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A novel mouse model for investigating α -synuclein aggregates in oligodendrocytes: implications for the glial cytoplasmic inclusions in multiple system atrophy

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Abstract

The aggregated alpha-synuclein (asyn) in oligodendrocytes (OLGs) is one of the pathological hallmarks in multiple system atrophy (MSA). We have previously reported that asyn accumulates not only in neurons but also in OLGs long after the administration of asyn preformed fibrils (PFFs) in mice. However, detailed spatial and temporal analysis of oligodendroglial asyn aggregates was technically difficult due to the background neuronal asyn aggregates. The aim of this study is to create a novel mouse that easily enables sensitive and specific detection of asyn aggregates in OLGs and the comparable analysis of the cellular tropism of asyn aggregates in MSA brains. To this end, we generated transgenic (Tg) mice expressing human asyn-green fluorescent protein (GFP) fusion proteins in OLGs under the control of the 2', 3'-cyclic nucleotide 3'-phosphodiesterase (CNP) promoter (CNP-SNCAGFP Tg mice). Injection of asyn PFFs in these mice induced distinct GFP-positive aggregates in the processes of OLGs as early as one month post-inoculation (mpi), and their number and size increased in a centripetal manner. Moreover, MSA-brain homogenates (BH) induced significantly more oligodendroglial asyn aggregates than neuronal asyn aggregates compared to DLB-BH in CNP-SNCAGFP Tg mice, suggestive of their potential tropism of asyn seeds for OLGs. In conclusion, CNP-SNCAGFP Tg mice are useful for studying the development and tropism of asyn aggregates in OLGs and could contribute to the development of therapeutics targeting asyn aggregates in OLGs.

Keywords Multiple system atrophy, MSA, α -synuclein, Glial cytoplasmic inclusions, GCI, Mouse model, Cellular tropism, Propagation, Prion hypothesis

Introduction

Multiple system atrophy (MSA) is a neurodegenerative disease that is clinically characterized by various symptoms, including parkinsonism, cerebellar ataxia, and autonomic dysfunction [1, 2]. The main pathological findings of MSA are the presence of alpha-synuclein (α syn) aggregates in oligodendrocytes (OLGs) called glial cytoplasmic inclusions (GCIs) and neuronal loss, but neuronal cytoplasmic inclusions (NCIs) are also observed [3, 4]. α syn aggregates in MSA are mainly formed in OLGs, but almost exclusively formed in neurons in Parkinson's

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disease (PD) and dementia with Lewy bodies (DLB) [5–13]. However, the mechanism by which α syn preferentially accumulates in OLGs in MSA remains unclear, even though the expression level of α syn in OLGs is much lower than that in neurons [14–17].

There is growing evidence suggesting that α syn has a prion-like property. The injection of α syn preformed fibrils (PFFs) into mouse brains resulted in the propagation of α syn in neurons [18–23], and injection of MSA brain lysates into mouse brains also caused the propagation of α syn exclusively in neurons, but not in OLGs [24, 25]. This result further deepened the mystery of how α syn accumulates in OLGs in MSA.

We previously reported that α syn accumulates not only in neurons but also in OLGs following long-term incubation (at least 7 months) in mice inoculated with α syn PFFs [26]. However, the process of early α syn aggregate formation in OLGs was difficult to be evaluated with the phosphorylated α syn antibody due to the overwhelming background signal generated by α syn aggregates in neurons.

The aim of this study was to create a mouse model that enables the sensitive and specific detection of α syn aggregates in OLGs and the analysis of their preferential accumulation in OLGs. To the best of our knowledge, this is the first study to precisely analyze the spatial and temporal development of oligodendroglial α syn aggregates in vivo.

Results

Oligodendrocyte-specific α synGFP expression in CNP-SNCAGFP Tg mice

We created transgenic (Tg) mice expressing human α syn-GFP fusion proteins in OLGs by the murine CNP promoter (CNP-SNCAGFP Tg mice) (Fig. 1A). The CNP promoter was employed because α syn has been reported to accumulate in mature OLGs as well as oligodendrocyte precursor cells (OPCs) in MSA, and CNP is known to be expressed in OPCs [27, 28]. OPCs were also reported to have a greater ability to incorporate α syn aggregates compared to mature OLGs and contribute to the development of GCIs [29]. In addition to the endogenous α syn expression, the human α syn-GFP fusion protein was expressed in CNP-SNCAGFP Tg mice (Fig. 1B). A histological study showed that α synGFP was robustly expressed in the corpus callosum, moderately expressed in the striatum and white matter of the cerebral cortex, and mildly expressed in the gray matter of the cerebral cortex (Fig. 1C). To examine the cells expressing α synGFP in CNP-SNCAGFP Tg mice, immunofluorescent staining with antibodies against markers of OLGs, neurons, astrocytes, and microglia was performed. GFP

co-localized with Olig2, CNP, TPPP, and GSTpi, but not with NeuN, GFAP, or Iba1, confirming that α synGFP is only expressed in OLGs (Fig. 1D). α synGFP was observed mainly in the cell bodies and also in the process of OLGs in CNP-SNCAGFP Tg mice (Fig. 1D). Note that the subcellular distribution of α synGFP in CNP-SNCAGFP Tg mice was the same to that of human α syn in CNP-SNCA Tg mice (Additional file 1: Fig. S1A). Furthermore, the phosphorylation or aggregation status of α synGFP did not change with aging (Additional file 1: Fig. S1B, Additional file 2: Fig. S2C left panel), and these mice did not show apparent motor impairment even at 12 months of age (Additional file 1: Fig. S1C).

Highly sensitive detection of α syn aggregates in OLGs in CNP-SNCAGFP Tg mice

We have previously shown that in mice inoculated with α syn PFFs, the accumulation of α syn in OLGs was observed long after inoculation [26]. In the present study, we tested whether the temporal development of α syn aggregates in OLGs could be monitored by the GFP signals in Tg mice according to the same paradigm.

Electron microscopy confirmed that α syn PFFs were fragmented to about 20–50 nm by sonication (Fig. 2A). α syn PFFs or PBS were inoculated in the left dorsal striatum of CNP-SNCAGFP Tg mice and histological analyses were performed at 1, 4, and 12 months after inoculation (Fig. 2B). Numerous phosphorylated α syn ($p\alpha$ syn)-positive aggregates were observed as early as 1-month post-inoculation (mpi), and these thread-like structures developed into round-shaped cytoplasmic inclusions (Fig. 2C). Although most of the $p\alpha$ syn-positive aggregates were colocalized with neuronal markers and of neuronal origin, $p\alpha$ syn and GFP positive α syn aggregates in OLGs were also identified. However, weak and rare $p\alpha$ syn-positive signals in OLGs were buried in strong background signals from neuronal aggregates in most of the brain regions, except for the corpus callosum where OLGs outnumber neurons (Fig. 2D, Additional file 3: Fig. S3).

To confirm that α synGFP aggregates, which we call GFP dots, were of oligodendroglial origin, immunofluorescent staining with antibodies against various cell-specific markers was performed. GFP dots colocalized with CNP and GSTpi, but not NeuN, GFAP, or Iba1, confirming that GFP dots were formed specifically in OLGs, and not in other cell types by the leakage or transfer of α synGFP from OLGs (Fig. 2E). GFP dots were immunopositive for ubiquitin (Ub), p62/sequestosome-1 (p62), aggregated form of α syn, and were PK resistant, suggesting that these are pathological aggregates (Additional file 2: Fig. S2).

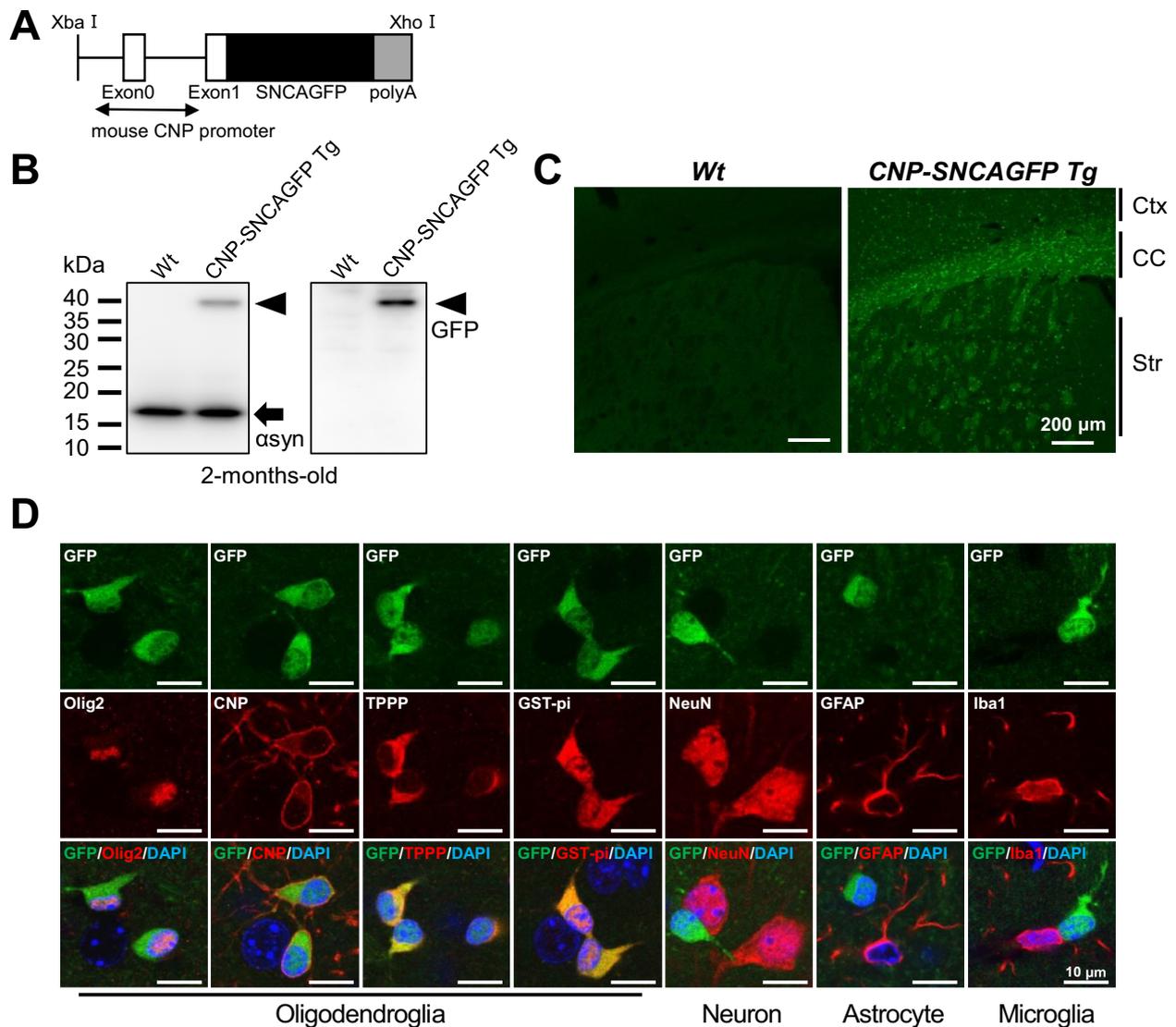


Fig. 1 The expression of human asynGFP in CNP-SNCAGFP transgenic (Tg) mice. **A** The transgenic construct. **B** Western blot images of whole-brain SDS-soluble fraction with anti-asyn and GFP antibodies in both genotypes (2 months old). The expression of endogenous asyn (arrow) and transgenic human asynGFP fusion protein (arrowhead) is shown. **C** Fluorescence microscopy showed the asynGFP expression in in both genotypes (2 months old). Scale bar = 200 µm. **D** Immunofluorescent staining of various cellular makers with asynGFP in the striatum of Tg mice (2 months old). The merged images include DAPI (blue). GFP signals were observed exclusively in OLGs. Note that anti-CNP antibodies strongly labeled the plasma membrane in OLGs. Scale bar = 10 µm. Wt, wild-type; asyn, alpha-synuclein; Ctx, cortex; CC, corpus callosum; Str, striatum

(See figure on next page.)

Fig. 2 The development of pasyn and asynGFP aggregates in CNP-SNCAGFP Tg mice inoculated with asyn PFFs. **A** Image of sonicated asyn PFFs used for injection in electron microscopy. **B** The schedule of asyn PFF injection and histological evaluation. Tg mice (age: 2–3 months) were inoculated with asyn PFFs or PBS, sacrificed at 1, 4, and 12 mpi, and histological examinations were performed ($n=6$). **C** Immunohistochemical staining for pasyn at the ipsilateral dorsal striatum of Tg mice inoculated with asyn PFFs or PBS. In Tg mice, numerous pasyn-positive neurites were observed as early as 1 mpi, and these thread-like structures grew into round-shaped cytoplasmic inclusions. Only a few pasyn-positive cells were observed in Tg mice inoculated with PBS. Scale bar = 50 µm. **D** Fluorescence micrographs of GFP and immunostaining with anti-pasyn (red) and anti-NeuN (gray) antibodies in the striatum of Tg mice. The merged images include DAPI (blue). Unaggregated asynGFP is denoted by dotted circles. Most of the pasyn-positive signals are of neuronal origin (arrowheads), and rarely merged with asynGFP aggregates (GFP dots) in OLGs (arrows). Scale bar = 20 µm. **E** Immunofluorescence micrographs of various cellular makers with GFP in Tg mice at 6 mpi of asyn PFFs. asynGFP aggregates were observed specifically in the OLGs. The merged images include DAPI (blue). Scale bar = 10 µm. mpi, month(s) post-inoculation; PFFs, preformed fibrils; pasyn, phosphorylated α -synuclein

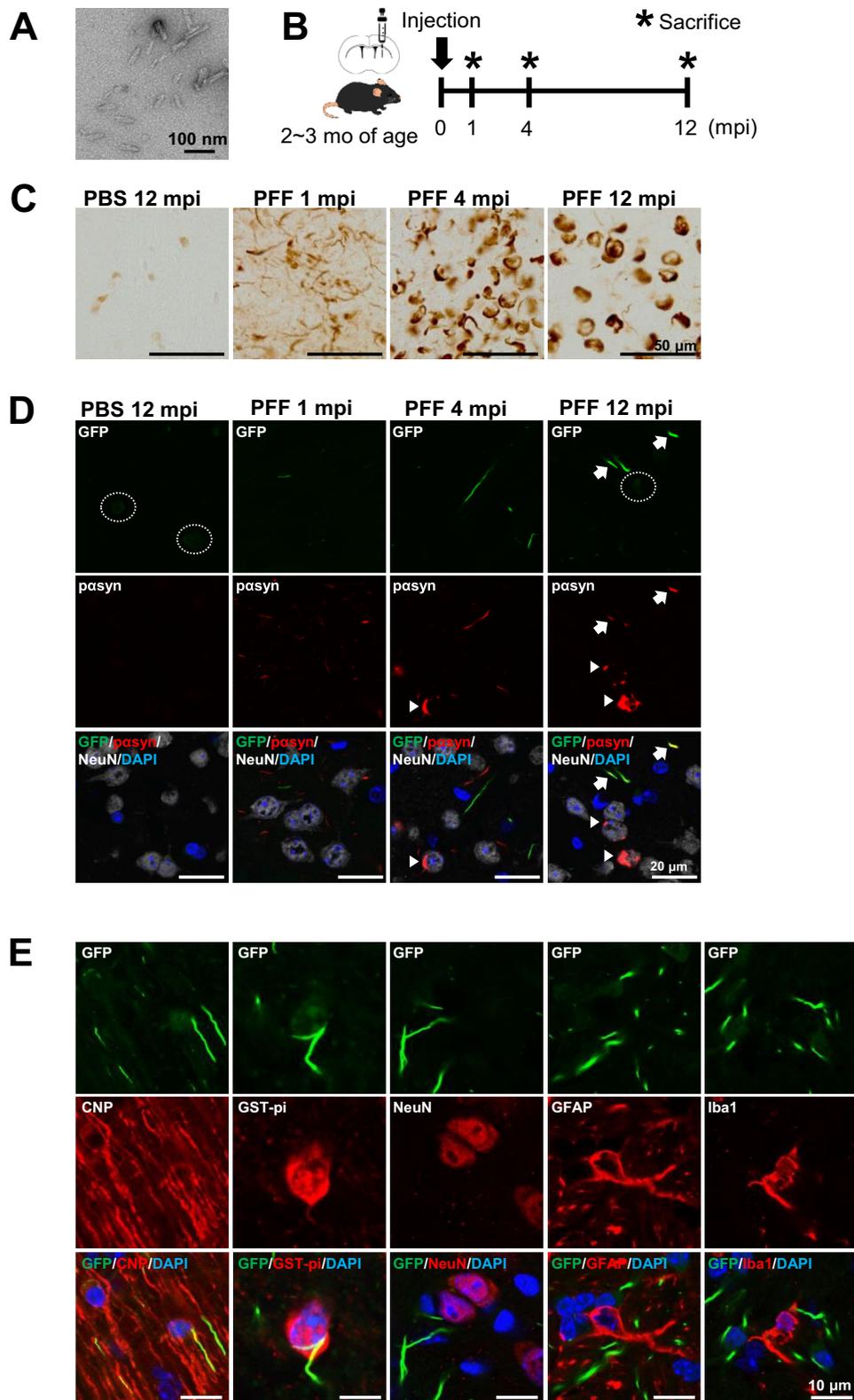


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Spatial and temporal distribution of asyn aggregates in neurons and OLGs in CNP-SNCAGFP Tg mice inoculated with asyn PFFs

The GFP dots in mice inoculated with asyn PFFs represented the asyn aggregates in OLGs. They were mainly

observed in the corpus callosum, followed by the striatum, and white matter of the cerebral cortex (Additional file 3: Fig. S3A). They first appeared as early as 1 mpi of asyn PFFs, and their signal intensity increased over time until 12 mpi (Figs. 2D, and 3A, Additional file 3: Fig.

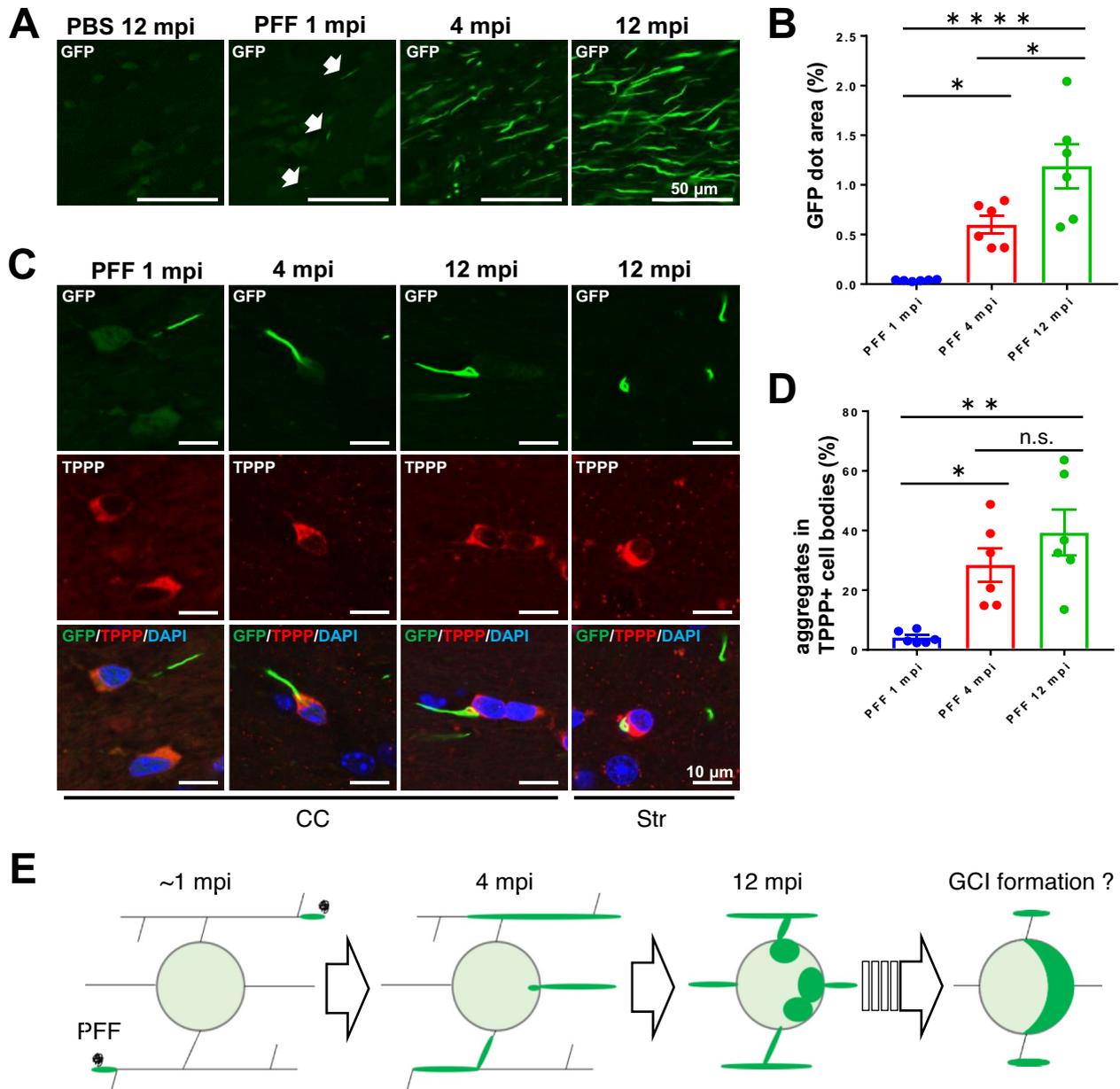


Fig. 3 asynGFP aggregates in OLGs developed in a centripetal manner in CNP-SNCAGFP Tg mice inoculated with asyn PFFs. **A** Fluorescence micrographs of the corpus callosum of Tg mice inoculated with asyn PFFs. Small asynGFP aggregates (GFP dots) were observed as early as 1 mpi (arrows), and their size and signal intensity increased over time. Scale bar = 50 μ m. **B** Quantification of the GFP dot area in **(A)** at 1, 4, and 12 mpi ($n=6$). **C** Fluorescence micrographs of GFP and immunostaining with anti-TPPP antibody in the corpus callosum of Tg mice at 1, 4, and 12 mpi. The merged images include DAPI (blue). GFP dots were initially observed in the OLG process and developed toward the cell body. Scale bar = 10 μ m. **D** Quantification of the number of aggregates in TPPP-positive cell bodies in **(C)** at 1, 4, and 12 mpi ($n=6$ at each time point). **E** The development of GFP dots is illustrated. Tukey's multiple comparisons test was performed in **(B)** and **(D)**; * $p < 0.05$, ** $p < 0.01$, **** $p < 0.0001$, n.s., not significant. Data indicate the mean \pm SEM. mpi, months post-inoculation; CC, corpus callosum; Str, striatum

S3A). In contrast, the signal intensity of p α syn reached a plateau at 4 mpi and remained stable until 12 mpi. These p α syn-positive aggregates rarely colocalized with the GFP dots (Additional file 3: Fig. S3B), and were sparse in OLG-rich regions, suggesting their neuronal origin (Additional file 3: Fig. S3A). Although majority of GFP-positive structures within OLGs also become p α syn-positive with increased sensitivity (Fig. S3C), this makes the analysis more difficult due to the strong p α syn signal of neuronal origin, especially in the striatum and cerebral cortex where neurons outnumber OLGs. These results suggest that, compared to α syn phosphorylation, GFP dot signals exhibit much higher sensitivity in detecting α syn aggregates in OLGs, especially in the early stage and regions abundant in neurons.

Development of α synGFP aggregates in a centripetal manner in CNP-SNCAGFP Tg mice inoculated with α syn PFFs

α synGFP aggregates (GFP dots) were first observed at 1 mpi, and their number, size, and signal intensity increased over time in CNP-SNCAGFP Tg mice inoculated with α syn PFFs (Fig. 3A, B). They were initially formed in the process far away from the TPPP-positive cell bodies in OLGs at 1 mpi (Fig. 3C). However, they developed in a centripetal manner and some of them became visible in TPPP-positive cell bodies in the striatum at 12 mpi (Fig. 3C, D). Figure 3E provides an illustrated summary of the development of GFP dots in these mice.

Thread-like α syn aggregates in the processes of OLGs in the brains of patients with MSA

The thread-like p α syn-positive structures were observed in the postmortem brains of patients with MSA (Fig. 4A), some of which co-localized with neuronal markers and were of neuronal origin. However, others did not co-localize with neuronal markers but co-localized with CNP, and were therefore of oligodendroglial origin (Fig. 4B). In contrast, in brains of patients with DLB, p α syn-positive aggregates co-localized with neuronal markers MAP2 and neurofilament, but not with CNP (Fig. 4B and Additional file 4: Fig. S4). These results suggest that some of the GCIs may developed from small α syn aggregates in the processes of OLGs in MSA, as shown in CNP-SNCAGFP Tg mice inoculated with α syn PFFs.

MSA brain homogenates (BH) induced more α syn aggregates in OLGs than DLB BH in CNP-SNCAGFP Tg mice
To investigate the tropism of α syn aggregates for OLGs in MSA, CNP-SNCAGFP Tg mice were inoculated with MSA and DLB BH and analyzed at 2 and 6 mpi. α syn

aggregates in OLGs were identified by GFP dots and those in neurons were identified by p α syn-positivity and GFP-negativity (Fig. 5A). When the numbers of neuronal and oligodendroglial α syn aggregates were compared between 2 and 6 mpi, there was no significant increase in the number of neuronal or oligodendroglial α syn aggregates in the DLB BH-treated group. In contrast, in the MSA BH-treated group, there was no significant increase in the number of neuronal α syn aggregates, but there was a significant increase in the number of oligodendroglial α syn aggregates, suggestive of the tropism for OLGs (Fig. 5B).

Discussion

How GCIs are formed and develop in MSA has remained a great mystery. Cases with preclinical MSA or incidental GCIs might contribute to answering this question [30, 31], but the number of such cases is limited and thus an animal model is an attractive option.

There are already excellent mouse models showing α syn pathology and associated motor symptoms. The model that expresses α syn under the PLP promoter show α syn pathology in OLGs and age-dependent motor symptoms as well as autonomic dysfunction [32–35]. Mouse models expressing α syn under the myelin basic protein (MBP) promoter are also useful models showing α syn pathology, motor symptoms, and neuroinflammation [36, 37]. M2 mice, which express human α syn under the CNP promoter, show phosphorylation of α syn in OLGs and age-dependent motor symptoms [38]. Furthermore, α syn seed administration to KOM2, a crossbreed between endogenous α syn knockout mice and M2 mice, induces α syn aggregate formation in the OLGs [39]. However, animal model specifically designed to explore the process and tropism of aggregate formation within OLGs is still lacking.

In this study, we succeeded in visualizing the aggregation process of endogenous α syn in OLGs from an early stage by employing newly generated CNP-SNCAGFP Tg mice inoculated with α syn PFFs. The sensitive, specific and early detection of α syn aggregates by GFP dots overcame the technical difficulties in identifying α syn aggregates, most of which are phosphorylated, in OLGs in the presence of numerous neuronal p- α syn aggregates. Indeed, we previously reported that α syn aggregates in OLGs were observed at 7 mpi of α syn PFFs in wild-type mice, as assessed by p α syn antibodies [26], whereas in the present study they could be detected as early as 1 mpi. In these mice, endogenous α syn aggregates were initially observed at the process and they developed toward the cell body in OLGs. This centripetal development of aggregates is in part consistent with findings in postmortem brains of MSA patients, and is not likely an effect of

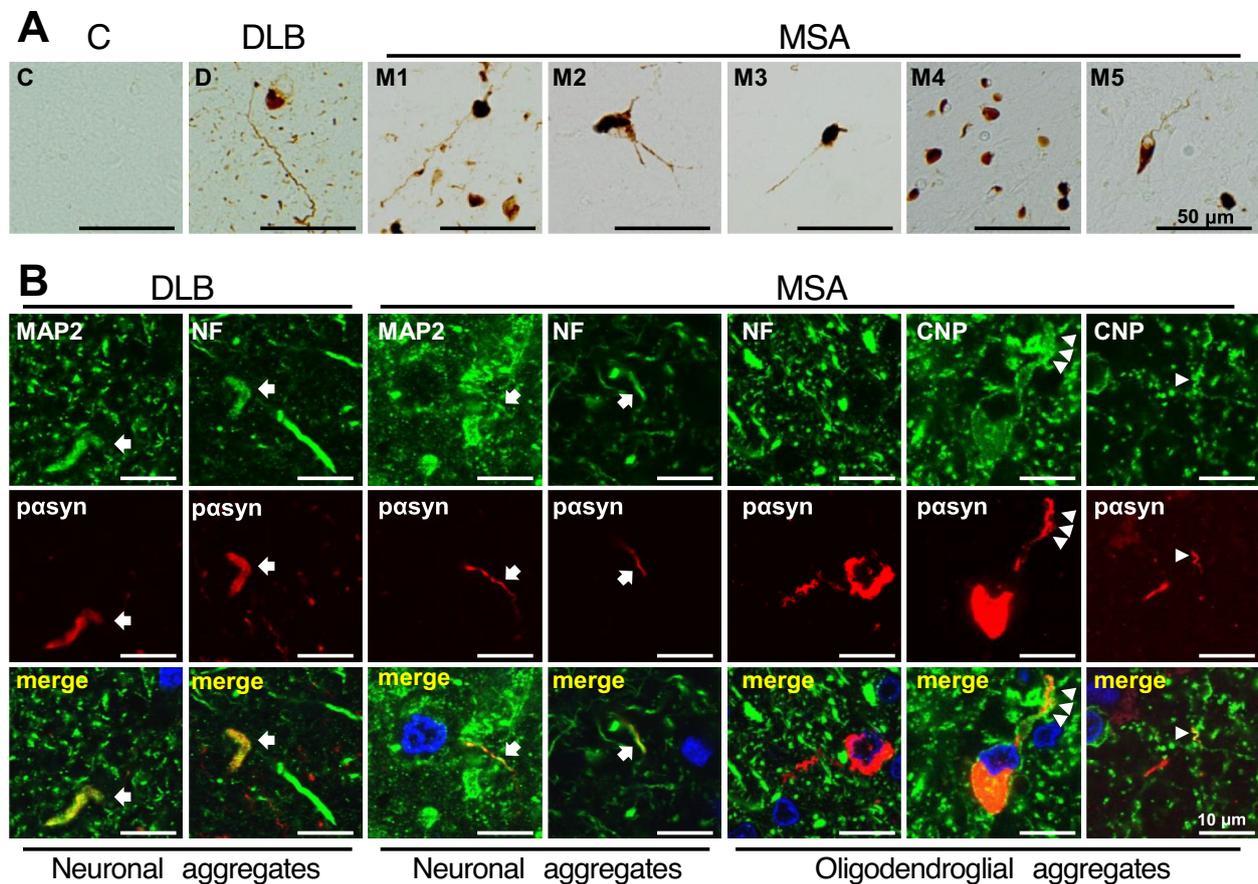


Fig. 4 Thread-like asyn aggregates in the process of OLGs in MSA brains. **A** Immunohistochemical staining of human autopsy brains with anti-pasyn antibody. Some of the pasyn-positive aggregates in MSA brains showed thread-like structures continuous with the cytoplasmic inclusions. The following regions were analyzed: the frontal lobes (C, D, M1-3), the putamen (M4), and the cerebellum (M5). Scale bar = 50 μ m. **B** Double immunofluorescent staining of thread-like aggregates with anti-pasyn antibody and various cellular markers in DLB and MSA brains. The merged images include DAPI (blue). Lewy neurites in DLB brains and some of the thread-like pasyn-positive aggregates in MSA brains co-localized with MAP2 and NF (arrows). Other pasyn-positive aggregates in MSA brains were not co-localized with neuronal makers but were co-localized with CNP and were continuous with GICs (arrowheads). Note that anti-CNP antibodies strongly labeled the plasma membrane in OLGs. Scale bar = 10 μ m. C, control; D, DLB (dementia with Lewy bodies); M, MSA. NF, neurofilament; pasyn, phosphorylated α -synuclein

the fused GFP protein because the distribution of human asynGFP did not differ from that of human asyn in CNP-SNCA Tg mice.

The origin of asyn aggregates in MSA has been considered to be OLGs, because in most cases GICs appeared

earlier and were more abundant than neuronal aggregates. Contrary to this notion that MSA is a primary oligodendroglialopathy, it has recently been reported that asyn aggregates in neurons are more widespread than previously thought [3, 5, 40, 41], and that in some cases

(See figure on next page.)

Fig. 5 Inoculation of brain homogenates from MSA samples induced distinct asyn aggregates in OLGs in CNP-SNCA GFP Tg mice. **A** Fluorescence micrographs of GFP and immunostaining with anti-pasyn (red) and anti-NeuN (gray) antibodies in the striatum of Tg mice treated with BH from DLB or MSA samples at 2 mpi and 6 mpi. The merged images include DAPI (blue). GFP dots (arrowheads) represent aggregates in OLGs. Aggregates that were immunopositive for pasyn and not co-localized with GFP were neuronal (arrows). **B** Comparison of the number of asyn aggregates in neurons and OLGs in the striatum (bregma + 0 mm) of Tg mice treated with BH from DLB or MSA. A single data point represents the average number of asyn aggregates of four mice at 2 mpi and two mice at 6 mpi, respectively. Neither the neuronal aggregate nor GFP dot exhibited a significant increase over time in the DLB BH-treated group. In contrast, the MSA BH-treated group displayed a notable increase only in GFP dots at 6 mpi. Scale bar = 20 μ m. Student's t-test was performed in **(B)**; * $p < 0.05$, n.s., not significant. Data indicate the mean \pm SEM. BH, brain homogenates; D, DLB (dementia with Lewy bodies); M, MSA; mpi, months post-inoculation; pasyn, phosphorylated α -synuclein

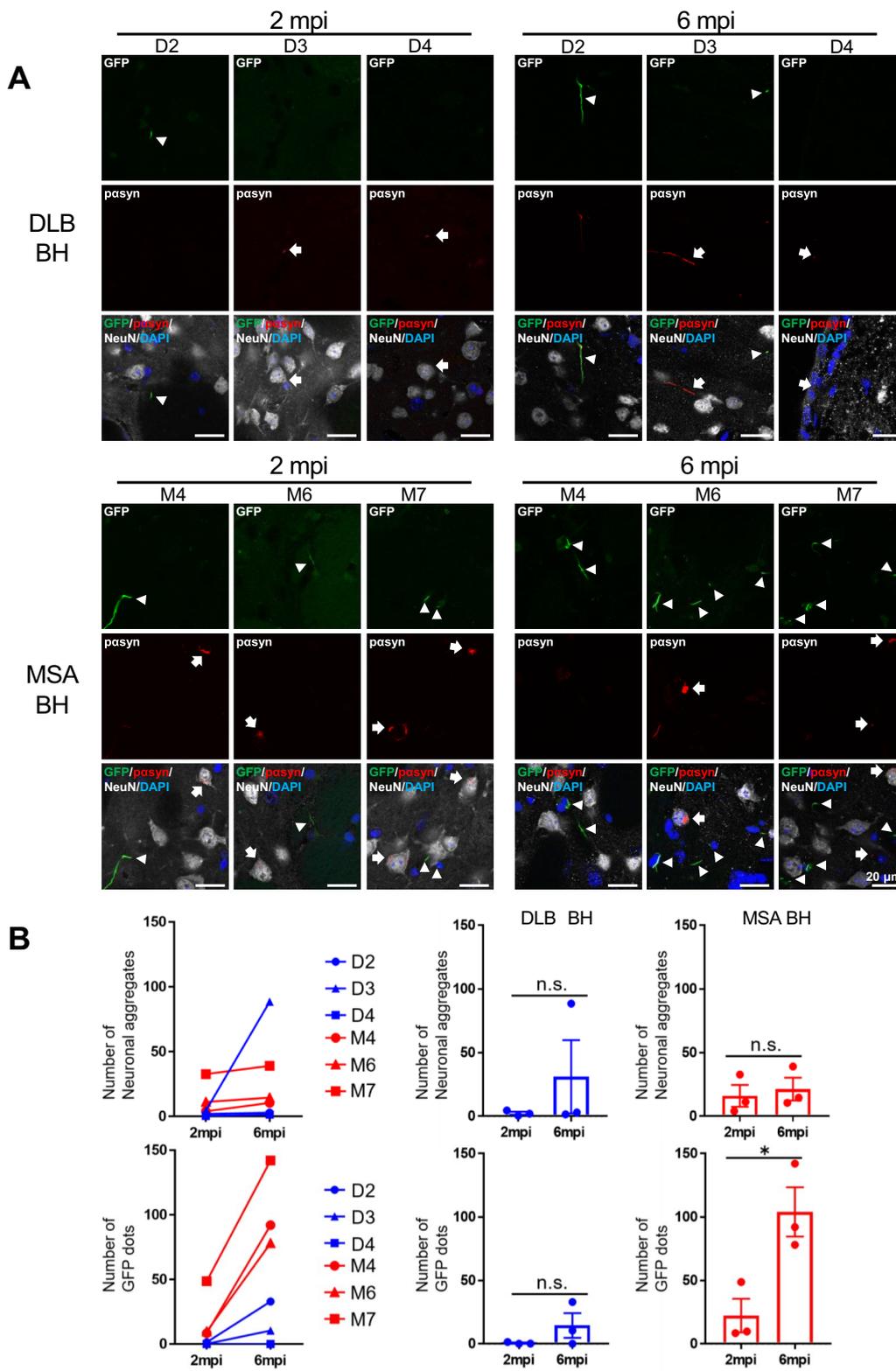


Fig. 5 (See legend on previous page.)

of MSA, they are predominantly found in neurons of specific brain regions [5]. α syn oligomers visualized by an α syn proximity ligation assay (PLA) are also reported to be observed in cortical neurons in the early stages of MSA [42]. Taken together with the fact that the expression level of α syn in OLGs is much lower than that in neurons, these studies strongly suggest the neuronal origin of α syn aggregates in GCIs [43]. There are also two possibilities for the origin of α syn seeds in OLGs in our mouse model. One is that α syn aggregates in OLGs directly seeded by inoculated α syn PFFs. This is supported by previous studies that upon treatment of α syn PFFs, α syn aggregates formed within primary OPCs [29] and in OLGs of mice that express human α syn exclusively in OLGs in a murine α syn null background [39]. These reports suggest that α syn aggregates can be taken up by and develop in OPCs/OLGs without neuronal α syn. The other possibility is that they propagated from α syn aggregates in neurons that were seeded by injected α syn PFFs. Although α syn aggregates were observed in OLGs at as early as 1 mpi of α syn PFFs, numerous α syn aggregates were already observed in neurons by this time. Therefore, it is difficult to determine whether the α syn aggregates in OLGs in this model are of neuronal origin or not. Another limitation is that the oligodendroglial aggregates observed in this model were less stained by antibodies for p α syn, possibly due to the fused GFP protein at the C-terminus of transgenic human α syn [44]. It is well known that the aggregation process of GFP fusion proteins is often different from that of naked proteins. In the case of α syn, it has been reported that α syn-GFP aggregates exhibit a fibril structure similar to that of naked protein aggregates by electron microscopic analysis [44], but the aggregation propensity of α syn-GFP is somewhat reduced compared to naked protein due to the presence of GFP at the C-terminus [44]. Therefore, physicochemical properties of α syn-GFP may be similar but not entirely identical to those of the naked α syn protein. Despite these limitations, this model may still be valuable for the sensitive and specific monitoring of the development of α syn aggregates in OLGs.

The properties of α syn aggregates have been shown to differ pathologically between PD, DLB, and MSA [45], and α syn aggregates amplified from patient blood samples also differ biochemically between PD, DLB, and MSA [46]. Recent studies showed that α syn seeds generated in the different cellular milieu cause different pathologies depending on their strains [39, 47, 48], and there may be α syn seeds that preferentially affect OLGs. To address this issue, we inoculated BH from MSA and DLB samples in CNP-SNCAGFP Tg mice. In contrast to many previous studies showing that MSA BH induced the α syn aggregates exclusively in neurons in wild-type

and α syn Tg rodents [24, 49, 50], not only MSA BH but also DLB BH induced α syn aggregates in OLGs. This can be explained by the overexpressed human α syn in OLGs and is consistent with the recent report showing that BH induced α syn aggregates in OLGs of P1 artificial chromosome (PAC) transgenic mice which overexpress human α syn in OLGs by the endogenous α syn promoter [51]. In our study, MSA BH induced more aggregates than DLB BH in neurons and OLGs at 2 mpi. This may be due to the higher seeding activities of MSA BH, but the greater increase in the number of aggregates in OLGs compared to neurons at 6 mpi can be explained by the tropism of α syn seeds in MSA for OLGs. These studies demonstrate that CNP-SNCAGFP Tg mice will be useful for investigating the cell-type preference of α syn seeds by sensitive detection in OLGs.

In conclusion, we generated a novel mouse model that enables the sensitive and specific detection of α syn aggregates in OLGs from an early stage. CNP-SNCAGFP Tg mice are expected to help investigate the formation of GCI and thus be useful as a preclinical model for the development of therapeutic agents, such as inhibitors of α syn aggregation in OLGs.

Methods

Animals

The mice used in this study were handled according to national guidelines. All procedures performed in this study were approved by the Institutional Animal Care and Use Committee of the Laboratory Animal Research Institute, Kyoto University Graduate School of Medicine (Med Kyo 18,215).

The transgenic construct of the CNP-SNCAGFP Tg mouse and CNP-SNCA Tg mouse were generated as follows. The coding sequence of human α synGFP or α syn was cloned into the pBSK plasmid harboring the murine 2',3'-cyclic nucleotide 3'-phosphodiesterase (CNP) promoter [52], which was a generous gift from Dr. Vittorio Gallo (Children's National Research Institute, DC). Next, the ligated plasmid was digested with XbaI and XhoI, purified and microinjected into C57BL/6J fertilized eggs. For genotyping, the following primer sets were used: Forward 5'-GGCTGGCTTTGAGGAGCC-3', Reverse 5'-GGGCTCCTTCTTCATTCTT-3'. Homozygous Tg was used in brain homogenate experiments and real-time PCR was performed to identify homozygous Tg using the following primer sets: Forward 5'-AAGTTCATCTGC ACCACCG-3', Reverse 5'-TCCTTGAAGAAGATG GTGCG-3'.

Human brain samples

Autopsied brains of patients with MSA, DLB, and non-neurodegenerative disease controls were used (listed in

Additional file 5: Table S1). Frozen tissue samples were obtained from Kyoto University Hospital, National Hospital Organization Hyogo-Chuo National Hospital, and the University of California, San Diego. Frozen human brain tissues were homogenized in PBS to 10% (wt/vol) and centrifuged at $2,000\times g$ for 10 min at 4 °C. The supernatant was used for inoculation. All procedures using human materials were performed in accordance with the ethical guidelines and approved by the Ethics Committee of Kyoto University Graduate School and Faculty of Medicine (R1038).

Immunohistochemical analysis of mouse and human brain samples

Mice were anesthetized with sevoflurane and transcardially perfused with phosphate-buffered saline (PBS), followed by 4% (w/v) paraformaldehyde in PBS. The brains were then removed and immersed in 4% (w/v) paraformaldehyde in PBS overnight at 4°C, replaced with 30% sucrose (w/v) in PBS, embedded in O.C.T compound (SAKURA Finetek), and frozen at -80°C. The mice brains were sliced in 40- μm -thick in a cryostat and stored in 0.02% NaN_3 (w/v) in PBS at 4°C. In addition, formalin-fixed, paraffin-embedded, 8- μm -thick brain sections were used for staining with antibodies for αsyn aggregates. For the immunohistochemical analysis of human brains, formalin-fixed, paraffin-embedded, 6- μm -thick sections were deparaffinized and heat-induced antigen retrieval was performed with Tris-EDTA buffer (pH 9.0).

Sections were immersed in 3% hydrogen peroxide in PBS for 30 min at room temperature (RT) to inactivate the endogenous peroxidase activity. After blocking with 10% goat serum and 0.02% Triton-X 100 in PBS for 1 h at RT, the sections were incubated overnight at 4°C with the primary antibody (listed in Additional file 5: Table S2). Then, the sections were incubated with peroxidase-labeled secondary antibodies (Histofine Simplestain Max PO, Nichirei Biosciences) and visualized by diaminobenzidine staining (Nacalai Tesque).

Immunofluorescence staining

Formalin-fixed, paraffin-embedded sections were deparaffinized, and heat-induced antigen retrieval was performed with Tris-EDTA buffer (pH 9.0). This process was not performed on free-floating mouse brain sections. After blocking with 10% goat or donkey serum and 0.02% Triton-X 100 in PBS for 1 h at RT, sections were incubated overnight at 4°C with primary antibodies (listed in Additional file 5: Table S2) (when MJFR-14-6-4-2 antibody was used, 0.5 M NaCl was added to the buffer). These sections were incubated with goat- or donkey-derived secondary antibodies (Alexa Fluor 488, 594, and 647, 1:400, Thermo Fisher) for 2 h at RT. The

sections were incubated with Vector TrueVIEW Auto-fluorescence Quenching Kit (Vector Laboratories) for 5 min to reduce autofluorescence, covered with Vibrance Antifade Mounting Medium with DAPI (Vector Laboratories), and observed with an FV1000 confocal microscope (OLYMPUS).

Proteinase K digestion

Proteinase K (PK) digestion was used to determine solubility of the human αsyn observed in OLGs with minor modifications [53]. Striatal sections were mounted onto FRONTIER-coated slide glass (Matsunami glass) and dried overnight at RT. After washing with PBS, the sections were digested with 200 $\mu\text{g}/\text{ml}$ PK in PBS for 30 min at RT. After several washes, the sections were immunostained as described above.

Generation of recombinant mouse αsyn monomers and fibrils

Recombinant αsyn monomers were generated as previously described, with minor modifications [26, 54, 55]. Briefly, mouse αsyn was expressed in *Escherichia coli* BL21 (DE3) (BioDynamics Laboratory) and purified by ion exchange using Q Sepharose Fast Flow (GE Healthcare). Endotoxin from *Escherichia coli* was removed using the ToxinEraser™ Endotoxin Removal Kit (Genscript), and the endotoxin levels were confirmed to be below the detection sensitivity with LAL Endotoxin Assay Kit (Genscript). After dialysis against 5 mM Tris-HCl and the addition of 10 \times PBS to make 1 \times PBS solution, αsyn monomers (5 mg/ml) were incubated at 37°C with constant agitation at 1,000 rpm for 7 days. The solution was sonicated for 5 min with a Bioruptor (Sonicbio) before injection.

Western blotting

Western blot analyses were conducted as described previously with minor modifications [56]. Briefly, mice brains perfused with cold PBS were homogenized in a tenfold volume of lysis buffer (150 mM NaCl, 1 mM ethylenediaminetetraacetic acid, 10 mM Tris-HCl, 1% (v/v) TritonX-100, 2% (v/v) SDS) or RIPA buffer (20 mM HEPES pH 7.4, 150 mM NaCl, 2 mM EDTA, 1% (v/v) Nonidet-P40, 0.5% (v/v) sodium deoxycholate, 0.1% (v/v) SDS). The homogenates were centrifuged at $100,000\times g$ for 30 min at RT. The supernatant was boiled in sample buffer (1% (w/v) SDS, 12.5% (w/v) glycerol, 0.005% (w/v) bromophenol blue, 50 mM dithiothreitol, 25 mM Tris-HCl, pH 6.8), and 5 μg of protein was loaded and separated by SDS-PAGE 5–20% (w/v) gradient gels (Wako). Proteins were transferred to polyvinylidene difluoride membranes using Trans-Blot SD Semi-Dry Transfer Cell (Bio-Rad). The membranes were treated with 4% (w/v)

paraformaldehyde in Tris-buffered saline (TBS) for 30 min at RT. After blocking for 1 h with 5% (w/v) skim milk in TBS with 0.1% (v/v) Tween 20 (TBST), each membrane was incubated with anti- α syn (BD Bioscience, clone 42, 1:2,000), anti-GFP (Abcam, EPR14104, 1:10,000), and anti-p α syn (Abcam, EP1536Y, 1:10,000) antibodies overnight at 4°C. These membranes were washed with TBST and incubated with horseradish peroxidase-conjugated secondary antibodies (NB7574 or NB7160, Novus Biologicals, 1:10,000) for 1 h. Images were acquired using an Amersham Imager 600 (GE Healthcare).

Transmission electron microscopy

Sonicated α syn PFFs were dropped onto a 200-mesh carbon-coated copper grid (Nissin EM), and PFFs adsorbed on the grid were negatively stained with 1% (w/v) uranyl acetate solution. Electron micrographs were obtained using a transmission electron microscope (H-7650, HITACHI) at 80 kV.

Intracerebral injection of α syn PFFs and brain homogenate (BH)

CNP-SNCAGFP Tg mice (age: 2–3 months) were anesthetized with isoflurane and inoculated with 2 μ l (10 μ g) of α syn PFFs ($N=20$), 2 μ l PBS ($N=3$) or 2 μ l BH ($N=18$) into the left dorsal striatum (A/P, +0.2 mm; M/L, +2.0 mm; D/V, -2.6 mm) using a 33-gauge Neurosyringe (Hamilton).

Statistical analysis

All statistical analyses were performed using the GraphPad Prism software program (version 7.04). All statistical tests performed are described in Figure Legends. P values of <0.05 were considered to indicate statistical significance.

Abbreviations

α syn	Alpha-synuclein
BGH	Bovine growth hormone
BH	Brain homogenate
CD68	Cluster of differentiation 68
CNP	2', 3'-Cyclic nucleotide 3'-phosphodiesterase
DLB	Dementia with Lewy bodies
GCLs	Glial cytoplasmic inclusions
GFAP	Glial fibrillary acidic protein
GFP	Green fluorescent protein
GSTpi	Glutathione S transferase pi
Iba1	Ionized calcium-binding adapter molecule 1
IHC	Immunohistochemistry
MAP2	Microtubule-associated protein 2
MSA	Multiple system atrophy
NeuN	Neuronal nuclear antigen
NF	Neurofilament
OLGs	Oligodendrocytes
p α syn	Phosphorylated alpha-synuclein
PBS	Phosphate-buffered saline
PD	Parkinson's disease
PFFs	Preformed fibrils
PK	Proteinase K

p62	P62/sequestosome-1
TBS	Tris-buffered saline
TBST	Tris-buffered saline with Tween 20
Tg	Transgenic
TPPP	Tubulin polymerization promoting protein
Ub	Ubiquitin
Wt	Wild type

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s13041-024-01104-7>.

Additional file 1: Fig. S1. The α synGFP in CNP-SNCAGFP Tg mice exhibits similar subcellular distribution to α syn in CNP-SNCATg mice and does not aggregate with age. (A) Fluorescence micrographs of GFP and immunostaining with anti-human α syn antibodies (red) and CNP (gray) of the striatum in CNP-SNCAGFP Tg mice and in CNP-SNCA Tg mice that express human α syn in OLGs. Scale bar = 10 μ m. (B) Western blot images of RIPA-soluble fraction with anti- α syn and p α syn antibodies in whole brains of Wt and Tg mice (2, 13, and 18 months old). The expression of endogenous mouse α syn (arrow) and transgenic human α synGFP fusion protein (arrowhead) are shown. (C) Latency to fall in the rotarod test at 12 months old. Student's t-test was performed (not significant). Data indicate the mean \pm SEM. Wt, wild-type; α syn, alpha-synuclein; p α syn, phosphorylated α -synuclein; PK, proteinase K; RT, room temperature.

Additional file 2: Fig. S2. GFP dot signals co-localized with markers of α syn aggregates. (A) Fluorescence micrographs of GFP and immunostaining with anti-ubiquitin, p62, and aggregated form of α syn antibodies (MJFR-14-6-4-2) in the striatum of Tg mice at 6 mpi of α syn PFFs. The merged images include DAPI (blue). Aggregates that were immunopositive for p62 and MJFR-14-6-4-2 that did not co-localize with GFP were neuronal (arrowheads). (B) Immunohistochemical staining with anti-GFP and aggregated form of α syn antibodies (5G4 and A17183A) in the striatum of Tg mice at 6 mpi of α syn PFFs (paraffin-embedded sample). The merged images include DAPI (blue). Scale bar = 10 μ m. (C) Immunohistochemical staining with human-specific α syn antibody (MJFR1) in the striatum of 2, 12, and 20 months-old Tg mice (left panel) and in the striatum of Tg mice at 6 mpi of PBS or α syn PFFs (right panel). The lower panel shows sections stained under the same conditions as the upper panel after PK digestion (200 μ g/ml, 30 min, RT). Arrows represent α syn aggregates in OLGs. Scale bar = 20 μ m. Ub, ubiquitin; PK, proteinase K; RT, room temperature.

Additional file 3: Fig. S3. Sensitive detection of oligodendroglial aggregates by α synGFP signals in CNP-SNCAGFP Tg mice treated with α synPFFs. Fluorescence micrographs of GFP and immunostaining with p α syn antibodies in Tg mice (low-power magnification). The signal intensity of GFP dots increased over time until 12 mpi, especially in the CC, whereas that of p α syn reached a plateau at 4 mpi in Tg mice inoculated with α syn PFFs. Scale bar = 200 μ m. (B) The high-power magnification of the area enclosed by the square in (A). Note that most of the α synGFP and p α syn-positive aggregates did not merge. Scale bar = 100 μ m. (C) With increased sensitivity, fluorescence micrographs of GFP and immunostaining with anti-p α syn (red) antibody in the striatum and corpus callosum of Tg mice inoculated with α syn-PFFs. The merged images include DAPI (blue). Arrows indicate weak but detectable p α syn signals of α synGFP aggregates in the striatum of the Tg mice at 1 and 4 mpi. Scale bar = 20 μ m. mpi, month(s) post-inoculation; Ctx, cortex; CC, corpus callosum; Str, striatum; PFFs, preformed fibrils; p α syn, phosphorylated α -synuclein.

Additional file 4: Figure S4. Immunohistochemical staining of human brains with anti-p α syn, CNP, MAP2, and NF antibodies. The merged images include DAPI (blue). Lewy neurites in the DLB brain did not co-localize with CNP. No p α syn-positive aggregates were observed in the control brain. Scale bar = 10 μ m. DLB, dementia with Lewy bodies; C, control; NF, neurofilament.

Additional file 5: Table S1. Clinical information for autopsy cases. Table S2. List of primary antibodies used for immunohistochemistry.

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Authors' contributions

T.I. performed the background studies, collected data, made the figures, wrote the manuscript, and arranged the financial support. H.Y. and R.T. designed the research, wrote the manuscript, made critical revisions of the manuscript for important intellectual content, and arranged the financial support. T.A., N.K., I.F. performed the pathological diagnosis of human brain samples. M.O., S.K., T.A., T.M., and S.M. supervised and made critical revisions to the manuscript for important intellectual content. All authors read and approved the final manuscript. All authors have approved the manuscript and agree with its submission.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author upon reasonable request.

Declarations

Ethics approval and consent to participate

All procedures involving the use of human materials were performed in accordance with the ethical guidelines set by Kyoto University. The Institutional Animal Care Committee of the Kyoto University Graduate School of Medicine approved all protocols in animal studies.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

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