaved sites detected in the seed-constituting αSyn. Protein

bands, Nos. 1–5, and control bands were subjected to mass spectrometric analysis (Fig. 2B),

and the results suggested the presence of 6 cleavage sites (Fig. 2C). The C-terminal cleavage

site at the 119/120 residue is a 20S proteasome-processing site[24]. To our knowledge,

cleavage sites detected at the N-terminal and NAC regions have not been reported previously.

Interestingly, conserved cleavage sites (KX|KXGV), the N-terminal sides of lysine residues

located at the N-terminal repeat regions (1–4) of αSyn, were detected (Fig. 2C). Although the

cleavage site at repeat 4 was detected in the results of the control sample in the mass

spectrometric analysis, this site was considered as a possible cleavage site, because the

primary amino acid sequence is completely conserved at repeat 3. These results suggested

that the N-terminus of the seed-constituting αSyn was partially processed by unknown

N-terminal lysine endopeptidase activity. This information on the conserved processing sites

is useful for identifying the processing enzyme. Therefore, we focused on these N-terminal

cleavage sites of αSyn.

This is the title and legend for Fig. 2

Fig. 2 Truncated αSyns are included in the pathological seeds. (A) Experimental design to

prepare samples for the detection of αSyn truncation using mass spectrometric analysis. **(B,**

C) Immunoprecipitated proteins were separated by SDS-PAGE and detected by silver

staining as in **(B)**. Gel bands indicated by the numbers 1–5 were dissected and subjected to

mass spectrometric analysis. Predicted truncation sites are shown in **(C)**. Truncated sites

detected in common amino acid sequence between human and mouse αSyn are indicated in

red, whereas sites detected in human or mouse αSyn-specific sequences are indicated in blue

or green, respectively. **(D)** Schematic representation of the predicted cleavage sites and the

corresponding insertion of each mutation to inhibit the cleavage.

αSyn is processed by SENP2

Based on the results of mass spectrometric analysis, candidate enzymes that cleave the

N-terminal side of lysine residues located at the N-terminal repeat regions of αSyn

(conserved cleavage sites in Fig. 2C) were screened using the Peptidase database, MEROPS,

which showed that carboxypeptidase B, USP7, and SENP2 were candidate enzymes that can

process the N-terminal side of the lysine residues. After screening these enzymes, each

enzyme's inhibitors were selected (Table 1). To examine the inhibitory effect of the inhibitors

on aggregate formation, primary neurons were cultured and treated with PFFs in the presence

of each inhibitor. In the inhibitor assay, NSC632839 could significantly inhibit the aggregate

formation (Fig. 3A–F), whereas the reagent inhibits the enzymatic activity of not only USP7

and SENP2 but also USP2 and UCHL1[25, 26]. Therefore, to examine whether αSyn is

directly processed by these enzymes, protease assays were performed using recombinant

active enzymes. Recombinant αSyn was cleaved by SENP2, and the resulting fragment was

detected clearly (Fig. 3G). However, cleaved fragments of αSyn were not detected in the

presence of USP7, USP2, and UCHL1 (S4 Fig. A–C). SENP2 preferentially cleaved the

soluble form of αSyn, rather than PFFs. The cleaved fragment of αSyn increased, depending

on the SENP2 dose (Fig. 3H, I). Interestingly, SDS-resistant high-molecular-weight signals of

αSyn were increased by high-dose SENP2 treatment. This cleavage pattern of αSyn by

SENP2 was diminished by the insertion of mutations at the predicted cleavage sites, as

represented in Fig. 2D (Fig. 3J).

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Fig. 3 Identification of processing enzyme. (A–F) Inhibition of aggregate formation by

NSC632839. Primary neurons were treated with PFFs and cultured in the presence of DMSO

(A) or NSC632839 **(B)**. The region marked by a white square in **(A)** and **(B)** is magnified in

(C) and **(D)**, respectively. Arrowheads in **(A)** and **(C)** indicate the formation of the aggregates.

Quantitative comparison of the phosphorylated-αSyn-positive area **(E)** and DAPI-positive

area **(F)** between DMSO-treated and NSC632839-treated neurons (n=3 fluorescent images

from 2 coverslips were analysed for each treatment; **$p < 0.01$, two-tailed, unpaired t-test;

two independent cultures were performed and confirmed reproducibility). **(G)** Cleavage of

αSyn by GST-fused recombinant SENP2 for 4 h at 37°C. A cleaved fragment is indicated by

the arrowhead. Mono, αSyn monomer. **(H–J)** Dose-dependent effect of SENP2 on

αSyn-cleavage. αSyn was processed for 24 h at 37°C. The cleaved fragment (red arrowhead)

was gradually increased in the presence of a higher amount of SENP2. Black and white

arrowheads show His-tagged full-length αSyn- and GST-fused SENP2 recombinant protein,

respectively. Note that αSyn-cleavage increased high-molecular-weight SDS-resistant

immunoreactivity (asterisk). This αSyn-cleavage by SENP2 for 4 h was inhibited by the

insertion of the N-terminal mutations **(J)**. Scale bar: 20 μm in **(A–D)**.

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Table 1. Screening of αSyn-processing enzyme

	Candidate enzyme	Inhibitor	Inhibition of aggregate formation	Activity of αSyn-cleavage
1	Carboxypeptidase B (B2, N)	Mergepta and CPI	(-)	
2	USP2			(-)

	(ubiquitin specific peptidase 2)			
3	USP7 (ubiquitin specific peptidase 7)	NSC632839	(+)	(-)
4	SENP2 (Sentrin-specific protease 2)			(+)
5	UCHL1 (ubiquitin carboxy-terminal hydrolase-1)			(-)

SENP2 was ordinarily expressed in the nucleus[27]. After aggregate formation, SENP2

was partially distributed in the aggregates (Fig. 4). In cultured neurons harbouring

phosphorylated-αSyn aggregates induced by PFFs, SENP2 was partially detected in those

aggregates (Fig. 4A). In the human PD brain, SENP2 was observed in the amorphous

aggregates composed of phosphorylated αSyn, but not in typical LBs, which are spheroidal

(Fig. 4B and S5 Fig. C, D).

This is the title and legend for Fig. 4

Fig. 4 Presence of endogenous SENP2 in αSyn-aggregates. (A) Presence of SENP2 in

PFF-induced αSyn-aggregates formed in primary cultured neurons. **(B)** Presence of SENP2 in

αSyn-aggregates formed in the neuron of the substantia nigra pars compacta of PD-brain.

Arrowheads indicate examples of the overlapping intense signals of SENP2 and

phosphorylated αSyn. Scale bar: 5 μm.

Inhibition of SENP2 suppresses aggregate formation and its propagation

Biochemical studies demonstrated that SENP2 cleaved the N-terminal repeat regions of αSyn

whereas its inhibitor, NSC632839, suppressed aggregate formation in cultured neurons. Next,

we studied the inhibitory effects of SENP2 through the inhibitor infusion on aggregate

formation and its propagation *in vivo*. The efficiency of the aggregate formation and

inhibitory effect of NSC632839 on the propagation in the brain was examined using

immunohistochemical methods. Injection of PFFs into the striatum (Fig. 5A) showed

aggregate formation, which was propagated from the striatum to other brain regions, such as

the piriform cortex and amygdala, whereas inhibitor infusion into the lateral ventricle

decreased aggregate formation and its propagation (Fig. 5B). The number of cell bodies

harbouring phosphorylated-αSyn-positive aggregates was significantly decreased in the

piriform cortex and amygdala (Fig. 5C, D). In this process, endogenous expression of soluble

αSyn was required for aggregate formation, because αSyn-KO mice did not show any

spreading of phosphorylated αSyn aggregates after the PFF injection (S6 Fig.). These results

indicated that the inhibition of the enzymatic activity attenuated aggregate formation and its

propagation efficiency.

This is the title and legend for Fig. 5

Fig. 5 Inhibition of aggregate formation and cell-to-cell seed transmission. (A) Schematic

representation of PFF injection in the WT mouse striatum, and DMSO or NSC632839

osmotic pump-based infusion into the lateral ventricle. C, caudal; LV, lateral ventricle; R,

rostral; Str, striatum. **(B)** Inhibitory effect of NSC632839 infusion on the propagation of

αSyn-aggregate formation in the ipsilateral sites of the striatum (injection site), piriform

cortex, and amygdala. The region marked by a white square in the amygdala is magnified in

each treatment. **(C, D)** Statistical analysis of the number of cell bodies harbouring

pathological aggregates of phosphorylated αSyn, with or without NSC632839 treatment (**C**,

n=5 NSC632839-treated and n=3 DMSO-treated control mice, the ipsilateral piriform cortex

was analysed by using the two-tailed, unpaired t-test with Welch's correction; **D**, n=3 mice in

each treatment for the ipsilateral amygdala analysed by the two-tailed, unpaired t-test; *$p <$

0.05). Bar: 100 μm in **(B)**.

Discussion
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Recent studies indicated that cell-to-cell seed transmission is a pivotal mechanism underlying

the propagation of Lewy-pathology in the PD brain[7]; however, the properties of seeds

remain unelucidated. In this study, SEC-HPLC analysis demonstrated that seeds released

from pathological cultured neurons were recovered as αSyn oligomers (approximately 1000

kDa), whereas native-PAGE assay showed a 140-kDa band. These results suggest that intact

seeds are composed of 140-kDa rigid oligomers as the elementary units. During

electrophoresis, the recovered seeds might become dissociated from the original

high-molecular structure into those elementary units. These seeds were preferentially

recognized by monoclonal antibody 5G4, but not O2. Furthermore, LBs and LNs were

recognized by these antibodies[28, 29]. Although LBs/LNs have epitopes against both 5G4

and O2 on their surfaces, seeds have only 5G4 epitopes on their molecular surfaces. The

different structural milieu between LBs/LNs and the seeds might be responsible for this

distinct immunoreactivity.

Biochemical and mass spectrometric analysis showed that seeds contained truncated forms

of αSyn. Interestingly, N-terminal sides of lysine residues located in the N-terminal repeat

regions of αSyn were processed by SENP2 – an enzyme that routinely removes SUMO from

various SUMOylated proteins[30]. However, in the present study, SENP2 directly cleaved

recombinant αSyn itself. After αSyn cleavage by high-dose SENP2, the formation of

SDS-resistant high-molecular oligomeric species was facilitated by this treatment (see, Fig.

3H, I). Furthermore, this enzymatic activity was crucial for aggregate formation. The

inhibitory effects of NSC632839 on aggregate formation and its propagation were

significantly observable *in vitro* and *in vivo*. Therefore, we further investigated the

knockdown effects of SENP2 using AAV-assisted shRNA expression system in primary

cultured neurons. However, SENP2 knockdown severely impaired normal neuronal

development. Moreover, neuronal viability was decreased by the knockdown treatment

during the neuronal maturation. On the other hand, significant knockdown efficiency was not

obtained using matured neurons. These results suggested that SENP2 was indispensable for

neuronal development, and its protein turnover rate in immature neurons might be higher than

that in matured neurons. In the present study, NSC632839-based inhibition of SENP2 did not

induce cellular damage, and this might be related to the neuronal maturation level.

Previous studies indicated that N-terminal repeat regions of αSyn implicate lipid

membrane association with conformational changes to α-helical structure[31, 32]. N-terminal

truncation causes β-sheet structure-enriched conversion within the truncated αSyn[31] and

the resultant decreased stability of PFFs[17]. Thus, the N-terminal truncation of αSyn might

promote pathogenesis, including αSyn oligomerization, and further provide an intracellular

milieu that readily accelerates seed production.

Interestingly, a recent study revealed the presence of various kinds of αSyn-PTMs, such as

acetylation, methylation, and carboxymethylation, in tissue extracts prepared from Lewy

body disease (LBD) and control brains[33]. According to this study, N-terminal lysine

residues at K12, K23, K34, and K45, which are SENP2-processing sites, are

post-translationally modified with acetylation. The seeding ability of soluble αSyn harbouring

K45Q, which is one of the acetylation mimic mutations, is significantly decreased. On the

other hand, lysine residues at K12 and K23 are modified with carboxymethylation in LBD

brains, but not in control brains. The seeding ability of αSyn might be regulated by the

balance of PTMs, including truncation, acetylation, and carboxymethylation.

SENP2 was partially distributed in aggregates of phosphorylated αSyn in cultured neurons

and PD brain. However, that was not detected in a typical LB, which might be attributable to

the maturation stages of the aggregates. In the PD brain, SENP2 was detected in amorphous

aggregates, rather than spheroidal aggregates, such as typical LBs, which are larger and more

rigid than amorphous aggregates. SENP2 cleaves soluble αSyn, and the resulting fragments

mediate aggregate formation. SENP2 may persist in immature aggregates but might be

excluded from matured aggregates.

Conclusion

382   Transmissible seeds contained the N-terminal truncated αSyn. SENP2 directly processed

383 N-terminal repeat regions of αSyn, and inhibition of its activity suppressed aggregate

384 formation and propagation in cultured neurons and mouse brains. However, these results,

385 especially the presence of the characteristic seeds in the human brain are unconfirmed. Owing

to the limited amount of seeds recovered, it is currently challenging to produce

structure-specific antibodies for *in vivo* seed detection using microfluidic devices. Further

studies are required for precise seed characterization. However, the present study revealed

that SENP2 activity is essential for Lewy-pathology formation. Thus, SENP2 constitutes a

potential novel therapeutic target, and fine-tuning the PTMs of αSyn, including N-terminal

truncation, could provide an innovative therapeutic strategy to prevent the progression of LB

diseases.

Materials and Methods

Animals

αSyn homozygous knockout (KO) mice (B6;129X1-*Snca^{tm1Rosl}*/J[34]) were purchased from

the Jackson Laboratory and maintained on a C57BL/6N background. Adult male C57BL/6N

wildtype (WT) and αSyn-KO (12- to 18-week-old) mice were used in the histological

experiments. KO mice were genotyped according to the manufacturer's protocol (Jackson).

The mice were housed under a 12-h light/dark cycle in a temperature-controlled environment

with ad libitum food and water in our animal facility. All animal experimental procedures

conformed to the Guidelines for Animal Experiments, Kyoto Prefectural University of

Medicine (approval numbers: M2024-148, M2024-154, and M2024-509) and national

regulations, and the number of animals used and their suffering were minimized.

Ethics approval and consent to participate

The use of postmortem brain tissue in this study was approved by the Institutional Review Boards of Kyoto Prefectural University of Medicine (approval no. ERB-C-1846-2). Written informed consent for cadaver surgical training and the use of tissue samples for medical research was obtained, both from patients themselves and their next of kin.

Consent for publication

Not applicable

Method details

Primary neuronal culture

Using a previously reported method[35], with slight modifications, primary cultured neurons were obtained. Briefly, dissociated cells obtained from the prefrontal cortices and hippocampi of WT mouse embryos (E15–18) were disseminated onto polyethyleneimine-coated coverslips or into the chambers of microfluidic devices, and cultured in Neurobasal Medium (Gibco, Thermo Fisher Scientific) supplemented with B-27 (Gibco, Thermo Fisher Scientific), L-glutamine (Nacalai Tesque), and penicillin/streptomycin (Nacalai Tesque). Half of the volume of the culture medium was changed every 3–4 days until use. To induce the formation

of intracellular phosphorylated αSyn aggregates, neurons were cultured in PFF-supplemented

complete Neurobasal Medium (final 20 μg/mL) for 4–5 days, unless specified otherwise.

Preparation of recombinant αSyn and PFFs

For the preparation of recombinant human αSyn from bacterial culture, plasmid vectors were

constructed as described previously[36]. Briefly, PCR fragments of human αSyn were

inserted into pTrc-His-TOPO vector (Thermo Fisher Scientific). His-tagged αSyn was

purified by cOmplete™ His-Tag Purification Column (Roche). To confirm the purity of the

protein by SDS-PAGE, the purified fraction was electrophoresed, and the protein band was

stained by Coomassie Brilliant Blue. The final protein yield was determined with Lowry's

method.

PFFs were prepared according to a previously reported method, albeit with some slight

modifications[11]. Briefly, PFFs were generated by incubating purified αSyn in sterile PBS

(final concentration: 2 mg/ml) at 37°C with constant agitation for 5–6 days, then

ultracentrifuging (at 200,000g) the PFF-containing solution for 2 h, and recovering the PFF

precipitate at the bottom of the centrifuge tube. The resulting pellet was dissolved in sterile

PBS and stored at −30°C until use.

To prepare recombinant αSyn harbouring mutations at the N-terminal repeat region (Fig.

2D), a DNA fragment of mouse full-length αSyn harbouring those mutations was artificially

synthesized and inserted into the pEX-A2J2 vector by Eurofins Genomics Japan. A DNA

fragment of the mutated αSyn was further cloned into pTrc-His2B vector (Thermo Fisher

Scientific). The presence of each mutation was confirmed by DNA sequencing using the ABI

PRISM3130 Genetic analyzer. The His-tagged mutated αSyn-recombinant protein was

purified using the abovementioned method described for WT αSyn.

Microfluidic devices

Transmission of pathological αSyn was reproduced *in vitro* using microfluidic devices[37].

To isolate cell-to-cell transmissible pathological seeds, two types of microfluidic devices

were used. For small-scale preparation of the seeds, three-compartment devices (TCND500)

were purchased from Xona and cell culture was performed according to the manufacturer's

protocol. Briefly, pathological neurons harbouring intracellular phosphorylated-αSyn

aggregates were cultured in the middle chamber, C2 (Fig. 1A). Seeds released from those

pathological neurons were recovered in the culture medium in the side chambers, C1 and C3,

which are connected to C2 through microgrooves. During cell culture, to prevent the

backflow of the PFF-containing medium into the C1 and C3 chambers, the medium in the C2

chamber was constantly maintained at a lower height than that of the C1 and C3 chambers.

For large-scale preparation of seeds to perform mass spectrometric analysis, a large

polydimethylsiloxane-based microfluidic device was developed (S2 Fig. A–C). Before the

dissemination of dissociated cells prepared from the embryonic brain, the large microfluidic

device was settled on a polyethyleneimine-coated glass dish that comprises two areas – a

central square area and a hollow square (outer) area – which are connected by microgrooves

through which the axons extend (S2 Fig. D, E). Pathological neurons were cultured in the

central area of this device, and the seeds released from these neuronal cells were recovered

from the medium in the outer area. During cell culture, the height of the medium in the

central area was constantly maintained lower than that in the outer area to prevent the

backflow of the PFF-containing medium.

Preparation of the seed fraction

Unless specified otherwise, the seeds were prepared using the microfluidic devices. Partly

seeds fraction was prepared using the batch method (S1 Fig. A–F). After the PFF treatment

on primary neurons cultured in the middle or central square chamber, half of the volume of

the seed-containing medium in the side (C1 and C3) or outer chambers was recovered every

3–4 days for 14 days. By ultrafiltration through a 3-kDa molecular-weight cutoff membrane

(Merck Millipore), this seed-containing medium was concentrated up to 10 folds. Next, a

discontinuous sucrose density gradient was prepared in a centrifuge tube purchased from

Beckman Coulter (Fig. 1C and S1 Fig. A). Each layer comprised phosphate-buffered saline

(PBS, pH 7.4) containing the indicated concentration of sucrose. The concentrated medium

was overlaid on a 0.8 M sucrose-containing PBS layer, ultracentrifuged at 200,000g for 12 h

at 4°C, and an equal volume (250-300 μL) was aliquoted and further subjected to

biochemical analysis. Seeds were recovered on the interface between the concentrated

medium and the 0.8 M sucrose-containing PBS layer (Fraction 4 in Fig. 1C).

For seed preparation using the batch method, the conditioned medium of pathological

neurons was recovered as the seed-containing medium. After 2-day culture under PFF

treatment, the PFF-containing medium was discarded, the cells were washed 4 times with

complete Neurobasal Medium and then cultured in a fresh Neurobasal Medium for 5–7 days.

This conditioned medium was collected and concentrated using ultrafiltration as described

above. The concentrated medium was fractionated using sucrose density-gradient

ultracentrifugation as well as the microfluidic method. Seeds were recovered at the interface

between the concentrated medium and 0.8 M sucrose-containing PBS and subjected to

biochemical analysis or aggregate-formation assay (S1 Fig.).

Immunoblotting assay

For SDS-PAGE, samples were denatured by heating with 1% 2-mercaptoethanol-containing

SDS-sample buffer (62.5 mM Tris-HCl, pH 6.8, 2% sodium dodecyl sulfate, 0.003% Pyronin

Y, and 10% glycerol) at 98°C for 3 min. Proteins were separated by 15% polyacrylamide gel

(Wako) and transferred to a polyvinylidene difluoride (PVDF) membrane (Bio-Rad

Laboratories). For native-PAGE, the samples were treated with sample buffer (15% glycerol,

0.5 M 6-aminocaproic acid, 0.001% Ponceau S, and 50 mM Bis-Tris/HCl, pH 7.0). Proteins

were separated by NativePAGE™ Bis-Tris Gel purchased from Thermo Fisher Scientific and

electrophoresed using anode (50 mM Bis-Tris/HCl, pH 7.0) and cathode (50 mM Tricine, 15

mM Bis-Tris/HCl, pH 7.0) buffers. After electrophoresis, polyacrylamide gel was immersed

in a transfer buffer containing 1% SDS, and the separated protein bands were also denatured

in this step. The denatured proteins were transferred to the PVDF membrane; the membrane

was blocked with 5% skim milk in Tris-buffered saline (TBS, pH 7.4) containing 0.1%

Tween-20 for 1 h at room temperature. After treatment with primary antibody, the membrane

was incubated with an alkaline phosphatase-conjugated secondary antibody. Protein bands

were detected using the NBT-BCIP system (Nacalai Tesque). For αSyn detection, mouse

monoclonal antibody Syn-1 (1:1000; BD Biosciences) or rabbit polyclonal antibody C-20

(1:2000; Santa Cruz Biotechnology) was used.

Immunocytochemistry

Cultured cells were fixed with 2% paraformaldehyde (PFA) in the culture medium for 10 min

at room temperature, permeabilized with 0.1% Triton X-100 in PBS for 10 min, and blocked

with 10% normal goat serum (NGS) in PBS for 30 min. Next, cells were incubated for 1–2 h

with primary antibodies in the blocking solution, washed with PBS, and treated with

secondary antibodies. The primary antibodies were detected with Alexa488- and

Alexa594-conjugated secondary antibodies (Thermo Fisher Scientific). After labelling with

the secondary antibodies, cells were washed with PBS and then with milliQ water (Merck

Millipore), before being mounted with FluorSave (Merck Millipore).

For αSyn detection, Syn-1 or C-20 antibodies were used. Phosphorylated αSyn was

detected by mouse monoclonal antibody pSyn#64 (1:1000; Wako) or rabbit polyclonal

antibody phospho-S129 αSyn (1:1000; Abcam). SENP2 was detected by rabbit polyclonal

antibody (1:250; GeneTex). Low-magnification images were acquired by using

immunofluorescent signals captured through a UPlanFl 10×/0.30 Ph1 objective with an

inverted fluorescent microscope IX71 (Olympus) equipped with a cooled charge-coupled

device camera (Olympus). For confocal observation, images were acquired as Z stacks (5–15

z-sections, 1-µm apart, 1024 × 1024 pixels) using a Plan-Apochromat 63×/1.40 Oil DIC

objective (Carl Zeiss) with an inverted laser-scanning confocal microscope, LSM900 (Carl

Zeiss) or using an HC PL APO CS2 63×/1.40 OIL objective (Leica) with a laser-scanning

confocal microscope, STELLARIS5 (Leica) for high-resolution imaging (Fig. 4A).

Atomic force microscopy

To acquire images of the seed structure, the seed fraction was dropped onto the surface of a

freshly cleaved mica plate (NILACO) and incubated for 5 min. Images were acquired using

atomic force microscopy, SPM-9700 (SHIMADZU), operating in Dynamic Mode. The

scanning parameters were as follows: operating point, 0.146–0.166 V; drive (tapping)

frequency, 323–325 kHz; and scan rate, 1.0 Hz.

Size-exclusion chromatography

To determine the molecular weight of the seeds, size-exclusion chromatography

(SEC)–high-performance liquid chromatography-based analysis was performed using the

SMART system (Pharmacia). Briefly, 50 μL seed fraction was injected into a sample tube and

separated using the Superose 6 column comprising filtered PBS (pH 7.4) as the mobile phase.

Each sample was eluted at a 40 μL/min flow rate; each 80 μL was collected as a fraction.

Protein elution was monitored by UV absorbance at 280 nm. To obtain a standard curve for

the molecular-weight determination, gel-filtration standards (Gel Filtration Calibration Kit

HMW) purchased from GE Healthcare were separated using the same SEC column and flow

rate. To detect the presence of the seed structure recovered in these SEC fractions, each SEC

fraction was further observed using immunoelectron microscopy.

Immunoelectron microscopy

For immunolabelling of seeds or PFFs, the seed fraction or sonicated PFFs were dropped

onto the surface of formvar-coated molybdenum grids, PVF-M15 STEM Mo150P

(Okenshoji), and incubated for 10 min. After incubation, the excess solution on each grid was

discarded; the grid surface was further blocked with 1% bovine serum albumin (BSA) in PBS

for 15 min and incubated with primary antibodies for 1 h. After washing in PBS, primary

antibodies were detected with 10-nm gold (BBI Solutions) and/or 6-nm gold

(Abcam)-conjugated secondary antibodies in PBS containing 1% BSA. After labelling, the

grids were washed with PBS, stained with uranyl acetate, and observed using a JEM-1220

electron microscope (JEOL) at 80 keV.

For αSyn detection, the C-20 antibody was used as a pan-αSyn antibody. For the detection

of oligomeric species of αSyn, the O2 (1:500; BioLegend) and 5G4 (1:500; Merck Millipore)

antibodies, which recognize soluble oligomers and fibril forms of αSyn, including LBs and

LNs[28, 29], were used.

Mass spectrometric analysis for screening of the αSyn-cleavage site

To detect the cleavage sites of the seed-constituting αSyn from the mass spectrometric

analysis sample, the seeds were concentrated by immunoprecipitation. Briefly, the seed

fraction was prepared using large microfluidic devices as the original fraction for

immunoprecipitation. The seeds were concentrated by immunoprecipitation after labelling by

a C-20 antibody and precipitation with protein A-conjugated magnet beads (Bio-Rad), and

then eluted by the SDS sample buffer at 56°C for 15 min. The eluted proteins were further

denatured by heating at 98°C with 1% 2-mercaptoethanol for 3 min, and separated by

SDS-PAGE. Protein bands were detected by silver staining using the SilverQuest™ Silver

Staining Kit (Thermo Fisher Scientific). The PFF-constituting αSyn was prepared as the

control for mass spectrometric analysis, because the results of the artificially cleaved sites

detected in the PFF-constituting αSyn were subtracted from those of the cleaved sites

detected in the seed-constituting αSyn. These protein bands (Nos. 1–5 and control bands, Fig.

2B) were subjected to mass spectrometric analysis.

For mass spectrometric analysis using Nano-liquid chromatography–tandem mass

spectrometry (LC–MS/MS), the proteins present in each gel slice underwent a series of

stepwise treatment as follows: reduction with 10 mM dithiothreitol (DDT) at 56°C for 1 h,

alkylation with 55 mM iodoacetamide at room temperature for 45 min in the dark, and

digestion with 10 µg/mL modified trypsin (Pierce, MS Grade) at 37°C for 16 h. The resulting

peptides were extracted using 1% trifluoroacetic acid and 50% acetonitrile, vacuum-dried,

and dissolved in a solution of 2% acetonitrile and 0.1% formic acid.

Mass spectra were acquired using an LTQ-Orbitrap Velos Pro (Thermo Fisher Scientific)

coupled with a nanoflow UHPLC system (ADVANCE UHPLC; AMR Inc.) featuring

Advanced Captive Spray SOURCE (AMR Inc.). The peptide mixtures were loaded onto a

C18 ID 0.1 mm × 20 mm, 5-µm (particle size) trap column (Acclaim PepMap 100 C18,

Thermo Fisher Scientific) and fractionated through a C18, ID 0.075 × 500 mm, particle size

3-μm column (CERI). The peptides were eluted at a flow rate of 300 nL/min with a linear

gradient of 5–35% solvent B over 60 min. Buffers A and B comprised 0.1% formic acid and

100% acetonitrile, respectively. The mass spectrometer was programmed for 11 successive

scans, including a full-scan MS over 350–2,000 m/z by orbitrap at a resolution of 60,000, and

automatic data-dependent MS/MS scans of the 10 most intense ion signals in the first

precursor scan by ion-trap detectors for the second and eleventh scans. Using a 2 m/z

isolation width and a 90-s exclusion time for molecules in the same m/z range, MS/MS

spectra were obtained with a normalized collision energy of 35% CID.

The raw files were searched against the mouse and human αSyn amino acid sequences,

including the N-terminal His-tag sequence, with Mus musculus dataset (Uniprot Proteome

UP000000589 2021_03_07 downloaded), Homo sapiens dataset (Uniprot Proteome

UP000005640 2022_05_25 downloaded), and the common Repository of Adventitious

Proteins (cRAP; ftp://ftp.thegpm.org/fasta/cRAP). The search was conducted through

MASCOT version 2.6 (Matrix Science) via Proteome Discoverer 2.4 (Thermo Fisher

Scientific). The following settings were used as the search parameters: Enzyme,

Semi-trypsin; false discovery rate (FDR), 0.01; precursor mass tolerance, 10 ppm; product

tolerance, 0.8 Da; fixed modification, carbamidomethylation of cysteine; variable

modification, oxidation of methionine, and acetylation of protein N-termini; and maximum

missed cleavages, 2.

Screening of the αSyn-processing enzyme and inhibitor assay

Based on the results of mass spectrometric analysis, the candidate enzymes which cleave the

N-terminal side of lysine residues located at the N-terminal repeat regions of αSyn

(conserved cleavage sites in Fig. 2C) were screened using the Peptidase database, MEROPS

(https://www.ebi.ac.uk/merops/); next, inhibitors for each enzyme were selected (Table 1),

dissolved with dimethyl sulfoxide (DMSO), and stored at −30°C until use. To ascertain the

inhibitory effect of the inhibitors on aggregate formation, primary neurons were cultured and

treated with PFFs in the presence of each inhibitor. The following inhibitors were purchased

from the manufacturers and used at the indicated final concentration: Mergepta (46 μM;

MuseChem), carboxypeptidase inhibitor derived from potato tuber (CPI) (1 μM; Sigma), and

NSC 632839 (9 μM; Selleckchem). As a control, primary neurons were cultured with diluted

DMSO as well as each inhibitor.

Image analysis and quantification of the inhibitory effect of NSC632839 on

aggregate formation in primary cultured neurons

The phosphorylated αSyn-positive and DAPI-positive areas were measured using WinRoof

2018 software (Mitani Corporation). As shown in S3 Fig., the phosphorylated-αSyn

immunoreactive area, excluding the DAPI-positive area, was quantitatively analysed because

the monoclonal antibody against phosphorylated αSyn #64 tends to react with nuclei. In

either the presence or absence of NSC632839, wherein the threshold intensity was kept

constant and set to include measurement profiles by visual inspections. Data represent the

mean ± SEM. Statistical analysis was performed using GraphPad Prism 5 with a two-tailed,

unpaired *t*-test and assessed using a significance level of $p < 0.05$ (n=3 fluorescent images

acquired from 2 coverslips were analysed for each treatment; two independent cultures were

performed, and reproducibility was confirmed).

Protease assay

To examine whether αSyn is processed by the several candidate enzymes summarized in

Table 1, a protease assay was performed using recombinant active enzymes: 7 µg soluble or

PFF forms of recombinant αSyn (substrate) was mixed with 0.5–5.2 µg of each recombinant

enzyme in the following buffer to a final volume of 100 µL: USP7 (50 mM HEPES (pH8.0),

0.1 M NaCl, and 1 mM TCEP), SENP2 (50 mM HEPES, pH 8.0; 0.15 M NaCl; and 1 mM

DTT), USP2 and UCHL1 (50 mM HEPES, pH 8.0, 0.15 M NaCl, 0.1 mM EDTA, and 1 mM

DTT). PFFs were sonicated using the Hielscher ultrasonic processor UP50H with sonotrode

MS1 (10 s on–10 s off, repeated 10 times on ice, at 100% amplitude setting) prior to use.

Each substrate was incubated with each enzyme at 37°C for the indicated duration (4–24 h).

The reaction was stopped by the addition of SDS sample buffer, and the samples were

immediately heated with SDS-sample buffer at 98°C for 3 min and subjected to

immunoblotting assay. Recombinant enzymes were purchased from the following

manufacturers: human USP2 catalytic domain, CF (R&D Systems), human His6-USP7

Protein, CF (R&D Systems), GST-tagged human SENP2 catalytic domain (Enzo), and mouse

active UCHL1 (ProSpec).

Human PD brain

The use of postmortem brain tissue in this study was approved by the Institutional Review

Boards of Kyoto Prefectural University of Medicine (approval no. ERB-C-1846-2). Written

informed consent for cadaver surgical training and the use of tissue samples for medical

research was obtained, both from patients themselves and their next of kin. The postmortem

brain specimen was obtained from a patient with PD (age: 85 years, female), fixed using

Thiel's method, with slight modifications. Briefly, a fixative containing 5% formalin was

perfused from the superior sagittal sinus. The brain was removed from the skull, immersed in

3.7% formalin for 14 days, and dissected into tissue blocks, including the substantia nigra

(SN). Tissue blocks were further fixed with 4% PFA in 0.1 M phosphate buffer (PB) for 2

days. After fixation, these tissue blocks were subjected to immunohistochemistry as described

below.

PFF injection and inhibitor infusion *in vivo*

To reproduce the propagation of LB-like pathology *in vivo*, WT mice were anesthetized using

a midazolam/medetomidine/butorphanol cocktail (4, 0.3, and 5 mg/kg, respectively), and 2

µL PFFs (7 mg/mL) was stereotaxically injected into the striatum, as reported previously[8].

A needle inserted into the left forebrain (+0.98 mm relative to Bregma, +1.70 mm from

midline) was used to direct the inoculum into the striatum (+3.5 mm beneath the dura) and

was injected using a 10-µL Hamilton syringe at a flow rate of 0.2 µL/min (total 14 µg PFFs

per site), with the needle left in place for >20 min.

The inhibitor (NSC 632839, final concentration 73.5 µM in sterile saline) was infused into

the left lateral ventricle using an Alzet osmotic pump (Model 2006) connected to a brain

infusion kit (Type 3) in accordance with the manufacturer's instructions (Alzet) for

controlled-release of the reagent (0.15 µL/h for 42 days). Subjects were monitored regularly

following their postoperative recovery, and sacrificed at 6 weeks after the surgery. The

efficiency of the aggregate formation and inhibitory effect of NSC632839 on aggregate

formation was determined using immunohistochemistry. Data from NSC635839-infused mice

(n=5) and DMSO-infused control mice (n=3) were included in the statistical analysis as

described below.

To study the effect of endogenous αSyn expression on aggregate formation, we injected

PFF into the striatum of WT and αSyn-KO mice (S6 Fig.), and the PFF injections were

690   performed as described above.

Immunohistochemistry and Hematoxylin and Eosin (HE) staining

Unless specifically mentioned, a standard immunohistological procedure was performed on

free-floating sections prepared from mouse and human brains.

For mouse brain, immunostaining was performed as previously described[38]. Briefly,

under deep anesthesia, adult male mice were intracardially perfused with PBS followed by

4% PFA in PBS. Brains were dissected into blocks and postfixed using the same fixative for

12 h at 4°C. Coronal sections (40 μm) were obtained using a vibratome (Dosaka-EM),

permeabilized with 0.3% Triton X-100 in PBS (PBST) for 1 h, and blocked with 10% NGS in

PBST for 20 h. These sections were then incubated with the primary antibody in the blocking

solution for 20 h, washed with PBST, and treated with a secondary antibody. For double

immunostaining, this staining procedure was repeated. For the detection of phosphorylated

αSyn, two types of antibodies were used: monoclonal mouse IgG antibody (#64) and

polyclonal rabbit IgG antibody (pS129 αSyn).

For the immunofluorescent method, the primary antibodies were detected with Alexa488-

and Alexa594-conjugated secondary antibodies (Thermo Fischer Scientific). Sections were

washed with PBST and then with 20 mM Tris-HCl buffer, and mounted with FluorSave

(Merck Millipore). For confocal observation, images were acquired as Z stacks (10–20

z-sections, 1-μm apart, 1024 × 1024 pixels) using a Plan-Apochromat 63x/1.40 Oil DIC

objective (Carl Zeiss) with an inverted laser-scanning confocal microscope, LSM900 (Carl

Zeiss).

For the immunoenzyme method, the primary antibody was detected by biotinylated

anti-rabbit IgG antibody (1:500; Vector Laboratory) for 4 h, followed by horseradish

peroxidase (HRP)-conjugated streptavidin (1:2; Nichirei Biosciences Inc.) detection for 1 h.

HRP was visualized with 3,3'-diaminobezidine tetrahydrochloride and H_2O_2 in 50 mM

Tris-buffered saline (pH 7.4). Immunosignals were acquired through the UPlanFl 60×

objective with an inverted microscope (IX71), equipped with a cooled charge-coupled device

camera (Olympus).

For the human PD brain, the dissected tissue block including the SN was cryoprotected

using 20% sucrose in 0.1 M PB (pH 7.4) and rapidly frozen in Tissue-Tek OCT compound

(Sakura Finetek Japan). Sections were obtained using a cryostat (Leica) at 30-μm thickness.

For HE staining, sections were stained on a slide glass using a Hematoxylin and Eosin Stain

Kit according to the manufacturer's standard protocol (ScyTek Laboratories). For

immunostaining with antigen retrieval, sections were immersed in formic acid for 10 min,

washed 4 times with PBS, and subjected to immunohistochemical analysis using

anti-phosphorylated αSyn (#64) and anti-SENP2 antibodies with the immunostaining

procedure described for mouse brain sections.

Quantification of the number of cell bodies harbouring pathological aggregates of phosphorylated αSyn

The number of cell bodies harbouring aggregates of phosphorylated αSyn in the amygdala and piriform cortex of NSC632839-treated and DMSO-treated control mouse brains was counted. For cell-counting in each region, 3–6 sections per animal for each treatment were subjected to the cell-counting procedure. The number of animals used in each experimental group is indicated in the figure legends (Fig. 5F, G). After the cell-counting procedure, the mean cell numbers per animal were individually calculated in each brain region; the mean values per experimental group were then used in the statistical analysis using GraphPad Prism 5 by the two-tailed, unpaired *t*-test, with or without Welch's correction. Data are expressed as the mean ± SEM. The significance levels were set at *$p < 0.05$.

Image processing

The digital images obtained on the LSM900 or IX71 were processed using Adobe Photoshop 7.0 (Adobe). Each figure set was prepared in Microsoft PowerPoint (Microsoft).

Statistical analysis and reproducibility

Statistical analyses between the two experimental groups were performed using the

two-tailed, unpaired *t*-test, with or without Welch's correction (GraphPad Prism 5, GraphPad

Software Inc.). The number of animals used in each experiment and the number of images

that were analysed are specified in the Methods and/or indicated in each figure legend. Data

are expressed as the mean ± standard error of the mean. The significance levels were set at *p*

< 0.05 (*) and $p < 0.01$ (**), as indicated. Reproducibility was confirmed by independent

experiments that were repeated at least two to three times, with similar results.

Resource availability

The experimental materials and data resources used in this study are available from the

corresponding authors, MT and KT, upon reasonable request.

Data and code availability

Mass spectrometric and proteomic analyses

The raw files of the mass spectrometric analysis were used to conduct searches against the

mouse and human αSyn amino acid sequences that were obtained from the Mus musculus

dataset (Uniprot Proteome UP000000589 2021_03_07 downloaded), Homo sapiens dataset

(Uniprot Proteome UP000005640 2022_05_25 downloaded), and the common Repository of

Adventitious Proteins (cRAP; ftp://ftp.thegpm.org/fasta/cRAP). The search was conducted by

using MASCOT version 2.6 (Matrix Science) via Proteome Discoverer 2.4 (Thermo Fisher

Scientific).

Based on the results of mass spectrometric analysis, the candidate enzymes which process

the N-terminal side of lysine residues located at the N-terminal repeat regions of αSyn were

screened using the Peptidase database, MEROPS (https://www.ebi.ac.uk/merops/).

All the remaining research data are available in this article and its Supplementary

Information. Furthermore, the source data have been provided with this paper.

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Author contributions

782 KT and MT conceived and designed the experiments. KT performed the experiments. YW

783 contributed to the analysis and discussion. KT and MT wrote the manuscript.

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Declaration of interests

The authors have no potential or actual conflicts of interest.

References

1.   Spillantini MG, Crowther RA, Jakes R, Hasegawa M, Goedert M. alpha-Synuclein

in filamentous inclusions of Lewy bodies from Parkinson's disease and dementia with lewy

bodies. Proc Natl Acad Sci U S A. 1998;95(11):6469-73. doi: 10.1073/pnas.95.11.6469.

PubMed PMID: 9600990; PubMed Central PMCID: PMCPMC27806.

2.   Dickson DW. Alpha-synuclein and the Lewy body disorders. Curr Opin Neurol.

2001;14(4):423-32. doi: 10.1097/00019052-200108000-00001. PubMed PMID: 11470957.

3.   Stefanis L. alpha-Synuclein in Parkinson's disease. Cold Spring Harb Perspect Med.

2012;2(2):a009399. doi: 10.1101/cshperspect.a009399. PubMed PMID: 22355802; PubMed

Central PMCID: PMCPMC3281589.

4.   Braak H, Del Tredici K. Neuroanatomy and pathology of sporadic Parkinson's

disease. Adv Anat Embryol Cell Biol. 2009;201:1-119. PubMed PMID: 19230552.

5.   Braak H, Del Tredici K, Rub U, de Vos RA, Jansen Steur EN, Braak E. Staging of

brain   pathology   related   to   sporadic   Parkinson's   disease.   Neurobiol   Aging.

2003;24(2):197-211. doi: 10.1016/s0197-4580(02)00065-9. PubMed PMID: 12498954.

6.   Li JY, Englund E, Holton JL, Soulet D, Hagell P, Lees AJ, et al. Lewy bodies in

grafted neurons in subjects with Parkinson's disease suggest host-to-graft disease propagation.

Nat   Med.   2008;14(5):501-3.   Epub   20080406.   doi: 10.1038/nm1746.   PubMed   PMID:

18391963.

7.   Volpicelli-Daley L, Brundin P. Prion-like propagation of pathology in Parkinson

disease. Handb Clin Neurol. 2018;153:321-35. doi: 10.1016/B978-0-444-63945-5.00017-9.

PubMed PMID: 29887143; PubMed Central PMCID: PMCPMC6625652.

8.   Luk KC, Kehm V, Carroll J, Zhang B, O'Brien P, Trojanowski JQ, et al. Pathological

alpha-synuclein transmission initiates Parkinson-like neurodegeneration in nontransgenic

mice. Science. 2012;338(6109):949-53. doi: 10.1126/science.1227157. PubMed PMID:

23161999; PubMed Central PMCID: PMCPMC3552321.

9.   Peng C, Gathagan RJ, Covell DJ, Medellin C, Stieber A, Robinson JL, et al. Cellular

milieu imparts distinct pathological alpha-synuclein strains in alpha-synucleinopathies.

Nature. 2018;557(7706):558-63. Epub 20180509. doi: 10.1038/s41586-018-0104-4. PubMed

PMID: 29743672; PubMed Central PMCID: PMCPMC5970994.

10.   Henderson MX, Cornblath EJ, Darwich A, Zhang B, Brown H, Gathagan RJ, et al.

Spread of alpha-synuclein pathology through the brain connectome is modulated by selective

vulnerability and predicted by network analysis. Nat Neurosci. 2019;22(8):1248-57. Epub

20190725. doi: 10.1038/s41593-019-0457-5. PubMed PMID: 31346295; PubMed Central

PMCID: PMCPMC6662627.

11.   Volpicelli-Daley LA, Luk KC, Patel TP, Tanik SA, Riddle DM, Stieber A, et al.

Exogenous alpha-synuclein fibrils induce Lewy body pathology leading to synaptic

dysfunction and neuron death. Neuron. 2011;72(1):57-71. doi: 10.1016/j.neuron.2011.08.033.

PubMed PMID: 21982369; PubMed Central PMCID: PMCPMC3204802.

12.   Suthar SK, Lee SY. Truncation or proteolysis of alpha-synuclein in Parkinsonism.

Ageing Res Rev. 2023;90:101978. Epub 20230605. doi: 10.1016/j.arr.2023.101978. PubMed

PMID: 37286088.

13.   Canever JB, Soares ES, de Avelar NCP, Cimarosti HI. Targeting alpha-synuclein

post-translational modifications in Parkinson's disease. Behav Brain Res. 2023;439:114204.

Epub 20221111. doi: 10.1016/j.bbr.2022.114204. PubMed PMID: 36372243.

14.   Fujiwara H, Hasegawa M, Dohmae N, Kawashima A, Masliah E, Goldberg MS, et al.

alpha-Synuclein is phosphorylated in synucleinopathy lesions. Nat Cell Biol. 2002;4(2):160-4.

doi: 10.1038/ncb748. PubMed PMID: 11813001.

15.   Rott R, Szargel R, Shani V, Hamza H, Savyon M, Abd Elghani F, et al.

SUMOylation and ubiquitination reciprocally regulate alpha-synuclein degradation and

pathological aggregation. Proc Natl Acad Sci U S A. 2017;114(50):13176-81. Epub 20171127.

doi: 10.1073/pnas.1704351114. PubMed PMID: 29180403; PubMed Central PMCID:

PMCPMC5740625.

16.   Savyon M, Engelender S. SUMOylation in alpha-Synuclein Homeostasis and

Pathology. Front Aging Neurosci. 2020;12:167. Epub 20200625. doi:

10.3389/fnagi.2020.00167. PubMed PMID: 32670048; PubMed Central PMCID:

PMCPMC7330056.

17.   Terada M, Suzuki G, Nonaka T, Kametani F, Tamaoka A, Hasegawa M. The effect of

truncation on prion-like properties of alpha-synuclein. J Biol Chem. 2018;293(36):13910-20.

Epub 20180720. doi: 10.1074/jbc.RA118.001862. PubMed PMID: 30030380; PubMed

Central PMCID: PMCPMC6130941.

18.   Sorrentino ZA, Giasson BI. The emerging role of alpha-synuclein truncation in aggregation and disease. J Biol Chem. 2020;295(30):10224-44. Epub 20200518. doi: 10.1074/jbc.REV120.011743. PubMed PMID: 32424039; PubMed Central PMCID: PMCPMC7383394.

19.   McGlinchey RP, Ni X, Shadish JA, Jiang J, Lee JC. The N terminus of alpha-synuclein dictates fibril formation. Proc Natl Acad Sci U S A. 2021;118(35). doi: 10.1073/pnas.2023487118. PubMed PMID: 34452994; PubMed Central PMCID: PMCPMC8536336.

20.   Legname G, Nguyen HO, Peretz D, Cohen FE, DeArmond SJ, Prusiner SB. Continuum of prion protein structures enciphers a multitude of prion isolate-specified phenotypes. Proc Natl Acad Sci U S A. 2006;103(50):19105-10. Epub 20061201. doi: 10.1073/pnas.0608970103. PubMed PMID: 17142317; PubMed Central PMCID: PMCPMC1748184.

21.   Tsujimura A, Taguchi K, Watanabe Y, Tatebe H, Tokuda T, Mizuno T, et al. Lysosomal enzyme cathepsin B enhances the aggregate forming activity of exogenous alpha-synuclein fibrils. Neurobiol Dis. 2015;73:244-53. Epub 20141022. doi: 10.1016/j.nbd.2014.10.011. PubMed PMID: 25466281.

22.   Mishizen-Eberz AJ, Guttmann RP, Giasson BI, Day GA, 3rd, Hodara R,

Ischiropoulos H, et al. Distinct cleavage patterns of normal and pathologic forms of

alpha-synuclein by calpain I in vitro. J Neurochem. 2003;86(4):836-47. doi:

10.1046/j.1471-4159.2003.01878.x. PubMed PMID: 12887682.

23.   Dufty BM, Warner LR, Hou ST, Jiang SX, Gomez-Isla T, Leenhouts KM, et al.

Calpain-cleavage of alpha-synuclein: connecting proteolytic processing to disease-linked

aggregation. Am J Pathol. 2007;170(5):1725-38. doi: 10.2353/ajpath.2007.061232. PubMed

PMID: 17456777; PubMed Central PMCID: PMCPMC1854966.

24.   Liu CW, Giasson BI, Lewis KA, Lee VM, Demartino GN, Thomas PJ. A

precipitating role for truncated alpha-synuclein and the proteasome in alpha-synuclein

aggregation: implications for pathogenesis of Parkinson disease. J Biol Chem.

2005;280(24):22670-8. Epub 20050419. doi: 10.1074/jbc.M501508200. PubMed PMID:

15840579.

25.   Yan C, Huo H, Yang C, Zhang T, Chu Y, Liu Y. Ubiquitin C-Terminal Hydrolase L1

regulates autophagy by inhibiting autophagosome formation through its deubiquitinating

enzyme activity. Biochem Biophys Res Commun. 2018;497(2):726-33. Epub 20180217. doi:

10.1016/j.bbrc.2018.02.140. PubMed PMID: 29462615.

26.   Nicholson B, Leach CA, Goldenberg SJ, Francis DM, Kodrasov MP, Tian X, et al.

Characterization of ubiquitin and ubiquitin-like-protein isopeptidase activities. Protein Sci.

2008;17(6):1035-43. Epub 20080418. doi: 10.1110/ps.083450408. PubMed PMID:

18424514; PubMed Central PMCID: PMCPMC2386736.

27.   Cubenas-Potts C, Goeres JD, Matunis MJ. SENP1 and SENP2 affect spatial and

temporal control of sumoylation in mitosis. Mol Biol Cell. 2013;24(22):3483-95. Epub

20130918. doi: 10.1091/mbc.E13-05-0230. PubMed PMID: 24048451; PubMed Central

PMCID: PMCPMC3826987.

28.   Kovacs GG, Wagner U, Dumont B, Pikkarainen M, Osman AA, Streichenberger N,

et al. An antibody with high reactivity for disease-associated alpha-synuclein reveals

extensive brain pathology. Acta Neuropathol. 2012;124(1):37-50. Epub 20120228. doi:

10.1007/s00401-012-0964-x. PubMed PMID: 22370907.

29.   Vaikath NN, Majbour NK, Paleologou KE, Ardah MT, van Dam E, van de Berg WD,

et al. Generation and characterization of novel conformation-specific monoclonal antibodies

for alpha-synuclein pathology. Neurobiol Dis. 2015;79:81-99. Epub 20150430. doi:

10.1016/j.nbd.2015.04.009. PubMed PMID: 25937088.

30.   Guo C, Henley JM. Wrestling with stress: roles of protein SUMOylation and

deSUMOylation in cell stress response. IUBMB Life. 2014;66(2):71-7. Epub 20140127. doi:

10.1002/iub.1244. PubMed PMID: 24470405.

31.   Kessler JC, Rochet JC, Lansbury PT, Jr. The N-terminal repeat domain of

alpha-synuclein inhibits beta-sheet and amyloid fibril formation. Biochemistry.

2003;42(3):672-8. doi: 10.1021/bi020429y. PubMed PMID: 12534279.

32.   Qin Z, Hu D, Han S, Hong DP, Fink AL. Role of different regions of alpha-synuclein

in the assembly of fibrils. Biochemistry. 2007;46(46):13322-30. Epub 20071027. doi:

10.1021/bi7014053. PubMed PMID: 17963364.

33.   Zhang S, Zhu R, Pan B, Xu H, Olufemi MF, Gathagan RJ, et al. Post-translational

modifications of soluble alpha-synuclein regulate the amplification of pathological

alpha-synuclein. Nat Neurosci. 2023;26(2):213-25. Epub 20230123. doi:

10.1038/s41593-022-01239-7. PubMed PMID: 36690898; PubMed Central PMCID:

PMCPMC10103650.

34.   Abeliovich A, Schmitz Y, Farinas I, Choi-Lundberg D, Ho WH, Castillo PE, et al.

Mice lacking alpha-synuclein display functional deficits in the nigrostriatal dopamine system.

Neuron. 2000;25(1):239-52. doi: 10.1016/s0896-6273(00)80886-7. PubMed PMID:

10707987.

35.   Taguchi K, Watanabe Y, Tsujimura A, Tatebe H, Miyata S, Tokuda T, et al.

Differential expression of alpha-synuclein in hippocampal neurons. PLoS One.

2014;9(2):e89327. Epub 20140225. doi: 10.1371/journal.pone.0089327. PubMed PMID:

24586691; PubMed Central PMCID: PMCPMC3934906.

36.   Watanabe Y, Tatebe H, Taguchi K, Endo Y, Tokuda T, Mizuno T, et al.

p62/SQSTM1-dependent autophagy of Lewy body-like alpha-synuclein inclusions. PLoS

One. 2012;7(12):e52868. Epub 20121231. doi: 10.1371/journal.pone.0052868. PubMed

PMID: 23300799; PubMed Central PMCID: PMCPMC3534125.

37.   Mao X, Ou MT, Karuppagounder SS, Kam TI, Yin X, Xiong Y, et al. Pathological

alpha-synuclein transmission initiated by binding lymphocyte-activation gene 3. Science.

2016;353(6307). doi: 10.1126/science.aah3374. PubMed PMID: 27708076; PubMed Central

PMCID: PMCPMC5510615.

38.   Taguchi K, Watanabe Y, Tsujimura A, Tanaka M. Brain region-dependent differential

expression of alpha-synuclein. J Comp Neurol. 2016;524(6):1236-58. Epub 20150929. doi:

10.1002/cne.23901. PubMed PMID: 26358191.

Figure legends for Supporting Information

S1 Fig. Isolation and characterization of pathological seeds using the batch method. (A)

Schematic diagram of fractionation by sucrose-density gradient ultracentrifugation of the

conditioned culture medium derived from pathological primary neurons harbouring Lewy

body- and Lewy neurite-like aggregates. CM, culture medium. **(B)** SDS-PAGE and an

immunoblotting assay of the fractions using anti-αSyn antibody. The numbers indicate the

recovered fractions shown in **(A)**. **(C)** Native-PAGE and immunoblotting assay of the

fractions. **(D)** The seeding ability of the pathological seeds recovered in Fraction 3. pS-αSyn,

phosphorylated αSyn. **(E)** Standard curve for HPLC (Log. molecular weight vs. elution

volume). **(F)** Immunoelectron microscopic observation of the αSyn-oligomer species

recovered in each HPLC-prepared fraction. **(G)** Immunoelectron microscopic observation of

PFFs as a control for αSyn immunolabeling. PFFs, preformed fibrils. Scale bar: 20 μm in **(D)**,

50 nm in **(F)**, and 100 nm in **(G)**.

S2 Fig. Development of the microfluidic device for large-scale preparation of

pathological seeds. (A–C) Overview of microfluidic device for large-scale preparation of

seeds. **(D, E)** Primary neuronal culture using the large-scale microfluidic device for 5 days

(D) and 12 days **(E)** *in vitro*. Arrowheads show the extending axons. The region marked by a

red square in **(D)** and **(E)** is magnified in each right panel, respectively. GC, growth cone.

Scale bar: 100 μm.

S3 Fig. Image analysis of the inhibitory effect of NSC632839 on aggregate formation.

Phosphorylated-αSyn immunoreactive area, excluding the DAPI-positive area surrounded by

dotted lines, was quantitatively analysed because the monoclonal antibody against

phosphorylated αSyn #64 tends to react with the nuclei. Arrowheads indicate examples of

phosphorylated αSyn-positive aggregates. Scale bar: 20 μm.

S4 Fig. Screening assay for possible αSyn-processing enzymes. (A–C) αSyn-processing

ability of possible NSC632839-inhibited enzymes was examined by a protease assay using

recombinant proteins. As indicated by asterisks, cleaved fragments were not detected in the

presence of USP-7 **(A)**, USP-2 **(B)**, or UCHL1 **(C)**. Both PFFs and a monomeric form of

αSyn were not processed as appropriate substrates by these enzymes. The reaction time and

temperature are indicated in each figure panel.

S5 Fig. Presence of SENP2 in Lewy pathology. (A, **B)** Lewy body, Lewy neurites, and

neuromelanin observed in the substantia nigra pars compacta of the PD-brain. **(C, D)**

Presence of endogenous SENP2 in amorphous aggregates **(C)**, but not in a typical Lewy body

(D). Arrowheads indicate the presence of both SENP2 and phosphorylated αSyn. LB, Lewy

body; LN, Lewy neurites; NM, neuromelanin. Scale bar: 20 μm in **(A, B)**, and 10 μm in **(C,**

D).

S6 Fig. Endogenous αSyn expression is indispensable for aggregate formation. (A)

Schematic diagram of PFF injection into the striatum of the mouse brain. C, caudal; R,

rostral; Str, striatum. **(B)** Intense formation of phosphorylated-αSyn aggregates was observed

in the WT, but not αSyn-KO, mouse brain. WT, wildtype; KO, knockout. Scale bar: 50 μm.

A

Medium volume

C1 C2 C3

↑
PFFs

B

pS-αSyn Pan-αSyn

C2 area

C3 area

20 μm

C

Frac. No.

CM 1
    2
    3
    4
0.8 5
    6
    7
    8
1.2 9
Suc. (M) 10

D

1 2 3 4 5 6 7

kDa
20 ← a
    ← b
15
10

IB: αSyn

E

kDa
669
440
140 ←
66

IB: αSyn

F

16.43
[nm]

1.00 μm   0.00
2.00 x 2.00 μm

G

100 nm

H

Pan-αSyn

I

O2 / pan-αSyn

J

5G4 / pan-αSyn

Bars: 50 nm

A

Transferred seed-containing culture medium

↓ Concentration using centrifugal filter

↓ Sucrose density gradient ultracentrifugation

Seed-containing fraction

↓ Immunoprecipitation using anti-αSyn antibody

SDS-PAGE (silver staining)

B

Mass sample | Antibody only | PFFs

kDa: 250, 150, 100, 75, 50, 37, 25, 20, 15, 10

←Heavy chain

←αSyn

C

Human αSyn

Repeat 1   Repeat 2   Repeat 3   Repeat 4   Repeat 5

¹MDVFMKGLS **K ¹¹AKEGV** VAAAE **²¹KTKQGV** AEAA ³¹G **KTKEGV** LYV ⁴¹GS **KTKEGV** VH ⁵¹GVATVAE **KTK ⁶¹EQV** TNVGGAV ⁷¹VTGVT**AV**AQK ⁸¹TVEGAGSIAA ⁹¹ATGFVKKDQL ¹⁰¹GKNEEGAPQE ¹¹¹GILEDMPVDP ¹²¹DNEAYEMPSE ¹³¹EGYQDYEPEA

Mouse αSyn

Repeat 1   Repeat 2   Repeat 3   Repeat 4   Repeat 5

¹MDVFMKGLS **K ¹¹AKEGV** VAAAE **²¹KTKQGV** AEAA ³¹G **KTKEGV** LYV ⁴¹GS **KTKEGV** VH ⁵¹GVTTVAE **KTK ⁶¹EQV** TNVGGAV ⁷¹VTGVT**AV**AQK ⁸¹TVEGAGNIAA ⁹¹ATGFVKKDQM ¹⁰¹GKGEEGYPQE ¹¹¹GILEDMPV**DP** ¹²¹GSEAYEMPSE ¹³¹EGYQDYEPEA

KX|KXGV repeat region

↑ Conserved cleavage site

D

Repeat region

1 — 65   NAC   95 — 140

11|12 A|K

22|23 T|K

33|34 T|K

44|45 T|K

76|77 A|V

119|120 D|P

Cleavage sites detected by MS analysis

11|12 G|A | 22|23 A|A | 33|34 A|A | 44|45 A|A   Mutations inserted in repeat region

A DMSO **B** NSC 632839

pS-αSyn

C **D**

20 µm

E

pS-αSyn-positive area

DMSO   NSC 632839
**

F

DAPI-positive area

DMSO   NSC 632839

G

Substrate   PFFs   Mono
GST-SENP2   −   +   −   +

kDa

IB: αSyn

H

GST-SENP2   −

kDa

*

IB: αSyn

I

GST-SENP2   −

kDa

CBB

J

Substrate   WT-αSyn   Mut-αSyn
GST-SENP2   −   +   −   +

kDa

IB: αSyn

A    SENP2       pS-αSyn      Merged

B    SENP2       pS-αSyn      Merged

5 µm

A

PFFs in Str

R

LV
DMSO or NSC 632839

C

B

	Striatum (injection site)	Piriform cortex	Amygdala

pS-αSyn

DMSO

NSC632839

100 μm

C

Piriform cortex

Number of cell bodies

DMSO   NSC632839

*

D

Amygdala

Number of cell bodies

DMSO   NSC632839

*