

RESEARCH ARTICLE

Biochemical Increase in Phosphorylated Alpha-Synuclein Precedes Histopathology of Lewy-Type Synucleinopathies

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Keywords

alpha synuclein, human brain, Lewy body disorder, Parkinson's disease, pathology staging, phosphorylation.

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Abstract

A key component in Lewy body (LB) pathology in LB disorders is α -synuclein phosphorylated at serine 129 ($p\alpha$ syn). However, it is not known if increase in the level of biochemically measurable $p\alpha$ syn precedes the presence of histologically identified Lewy-type synucleinopathy (LTS). To gain sights into possible temporal sequence, we measured levels of $p\alpha$ syn in cingulate and temporal cortices that develop LTS pathology at later stages of LB disorders. Brain homogenates from 128 autopsy cases including normal controls and subjects classified by Unified LTS histopathology staging system were studied. We found that biochemically measurable $p\alpha$ syn levels in cingulate and temporal cortices were significantly increased at Unified stages III and IV. When $p\alpha$ syn levels were compared between LTS density scores instead of Unified stages, significant increases were detected even as LTS density scores increased from 0 to 1 in olfactory bulb and substantia nigra. Therefore, our findings demonstrated that changes of $p\alpha$ syn levels in cingulate and temporal cortices coincided with the early appearance of the LTS pathology in olfactory bulb and substantia nigra, even though histologically demonstrable LTS was lacking in the cortical region. Therefore, identifying the underlying mechanisms driving these changes could be crucial to understanding the pathogenesis of LB disorders.

INTRODUCTION

Soluble alpha synuclein (α syn) protein is normally concentrated in presynaptic terminals in the brain (28). Deposition of aggregated α syn in neuronal cytoplasm, processes and presynaptic terminals is believed to be a central event in the pathogenesis of Lewy body (LB) disorders, which include Parkinson's disease (PD), Alzheimer's disease with a restricted distribution of Lewy bodies (ADLB) and dementia with Lewy bodies (DLB) (4, 26, 44, 53). Subsequently, α syn-immunoreactive Lewy-type synucleinopathy (LTS) has become a useful criterion for the neuropathological diagnosis of these heterogeneous diseases and it has been shown that the abundance and distribution of α syn-containing pathology is associated with the severity of LB disorders. Several staging systems have been developed to classify the stages of PD and DLB based on the regional brain distribution of LTS histopathology (9, 10, 11, 35, 36, 41, 50). Among these, the system developed by Braak and colleagues was specifically designed to classify PD subjects (10, 12). They proposed that α syn pathology progresses in six sequential stages, beginning in the dorsal motor nucleus of vagus

following a caudal-to-rostral direction in the order of medulla, pons, midbrain, nucleus basalis of Meynert, limbic regions and neocortex (10, 12). In the Braak's scheme, the involvement of olfactory bulb was thought to be inconsistently present, coincidental to medullary affectation. Another staging system was specifically devised for DLB by the Dementia with Lewy Bodies Consortium (third version), which grades α syn-immunoreactive pathologies in multiple brain regions to classify three types of Lewy pathology development: brain stem predominant, limbic/transitional and diffuse neocortical types (40, 41); in this system, there is no consideration of olfactory bulb involvement.

The efficacy of these classificatory systems has recently been assessed by several groups (8, 14, 20, 23, 29, 31). It has been reported that 6.3% to 47% of clinical PD cases did not follow the proscribed caudal-to-rostral pattern of LTS pathological development (13, 30, 46). A similar situation was found to exist for the DLB Consortium system where again, the dictated caudal-to-rostral progression was often not followed. Additionally, both systems failed to identify the presence of a relatively common olfactory bulb-only stage, which may be the first stage common to

all LB disorders (8). Because of these deficiencies, we proposed “Unified Lewy Body Staging System,” which can be used to classify all subjects regardless of types of LB disorders (Figure 1) (7). This system was established by the abundance and distribution of phosphorylated serine 129 alpha synuclein ($p\alpha$ syn) immunoreactive pathologies in a large series of autopsy cases from the Banner Sun Health Research Institute Brain and Body Donation Program (7).

Although there are other serine and tyrosine residues on α syn, phosphorylation at serine 129 is the most prominent pathological species in LB disorders (2, 21, 22). Under normal condition, the brain expression of $p\alpha$ syn is very low, but in LB disorders, its levels are increased in synapse-enriched fractions as well as in the hallmark pathologies, Lewy neurites and Lewy bodies (2, 43). Phosphorylation of α syn promotes oligomerization, aggregation, neurotoxicity and formation of LTS pathology, and reduces the ability to regulate tyrosine hydroxylase (17, 25, 27, 33, 44, 45, 49, 52). In α syn transgenic mouse model, enhancing phosphatase

activity has been shown to reduce the levels of $p\alpha$ syn and improve neuronal activity and motor performance (32).

In the present study, we explored the temporal and anatomical relationship between the presence of histologically identified $p\alpha$ syn-containing LTS and biochemically measured $p\alpha$ syn. We hypothesized that increase in biochemically measurable serine 129 $p\alpha$ syn precedes the presence of histologically identified LTS pathologies. The hypothesis was tested in cingulate and temporal cortices of 128 subjects (normal controls as well as cases from all LB stages). Because LTS pathology is absent or rare in the cingulate and temporal cortices at early disease stages, such as stages I and II, they can be used to determine the temporal relationship between the biochemical and histological changes of $p\alpha$ syn occurring at early stages of the diseases.

We measured both $p\alpha$ syn and the combined amounts of unmodified (nonphosphorylated) and phosphorylation-modified α syn, defined as total α syn ($t\alpha$ syn) and determine the ratios of $p\alpha$ syn to $t\alpha$ syn in individual samples. Based on this ratio, we determined whether increases in $p\alpha$ syn levels were selective or due only to increases in total α syn. We also determined if the increase in biochemically measurable $p\alpha$ syn in cingulate and temporal cortical regions had already occurred when LTS histopathologies began to appear in olfactory bulb, and spread to substantia nigra and amygdala.

Our results showed a significant increase in the biochemical level of $p\alpha$ syn when LTS density score in each given region becomes detectable, and the levels of $p\alpha$ syn in cingulate and temporal cortices were significantly and selectively elevated when histological LTS density scores increased from 0 to 1 in either the olfactory bulb or the substantia nigra, even when LTS scores were zero in cingulate and temporal cortices. These results indicate that aberrant phosphorylation of α syn is an early and thus possibly important phenomenon during the pathogenesis of the LB disorders.

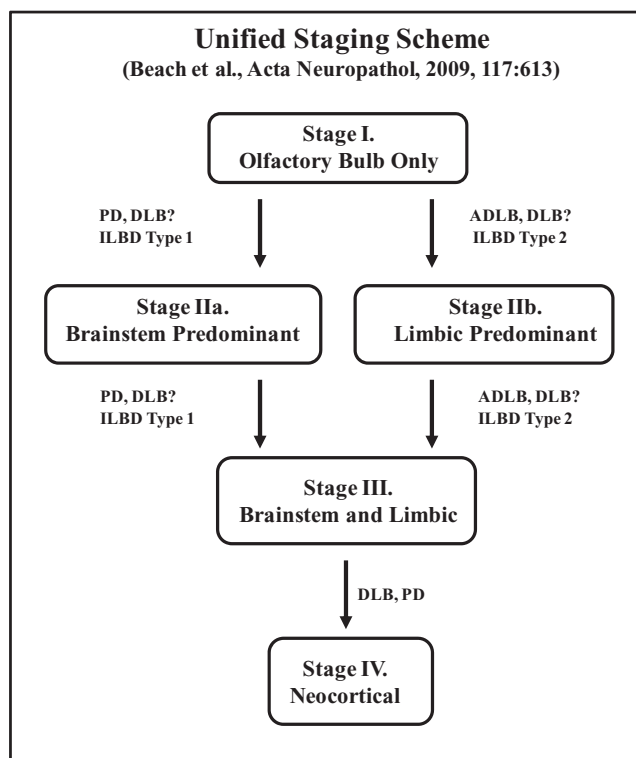


Figure 1. Unified staging scheme. This Lewy body (LB) pathology classification system is based on a scoring system grading the anatomical distribution of phosphorylated α -synuclein ($p\alpha$ syn) published previously (7). The system proposed that the development and progression of $p\alpha$ syn-containing LB pathology is initiated in olfactory bulb as stage I, followed by brain stem-predominant (stage IIa) or limbic-predominant stages (stage IIb), then merged as both brainstem and limbic stages (stage III) and eventually advancing to neocortex (stage IV). The diseases which are likely to be observed at different stages are shown also in the scheme: PD = Parkinson’s disease; ILBD = Incidental Lewy body disease; ADLB = Alzheimer’s disease with Lewy bodies; DLB = dementia with Lewy bodies.

MATERIALS AND METHODS

Human subjects for Brain Donation Program

Subjects who enrolled in the Brain and Body Donation Program (BBDP) of Banner Sun Health Research Institute (BSHRI) were consented according to the approved protocols of the Banner Health Institutional Review Board. Most subjects receive annual standardized batteries of neurological and neuropsychological assessments (1, 15). For those subjects lacking standardized ante-mortem evaluations, information was obtained, for some subjects, from a post-mortem telephone interview with a contact, including an adaptation of the Clinical Dementia Rating Scale (CDR). The demographic features of the subjects used in this study are summarized in Table 1.

An overview of the BSHRI Brain Donation Program (now the Brain and Body Donation Program, BBBDP) has been published previously (5). Autopsy cases for this study were chosen by searching the BBBDP database for cases in order to establish adequate subject numbers at each Unified stage. A secondary objective was to include subjects from all types of Lewy body disorders, including PD, DLB and ADLB. The diagnostic criteria used for AD, PD and DLB have been previously described (5). For AD and DLB, cases received the diagnosis if they were classified as “intermediate” or “high” probabilities in their respective classification

Table 1. Demographic and neuropathological features of the patients grouped by Unified Staging System.

Features/stages	Stage 0	Stage I	Stage IIa	Stage IIb	Stage III	Stage IV
Number of subjects	17	11	5	16	34	45
Expired age (years)	84.1 \pm 1.5	85.8 \pm 1.2	87.0 \pm 1.4	81.6 \pm 1.9	80.6 \pm 1.2	78.8 \pm 0.8
Mean \pm SE						*(0, I, IIa)
Post-mortem delay (PMI, hours)	2.5 \pm 0.1	3.0 \pm 0.2	2.4 \pm 0.3	4.3 \pm 0.7	3.3 \pm 0.2	3.0 \pm 0.1
Mean \pm SE				*(0, I, IIa, III, IV)		
Rates of apoE ϵ 4 carrier in each stage	4/17	7/11	2/5	9/16	12/34	24/45
Gender ratios (male to female)	12/5	5/6	3/2	7/9	21/13	26/19
Total amyloid plaque scores	3.8 \pm 1.0	12.5 \pm 0.5	9.8 \pm 1.6	13.3 \pm 0.5	6.9 \pm 0.9	10.8 \pm 0.6
Mean \pm SE	*(I, IIb, IV)	*(0, III)		*(0, III)	*(I, IIb, IV)	*(0, III)
Braak's scores	2.8 \pm 0.2	4.8 \pm 0.4	3.7 \pm 0.5	5.3 \pm 0.2	2.9 \pm 0.2	3.8 \pm 0.2
Mean \pm SE	*(I, IIb, IV)	*(0, IIa, III, IV)	*(I, IIb)	*(0, IIa, III, IV)	*(I, IIb, IV)	*(0, I, IIb, III)
Substantia nigra depigmentation scores	0.5 \pm 0.2	1.5 \pm 0.2	2.0 \pm 0.3	0.9 \pm 0.2	2.8 \pm 0.1	2.4 \pm 0.1
Mean \pm SE	*(I, III, IV)	*(0, IIb, III, IV)	*(III)	*(I, III, IV)	*(0, I, IIa, IIb)	*(0, I, IIb)

Symbol * in front of parenthesis denotes statistical significance level at $P < 0.05$.

The Roman numerals within the parenthesis represent the stages that differed significantly with the stage indicated at the top row of this table.

schemes. Cases were classified as ADLB if they had AD and Lewy bodies in any brain region but failed to meet criteria for DLB or PD. The PD cases contained variable plaque and tangle scores and some of them met criteria for the diagnosis of AD. We called these cases PDAD. The cases, which did not contain AD pathology, were called PD when they were not demented and PDD when they were demented. Cases with DLB also often have AD pathology; in this study, 24 met intermediate or high criteria for AD while three did not. Subjects without dementia, Parkinsonism or neuropathological evidence of a LB disorder were chosen as control groups for the study.

Histological methods

Neuropathological diagnosis was made from brain tissue blocks fixed with 4% neutral-buffered formaldehyde and cut from both paraffin embedded and cryoprotected blocks of various brain regions. Tissue sections were processed with our standard histological procedure (7, 6). Histopathological grading of AD followed Braak neurofibrillary tangle staging and Consortium to Establish a Registry for Alzheimer's Disease (CERAD) templates (42). Microscopic observations of α syn-stained histological slides were performed by a single experienced observer (TGB) without prior knowledge of the patient's medical record or neurological status. Each case was first staged according to the Unified Staging System for LB disorders using a standard set of brain sections stained with an immunohistochemical method for α syn as previously described (7). In this methodology, proteinase K pretreatment was used to unmask the epitopes for α syn in paraformaldehyde-fixed brain tissues. For staging purposes, 10 brain regions were examined, including olfactory bulb and tract, anterior medulla, anterior and mid-pons, mid-amygdala and adjacent transentorhinal area, anterior cingulate cortex, middle temporal gyrus, middle frontal gyrus and inferior parietal lobule. The density of α syn-immunoreactive neuronal cytoplasmic inclusions as well as punctate and neuritic profiles was graded in each region according to the templates published by the DLB Consortium (41). The scores were assigned as none, sparse, moderate, frequent and very frequent.

Based on these results, the autopsy cases were assigned to one of the following stages: I, IIa, IIb, III and IV, or 0 when there was no LB pathology present. Stage I is also called "olfactory bulb only;" stage IIa, "brain stem predominant;" stage IIb, "limbic predominant;" stage III, "brain stem and limbic;" and stage IV, "neocortical" as shown in Figure 1 (7).

Brain tissue homogenization and protein extraction

A collection of 128 autopsy cases received at the BBDP of BSHRI between years 1998 and 2008 with LB pathology classified as Unified stage 0 to IV were used for the study. The numbers of cases in each stage are stage 0, 17 normal control cases; stage I, 11 ADLB cases; stage IIa, one PDAD and four PD cases with mild cognitive impairment (MCI); stage IIb, one PDD and 15 ADLB cases; stage III, five PD, 12 PDD, eight PDAD and nine DLB cases; and stage IV, seven PDD, 10 PDAD and 28 DLB cases. Gray matter from cingulate and temporal cortices were dissected out from 1-cm thick brain slices that were snap frozen between dry ice slabs at autopsy and then stored at -70 to -80°C . Samples were further cut into 2 mm cubes, weighed and homogenized with a sonicator on ice in four volumes of prechilled 1X radio-immunoprecipitation assay (RIPA) buffer consisting of 25 mM Tris-HCl (pH 7.6), 150 mM NaCl, 1% NP-40, 1% sodium deoxycholate, 0.1% sodium dodecyl sulfate (SDS), 1X cocktail protease and phosphatase inhibitors and 1 mM EDTA (Pierce, Thermo Scientific, Rockford, IL, USA). This buffer is commonly used for extraction of cytosolic, membrane-associated and extracellular matrix-associated proteins. After sonication, the extraction of proteins continued for 30 minutes on ice with gentle shaking. To remove unextracted nuclear components and debris, homogenates were centrifuged at 20 000 rpm for 30 minutes in a refrigerated table top centrifuge. A fraction of supernatant was removed for protein assay using Micro BCA™ method (Thermo Scientific). The rest of the supernatant was aliquoted and stored at -80°C until preparation for Western blot procedure.

Western blotting

Frozen RIPA buffer extracts of cingulate and temporal cortical tissues were thawed and mixed with 4X lithium dodecyl sulfate (LDS) Western blot sample buffer (BD Biosciences, San Jose, CA, USA) and water in the absence of reducing reagent dithiothreitol to obtain protein concentration at 1 μ g/ μ L in 1X LDS buffer. Samples are heated immediately before loading to NuPAGE® 10%–14% Bis-Tris Mini gels (BD Biosciences). Ten μ g of protein was loaded to each lane. The order of the samples was arranged to ensure that samples from cingulate and temporal cortices of the same autopsy case were loaded in adjacent lanes. We also made sure that samples loaded on the same gel contained samples from subjects of stage 0 to stage IV. Some of the Western blot samples were used as across-gel loading and processing controls. These were used to assess the extent of discrepancy between gels at the end of procedure.

Samples were electrophoresed at 200 V for 40 minutes using 2-(*N*-morpholino)ethanesulfonic acid (MES) buffer (BD Biosciences), and then transferred to nitrocellulose membrane at 30 V for 75 minutes. Blots were air-dried before blocking with tris-buffered saline consisting of 0.05% Tween® 20 (VWR International, West Chester, PA, USA) (TBS-T) and 5% nonfat dry milk. A rabbit polyclonal antibody raised against p α syn (S129P), characterized previously, was used for immunodetection (3, 7, 22). The specificity of the antibody was determined in a separate investigation (unpublished observation) by showing decreased immunoreactivity after treating the brain protein on the blots with phosphatase and after detection of p α syn protein generated by co-incubating α syn with casein kinase.

After incubation with this antibody at 1:5000 dilution overnight at room temperature, blots were washed four times with TBS-T and then probed with a goat anti-rabbit IgG conjugated with horseradish peroxidase (HRP; Thermo Scientific) at 1:20 000 concentration, and washed four times with TBS-T before detection with Western Extended Dura chemiluminescence substrate for HRP (Thermo Scientific).

Parallel blots were probed with a mouse anti- α syn antibody (No. 610786, BD Biosciences) at 1:1000 dilution. This antibody, raised against nonphosphorylated epitopes between amino acid 15–123, does not distinguish p α syn from unmodified form, thus its immunoreactive products consisted of both phosphorylated and nonphosphorylated α syn. We designated this measurement as t α syn. After overnight incubation with primary antibody, anti-mouse IgG conjugated with HRP (Thermo Scientific) was used at 1:10 000. The level of β actin in each sample was detected in each blot with a mouse anti- β actin (Clone AC-15, Sigma, St. Louis, MO, USA) antibody at 1:5000. Finally, the immunoreactivities of the bands were detected with Western Extended Dura chemiluminescence kits.

Images of immunoreactive bands were captured using CCD camera in the FluorChem® Q (Cell Biosciences, Santa Clara, CA, USA) ChemiImager™. Images from three durations of exposure were taken for each protein, ranging from 5 s to 1.5 minute; and the intensity of individual band was quantified with the software provided with FluorChem® Q. The results of densitometry were compared between exposure durations for each protein in order to determine if the development of immunoreactivity was within linear range. Although three exposures were taken for each blot, only one set of densitometry results was used for normalization by β actin and calculation of the ratio of p α syn to t α syn.

Statistical analysis

Statistical analysis was performed with MedCalc software version 9.8 (Mariakerke, Belgium). The significance level for all comparisons was set at $P < 0.05$. One-way analysis of variance (one-way ANOVA) was used to analyze the stage-related differences between p α syn, total α syn and their ratios in each brain region followed by Student Newman–Keuls pairwise comparisons. To assess the effects of pathological stages and brain regions on the levels of p α syn, total α syn and their ratios, two-way ANOVA was used, followed by Student Newman–Keuls analysis for pairwise comparison. The *F*-ratio is considered to be significant when $P < 0.05$. Spearman Rank correlation was used to relate the biochemical levels of p α syn to pathological features including a standard set of inputs, namely, the LB-related density scores from olfactory bulb, substantia nigra, amygdala, cingulate cortex, temporal cortex and frontal cortex.

RESULTS

Characteristics of the autopsy cases in this study

Demographic and neuropathological features

Demographic and neuropathological features including age at death, post-mortem interval, frequency of the apolipoprotein E ϵ 4 (ApoE ϵ 4) genotype, total amyloid plaque density scores, Braak's stage and substantia nigra pigmented neuron loss scores are shown in Table 1.

Subjects at Unified stage IV died at younger age (mean \pm SE = 78 \pm 0.8) than subjects at stages 0, I and IIa ($P < 0.05$). Post-mortem intervals (PMI) were significantly longer for subjects at Unified stage IIb but did not otherwise differ between groups. The frequency of ApoE ϵ 4 allele carriers was not significantly different between subjects at different Unified stages. The substantia nigra pigmented neuron loss score was significantly different from controls for all groups except for those in stage IIb. In terms of AD-related pathology, total plaque scores were not significantly different between stages I, IIb and IV, but were significantly higher than stages 0 and III. Braak's stages were not significantly different between Unified stages I and IIb, but were significantly higher than Unified stages 0, IIa, III and IV.

Lewy-type synucleinopathy (LTS) histopathological density scores in various brain regions

A grading system with a scale of 0 to 4 was used to assess the abundance of p α syn-immunoreactive Lewy-type synucleinopathy (LTS) in olfactory bulb, brain stems, amygdala, trans-entorhinal cortex, cingulate cortex and neocortical cortical regions, as described in previous publication (41). The means and standard errors of density scores of p α syn-immunoreactive LTS in 10 brain regions are listed according to Unified stages of this studied population in Table 2.

We examined the p α syn-immunoreactive LTS density scores in cingulate and temporal cortex, the two brain regions that we studied with biochemical methods. At stage IIb, the density scores were lower than 1 in both brain regions; 0.60 \pm 0.13 (mean \pm SE)

Table 2. Lewy pathology density scores in 10 brain regions of the studied patients grouped by Unified stages [mean \pm standard errors (subject numbers)].

Unified stages/brain regions	I	Ila	Ilb	III	IV
Olfactory bulb	1.71 \pm 0.19 (11)	1.40 \pm 0.68 (5)	3.31 \pm 0.18 (16)	2.17 \pm 0.24 (30)	2.92 \pm 0.20 (39)
Substantia nigra	0.00 \pm 0.00 (11)	0.50 \pm 0.29 (5)	0.50 \pm 0.18 (16)	2.27 \pm 0.18 (33)	2.67 \pm 0.13 (42)
Pons	0.00 \pm 0.00 (11)	0.75 \pm 0.48 (4)	0.25 \pm 0.18 (12)	2.74 \pm 0.18 (34)	2.78 \pm 0.16 (40)
Medulla	0.00 \pm 0.00 (11)	1.00 \pm 0.70 (4)	0.83 \pm 0.21 (12)	2.88 \pm 0.19 (32)	2.81 \pm 0.17 (42)
Amygdala	0.00 \pm 0.00 (11)	0.60 \pm 0.40 (5)	3.60 \pm 0.13 (15)	3.44 \pm 0.11 (32)	3.73 \pm 0.09 (41)
Trans-entorhinal cortex	0.00 \pm 0.00 (11)	0.40 \pm 0.40 (5)	2.93 \pm 0.31 (14)	2.36 \pm 0.17 (33)	3.31 \pm 0.13 (42)
Cingulate cortex	0.00 \pm 0.00 (11)	0.20 \pm 0.20 (5)	0.60 \pm 0.13 (15)	1.88 \pm 0.13 (32)	3.11 \pm 0.12 (44)
Temporal cortex	0.00 \pm 0.00 (11)	0.00 \pm 0.00 (5)	0.33 \pm 0.13 (15)	0.97 \pm 0.05 (32)	2.50 \pm 0.12 (44)
Parietal cortex	0.00 \pm 0.00 (11)	0.00 \pm 0.00 (5)	0.06 \pm 0.06 (15)	0.77 \pm 0.09 (34)	1.73 \pm 0.11 (45)
Frontal cortex	0.00 \pm 0.00 (11)	0.00 \pm 0.00 (5)	0.06 \pm 0.06 (15)	0.77 \pm 0.07 (34)	1.67 \pm 0.11 (45)

in cingulate cortex and 0.33 ± 0.13 in temporal cortex, demonstrating a limited presence of LTS histopathology in these two brain regions at this stage of LB disorders. In fact, 10 of 15 stage IIb cases had no LTS pathology in temporal cortex, whereas six cases had no LTS pathology in cingulate cortex. The LTS scores remained no greater than 1 in temporal cortex as the disease progressed to Unified stage III. Conversely, the majority of stage III cases had LTS density scores greater than 1 in cingulate cortex. These data show that LTS pathological development in temporal cortex is behind cingulate cortex, which is consistent with our previously published observations and classification system that showed appearance of LTS pathology in cingulate cortex preceding neocortical LTS in a large studied population (7).

Levels of $p\alpha$ syn and $t\alpha$ syn, and their ratios in cingulate and temporal cortices

The main goal of this study was to determine the temporal relationship between the spread of LTS histopathology and changes in the levels of $p\alpha$ syn and $t\alpha$ syn in RIPA buffer brain extracts. These α syn proteins were measured by Western blotting with a serine 129 phosphorylation-specific antibody and an antibody-detecting epitopes before serine 129, respectively; then normalized by the levels of β actin in each sample. A panel of Western blot images of $p\alpha$ syn, $t\alpha$ syn and β actin from representative cases across all Unified LTS stages is shown in Figure 2A. This figure shows the arrangement of cingulate ("C" in the graph) and temporal cortical ("T" in the graph) samples from the same autopsy case in adjacent lanes on the same gel. Strong immunoreactivity for $p\alpha$ syn protein was observed in the samples from most of stage IV and some stage III cases.

The intensities of full-length $p\alpha$ syn- and $t\alpha$ syn-immunoreactive bands at 18 kD, based on the molecular weight marker run simultaneously, were quantified. The $p\alpha$ syn and $t\alpha$ syn proteins were detected on separate membranes to avoid overlapping of the band images caused by minor differences in molecular weights between $p\alpha$ syn and $t\alpha$ syn proteins. To assess if there would be a gel-to-gel discrepancy occurred at different steps of the procedure, same samples were repeatedly run on different gels. Our results showed no evidence of gel-to-gel differences.

We first used one-way ANOVA to determine if the levels of $p\alpha$ syn and $t\alpha$ syn were statistically different between samples from different Unified stages. The results were shown in Figure 2B–G. There

were no significant changes detected between stages zero, I, IIa, IIb and III. A drastic increase in the levels of $p\alpha$ syn and the ratio of $p\alpha$ syn to $t\alpha$ syn was detected between stage IV samples and the samples of other stages ($P < 0.05$). From stage III to stage IV, there were approximately 1.9-fold and 2.1-fold increases in $p\alpha$ syn, and 1.4-fold and 1.1-fold increases in the ratios in cingulate and temporal cortices, respectively. Because we were interested in determining the changes in $p\alpha$ syn levels at earlier disease stages, we reanalyzed the data by excluding stage IV cases. This time, the analysis showed significant twofold and 1.5-fold increases in cingulate and temporal cortices in stage III, respectively, from the earlier stages.

Because our stage IV subjects consisted of more DLB cases than other disease categories (27 DLB, nine PDAD and seven pure PDD), we determined whether there were disease-associated differences in the levels of $p\alpha$ syn in this particular stage. One-way ANOVA results showed that the levels in DLB cases were significantly greater than the levels in other two disease groups, and this observation was found in cingulate (DLB: 2.33 ± 0.21 ; PDAD: 1.08 ± 0.33 , pure PDD: 1.12 ± 0.27 ; $P < 0.05$) and temporal cortices (DLB: 1.88 ± 0.24 ; PDAD: 0.908 ± 0.20 ; Pure PDD: 0.767 ± 0.22 , $P < 0.05$). We then examined whether this would also occur in stage III subjects with DLB. There were eight DLB cases in stage III group, in addition to 12 pure PDD, eight ADPD and five PD cases. There were no disease group-associated differences detected in both brain regions in stage III cases. To determine whether DLB cases were responsible for the statistical significances in the $p\alpha$ syn levels analyzed by Unified stages, we analyzed the data again by excluding all of the DLB cases from stages III and IV. The results showed that the significant increases in $p\alpha$ syn in these stages remained.

The levels of $t\alpha$ syn did not change with stages in both brain regions. Thus, the changes in the levels of $p\alpha$ syn are accountable for the changes in the ratios. When two-way ANOVA was used to assess whether there were effects of LTS histopathological stages and brain regions on the levels of $p\alpha$ syn, the results showed significant effects caused by the LB stages, but not by brain regions. No interaction between the effects of LB stages and brain regions were detected.

Relationship between the biochemical measures in cingulate and temporal cortices

We then determined if the levels of $p\alpha$ syn and $t\alpha$ syn, and the ratios of $p\alpha$ syn to $t\alpha$ syn were correlated between two cortical regions

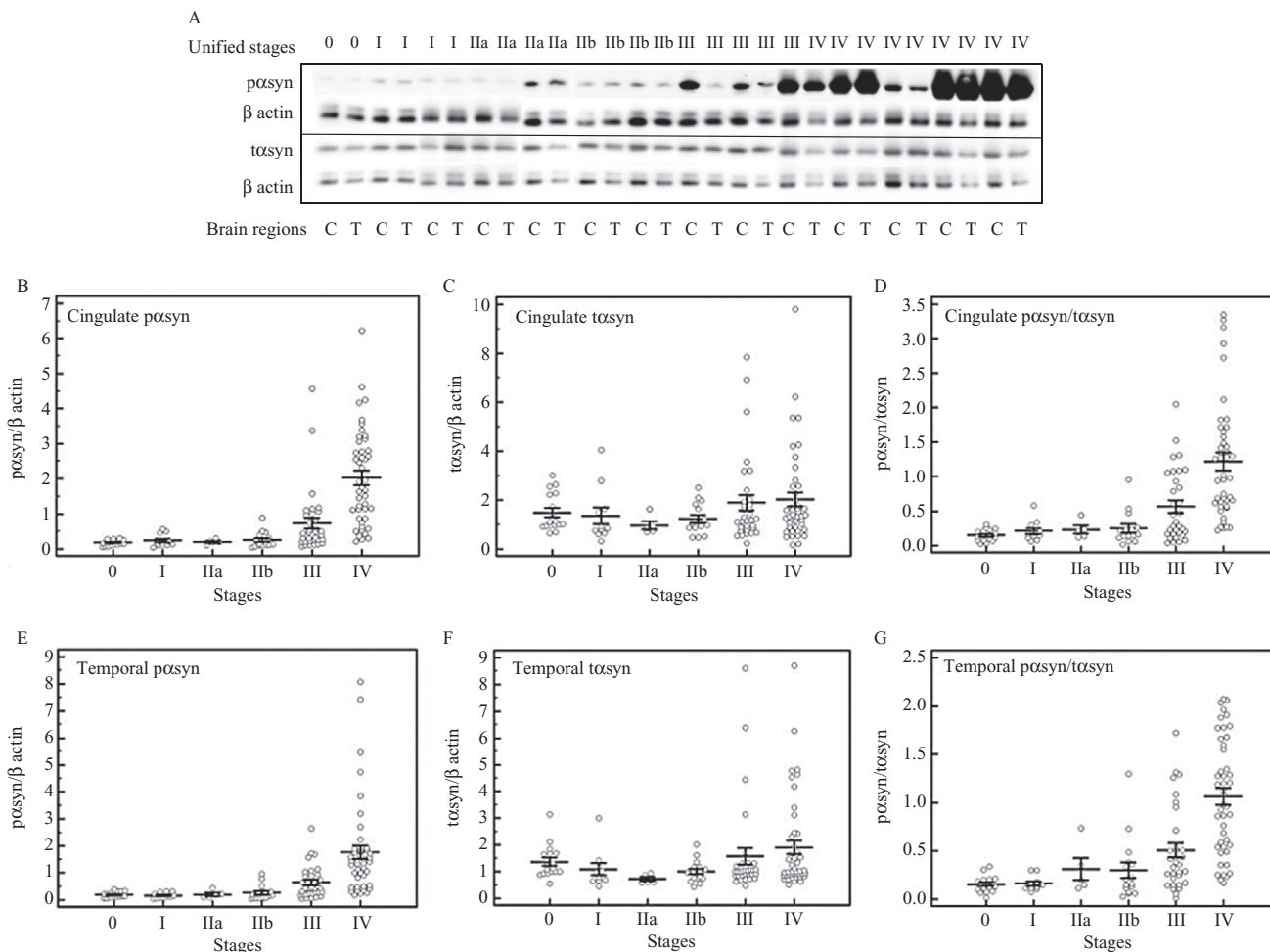


Figure 2. Levels of $p\alpha\text{syn}$ and total αsyn ($t\alpha\text{syn}$), and their ratios according to Unified stages. **A.** Immunoblot images of $p\alpha\text{syn}$ and $t\alpha\text{syn}$ from cingulate (C) and temporal (T) cortical samples across Unified stages (0, I, IIa, IIb, III and 4), the bars in the graphs represent mean values and standard errors of $p\alpha\text{syn}$ in cingulate and temporal cortices (**B** and **E**) $t\alpha\text{syn}$ in cingulate and temporal cortices (**C** and **F**); and the ratios of $p\alpha\text{syn}$ to $t\alpha\text{syn}$ in cingulate and temporal cortex (**D** and **G**).

using a linear regression model. Highly significant ($P < 0.001$) positive linear relationships were detected between these two brain regions in all of the three measures ($R^2 = 0.4643$ for $p\alpha\text{syn}$, $R^2 = 0.7497$ for total αsyn and $R^2 = 0.4295$ for the ratios).

To determine how well the $p\alpha\text{syn}$ levels in cingulate and temporal cortices correlated with LTS density scores, we performed Spearman’s rank correlation analysis. The correlation coefficients Rho showed that across four Unified stages, cingulate and temporal $p\alpha\text{syn}$ levels correlated very well with LTS scores not only within the same brain regions, but also in other brain regions that are affected at earlier stages, including the olfactory bulb, substantia nigra and amygdala (Table 3). The correlation coefficients between expression levels of $p\alpha\text{syn}$ and LTS scores are 0.732 ($P < 0.0001$) in cingulate cortex and 0.711 ($P < 0.0001$) in temporal cortex.

We have already shown that $p\alpha\text{syn}$ levels in cingulate and temporal cortical regions significantly elevated at more advanced Unified stages. We then analyzed the levels of $p\alpha\text{syn}$ by LTS histopathology density scores (0 to 4) in cingulate and temporal cortices. The results were shown in Figure 3A (cingulate cortex) and 3B

(temporal cortex). One-way ANOVA showed that in the subjects with cingulate LTS density scores = 3 and 4, the levels of $p\alpha\text{syn}$ within the same brain region were significantly higher than lower LTS density scores ($P < 0.05$), while there were no difference in levels of $p\alpha\text{syn}$ between the samples with LTS scores = 3 and 4. There was also no difference in $p\alpha\text{syn}$ levels between LTS scores = 0 and 1. However, when LTS score = 2, the levels of $p\alpha\text{syn}$ were not different from the samples with LTS = 0 and 1, but significantly lower than the samples with LTS = 3 and 4. In temporal cortex, the levels of $p\alpha\text{syn}$ were significantly elevated in the samples with LTS scores > 0 than in the samples with LTS scores = 0 ($P < 0.05$).

The expression levels of biochemically measurable and the histologically detectable $p\alpha\text{syn}$ do not always coincide in the studied subjects. For examples, 25%–30% of stage III cases and 5% of stage IV cases had $p\alpha\text{syn}$ levels lower than the mean values of $p\alpha\text{syn}$ in cingulate and/or temporal cortices in stage II subjects. We showed the LTS histopathological images from some of the representative cases, which had discrepancy between biochemical and

Table 3. Spearman's rank correlation coefficients between Unified staging LTS density scores and the levels of phosphorylated α synuclein in cingulate and temporal cortices.

Brain regions	Phospho- α synuclein levels in cingulate cortex (Rho, significance level)	Phospho- α synuclein levels in temporal cortex (Rho, significance level)
Olfactory bulb		
LTS density scores	0.412, $P < 0.0001$	0.411, $P < 0.0001$
Substantia nigra		
LTS density scores	0.623, $P < 0.0001$	0.589, $P < 0.0001$
Amygdala		
LTS density	0.476, $P < 0.0001$	0.477, $P < 0.0001$
Trans-entorhinal		
Cortex LTS density scores	0.548, $P < 0.0001$	0.560, $P < 0.0001$
Cingulate		
Cortex LTS density scores	0.732, $P < 0.0001$	0.660, $P < 0.0001$
Temporal		
Cortex LTS density scores	0.745, $P < 0.0001$	0.711, $P < 0.0001$
Parietal		
Cortex LTS density scores	0.722, $P < 0.0001$	0.677, $P < 0.0001$
Frontal		
Cortex LTS density scores	0.701, $P < 0.0001$	0.660, $P < 0.0001$

histological abundance in Figure 4. LTS histopathological profiles detected with p α syn antibody were photographed from 5- μ m paraffin sections cut from cingulate and temporal cortices from the representative cases that showed agreement or discrepancy in two types of measures of p α syn (Figure 4A–H). The p α syn immunoreactivities appeared in dark blue color in the images. The LTS scores and p α syn level were written side by side with each histological image. Case 1 had LTS histopathology scores = 4 in cingulate cortex and LTS scores = 3 in temporal cortex, case 2 had LTS score = 4 in cingulate cortex and LTS score = 3 in temporal cortex,

whereas case 3 and case 4 all had LTS score = 1 in both cingulate and temporal cortices. Although case 1 had higher biochemically measurable p α syn level than case 2, there is hardly any difference in the abundance of LTS pathology (Figure 4A–D). Cases 3 and 4 had the same LTS scores in both cingulate and temporal cortices, but case 3 had higher biochemically measurable p α syn than case 4. Nevertheless, there is no observable difference in the presence of LTS pathology between the cases 3 and 4 (Figure 4E–H).

We then determined whether the levels of p α syn in cingulate and temporal cortices is associated with the presence of AD pathology

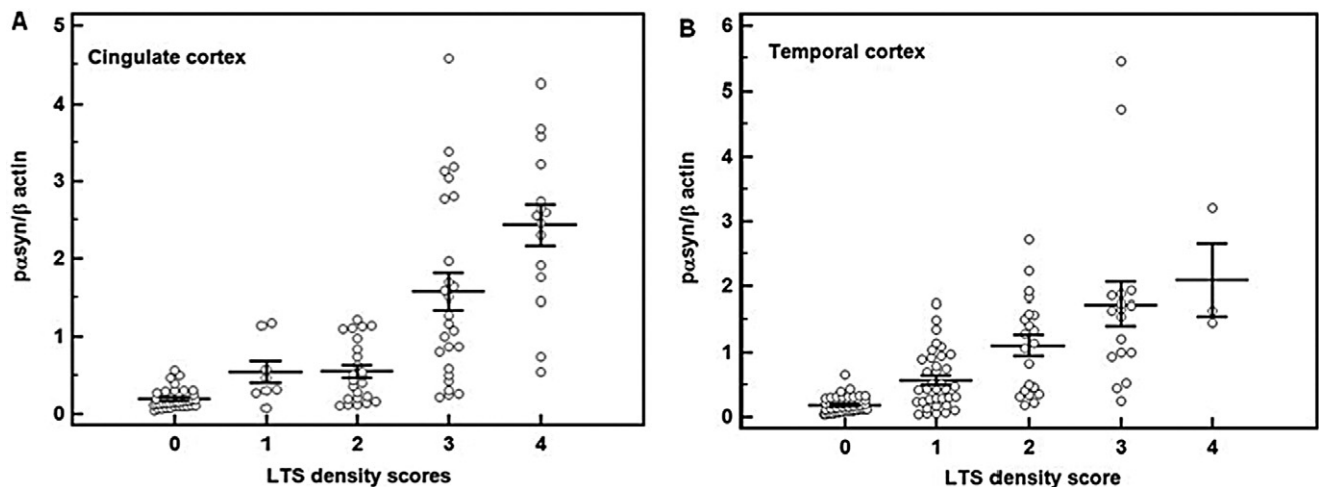


Figure 3. Increases in the levels of p α syn with LTS histopathological scores. The biochemical levels of p α syn are grouped by LTS histopathological scores 1 to 4. Horizontal lines indicated means and the vertical bars represent standard errors. Each circle in the graph represents the data from one individual. One-way ANOVA showed that in the cases that cingulate LTS density scores = 3 and 4, the levels of p α syn within the same brain region were significantly higher than lower LTS density scores ($P < 0.05$), while there were no difference in levels of p α syn

between the samples with LTS scores = 3 and 4. There was also no difference in p α syn levels between LTS scores = 0 and 1. When LTS score = 2, the levels of p α syn were not different from the samples with LTS = 0 and 1, but significantly lower than the samples with LTS = 3 and 4. In temporal cortex, the levels of p α syn were significantly elevated in the samples with LTS scores > 0 than the samples with LTS scores = 0 ($P < 0.05$).

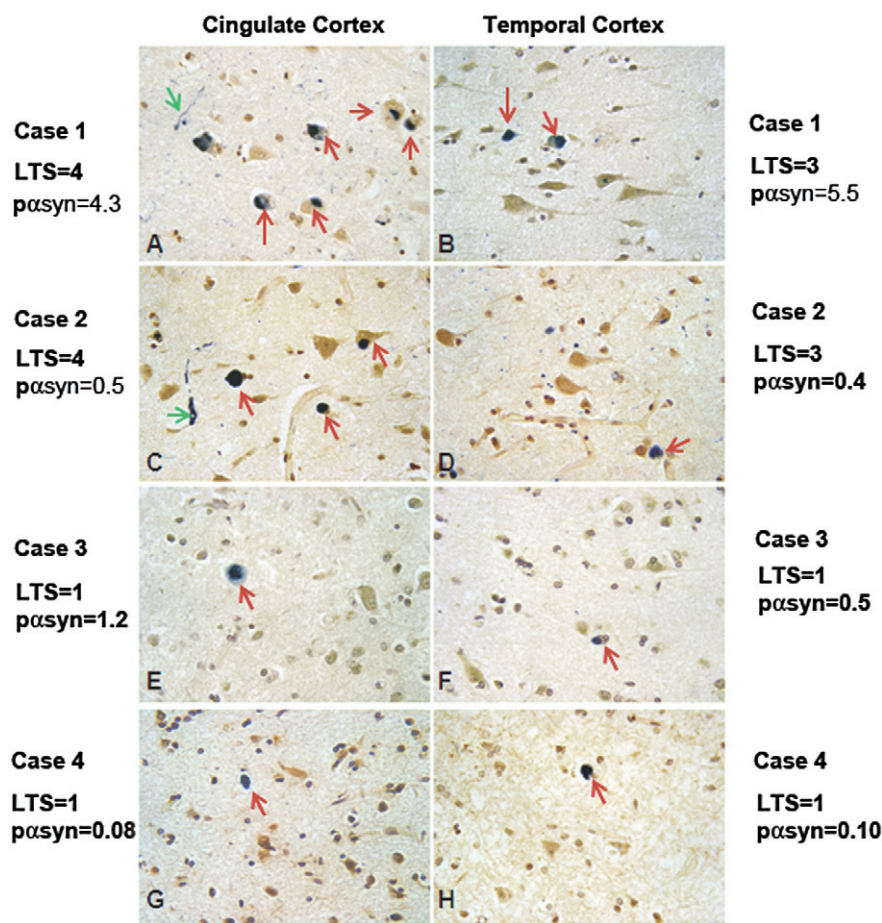


Figure 4. Representative images of $p\alpha syn$ immunoreactivities in cingulate and temporal cortices. Four representative cases were documented for the abundance of $p\alpha syn$ immunoreactivities shown in dark blue color. Images from 5 μm paraffin sections from cingulate and temporal cortices were taken with CCD camera and a 40X objective on a light microscope (Olympus). The LTS scores and $p\alpha syn$ level were written at the sides of each histological image. Case 1 had LTS histopathology scores = 4 in cingulate cortex and LTS scores = 3 in temporal cortex, case 2 had LTS score = 4 in cingulate cortex and LTS score = 3 in temporal cortex, case 3 and case 4 all had LTS score = 1 in both cingulate and temporal cortices. Although case 1 had higher biochemically measurable $p\alpha syn$ level than case 2, there is hardly any difference in the abundance of LTS pathology in the tissues as shown in Figures A–D. Cases 3 and 4 had the same LTS scores in both cingulate and temporal cortices, but case 3 had higher biochemically measurable $p\alpha syn$ than case 4. Nevertheless, there is no observable difference in the presence of LTS pathology between the cases 3 and 4 (Figures E–H).

especially in the cases that LTS pathology is absent. The AD pathology was assessed by total tangle scores, Braak’s stages and plaque density scores obtained according to our standard histological procedure. We performed Spearman’s rank correlation analysis in the subset of the cases that has no LTS pathology. We found that there was no association between the levels of $p\alpha syn$ in either cingulate or temporal cortices and the AD pathologies in these cases. Thus, it is likely that the expression of $p\alpha syn$ in these cases were not affected by AD neuropathology.

Increases in the levels of $p\alpha syn$ occur at early stages of Lewy pathology formation

To further investigate how early the levels of $p\alpha syn$ in cingulate and temporal cortices are altered during disease progression, we focused on the relationship of $p\alpha syn$ with the LTS density scores in brain regions and stages that are selectively affected early in the disorders, namely, olfactory bulb and substantia nigra. We compared the $p\alpha syn$ levels from subjects with LTS density scores graded as 0 or 1 in olfactory bulb and substantia nigra, and 0 in cingulate and temporal cortices.

A nonparametric test was performed to compare the samples derived from subjects who were at Unified LB stages 0–II. These subjects were grouped by LTS density scores, 0 or 1 in olfactory bulb or substantia nigra, and 0 and 1–2 in amygdala. As shown in

Figure 5A–F, the levels of $p\alpha syn$ were significantly elevated in cingulate and temporal cortices when LTS density scores increased from 0 to 1 in olfactory bulb and substantia nigra. The change in $p\alpha syn$ levels between the cases with amygdala LB score 0 and greater than 1 were significantly different in the temporal cortex samples, but not in the cingulate samples. This trend continues as the LTS scores in olfactory bulb and substantia nigra increased to 2; there were more than twofold increases observed in $p\alpha syn$ levels in both brain regions (data not shown).

DISCUSSION

Previously, the distribution patterns and abundance of histological deposits of immunoreactive αsyn in olfactory bulb, brain stem, limbic and neocortex have been used to describe the progression of PD and DLB, two major diseases caused by Lewy-type α -synucleinopathy. However, a new Unified Staging System for Lewy Body Disorders, based on $p\alpha syn$ immunohistochemistry, has demonstrated an improved efficacy for the classification of LB disorders. In this system, the grading of $p\alpha syn$ -containing LTS histopathology was conducted in 10 brain regions. The LTS histopathology begins in olfactory bulb at Unified stage I, followed by either brain stem (stage IIa) or amygdala (stage IIb) and by stage III, the LTS histopathology coexists in both brain stem and limbic regions. The LTS histopathology spread to neocortical

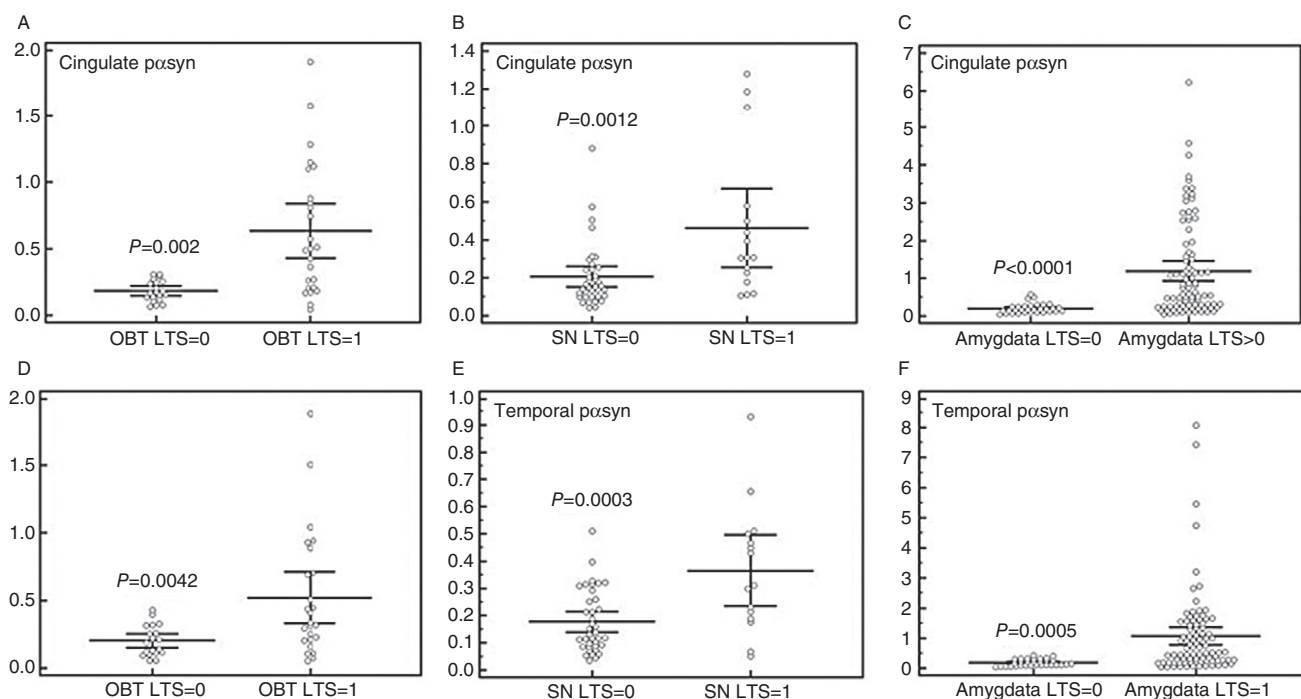


Figure 5. Levels of $p\alpha$ syn in cingulate and temporal cortices according to Lewy-type synucleinopathy (LTS) density scores in olfactory bulb, substantia nigra and amygdala. Bar graphs represents mean values and standard errors of $p\alpha$ syn in cingulate (A, B, C) and temporal (D, E, F) cortices according to LTS density scores in olfactory bulb (OBT) (A and D), substantia nigra (SN) (B and E) and amygdala (C and F).

regions in stage IV. This new system has demonstrated that the deposition of α syn phosphorylated at serine 129 is a valid and sensitive marker of LTS pathological progression in LB disorders (7).

The present study analyzed RIPA buffer-soluble α syn and $p\alpha$ syn in cingulate and temporal cortices from subjects across Unified LTS stages to determine whether the levels of $p\alpha$ syn are altered prior to its presence in LTS histopathology in LB disorders. Biochemical methodology is effective for detection of $p\alpha$ syn protein, which is not in LB pathologies (2, 43). The increase in $p\alpha$ syn expression could be independent of the presence of LTS histopathology as nonaggregated $p\alpha$ syn could be localized in synapse-enriched fraction (43).

We chose to study cingulate (belong to limbic system) and temporal cortices because these two brain regions have absent or rare LB pathology at early disease stages such as stages I and II, thus they are suitable for inferring a possible temporal relationship between the biochemical and histological changes of $p\alpha$ syn at early stages of diseases. Our results showed that $p\alpha$ syn could be detected in cingulate and temporal cortices in normal controls and across four Unified stages. Our results also showed that $p\alpha$ syn expression levels in both cingulate and temporal cortices are significantly elevated as Unified stages advance from II to III, and also as early as the LTS histopathology density score increased from 0 to 1 in temporal cortex as well as from 2 to 3 in cingulate cortex. Approximately 70% of our stage III cases have cingulate LTS density scores 2 and 3, while their temporal LTS scores are 1. These results suggested that biochemically measurable $p\alpha$ syn in temporal cortex could be more sensitive than cingulate cortex for reflecting the

increases in LTS histopathology within same brain region during these stages of LTS development.

Our second major finding was obtained from exploring how early the biochemical levels of $p\alpha$ syn in cingulate and temporal cortices were altered in relation to LTS pathology developed in the brain regions that are affected at early stages of LB disorders. We compared the $p\alpha$ syn levels from subjects with LTS density scores graded as 0 or 1 in olfactory bulb and substantia nigra, and 0 in cingulate and temporal cortices. Patients with olfactory bulb LTS density score = 1 were either at Unified stages I or II, whereas patients with substantia nigra LTS score = 1 were usually at stage IIa. Our results showed biochemically measurable $p\alpha$ syn levels were significantly elevated as LTS density scores in olfactory bulb and substantia nigra increased from 0 to 1, indicating that changes of $p\alpha$ syn levels in cingulate and temporal cortices coincided with the early appearance of the LB pathology in olfactory bulb and substantia nigra, even though histologically demonstrable LTS was lacking in these two cortical regions. This finding raises the question whether phosphorylation modification is a widespread event across brain regions, even when LTS histopathology has only limited appearance in olfactory bulb, brains stem or amygdala.

Although our data demonstrated phosphorylation of α syn in cingulate and temporal cortices as an early abnormality in LTS-associated disorders, we need to point out the existence of the variability in the data obtained in this study. For example, approximately 25%–30% of stage III cases had $p\alpha$ syn lower than the mean values of stage II samples in one or both of the brain regions. From demographical features and histopathological measures of AD and PD that were available from these cases, we could not identify the

specific causes that could have made these cases to express lower levels of p α syn. Current knowledge on the dynamics between increases in p α syn compartmentalized in cytoplasm and synapses, and formation of Lewy bodies and neurites are still not clearly understood. The possibility for the coexistence of biochemical and/or pathological modifiers in favor of pathological formation in these cases could not be excluded.

It has been demonstrated that AD pathologies have synergistic effects on the formation of LTS pathologies and it is frequent to find concurrent AD pathology in LB disorders (7, 9, 18, 19, 41). Among the molecules that have been shown to interact with α syn and p α syn, amyloid beta (A β) peptide and phosphorylated tau are on top of the list. Phosphorylated tau was shown to be increasingly co-localized with p α syn in the synaptic-enriched fractions (43). A β peptide increased intraneuronal accumulation of α syn in cell culture and transgenic mouse model (37). Transgenic mice over-expressing α syn increased accumulation of A β as well as ApoE (24). We have assessed the cases that did not have LTS pathology in cingulate and temporal cortices and the cases that had higher LTS but low biochemical values of p α syn for the effects of AD pathology. In general, we did not detect an association of AD hallmark pathologies with the levels of p α syn in these cases. However, we did not measure the biochemical levels of A β and phosphorylated tau in these brain samples, we could not conclude if these proteins had influence on the levels of p α syn in our studied subjects.

Evidence has indicated that phosphorylation at serine 129 could play an important role in regulation of oligomerization, fibrillogenesis and aggregation of α syn (17, 25, 27, 33, 44, 45, 49, 52). Increased expression of monomeric p α syn promotes oligomerization of p α syn, which causes neurotoxicity and synaptic damages (16, 51). Although the effects of p α syn on cellular functions have not been completely characterized, a recent study using a molecular pull-down technique for proteomic analysis showed that p α syn interacted with a vast range of functional and structural proteins, including cytoskeletal proteins, cell-signaling proteins and molecules associated with synaptic vesicle trafficking (39).

If phosphorylation of α syn at serine 129 is indeed a critical step in the pathogenesis of LB disorders, then possible mechanisms leading to this, including increased activity of the specific kinases, reduced dephosphorylation capability and reduced degradation and clearance, could be important processes for therapeutic targeting. To this date, three families of kinases have been shown capable of *in vitro* phosphorylation of nonaggregated α syn at serine 129, including casein kinases (CK) 1 and 2, G-protein-coupled receptor kinases (GRK) leucine-rich repeat kinase 2 (LRRK2) and polo-like kinases (PLK) 1, 2 and 3 (38, 48, 54). Among these, PLK1, CK1 and CK2 have also been shown to phosphorylate aggregated α syn (41, 54, 55). *In vitro*, phosphorylation of α syn by PLK2 was shown to compromise the ability of α syn to regulate the activity of tyrosine hydroxylase (TH), the rate-limiting enzyme of dopamine synthesis (33). By inhibition of the enzyme that dephosphorylates TH, called phospho-protein phosphatase 2A (PP2A), α syn promotes the activity of the TH (47). The phosphorylation at serine129 of α syn reduced this function (33). One recent study in a transgenic mouse model over-expressing α syn showed that by enhancing phosphatase activity of the PP2A, the phosphorylation and aggregation of α syn could be reduced (32). As understanding of the mechanisms of phospho-

rylation and dephosphorylation of α syn is advancing, to clarify the mechanisms contributing to the accumulation of p α syn in LB pathologies, it is important also to examine the mechanisms for degradation of p α syn, which are currently unclear; the ubiquitin-independent proteasome pathway or a ubiquitin-dependent pathway after dephosphorylation have both been implicated (34).

Taken together, we have shown significant, Unified stage-associated increases in RIPA buffer-soluble p α syn in cingulate and temporal cortices; and we have also shown significant increases in these measures coincided with the appearance of the LTS histopathology in olfactory bulb and substantia nigra. This study support that the mechanisms driving the increases in phosphorylation of α syn are at work at early stages of LB disorders and the effects are more widespread than what have known previously, thereby justifies the importance of identifying the causes and mechanisms leading to aberrant phosphorylation of α syn in these conditions.

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