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α -Synuclein and astrocytes: tracing the pathways from homeostasis to neurodegeneration in Lewy body disease

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Abstract

α -synuclein is a soluble protein that is present in abundance in the brain, though its normal function in the healthy brain is poorly defined. Intraneuronal inclusions of α -synuclein, commonly referred to as Lewy pathology, are pathological hallmarks of a spectrum of neurodegenerative disorders referred to as α -synucleinopathies. Though α -synuclein is expressed predominantly in neurons, α -synuclein aggregates in astrocytes is a common feature in these neurodegenerative diseases. How and why synuclein ends up in the astrocytes and the consequences of this dysfunctional proteostasis in immune cells is a major area of research that can have far-reaching implications for future immunotherapies in α -synucleinopathies. Accumulation of aggregated α -synuclein can disrupt astrocyte function in general and, more importantly, can contribute to neurodegeneration in α -synucleinopathies through various pathways. Here, we summarize our current knowledge on how astrocytic α -synucleinopathy affects CNS function in health and disease and propose a model of neuroglial connectome altered by α -synuclein proteostasis that might be amenable to immune based therapies.

Keywords

α -synuclein; Lewy body; glial cytoplasmic inclusion; neurodegeneration; exosome; tunneling nanotube; transmission; astrocyte heterogeneity; NAC domain; therapy

Introduction

Intraneuronal inclusions of α -synuclein (α Syn) protein, commonly referred to as Lewy bodies (LB) or Lewy neurites (LN), are hallmark pathologies in a group of neurodegenerative disorders collectively known as α -synucleinopathies [13, 164]. The gene coding for the α Syn protein, *SNCA*, is a genetic risk factor for both sporadic and familial forms of multiple α -synucleinopathies, including Parkinson's disease (PD) and Dementia with Lewy Bodies (DLB) [70, 139]. α Syn aggregates are found in different cell types as well as in different brain regions across the spectrum of α -synucleinopathies. For example,

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intraneuronal LB and LN pathology are frequently observed in PD, PD with dementia (PDD), incidental Lewy body diseases (iLBD) and DLB (Figure 1a–d). In multiple system atrophy (MSA) patients, in addition to few neuronal cytoplasmic inclusions (NCI) of α Syn, α Syn inclusions are primarily found in oligodendrocytes and are referred to as glial cytoplasmic inclusions (GCI) or Papp-Lantos bodies [75] (Figure 1e). On the other hand, astrocytic inclusions of α Syn are invariant features in PD and DLB but observed less frequently in MSA (Figure 1a–c) [145]. The clinical and neuropathological heterogeneity in the spectrum of α -synucleinopathies is believed to result from a combination of various factors, including unique conformational variants of α Syn protein that might contribute to distinct clinical phenotypes [118, 121].

α Syn was originally identified from an immunosera raised against purified cholinergic vesicles of the Torpedo electric organ and was also detected in the nucleus [101], resulting in the portmanteau designation of ‘synuclein’ (protein present in the synapse and nucleus). α Syn has been shown to exist as an inherently disordered monomer [161] or more controversially, as tetramers [15]. The N terminus of α Syn forms amphipathic α helix structure, whereas the highly acidic C terminus, which exists as a random coil, stabilizes the structure by transiently interacting with the N terminus [18, 55]. The non-amyloid β component (NAC) domain found in the middle of the protein is hydrophobic and is required for α Syn to polymerize into amyloid fibrils [89]. The N terminus has a unique repeated motif of seven 11 amino acid residues and all the known PD-associated mutations (A30P, E46K, H50Q, A53E, A53T, G51D) tend to cluster in this domain. This motif is key to interaction between α Syn and membranes. The physiological function of α Syn remains highly debated, though several studies have shown that it plays a highly diverse role in the CNS - synaptic vesicle trafficking [116], lipid metabolism and membrane remodeling [115, 172], synaptic plasticity [59], molecular chaperoning [129] and mitochondrial membrane remodeling [111]. The natively unfolded structure of α Syn is determined by its low hydrophobicity and high net charge (pI of 4.7 at neutral pH). Destabilization of the native structure, possibly through post-translational modification or interactions with proteins or cell membranes, induces formation of insoluble oligomers or fibrils with highly ordered β -sheet conformation [62]. Phosphorylation at Serine 129 and C terminal truncation products have both been shown to be common species in insoluble LB, LN and GCI [5] and though controversial, some studies have shown that phosphorylation at Ser129 and C-terminal truncation can modulate α Syn fibrillization [54, 147].

Clinical research suggests that the predilection for astrocytic α Syn buildup in the temporal lobe of DLB and advanced PD patients may play a role in the symptomatic progression of these diseases and warrants further study to understand associated functional impairments [22]. It is generally believed that specific clinico-pathologic outcomes, representative of the heterogeneity observed in α -synucleinopathies, may be traced to the involvement of neuronal as well as glial α Syn pathology [119, 121, 122]. The biological mechanisms that trigger the normally soluble protein α Syn to aggregate into pathological inclusions, such as GCI, LB and LN, remains highly debated. Factors relating to the cellular environment, such as autophagic impairment, mitochondrial dysfunction, vesicular trafficking, oxidative stress, ER stress and inflammatory stress as well as factors related to α Syn metabolism, such as post translational modifications (nitrosylation or proteolytic processing) may promote α Syn

aggregation in neurons [26, 60, 82, 94, 109, 137, 147]. On the other hand, the provenance of astrocyte-resident α Syn remains unclear as α Syn is expressed predominantly from neurons in both human and mice. It is thought that astrocytes accumulate α Syn by scavenging extracellular α Syn released from the neurons through an active process or following the demise of neurons containing these pathological α Syn inclusions [93, 97, 130]. In some cases, astrocytes containing α Syn inclusions are found distal to extracellular LBs or dead neurons and sometimes even outnumber LB positive neurons, which may also suggest a de novo mechanism [22]. Observations in experimental animal models have also suggested that distinct α Syn conformers formed in the neurons are capable of templating normal soluble α Syn through a prion-like process [165]. Following a prion-like seeding process, pathological α Syn conformers can potentially be passaged through inter-neuronal or even neuroastroglial connections and result in astrocytic accumulation of pathologic α Syn [1, 98]. However, overall very little mechanistic details are known regarding the role of these ubiquitous cells in the pathogenesis of α -synucleinopathies. To understand whether astrocytes play a beneficial or detrimental role, we examined the literature on interaction between pathologic α Syn and astrocytes in the context of cellular models, in vivo experimentation, and human disease. In the next sections, we will discuss how α Syn broadly affects the immune system and proceed to understanding the specific ways that α Syn accumulation alters astrocyte function and CNS homeostasis.

α Syn activates the immune system

Innate immune activation is a cardinal neuropathological feature accompanying end-stage α -synucleinopathy in patients and preclinical models (reviewed in [4, 27]). Curiously, the association of viral infection with Parkinsonism symptoms in von Economo's and postencephalitic PD patients spurred the idea that peripheral immune activation can trigger PD type neurodegeneration [128]. One of the earliest neuropathological demonstrations of CNS immune activation in PD patients showed increased HLA-DR (human leukocyte antigen DR isotype) positive reactive microglia [102], though PD patients often do not show typical hypertrophic reactive astrocytes [145]. Interestingly, as discussed in more details in later sections, while subcortical protoplasmic astrocytes accumulate α Syn in PD, MSA patients do not typically show α Syn immunoreactive astrocytes [145]. Later studies showed that PD patients show elevated CSF levels of inflammatory cytokines such as IL-1 β and IL-6 [19]. More recent studies have used cutting edge imaging and immunophenotyping studies to explore the link between inflammation and α -synucleinopathies. DLB patients with mild impairment displayed increased gliosis (using [(11)C]PK11195 PET tracer) in specific areas of the brain such as the caudate nucleus and cuneus compared to patients with more severe impairment, suggesting that inflammatory changes may occur in the early stages of disease [153]. Concomitant with CNS changes, these patients showed elevated peripheral blood inflammatory cytokines, such as MIP-3, IL-17A and IL-2 [153]. This is consistent with a recent seminal demonstration that PD patients have a unique T cell signature [151]. This study identified IL5-producing CD4+ T cells and IFN- γ producing CD8+ cytotoxic T cells in PD patients that were reactive to native as well as protofibrillar forms of α Syn. Several genome wide association studies have unearthed a link between HLA locus (human leukocyte antigen locus encoding the major histocompatibility complex class II (MHC-II))

and PD, strongly linking immune response as a risk factor in α -synucleinopathies [34]. Together these results show a functional link between activation of innate and adaptive immunity and α -synucleinopathies.

As a typical damage associated molecular pattern (DAMP) containing moiety, α Syn aggregates or conformers has been shown to directly interact with astrocytes and microglia via pattern recognition receptors, including Toll like receptor (TLR) 4 in cell culture [35, 93, 127, 179]. While α Syn aggregates have been specifically identified in hypertrophic ‘activated’ astrocytes in patient brains, whether the astrocytic inclusions themselves directly compromise these cells and trigger further cell- and non-cell autonomous damage and intercellular α Syn transmission is hotly debated. α Syn induced chronic astrocyte activation can also lead to bystander pathologies, such as increased cytokine and chemokine expression, antigen presentation, Ca^{2+} flux and oxidative stress that can ultimately lead to maladaptive neuronal activity and neurodegeneration by non-cell autonomous mechanisms [6, 35, 93].

Based on neuropathological findings in limited cohort of PD patients [14, 24, 63, 170] and the abundance of non-motor prodromal symptoms in PD patients [12], the gut-brain axis in PD was proposed which hypothesizes that initial α Syn aggregation can occur in the myenteric plexus of the gut (and other mucosal surfaces) which then propagates along vagal projections to the brain in a prion-like manner. There is indirect evidence of the involvement of peripheral immunity and the gut microbiome composition in induction and propagation of PD pathologies (reviewed in [80]). Indeed, the composition of PD patient gut microbiome is different from age-matched controls [77, 138] and transplantation of PD patient gut microbiota in a mouse model enhances motor deficits [135]. This latter study also showed for the first time that altering the gut microbiome composition modifies α Syn pathophysiology in the CNS of mice [135]. Since α Syn is expressed in enteric neurons [16] and can be physiologically secreted via a conventional, endoplasmic reticulum/Golgi-dependent exocytosis in a neuronal activity-regulated manner [117], it is easy to envisage that α Syn present in the gut can promote gut inflammation and contribute to α Syn pathogenesis along the gut-brain axis. However, it remains to be seen whether astrocytes (or other immune cells) in the enteric plexus play additional role in gut inflammation or the induction and spread of α Syn pathologies along the gut brain axis.

Recent advances in single cell transcriptomics and mathematical modeling have revealed regional, functional and disease state-specific heterogeneity in brain resident astrocytes [78], suggesting that astrocyte diversity in different brain regions may also affect neural circuits differentially depending on the spatio-temporal context of the underlying α Syn pathology. Interestingly, identification of MHCI in a subset of neurons also infers that the neurons may not be immune privileged after all and may dynamically interact with alterations in glial cell function [42]. Thus it is possible that a specific combination of the cellular context [121], immune milieu [176] and/or α Syn conformers [118] will determine region-specific immune response including astrocyte function [154]. This complex inter-relationship between innate immunity, peripheral immunity and dysfunctional CNS proteostasis can likely alter disease progression and the phenotypic manifestation in different α -synucleinopathies. A mechanistic knowledge of the interaction between cell autonomous (intraneuronal protein

aggregation and related sequela) and non-cell autonomous (inflammation and glial response to α Syn accumulation) pathways in α -synucleinopathies can lead to potential disease modifying immunobiotherapies.

While there are excellent reviews that capture this complex interaction of immunity in α -synucleinopathies [4, 27], very little is known about how astrocytes specifically are involved in this process. In the next sections, we focus specifically on the role of astrocytes in the pathogenesis of α -synucleinopathies by examining the literature on interaction between pathologic α Syn and astrocytes.

Astrocytes: role in homeostasis and diseased brain

Astrocytes are the most numerous cell type in the brain; these “star” shaped cells use their thin processes to directly ensheath up to two million synapses each for the purpose of maintaining synaptic function [3]. At the synapse, astrocytes shuttle ions, neurotransmitters, and other species across their membranes to ensure optimal conditions for neurotransmission. Their additional functions include protection against metabolic and oxidative stress, nutrient transport across the blood brain barrier, production of a quasi-lymphatic system draining waste into the CSF, resolution of neural injury via glial scar formation, and even higher executive functions related to synaptic plasticity [47, 166]. Impairment of these processes is evident in post mortem brains of neurodegenerative proteinopathies in general, with preclinical studies further suggesting that loss of astrocytic support may play a direct causal role in the neurodegeneration in these diseases [86, 120].

Astrocytes are a heterogenous group of cells that arise from neural stem cells controlled by complex series of developmental cues, much of which is still undetermined [107]. Broadly, there are two classes of astrocytes – protoplasmic astrocytes found in grey matter and fibrous astrocytes found in the white matter. There are other populations of astrocytes with specialized functions, for example, retinal Muller glia and cerebellar Bergman glia. In the context of α -synucleinopathies, the primary role of astrocytes in relieving oxidative stress, generating neurotrophic factors, modulating inflammation, and reducing pathogenic α Syn have been demonstrated in several preclinical studies, as we will discuss in details in the next sections. Developing the concept of immunoproteostasis, where immunity and α Syn protein homeostasis pathways are continually interacting with each other, we can envision a double-edged sword scenario where this interaction can result in harmful or beneficial outcomes. Further, it can be argued that intraneuronal Lewy aggregates are relatively harmless [88] and is a simple measure of cellular response to proteostasis imbalance [39] whereas the inappropriate astrocyte response to these dysfunctional neurons is what drives the inexorable pathological sequela including synaptic failure, neuronal death and clinical symptoms. This is consistent with recently described, though controversial, paradigm called gliotransmission by which neuronal activity induces astrocytes to secrete neuroactive substances into the synaptic cleft that might ultimately regulate brain organ health. It is also tempting to speculate that the subcellular localization of α Syn aggregates within astrocytes, compared to intraneuronal α Syn found within LBs and GCIs, may result in a new form of α Syn with unique properties such as different aggregation potential, prion-like conformational templating, and toxicity. Such unique forms or α Syn ‘strains’ are

increasingly being recognized as important determinants of disease pathogenesis [118, 121]. Another factor to consider is that the inherent heterogeneity in astrocytes (regional, developmental and functional; [78]) may play a role in generation of different α Syn strains, leading to specific disease progression scenarios in the brain. In the next sections, we will discuss the implications of these hypotheses and further introduce some recent paradigms that can have critical impact on therapeutic strategies.

Astrocytic α Syn inclusions are distinct from intraneuronal Lewy pathology

Aggregated α Syn have been reported in astrocytes of PD, DLB and MSA patients [79, 145] (Figure 1). On the whole, post-mortem studies show that the distribution of astrocyte α Syn pathology closely mirrors the appearance of intraneuronal LB pathology in the temporal and insular cortex in PD and DLB patients [22, 52]. In PD and DLB patients, most of the aggregated α Syn is observed in astrocytes in the white matter and in Bergman glia [110, 124] while some are noted in grey matter areas such as the temporal lobe, limbic areas, cortex and substantia nigra (Table 1). These astrocytic α Syn inclusions are characteristic of PD and DLB and rarely observed in MSA while being absent in other dementias and healthy controls (Table 1). Some studies have reported at least 60–100+ α Syn positive astrocytes within a single unilateral section from the midbrain or cortex of PD/DLB or other α -synucleinopathy patients, which was comparable to the levels of intraneuronal LB pathology within the same field of interest [66, 160, 168, 177]. The presence of α Syn positive astrocytes largely paralleled both the location and severity of neuronal LBs within the cortex or midbrain as noted by 5 studies examining LBD or PD cases [22, 66, 141, 145, 168]. In many cases, these astrocytes do not show hypertrophic morphology as would be expected in a typical reactive state, but instead are characterized as having many fine radiating processes [145, 159, 160, 168, 169]. In comparison with MSA where astrocytic α Syn is sparse but there is widespread gliosis [74], the presence of α Syn laden but non-reactive astrocytes in LBD/PD may suggest that the accumulated α Syn is impairing glial response. These astrocytic α Syn inclusions are ultrastructurally different from the typically densely packed neuronal LB pathology; astrocytic α Syn is diffusely distributed through the cell body and processes, with granular staining pattern reminiscent of vesicles or lysosomes [22]. Moreover, unlike the neighboring neurons with LBs that are readily characterized by H&E histology, silver staining or phosphorylated Serine 129 (pSer129)- α Syn immunoreactivity, astrocytic LBs do not share these histological properties. Indeed, the most critical roadblock in these post-mortem analysis appears to be accessing and identifying the specific α Syn species, especially when using formalin-fixed and paraffin-embedded human brain sections. A clear breakthrough was achieved when using antibodies to the central domain of α Syn, several groups could successfully demonstrate α Syn inclusions in astrocytes of PD/DLB patients (Table 1) [2, 7, 22, 65, 87, 110, 125, 143, 147, 159, 162, 163, 165, 171, 172, 181]. In addition, because of epitopes being potentially masked or modified in fixed brain tissues, astrocytic α Syn can be visualized only after using harsh antigen retrieval methods. Several studies used formic acid pretreatment to uncover epitopes in the central portion of α Syn (including the NAC domain) that seem to be specifically present in astrocytic α Syn inclusions in PD/DLB patients but absent in MSA patients or controls (Table 1) [22, 66, 87, 93, 141, 145, 156, 160, 177]. These same studies observed that there was little to no labeling

of astrocytic α Syn when N or C-terminal antibodies against α Syn were used, or when formic acid pretreatment was excluded [22, 87, 156, 160, 177]. This pattern was not seen with neuronal α Syn aggregates where all antibodies seemed to detect LBs equally [22, 86, 156, 160, 177]. Further, while neuronal LBs are positive for Gallyas-Braak silver staining, astrocytic α Syn inclusions were reported to be reactive to Gallyas-Braak silver only in a few studies [159, 160, 168, 169] (Table 1). In addition to being unreactive to N- and C-terminal α Syn antibodies, astrocytic α Syn does not colocalize with p62/SQSTM1 or ubiquitin, as is commonly observed with neuronal LBs [22]. Collectively, these data suggest that astrocyte-resident α Syn may have unique ultrastructure and have extensive modifications, including post-translational modifications and truncations. Such unique structural properties of astrocytic α Syn can have consequences relating to α Syn aggregation and prion-like transmission properties.

Based on the cumulative findings of the aforementioned studies, we generated a monoclonal antibody raised against the central portion of α Syn (antibody 3H11; residues 43–63 of α Syn abutting the NAC) and used it to demonstrate robust astrocyte-resident α Syn pathology in two LBD cases following antigen retrieval using formic acid (Figure 1a–c). In these LBD cases, dense astrocytic α Syn was particularly prominent in limbic structures of the temporal lobe such as the hippocampus and amygdala, where double labeling using the GFAP antibody with the 3H11 antibody confirmed the finding (Fig. 1c). This astrocytic α Syn is thought to be pathologic in nature, as no astrocytic α Syn staining was detected in the hippocampus of an Alzheimer's case (Figure 1f). Interestingly, by immunoblotting, 3H11 antibody recognized different PD-associated α Syn mutants differentially, indicating that some of these mutations disrupts the 3H11 epitope in α Syn (Fig. 1g–i). We observed α Syn reactive astrocytes not only in the direct vicinity of cortical LBs, but even extending further into the parenchyma where the majority of α Syn reactive cells appear to be astrocytes (Figure 1a). Whether astrocytic α Syn accumulation precedes the development of cortical LBs is still unknown but knowledge about the temporal aspect of cellular vulnerability to pathological α Syn will inform us on the etiology of α -synucleinopathies, including the prion-like properties of α Syn. Whether 3H11-immunoreactive astrocyte inclusions in the enteric or central nervous system track with the Braak pathological staging [23] will need to be empirically determined to understand how astrocytes contribute to α Syn pathogenesis. One feasible theory is that astrocytes initially function as scavengers of extracellular pathologic α Syn preceding the formation of neuronal LBs. It is possible that astrocytes containing these pathologic α Syn, having processes extending to hundreds of thousands of synapses, can subsequently contribute to spreading of α Syn fibrils themselves. This will be elaborated in subsequent sections.

In rodent transgenic models of α -synucleinopathy, α Syn immunoreactivity has been readily observed in neurons as well as astrocytes [82, 132, 133, 146, 148] (Figure 2). These astrocytic inclusions even occur when α Syn is expressed under a neuronal promoter ((Thy-1)-human [A30P] α Syn), suggesting that α Syn can probably be internalized by astrocytes and subsequently accumulate [97]. An important difference between these mouse models and human disease is that full-length aggregated α Syn is detectable in mouse astrocytes even without formic acid pretreatment, although it remains to be seen whether different pre-treatment(s) would uncover additional novel pathologies in mice. Therefore,

these transgenic mouse models likely represent a mix of features from both PD/DLB and MSA, but overall, they may prove useful in investigating differential evolution of inclusions in astrocytes compared with neurons.

Astrocytic α Syn pathology: questions on its origin and fate

The etiology and functional consequence of astrocytic α Syn remains unresolved. The field is generally constrained by a lack of systems level insights into neuronal function and cell type specific gene expression patterns that can shed light into astrocyte-mediated functional impairment, neurodegeneration and symptomatic progression of PD/DLB. A critical question in this context is whether astrocytes facilitate removal of toxic α Syn by phagocytosis [53] or do astrocytes directly contribute to the increasing neuronal proteostasis via uptake and subsequent prion-like propagation of the phagocytosed pathologic material? The latter premise that astrocytes can facilitate α Syn transmission comes from a transgenic mouse line that overexpresses human A53T α Syn from the astrocyte-specific GFAP promoter (GFAP-tTA/tetO- α -syn mice). In these mice, diffuse intraneuronal LBs are observed throughout the CNS suggesting that astrocyte to neuronal spread of pathologic α Syn had taken place [64]. Some plausible theories behind the origin of astrocytic α Syn pathology are 1) phagocytic or pinocytotic uptake of extracellular α Syn, 2) direct transfer from neurons through an active process, such as tunneling nanotubes or exosomal transfer or 3) de novo induction of α Syn pathology during pathogenesis. In this section, we will discuss the literal yin-yang of astrocyte scavenger function vis-à-vis its role in propagation of α Syn pathology.

Several in vitro studies have reported that astrocytes can rapidly ingest extracellular α Syn, within minutes of being exposed [25, 33, 83, 97]. All of these studies concluded that astrocytes internalize pathologic α Syn more readily than neurons, suggesting that astrocytes are primed to scavenge DAMPs, such as α Syn conformers. In vivo evidence of this phenomenon was reported in mice where overexpression of α Syn under neuronal Thy-1 or PDGF β promoters (A30P α Syn and wild type α Syn respectively) resulted in increased astrocytic α Syn, presumably following endocytosis from extracellular sources [93, 97]. Following endocytosis, the α Syn can be subsequently trafficked to the lysosomes [93, 97]. The process of α Syn internalization and cellular trafficking is thought to be different between neurons, microglia and astrocytes, which suggests that this might potentially lead to formation of unique α Syn conformers within different cell types. Neurons can internalize α Syn by interacting with cell surface heparin sulphates [67] or specific receptors such as Lag3 [100] or Na⁺/K⁺ transporting ATPase subunit α 3 [142], following which it can be either transported along microtubules as naked proteins or inside vesicles into the lysosomes. Microglia can engulf α Syn using TLR4 [53], resulting in NF κ B activation, release of pro-inflammatory cytokines and production of reactive oxygen species. Astrocytes, however, do not seem to require TLR4 for α Syn phagocytosis [53]. Further, astrocytes seem to process endocytosed α Syn differentially than neurons, as shown by a unique interactome revealed by mass spectrometry [143] and presence of partial breakdown products of α Syn that are reactive specifically to α Syn central domain antibodies (Figure 1, Table 1). It is possible that extracellular α Syn is cleaved by matrix metalloprotease MMP3 before astrocytic phagocytosis or processed by lysosomal cathepsins after ingestion, leading

to generation of truncated species [125, 152]. In vitro studies have shown α Syn aggregation follows different kinetics at normal pH (7.4) and at acidic pH (typically found in lysosomes and endosomes). At pH values below 6, α Syn aggregation is predominantly triggered via secondary nucleation changes (such as fragmentation and surface-assisted nucleation) rather than a simple elongation process [30]. Such secondary nucleation processes can lead to faster multiplication of aggregates and presumably aggravate disease pathogenesis. Therefore, endosomal trafficking of α Syn may result in an exacerbated neurotoxic phenotype since C-terminal truncated α Syn is known to fibrillize more rapidly than full length variants and cause more cytotoxicity [147].

Astrocytes and neurons are capable of undertaking bidirectional or unidirectional inter-cellular transport of α Syn through various mechanisms [167]. Both in vitro and in vivo studies have shown the presence of tunneling nanotubes (TNT) that can transfer α Syn between neurons and astrocytes [1, 46, 130]. TNTs are thin (50 – 200 nm in diameter), actin-rich membranous channels that can connect the cytoplasm of cells as far apart as 100 μ m. Lysosomal resident α Syn can traverse between neurons [1] or more intriguingly, stressed astrocytes with defective lysosomal function can barter ‘undigested’ α Syn with a healthy astrocyte in exchange for healthy functional mitochondria [130]. This could result in a double-edged sword scenario: infection of healthy astrocyte with potentially toxic α Syn conformers leading to prion-like transmission and functional recuperation of α Syn-damaged astrocytes with fresh mitochondria. Another mode of intercellular communication is through exosomes. Exosomes are small (40–100nm) vesicles that originate from multivesicular bodies of the endocytic pathway of neurons, astrocytes and microglia and are thought to be involved in clearance of cell-derived debris [29]. α Syn carrying exosomes can readily cause neuronal death in vitro [50]. Recent data also shows that brain-derived exosomes from DLB patients contain pathologic α Syn that can be internalized by both neurons and astrocytes leading to increased pathogenicity in mouse brains [112]. Interestingly, exosomal α Syn release is not reduced by blocking ER-Golgi transport mechanism, suggesting that α Syn containing exosomes can be extruded through non-conventional pathways [65, 71], such as exosome-associated exocytosis and/or exophagy [49]. Thus, so far, the data suggests that uptake of exosomes or vesicular α Syn by astrocytes can potentially have detrimental effects on disease pathogenesis in α -synucleinopathies by allowing transcellular propagation of pathologic α Syn and altering astrocyte homeostasis.

Among these theories, the last one, i.e., reactive de novo aggregation of α Syn is not fully supported by direct experimental evidence in vivo. α Syn immunoreactive protoplasmic astrocytes have been found in abundance in human brain regions typically devoid of LB, such as striatum and dorsal thalamus [22] which anecdotally supports this theory. However, the amount of endogenous α Syn protein that is typically expressed by glia is far less than what would be expected to be needed for successful de novo templated aggregation and transmission of pathology. In the healthy brain, α Syn is detectable in white matter astrocytes following immunohistochemical staining; however, this occurs only at very low levels and the tissue must be extensively pre-treated [108]. Immuno-EM studies have shown that the subcellular distribution of α Syn in these astrocytes tend to be diffuse throughout the cell body and processes and remain in association with mitochondrial membranes, ribosomes, and small vesicles [108]. In vitro evidence from astrocytic cell lines showed that only small

amounts of α Syn can be detected by both mRNA and protein assays [158]. Interestingly, later studies have shown a clear association between upregulation of synuclein levels and astrocyte activation status, at least in vitro. For example, α Syn is upregulated in cultured astrocytes when exposed to inflammatory cytokines [158] or during oxidative stress [37]. Therefore, although upregulation of endogenous astrocytic α Syn may contribute to disease, it is more likely that the majority of α Syn found within astrocytes in the diseased brain is mostly derived from neuronal origins.

Functional consequences of astrocytic α Syn: lessons from in vitro studies and transgenic animal models

The belief that astrogliosis is a passive bystander response to increasing intraneuronal α Syn proteostasis and the neurodegenerative cascade is now under debate. Recent data showing that astrocytes can actively communicate with neurons and alter neuronal plasticity and homeostasis has been critical in establishing the hypothesis that astrocytes may be more directly involved in α Syn pathogenesis [61].

The function of endogenous α Syn within normal astrocytes is unclear, although there is some evidence that α Syn is involved in astrocytic fatty acid metabolism [32]. However, later studies have shown a clear association between upregulation of synuclein levels and astrocyte activation status, at least in vitro [161]. Although upregulation of endogenous astrocytic α Syn may contribute to disease, it is more likely that the majority of α Syn found within astrocytes in the diseased brain is derived from neuronal origins, as has been discussed in the previous section. Studies in PD brains have shown that midbrain astrocytes undergo apoptosis in addition to nigral neurodegeneration [73, 85, 106], suggesting that non-cell autonomous events have a profound effect on the neurodegenerative cascade. In cultures treated with pathologic α Syn or human brain-derived LBs, astrocytes display morphological changes consistent with activation, as well as mitochondrial fragmentation, autophagic impairment, impaired Ca^{2+} flux, sensitization to oxidative stress and death [6, 25, 33, 64, 83, 97, 104, 149]. The mechanism by which aggregated α Syn induces toxicity in astrocytes appears to be through a combination of mitochondrial dysfunction and impaired autophagy and mitophagy leading to oxidative stress and apoptosis, as demonstrated by in-vitro studies (Figure 3) [25, 33, 51, 97]. It is possible that several familial PD risk genes that lead to Parkinsonism without necessarily causing α Syn accumulation have important roles in mitochondrial function and oxidative stress, suggestive of commonalities in etiology of Parkinsonian syndromes (Figure 3).

In animal models, neuronal overexpression of human α Syn or injection of human brain materials leads to astrocyte activation, as evident by increased GFAP, S100 β or vimentin immunoreactivity (reviewed in [84]). In a peripheral to central transmission model of α Syn pathology expressing human [A53T] α Syn from prion promoter (Line M83; B6;C3-Tg(Prnp-*SNCA**A53T)83Vle/J), we observed motor neuron death and pSer129- α Syn pathology prior to the appearance of astrogliosis (Figure 2a) [132, 148] as in human PD where astrocytic hyperactivation usually follows nigral degeneration [163]. Intriguingly, in Line M83 mice, we observed widespread nigral astrocytic pSer129- α Syn pathology while

sparing the dopaminergic neurons following intracranial seeding of α Syn aggregates (Figure 2c). Therefore, the relationship between astrocytic α Syn toxicity and neuronal dysfunction remains unclear. This aspect was also investigated in transgenic mice expressing α Syn exclusively from astrocytes. In an inducible transgenic mouse model expressing human [A53T] α Syn specifically from astrocytes (GFAP-tTA/tetO- α -syn line), extensive astrocytic death, mitochondrial dysfunction and dopaminergic neurodegeneration was observed [64]. However, another transgenic model with dramatic levels of astrocytic expression of α Syn (human [A30P+A53T] α Syn driven by chicken β actin promoter; BAsyn/PaKO line) did not show overt neurodegeneration in the substantia nigra but had profound motor deficits, suggesting that impairment of astrocytes can lead to neuronal dysfunction even without overt cell death [104]. Interestingly, a specific subtype of astrocyte, A1 astrocyte, has been identified in the nigra of PD patients [95] and a prion promoter driven human [A53T] α Syn transgenic mouse model (C57BL/6; Prnp-*SNCA**A53T [182]) that has been postulated to drive neurodegenerative cascade in α -synucleinopathies. These A1 astrocytes are devoid of their usual ability to promote neuronal outgrowth, synaptogenesis and phagocytosis, and further can lead to death of neurons and oligodendrocytes. In addition, blocking the formation of these reactive astrocytes leads to elongation of life in the human [A53T] α Syn mouse model of α -synucleinopathy [182]. Overall, though the A1/A2 characterization of astrocyte functionality has provided a simplistic and convenient platform for initiating our understanding into astrocyte involvement in α -synucleinopathies, future transcriptomic and functional studies should be conducted to reveal physiologically-relevant disease-associated profiles that can shed light on astrocyte heterogeneity and selective vulnerability in α -synucleinopathies.

Recent work has illuminated that one of the pathways underlying the stereotypical spread of α Syn pathology is by prion-like transmission of α Syn in transgenic mouse models of α -synucleinopathy injected with α Syn fibrils or disease associated brain extracts (reviewed in [165]). While some groups reported mostly intraneuronal α Syn pathology during this transmission process in human [A53T] α Syn transgenic mice [99], other groups including ours reported robust astrocytic α Syn pathology in these mice suggesting that astrocytes can also contribute to α Syn transmission [131, 146, 148]. Interestingly, we observed that the majority of the pathological α Syn in the substantia nigra of these mice were localized to astrocytes, and none to dopaminergic neurons (Figure 2C). This might imply that astrocytes can modify disease pathogenesis in this transmission model either by scavenging and having a neuroprotective function or as an unwitting accomplice in α Syn pathogenesis. Other groups have shown that astrocytes can take up these α Syn assemblies and direct these to lysosomes for degradation in organotypic slice cultures from wild type mice [98], though chronic uptake may also lead to cellular damage and inclusion formation in primary cultures [97]. Importantly, whether astrocyte resident pathological α Syn has unique strain-like properties distinct from neuronal α Syn inclusions, as observed in primary astrocytes in vitro [131], remains a matter of debate [103].

Astrocytes can protect dopaminergic neurons by improving mitochondrial function [48] and simultaneously can also regulate α Syn pathogenesis in the synaptic clefts [97]. Additionally, astrocytes participate in dopamine metabolism through uptake from the synaptic cleft [157] and impairment of this process may lead to increased cellular stress through buildup of

reactive dopamine products [105, 173]. However, whether astrocytic processes can potentially modulate clinical symptoms by regulating formation and clearance of dopamine- α Syn adducts or α Syn induced dopamine oxidation in the extrasynaptic space, remains to be investigated.

Cumulatively, the current evidence suggests that various pathologic forms of α Syn are toxic to astrocytes through a pathway incorporating impaired autophagy and resulting mitochondrial dysfunction from what may be mitophagy dysregulation [51]. Overall, an idea that is supported by clinical and preclinical evidence is that pathologically modified forms of α Syn may slowly accumulate in astrocytes following passive uptake or through some form of extrusion from neurons. Ultimately, this might reach a threshold where astrocytes are too impaired to respond to toxic build-up of LB pathology. One might speculate that α Syn overload of astrocytes would theoretically result in the loss of the preferential uptake of α Syn by astrocytes causing a shift towards trans-synaptic propagation and microglial involvement which would lead to dysfunctional immune function and worsening of disease.

Gliotransmission in α -synucleinopathies: possible co-morbid factor in α Syn pathogenesis

Metabolic exchanges across the astrocyte-neuron synapses or tripartite synapses are critical in maintaining CNS homeostasis [123]. Astrocytes are responsible for scavenging glutamate from the synaptic cleft as well as supplying lactate for neuronal activity through a process called gliotransmission. α Syn, being an abundant pre-synaptic protein, can potentially alter the function of the tripartite synapse formed between astrocytes and the pre- and post-synaptic densities. Recent studies have reported toxic gain of function of α Syn at the synapse by redistribution of SNARE protein, affecting dopamine transporter function and increasing glutamate excitotoxicity [17, 58, 136]. Overexpression of human [A53T] α Syn in the GFAP-tTA/tetO- α -syn transgenic mice led to reduced glutamate transporter expression from the astrocytes, suggesting a direct role of α Syn in neurometabolic coupling to astroglia [64]. Whether this altered gliotransmission plays a role in α Syn pathogenesis remains an intriguing premise that needs to be validated in preclinical mouse models, but could be a modality by which impaired astrocytes potentially contribute to clinical symptoms (Figure 3).

Astrocyte function can be altered by PD-associated genetic risk factors

Aside from *SNCA*, several other genes have been identified to be key risk factors in PD (reviewed in [21]). Among these, *PARK2*, *PARK7*, *GBA*, *LRRK2*, *PINK1* and *NR4A2* are expressed from mature astrocytes and unsurprisingly, most of these have been linked to neuroinflammatory response in preclinical models [21]. Therefore, it is possible that aside from disrupting neuronal function, these α Syn variants also compromise astrocyte homeostasis. Some of these gene variants do not necessarily cause α Syn pathology in all patients; however, by regulating mitophagy or redox function, these genes may cause generalized proteostasis imbalance or dysfunctional autophagic response in astrocytes leading to clinical symptoms reminiscent of Parkinsonism [21].

A common major cellular pathways affected by these genes is induction of oxidative stress causing widespread damage to cellular lipids, proteins, and DNA [41, 96, 114]. Indeed, accumulation of reactive oxygen species (ROS) can induce formation of TNTs [182], which has been shown by various groups to be conduits for α Syn propagation between astrocytes and neurons [1, 132, 146]. A homeostatic response of astrocytes to mitigate ROS is by a redox-sensitive decrease in KEAP1 mediated degradation of a transcriptional regulator, Nrf2. This induces expression of anti-oxidant response elements (ARE) leading to expression of detoxification and antioxidant enzymes such as glutathione, metallothioneins, NAD(P)H Quinone Dehydrogenase 1 (NQO1) and heme oxygenase. Several preclinical studies have shown a protective effect of Nrf2 on α -synucleinopathy [36, 57, 69, 72, 90, 171]. Interestingly, astrocytic Nrf2 overexpression rescues the phenotype and extends life span of human [A53T] α Syn transgenic mice [57]. This has spurred multiple studies testing the neuroprotective effects of Nrf2 activators as potential PD therapeutics [44, 56, 76]. Impairment of these protective functions of astrocytes by α Syn accumulation may contribute to neuronal demise, and restoration of them could represent avenues for therapeutic development (Figure 3).

DJ-1 (encoded by *PARK7*) that has been implicated in oxidative stress in PD [20, 155] as well as modulating α Syn aggregation and toxicity in preclinical models [140, 183]. DJ-1 is redox sensor of the cell and upon identifying ROS, it utilizes multiple defense mechanisms including direct neutralization of ROS, stabilization of Nrf2 and increased mitochondrial association [31, 40, 81]. Upregulating DJ-1 reduces α Syn pathology and rescues motor symptoms [181]. Additionally, DJ-1 can physically interact with α Syn to reduce its aggregation propensity [183]. Disease-associated variants of other astrocytic specific genes, such as *PARK2*, *LRRK2* and *GBA1*, lead to increased α Syn pathology in patient-derived induced pluripotent stem cells [68, 113, 174], suggesting a close association between astrocyte homeostasis and α Syn pathogenesis. In spite of reports that PINK1 is absolutely critical for astroglialogenesis [38] and that it phenocopies PARKIN (encoded by *PARK2*) function [8], there are no direct reports associating PINK1 with α Syn pathogenesis.

NURR1/NR4A2, enriched in mouse astrocytes and human microglia, is essential for dopaminergic neuronal development. Mutations in this gene has been linked to familial late-onset PD [92]. The primary function of Nurr1 is to suppress NF- κ B, leading to neuroprotection of dopaminergic neurons [134]. A second, more controversial, study showed that Nurr1 can also remediate α Syn toxicity in dopaminergic neurons by stimulating GDNF signaling [43]. Interestingly, Nurr1 was also shown to negatively regulate α Syn transcription [175], suggesting a common link between glial function and α Syn levels.

In PD and LBD, evidence of oxidative stress in areas of neuronal loss is seen in post mortem samples with increased lipid and protein peroxidation products, and altered expression of antioxidants within astrocytes (reviewed in [126]). Inadequate astrocytic production of glutathione, metallothionein 1/2, NQO1, DJ-1 and other ROS neutralizing factors in times of dopaminergic neuronal stress may contribute to their selective neurodegeneration in α -synucleinopathies. Coupled with the possibility that α Syn can induce astrocyte toxicity, the impairment of supportive functions may represent a new dimension for α Syn toxicity. Many of the PD risk genes discussed here function in ROS regulatory pathways within astrocytes,

and their loss may herald the onset of Parkinsonism either synergistically with α Syn pathology or in a parallel pathway by causing functional impairment of astrocytes (Figure 3).

Immune based biotherapies in α -synucleinopathies

A major unmet need in α -synucleinopathies is the lack of disease modifying therapies. Clinical and neuropathological heterogeneity in the spectrum of α -synucleinopathies, unclear etiologies, as well as the absence of validated biomarkers presents critical challenges regarding the development of effective therapeutics. The two major broad therapeutic approaches that are being currently tested in patients are anti- α Syn immunotherapy (passive as well as active) and small molecule drugs that inhibit α Syn accumulation [28], neither of which directly target astrocyte or innate immune mediators. The fact that astrocytes accumulate α Syn aggregates and may be involved in transmission of α -synucleinopathy opens up possibilities that targeting these specific pathologic inclusions or attenuating their consequences through astroglia targeted therapies can have disease modifying effects. However, it is not yet clear how best to harness the immune system towards a beneficial outcome on astrocyte function. In this section, we will first describe the trophic factor therapies that have been used in patient trials, which among many factors, alters neuroinflammation but does not necessarily modulate astrocyte function directly.

In previous clinical trials, several neurotrophic factors (that are produced by astrocytes) were tested as neuroprotective pharmacotherapeutics [150]. Of these, glial cell line-derived neurotrophic factor (GDNF) and Neurturin (NRTN) were well-tolerated in patients but, unfortunately, did not meet predetermined end-points in efficacy. More recently, other neurotrophic factors that can be produced by astrocytes, such as cerebral dopamine neurotrophic factor (CDNF) [91] and mesencephalic astrocyte-derived neurotrophic factor (MANF) [180], have shown promise in preclinical models of α -synucleinopathy. Since the receptors and cellular signaling pathways for many of these neurotrophic factors are relatively unknown, more work needs to be done to forward these into the clinical realm.

Several preclinical strategies that work partly by targeting astrocytes have been recently described. Glucagon-like peptide-1 receptor (GLP1R) agonists have shown efficacy in multiple models of neurodegenerative proteinopathies by modulating mitochondrial biogenesis, suppressing microglial activation and inflammation, enhancing autophagy, and clearance of aggregated proteins. One such GLP1R agonist, NLY01, was shown to be neuroprotective in PD models by specifically regulating NF- κ B mediated inflammatory signaling which is downstream of the GLP-1R/PI3K/AKT pathway [9]. More recently, NLY01 was effective in blocking microglia-mediated conversion of astrocytes to the neurotoxic A1 phenotype in a rodent model of α -synucleinopathy [178]. Indeed, a limited open-label trial and a subsequent double blind trial of the GLP1R agonist exenatide appeared to improve motor and cognitive function in PD patients [10, 11].

A direct line of preclinical research targeting astrocyte function was done by using midbrain derived astrocyte grafts into a rat model of PD. These astrocytes, following grafting into the striatum, resulted in remarkable recovery of dopaminergic neurons, especially in the

presence of Nurr1 and Foxa2 co-expression. Through a paracrine pathway, these grafted astrocytes were found to exert marked trophic action on the glutamatergic and GABAergic neurons and stimulate endogenous dopaminergic neurons to secrete dopamine [144]. Such therapies have the potential to be the most effective, as instead of isolating one single astrocytic product for supplementation, this would generate the entire spectrum of trophic factors, anti-oxidant compounds, and other astrocytic protective functions needed to assist neurons for overcoming cellular stresses induced by pathologic α Syn.

Conclusion

Astrocytes are the most predominant glial cells in the brain and play critical role in neuronal development, neural activity, synaptic transmission and learning and memory. However, their role in the etiology of neurodegenerative α -synucleinopathies is still mostly sheathed in mystery (Box 1). Several lines of evidence, neuropathological, functional and genetic, indicate that astrocytes are functionally involved in α -synucleinopathies (Figure 3). How and why these astrocytes synergize with the evolution of proteostasis failure and heterogeneous clinical symptoms need to be charted out, along with the contribution that regional astrocyte heterogeneity may play a role in selective neurodegeneration. Interestingly, the underlying mechanisms of astrocytic modulation of α -synucleinopathy may be divergent – in some cases, astrocytic activity may need to be enhanced whereas in some cases, its pathological function will need to be attenuated based on the context and timing of intervention. The fact that astrocytes are a heterogeneous group of cells, in terms of functionality and region-wide distribution, remains under-appreciated in the field of neurodegenerative α -synucleinopathies. Whether this physiological heterogeneity has a role to play in selective vulnerability of neurons along the spectrum of α -synucleinopathy (PD, DLB, MSA) that might ultimately be informative of the spectrum of clinical heterogeneity seen in the patients also need to be examined. Genetic predisposing factors that alter risk for α -synucleinopathies such as *PARK7*, *GBA*, *LRRK2*, *PINK1* and *NR4A2* may also affect astrocyte function and alter disease etiology through altering redox stress or other pathways (Figure 3). Whether mining such genetic and risk factor data can enable stratification of patients for personalizing treatment options, based on their unique clinical presentation, should be examined in future therapeutic studies. Current strategies of harnessing immunoproteostasis by either using small molecule drugs or astrocyte-derived neurotrophic molecules or astrocyte transplantation in combination with immunotherapy remain intriguing and will have to be tested in multiple models of α -synucleinopathies (Figure 3). Overall, bilateral interactions or non-cell autonomous interactions between astrocytes and neurons or other glia in specific regions of the brain may provide us insights into neuronal dysfunction and death (Figure 3). Deeper knowledge of astrocyte biology and its functional alterations will be necessary before we can successfully embark on such disease modifying therapies in α -synucleinopathies.

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Abbreviations

αSyn	<u>α</u> - <u>syn</u> uclein
AKT	' <u>A</u> <u>K</u> ' thymoma
ARE	<u>A</u> nti-oxidant <u>r</u> esponse <u>e</u> lement
CNS	<u>C</u> entral <u>N</u> ervous <u>S</u> ystem
CDNF	<u>C</u> erebral <u>d</u> opamine <u>n</u> eurotrophic <u>f</u> actor
CSF	<u>C</u> erebrospinal <u>f</u> luid
DAMP	<u>D</u> amage <u>a</u> ssociated <u>m</u> olecular <u>p</u> attern
DLB	<u>D</u> ementia with <u>L</u> ewy <u>B</u> odies
EM	<u>E</u> lectron <u>M</u> icroscope
FA	<u>F</u> ormic <u>A</u> cid
Foxa1	<u>F</u> orkhead Box <u>A</u> 1
GABA	<u>G</u> amma- <u>A</u> mino <u>B</u> utyric <u>a</u> cid
GDNF	<u>G</u> lial cell line- <u>d</u> erived <u>n</u> eurotrophic <u>f</u> actor
GCI	<u>G</u> lial <u>c</u> ytoplasmic <u>i</u> nclusions
GFAP	<u>G</u> lial <u>F</u> ibrillary <u>A</u> cidic <u>P</u> rotein
GLP1R	<u>G</u> lucagon- <u>l</u> ike peptide- <u>1</u> receptor
GBA	<u>G</u> lucosidase <u>β</u> eta <u>a</u> cid
H&E	<u>H</u> ematoxylin & <u>E</u> osin staining
HLA-DR	<u>H</u> uman <u>l</u> eukocyte <u>a</u> ntigen DR isotype
MHC-II	<u>M</u> ajor <u>h</u> istocompatibility <u>c</u> omplex class <u>I</u> I
MMP	<u>M</u> atrix <u>m</u> etalloproteinase
MANF	<u>M</u> esencephalic <u>a</u> strocyte-derived <u>n</u> eurotrophic <u>f</u> actor
HLA-DR	<u>H</u> uman <u>L</u> eukocyte <u>A</u> ntigen – <u>D</u> R isotype
iLBD	<u>I</u> ncidental <u>L</u> ewy <u>b</u> ody <u>d</u> iseases
Keap1	<u>K</u> elch-like <u>e</u> ch-associated protein 1
LRRK2	<u>L</u> eucine- <u>r</u> ich <u>r</u> epeat <u>k</u> inase <u>2</u>
LB	<u>L</u> ewy <u>b</u> odies
LN	<u>L</u> ewy <u>n</u> eurites

MSA	<u>M</u> ultiple <u>S</u> ystem <u>A</u> trophy
NQO1	<u>N</u> AD(P)H <u>Q</u> uinone Dehydrogenase <u>1</u>
NCI	<u>N</u> euronal <u>c</u> ytoplasmic <u>i</u> nclusions
NRTN	<u>N</u> eurturin
NAC	<u>N</u> on-amyloid β <u>c</u> omponent
Nrf2	<u>N</u> uclear factor erythroid <u>2</u> -like <u>2</u>
NFκB	<u>N</u> uclear factor κ -light-chain-enhancer of activated <u>B</u> cells
Nurr1/NR4A2	<u>N</u> uclear receptor subfamily <u>4</u> , group <u>A</u> , member <u>2</u>
PD	<u>P</u> arkinson's <u>d</u> isease
PARK	<u>P</u> arkinson's disease associated gene
PDD	<u>P</u> D with <u>d</u> ementia
PI3K	<u>P</u> hosphatidylinositol <u>3</u> - <u>k</u> inase
PDGF β	<u>P</u> latelet-derived growth factor <u>β</u>
PET	<u>P</u> ositron-emission <u>t</u> omography
pSer129	<u>P</u> hosphorylated <u>S</u> erine <u>129</u>
PINK1	<u>P</u> ten-induced putative <u>k</u> inase <u>1</u>
ROS	<u>R</u> eactive <u>o</u> xygen <u>s</u> pecies
SQSTM1	<u>S</u> questosome <u>1</u>
Thy-1	<u>T</u> hymocyte differentiation antigen <u>1</u>
TLR	<u>T</u> oll-like <u>r</u> eceptor
TNT	<u>T</u> unneling nanotube

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BOX 1:**CRITICAL OUTSTANDING QUESTIONS**

What is the source of astrocytic α -synuclein?

Does a bilateral communication between neurons and astrocytes lead to exacerbated disease?

What is the mechanism of bilateral transmission of α Syn – direct physical contact or passive uptake?

How do the astrocyte-resident PD risk genes lead to proteostasis failure in glia – through direct or non cell autonomous pathways?

Does astrocytic dysfunction precede α Syn proteinopathy or is the dysfunction a result of neuronal α Syn proteinopathy?

Does astrocytic metabolism of internalized α Syn result in toxic by-products that contribute to disease pathogenesis?

Is restoring astrocytic function sufficient to rebalance proteostasis as a monotherapy?

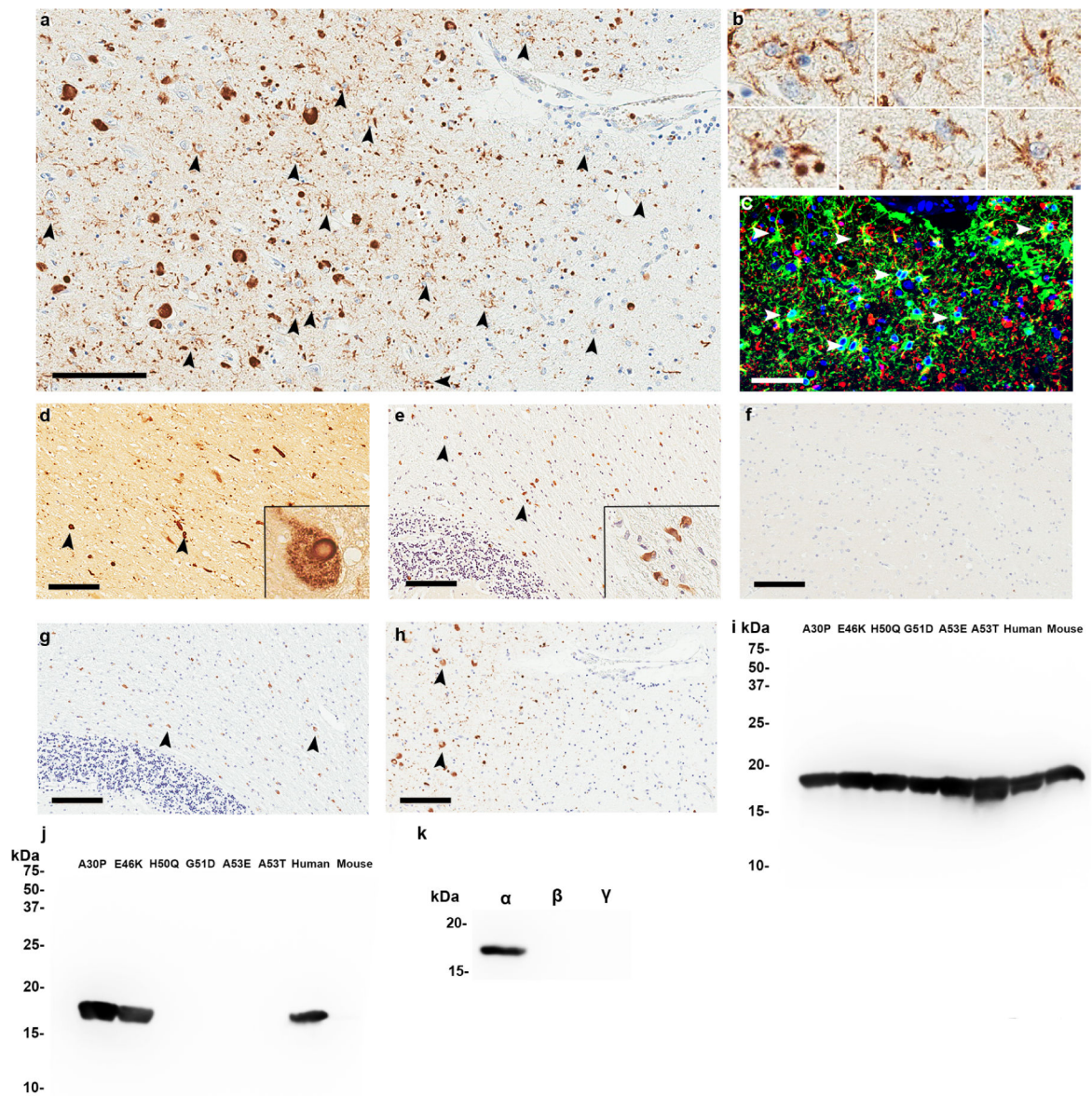


Figure 1. Astrocytic α Syn inclusions in DLB patients specifically detected by an antibody in the middle domain of α Syn.

a. Immunohistochemical staining with 3H11 antibody raised against residues 43–63 of α Syn with formic acid retrieval. α Syn is extensively detected within astrocytes in the hippocampus of a DLB patient. b. Higher magnification from Panel A showing astrocytes immunopositive for α Syn antibody 3H11. c. Immunofluorescent analysis with antibodies to GFAP (green) and α Syn 3H11 (red) demonstrating that α Syn is present in astrocytic processes in the DLB hippocampus. d. Detection of α Syn within classical LBs in the substantia nigra of a DLB patient with the 3H11 antibody. Inset shows magnified view of a typical intraneuronal LB. e. Detection of α Syn within GCI inclusions in the cerebellum of an MSA individual using the 3H11 antibody. Inset shows magnified view of typical GCI. f. α Syn antibody 3H11 fails to show any immunoreactivity in the hippocampus of an Alzheimer's disease patient negative for pSer129 α Syn inclusions. g. Immunohistochemical

staining of the DLB hippocampus from panel a using antibody EP1536Y against pSer129 α Syn; extensive astrocytic inclusions are not seen using this antibody. h. Immunohistochemical staining of the MSA cerebellum from panel e using antibody EP1536Y against pSer129 α Syn; 3H11 detects GCIs similarly to this antibody but no astrocytic staining is seen using either. i. A western blot of 200 ng recombinant α Syn proteins harboring various familial mutations probed with antibody 94–3A10 (residues 130–140) [45]. j. A western blot using the same proteins from panel G but probed with antibody 3H11; the antigenic region is residues ~47–55. k. Western blot showing that 3H11 specifically recognizes human α Syn but not β Syn or γ Syn. Scale Bar: 100 μ m (A, D, E, F, G, H), 50 μ m (C).

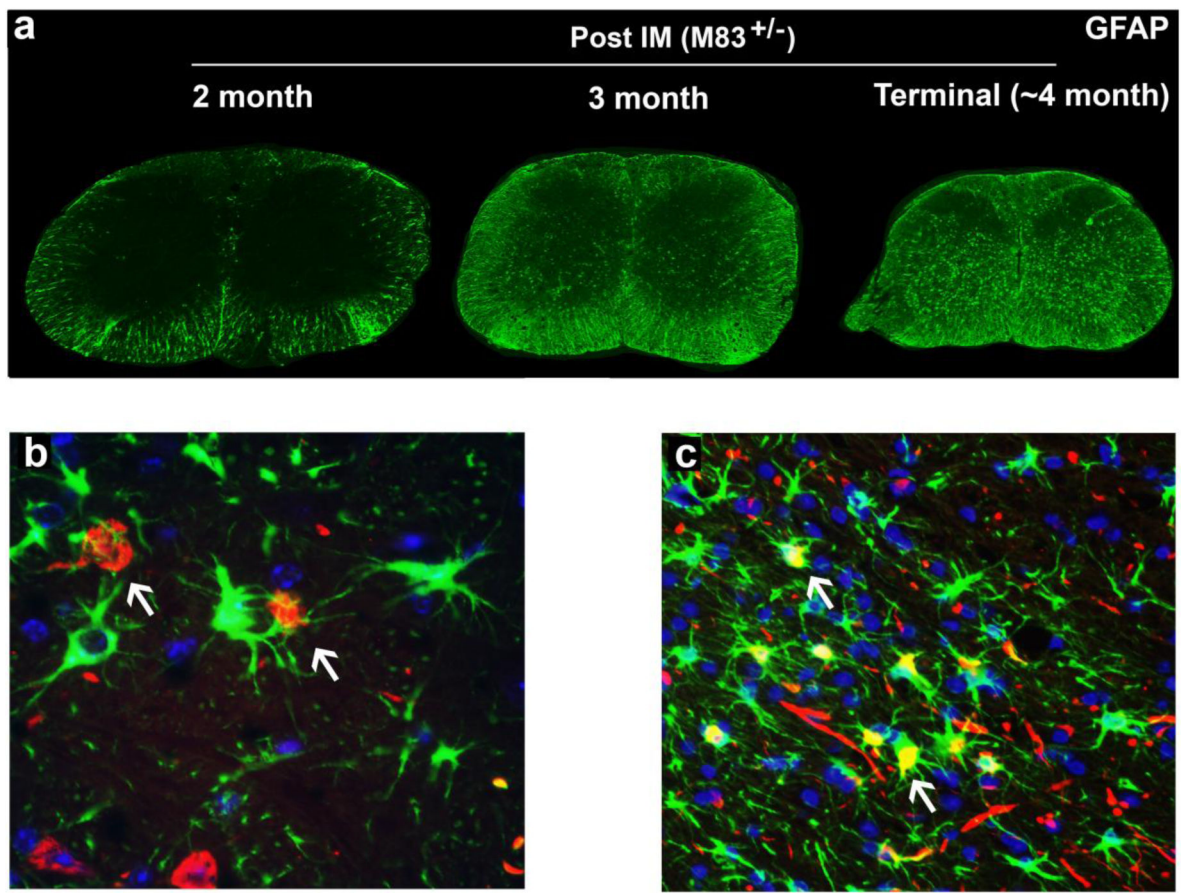


Figure 2. Robust astrogliosis and astroglial α Syn accumulation in transgenic mouse models of α -synucleinopathy.

a. Immunofluorescent detection of GFAP in spinal sections from M83^{+/-} transgenic mice overexpressing human [A53T] α Syn that were intramuscularly injected with preformed wild type mouse α Syn fibrils. In this model, pSer129- α Syn pathology and motor neuron death is apparent at 2 months post injection whereas increase in GFAP, representative of astrocyte activation, is observed at later time points. b. Immunofluorescent detection of GFAP (green) and pSer 129 α Syn (red) in the spine of terminal M83^{+/-} transgenic mice described in panel a; in regions of neuronal death, astrocytes closely interact with extracellular aggregated α Syn. c. Immunofluorescent detection of GFAP (green) and pSer 129- α Syn (red) in the midbrain of M20 transgenic mice overexpressing human α Syn that were seeded with preformed α Syn fibrils in the striatum showing the presence of abundant astrocytic α Syn pathology. Scale Bar: 500 μ m (B), 100 μ m (C).

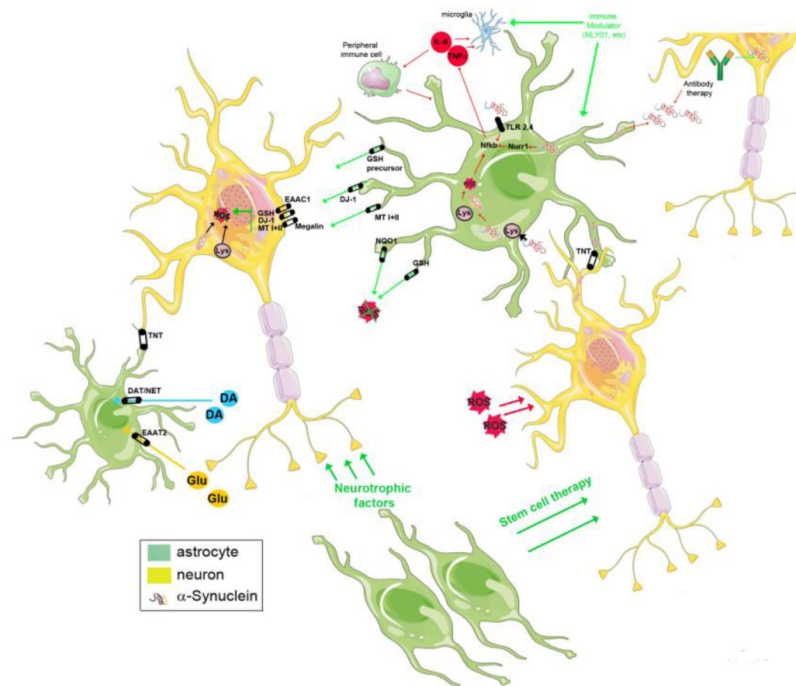


Figure 3. αSyn mediated alterations in neuro-glia homeostasis in health and disease.

Accumulation of αSyn leads to Lewy pathology through a variety of cellular processes. In neurons and astrocytes, αSyn has been implicated in dysfunction of lysosomes (LYS) and mitochondria (MITO), which enhances ROS production and toxic sequela. Neuronal αSyn can be transported into neighboring astrocytes via tunneling nanotubes (TNT), receptors such as Toll like receptors (TLR), or several indirect methods such as internalization of exocytosed material from the synaptic space. We hypothesize that astrocytic lysosomes preferentially cleave full length αSyn into smaller toxic truncation products that might have additional pathological functions with astrocytes. Toxic accumulation of αSyn within astrocytes may damage their normal functions and have adverse results such as synaptic accumulation of neurotransmitters including glutamate and dopamine (DA) that are in part cleared by astrocytes through EAAT2 and DAT/NET respectively. Astrocytes normally protect against oxidative stress in the CNS through production of multiple products including ARE factors, KEAP1/NFR2, Glutathione, NURR1 and NQO1; loss of these factors due to toxic accumulation of αSyn may contribute to neurodegeneration. Astrocytic uptake of αSyn and oxidative stress may impinge on the NF-κB pathways that can result in increased inflammatory cytokines, such as IL-6 or TNFα that will affect the function of neighboring microglia or peripheral immune cells which represents another modality by which astrocytes may contribute to neurodegeneration. Additionally, common risk factor genes such as DJ-1, Parkin and Pink1 are highly expressed in astrocytes and their dysfunction may affect mitophagy in a parallel pathway to that of toxic αSyn resulting in similar degenerative features. Red arrows denote pathological processes, green arrows denote putative therapies or beneficial pathways and black arrows denote normal physiological processes.

Table 1.

Astrocytic α Syn inclusions: a brief review of literature.

# subjects with astrocyte inclusions (total #)	Location of astrocyte inclusions	Antigen Retrieval	α Syn antibody (astrocyte labeling)	Comments	Ref
24 (30) PD 0 (7) iLBD 0 (30) control	13/24 SNpc 24/24 midbrain/pontine tegmentum	NA	Polyclonal α Syn (114–131) (+)	<ul style="list-style-type: none"> number of positive astrocytes parallels severity of LBs amount of astrocyte inclusions correlated with neuronal loss double labeling with GFAP and vimentin to identify astrocytes 	168
8 (8) DLB	8/8 temporal lobe 8/8 frontal cortex 7/8 basal ganglia 6/8 midbrain/pontine tegmentum 5/8 SNpc	FA	α Syn NAC 60–75 (+) α Syn N terminal 1–15 (–) α Syn C terminal 108–122 (\leftrightarrow)	<ul style="list-style-type: none"> “star like astrocytes” with α-syn positivity throughout perikarya and processes densest amount of α-syn positive astrocytes in temporal lobe with 100–200 SLA within visual field 	160
x (7) PD x (5) DLB 0 (6) MSA 0 (25) other disease 0 (15) control	PD, DLB: cerebral cortex, basal ganglia	FA	α Syn N terminal 1–15 (+) α Syn NAC 60–75 (+) α Syn 110–140 (+)	<ul style="list-style-type: none"> more positive astrocytes in DLB compared with PD number of positive astrocytes parallels severity of LBs grey matter astrocyte staining only 	141
9 (9) LBVAD 2 (2) DLB 1 (1) PD 0 (17) other disease 0 (4) control	PD, DLB, LBVAD (distinction not made): neocortex, hippocampus	FA	α Syn N terminal 1–9 (–) α Syn NAC 61–75 (+) α Syn C terminal (101–110&131–140) (–)	<ul style="list-style-type: none"> grey matter and cortical astrocyte staining astrocyte inclusions tau negative 	156
10 (10) PD 0 (2) iLBD 0 (5) control	10/10 striatum 10/10 SNpc 10/10 midbrain and pontine tegmentum 8/10 cingulate and other cortex	HCl	Polyclonal α Syn full-length (+)	<ul style="list-style-type: none"> 60+ glial inclusions per area in cingulate compared with 50+ LBs in same area glial inclusions correlate with neuronal loss and severity of LB 	66
14 (14) PD 0 (6) control	14/14 cortex 12/14 striatum 11/14 thalamus	FA	α Syn NAC 91–99 (+) α Syn C terminal 116–131 (–)	<ul style="list-style-type: none"> astrocytic staining parallels severity of LB pathology no correlation between plaques and astrocytic inclusions astrocytic staining is diffuse and granular throughout cell densest in temporal lobe 	22
0 (10) PD 0 (8) DLB 0 (9) other disease 0 (3) control	N/A	NA	α Syn N terminal 1–15 (–)	<ul style="list-style-type: none"> silver positive glial inclusions in all PD and DLB cases most abundant in SN 	7
17 (20) PD 0 (26) other disease 0 (30) control	8/17 SNpc 17/17 other midbrain region	NA	No antibodies used	<ul style="list-style-type: none"> silver positive, tau negative inclusions in astrocytes in PD positive astrocytes had many fine radiating argyrophillic tendrils; differ from tufted astrocytes of PSP 	169

# subjects with astrocyte inclusions (total #)	Location of astrocyte inclusions	Antigen Retrieval	αSyn antibody (astrocyte labeling)	Comments	Ref
6 (15) MSA 0 (20) DLB 0 (20) control	3/6 periventricular astrocytes in cerebrum 6/6 subpial astrocytes in midbrain, brainstem 1/6 bergmann glia in cerebellum	NA	αSyn pSer129 (+)	<ul style="list-style-type: none"> 20+ astrocyte inclusion in one unilateral PD midbrain section not correlated with amount of remaining neurons or LBs in SN amount of positive astrocytes correlated with MSA of longer duration 	110
1 (1) PD (fetal transplant)	Within fetal transplant	NA	αSyn pSer129 (↔)	<ul style="list-style-type: none"> glial inclusions throughout the graft and outside the transplant region 	2
6 (8) DNTC 0 (6) control	6/6 temporal lobe most densely within entorhinal cortex	FA	αSyn N terminal 1–15 (-) αSyn NAC 60–75 (+) αSyn 108–122 (-)	<ul style="list-style-type: none"> α-syn pathology confined to temporal semiquantitative comparison indicated near equivalent levels of astrocytic and neuronal α-syn pathology 	177
4 (6) PD 5 (7) DLB 3 (7) MSA 0 (5) control	PD, DLB, MSA: 12/12 bergmann glia molecular layer cerebellum	NA	αSyn C terminal 114–131 (+)* αSyn full-length (+)*	<ul style="list-style-type: none"> LB like inclusions and doughnut shaped structures; also detectable by H&E no correlation between Bergmann glia α-syn positivity and disease outcome 	124
x (34) PD/DLB x (14) MSA 0 (x) Control	PD,DLB: Temporal lobe, cingulate cortex	FA	αSyn N terminal 46–53 (+)	<ul style="list-style-type: none"> tissue microarray analysis 	87
13 (13) PD 0 (29) MSA 0 (44) other disease 0 (13) control	13/13 subcortical grey matter	FA	αSyn 15–123 (+) αSyn full-length (+)	<ul style="list-style-type: none"> 40–45% of subcortical astrocytes in PD accumulated α-syn α-syn positive astrocytes and neurons localized in same regions protoplasmic astrocytes affected more than fibrous astrocytes 	145

Note: FA, Formic Acid; +, positive staining; -, negative staining; ↔, weak staining;

* , labeled Bergman Glia only; x, n number not specified; NA, information not available. Additional studies have noted glial reactivity for αSyn in PD/DLB, however the lack of differentiation between microglia, astrocytes, and oligodendrocytes led to them being omitted from this Table.