



Expression of α -synuclein is regulated in a neuronal cell type-dependent manner

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Abstract

α -Synuclein, the major component of Lewy bodies (LBs) and Lewy neurites (LNs), is expressed in presynapses under physiologically normal conditions and is involved in synaptic function. Abnormal intracellular aggregates of misfolded α -synuclein such as LBs and LNs are pathological hallmarks of synucleinopathies, including Parkinson's disease (PD) and dementia with Lewy bodies (DLB). According to previous studies using pathological models overexpressing α -synuclein, high expression of this protein in neurons is a critical risk factor for neurodegeneration. Therefore, it is important to know the endogenous expression levels of α -synuclein in each neuronal cell type. We previously reported differential expression profiles of α -synuclein in vitro and in vivo. In the wild-type mouse brain, particularly in vulnerable regions affected during the progression of idiopathic PD, α -synuclein is highly expressed in neuronal cell bodies of some early PD-affected regions, such as the olfactory bulb, the dorsal motor nucleus of the vagus, and the substantia nigra pars compacta. Synaptic expression of α -synuclein is mostly accompanied by expression of vesicular glutamate transporter-1, an excitatory synapse marker protein. In contrast, α -synuclein expression in inhibitory synapses differs among brain regions. Recently accumulated evidence indicates the close relationship between differential expression profiles of α -synuclein and selective vulnerability of certain neuronal populations. Further studies on the regulation of α -synuclein expression will help to understand the mechanism of LB pathology and provide an innovative therapeutic strategy to prevent PD and DLB onset.

Keywords Dementia with Lewy bodies · Excitatory neuron · Inhibitory neuron · Parkinson's disease · Synapse

Introduction

α -Synuclein is a major constituent of Lewy bodies (LBs) and Lewy neurites (LNs), which are pathological hallmarks of synucleinopathies, including Parkinson's disease (PD) and dementia with Lewy bodies (DLB) (Spillantini et al. 1998; Dickson 2001; Stefanis 2012). Several missense mutations, as well as duplicate and triplicate regions of the α -synuclein gene are responsible for familial PD (Polymeropoulos et al. 1997; Zarranz et al. 2004; Kruger et al. 1998; Singleton et al.

2003; Chartier-Harlin et al. 2004). In studies of α -synuclein pathogenicity, it was demonstrated that overexpression of α -synuclein in neurons results in the formation of inclusion bodies and neuronal loss (Masliah et al. 2000; Van der Perren et al. 2015; Singleton et al. 2003). Therefore, an increase in the intracellular amount of α -synuclein is a probable risk factor for neurodegeneration.

Expression of α -synuclein is regulated by various transcription factors, such as zinc finger and SCAN domain containing 21 (ZSCAN21) (Clough et al. 2009; Dermentzaki et al. 2016), GATA-1 and GATA-2 (Scherzer et al. 2008), Nurr1 (Yang and Latchman 2008), TRIM32 (Pavlou et al. 2017), and p27^{Kip1} (Gallastegui et al. 2018). These transcription factors interact directly with the promoter region of α -synuclein, and this critical link between these transcription factors and α -synuclein may enable the design of therapies to lower production of α -synuclein. Further studies of the regulation machinery of α -synuclein expression via these transcription factors will help develop novel therapeutic strategies for synucleinopathies.

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According to a recent study, β 2-adrenoreceptor (β 2AR) was identified as a novel regulator of the α -synuclein gene (Mittal et al. 2017). β 2AR activation by selective agonists reduces α -synuclein expression in mouse substantia nigra. Conversely, suppression of β 2AR expression or chemical inhibition of β 2AR activity increases α -synuclein expression. Furthermore, longitudinal studies of incident PD throughout Norway showed that the β 2AR antagonist, propranolol, is associated with a markedly increased risk of PD. However, salbutamol, a β 2AR agonist, is associated with a decreased risk of PD. Thus, fine-tuning of intrinsic α -synuclein expression levels could constitute an innovative therapeutic strategy to prevent PD onset.

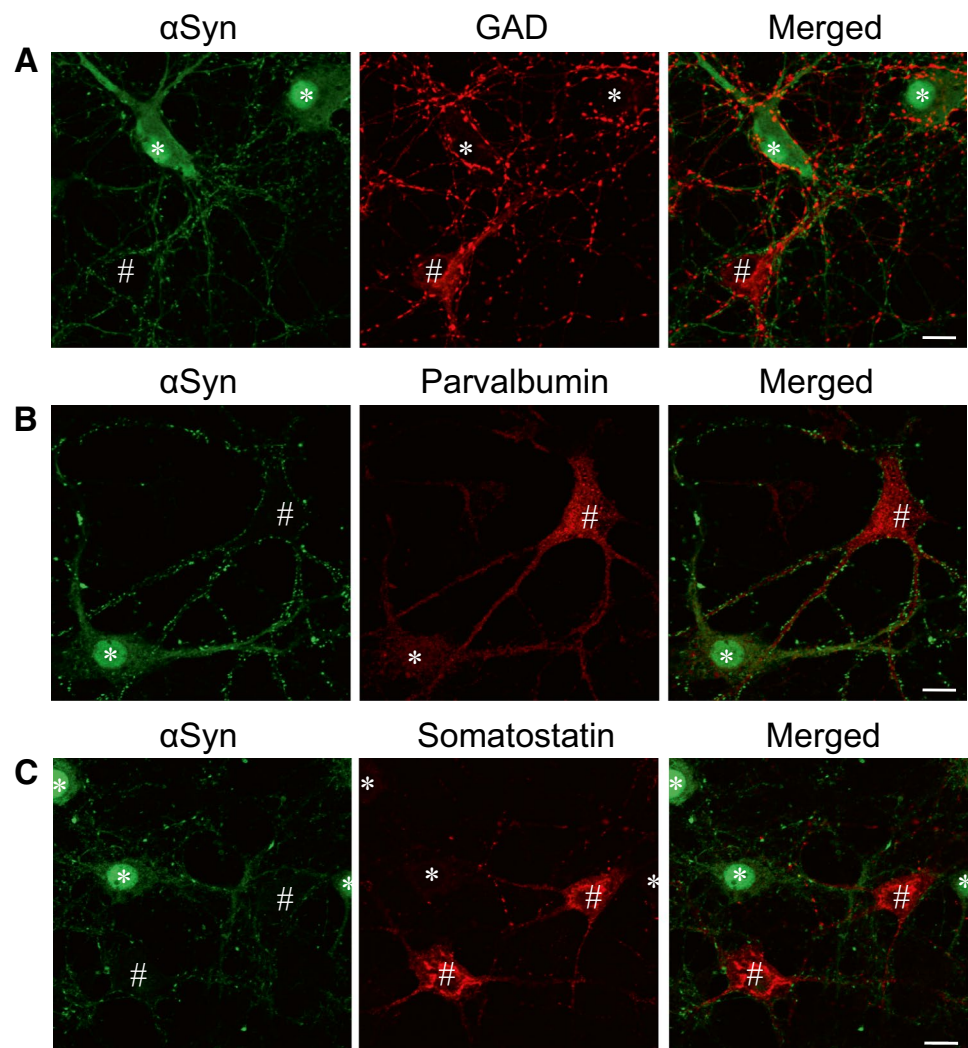
Expression levels and subcellular distribution of α -synuclein in each neuronal cell type are related closely to the pathogenicity and the physiological function of α -synuclein. In this review, we focus on the characteristic profile of α -synuclein expression in vitro and in vivo, and further discuss new findings obtained from recent studies on this protein.

Differential expression of α -synuclein under physiological conditions

α -Synuclein is enriched in brain and is localized at presynapses under physiologically normal conditions in vitro and in vivo (Withers et al. 1997; Totterdell et al. 2004; Totterdell and Meredith 2005; Vivacqua et al. 2011). It has been suggested that α -synuclein plays a role in the generation and maintenance of synapses because this protein appears earlier than synaptophysin—a synaptic vesicle protein—during development of the central nervous system (CNS) and is localized to axon terminals throughout the adult mammalian brain (Hsu et al. 1998; Petersen et al. 1999).

α -Synuclein binds directly to synaptobrevin-2 in pre-synaptic regions and functions to sustain soluble *N*-ethylmaleimide-sensitive factor attachment protein receptor (SNARE)-complex assembly in vivo and in vitro (Burre et al. 2010). According to a more recent study, α -synuclein modulates the kinetics of exocytotic events, promoting

Fig. 1 a–c Low expression of α -synuclein (α Syn) in hippocampal inhibitory neurons. # Cells expressing the inhibitory neuronal marker proteins, glutamic acid decarboxylase (*GAD*) (a), parvalbumin (b), and somatostatin (c), show low expression of α Syn. Cells with high expression of α Syn are labeled with asterisks. Bars 10 μ m (Taguchi et al. 2014)



cargo discharge and reducing pore closure (Logan et al. 2017). Thus, these studies indicate a synaptic function of α -synuclein during the neurotransmission process.

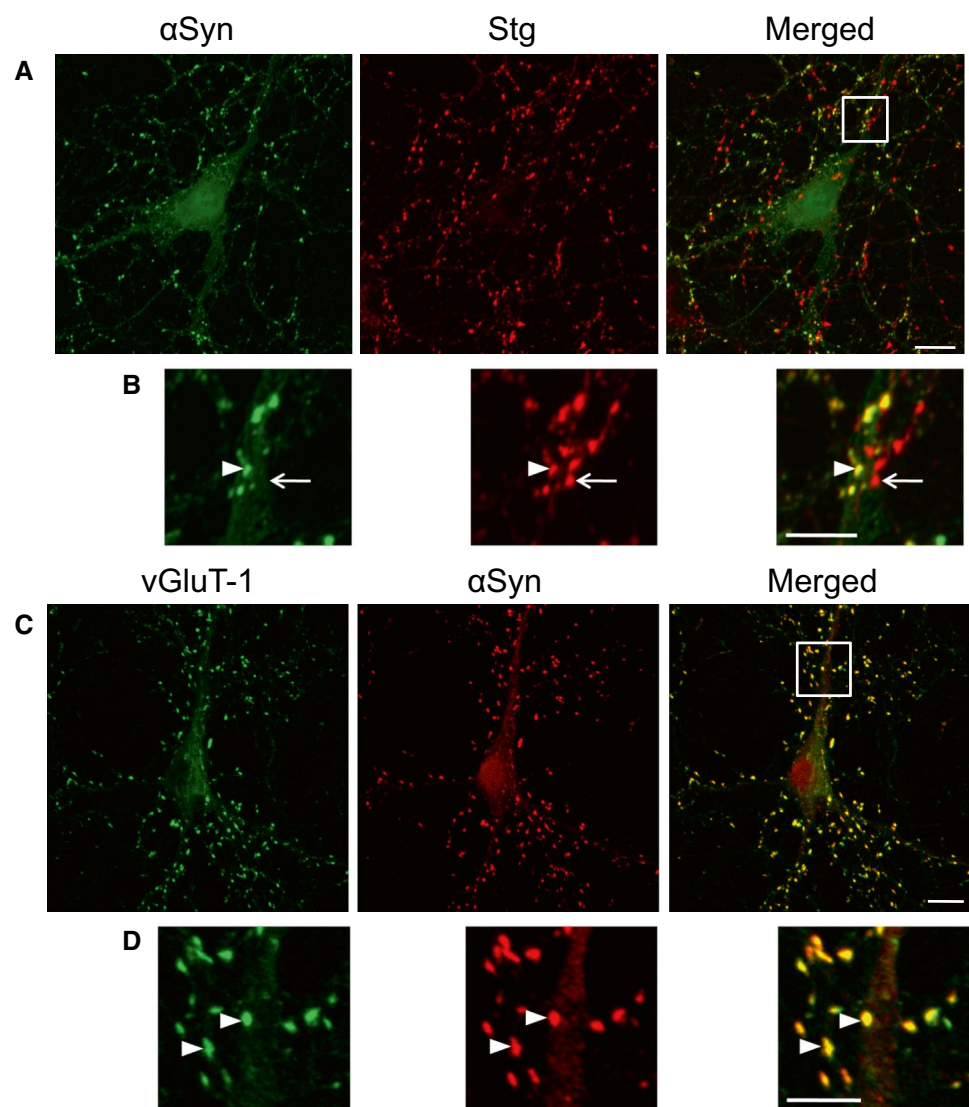
It is believed that α -synuclein is generally expressed in synapses, whereas the amount of α -synuclein protein expressed in each cell differs between different neuronal cell types. Li and colleagues indicated that α -synuclein is present abundantly in central catecholaminergic systems (Li et al. 2002), whereas α -synuclein expression is comparatively weak in many cholinergic brain regions.

We previously investigated the subcellular distribution of α -synuclein in normal and pathological conditions using primary cultured hippocampal neurons (Taguchi et al. 2014). While some neurons expressed high levels of α -synuclein in presynapses expressing synaptotagmin (Stg)—a presynaptic marker protein—and cell bodies, other neurons either did not express the protein, or only very weakly (Figs. 1, 2a,b). These α -synuclein-negative cells were identified as

inhibitory neurons expressing glutamic acid decarboxylase (GAD) (Fig. 1a), parvalbumin (Fig. 1b), or somatostatin (Fig. 1c). In contrast, α -synuclein-positive synapses were clearly colocalized with vesicular glutamate transporter-1 (vGluT-1)—an excitatory synapse marker protein (Fig. 2c,d). This characteristic expression pattern of α -synuclein is conserved in the hippocampus in vivo.

According to a previous study, α -synuclein plays a role in maintaining the overall size of the recycling pool of synaptic vesicles (Scott and Roy 2012). The sizes of both the recycling pool and total vesicular pool are more variable at glutamatergic synapses than at gamma-aminobutyric acid (GABAergic) synapses (Moulder et al. 2007). This heterogeneity of the size of the recycling pool at glutamatergic synapses may provide a dynamic range for synaptic strength that is not present at GABAergic synapses (Moulder and Mennerick 2005; Moulder et al. 2007). Therefore, α -synuclein

Fig. 2 Presynaptic localization of α -synuclein in hippocampal excitatory neurons. **a, b** Confocal images of double immunostaining for α -synuclein (α Syn) and synaptotagmin (Stg). The region marked by a white square in **a** is magnified in **b**. Arrowhead in **b** indicates the presynapse, expressing both α Syn and Stg. However, there are some Stg-positive synapses lacking α Syn (arrow). **c, d** Confocal images of double immunostaining for α Syn and vGluT-1. The region marked by a white square in **c** is magnified in **d**. α Syn is clearly colocalized with vGluT-1 in **d** (arrowheads). Bars **a, c** 10 μ m; **c, d** 5 μ m (Taguchi et al. 2014)



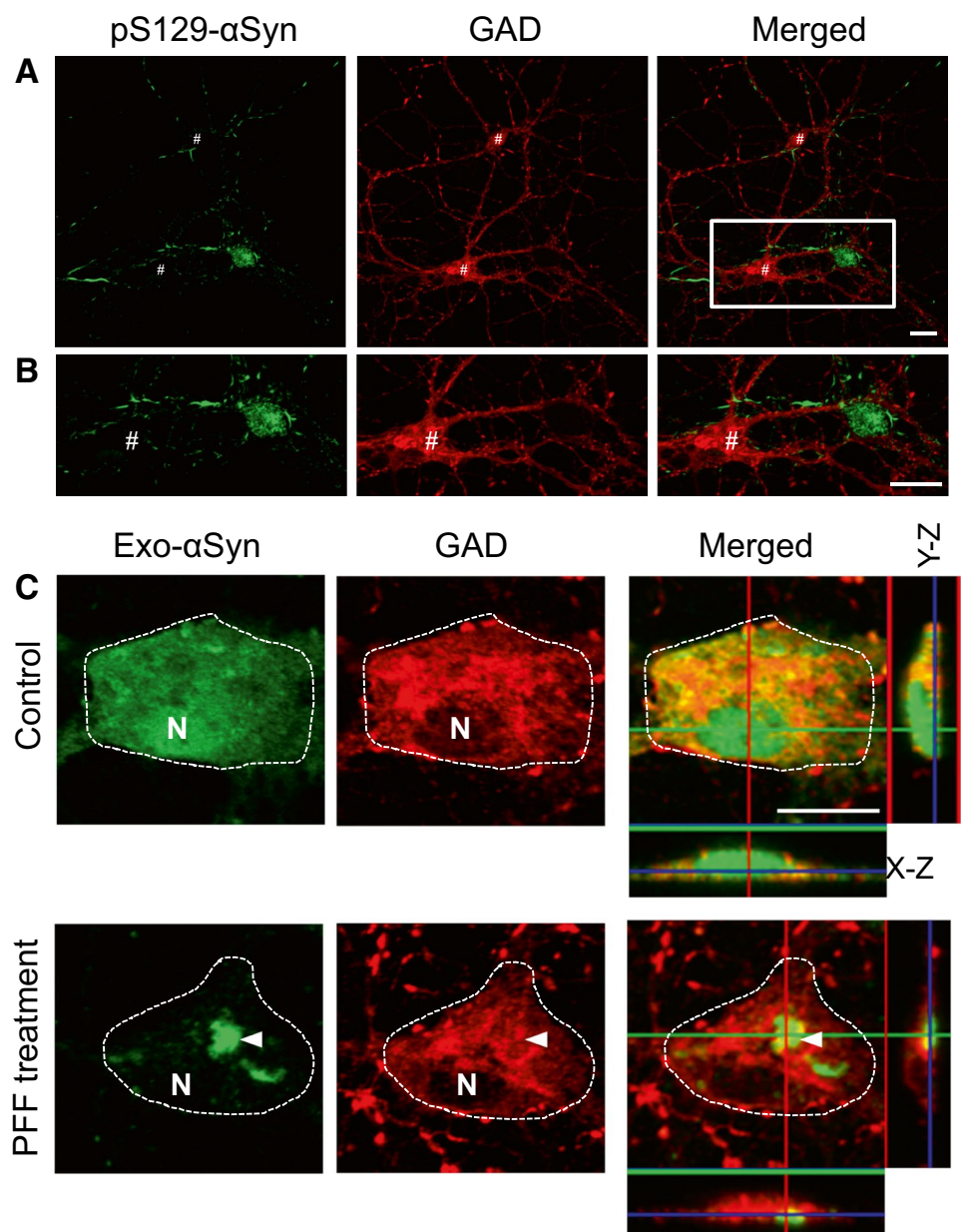
might act as a modulator of the size of the recycling pool at excitatory synapses.

Cell-type dependent aggregate formation and synapse impairment

Intracellular aggregates such as LBs and LNs are composed mainly of α -synuclein. These aggregates are formed by recruitment of the intrinsic soluble α -synuclein into the insoluble aggregate core. Therefore, endogenous expression of α -synuclein is required for aggregate formation (Volpicelli-Daley et al. 2011). Preformed fibrils (PFFs) prepared from recombinant α -synuclein can induce the formation of LB-like aggregates. If inhibitory hippocampal neurons

have low expression levels of endogenous α -synuclein as described above, they may not be able to form LB-like aggregates. As expected, most of the GAD-positive inhibitory hippocampal neurons were free of LB-like aggregates (Fig. 3a,b) (Taguchi et al. 2014). However, LB-like aggregate formation was induced successfully by PFFs in hippocampal inhibitory neurons overexpressing exogenous human α -synuclein (Fig. 3c). The low frequency of LB-like aggregate appearance is probably due to the lower amount of endogenous α -synuclein expressed in the inhibitory neurons. As described below, inhibitory neurons in the cerebral cortex and some other regions including the hippocampus show low expression of α -synuclein in adult mouse brain (Taguchi et al. 2016). A pathological study of DLB patients

Fig. 3a–c Formation of intracellular aggregates of α -synuclein. **a, b** Confocal images of double immunostaining for phosphorylated α -synuclein (pS129- α Syn) and GAD after treatment with preformed fibrils (PFFs) of α -synuclein. The region marked by a *white square* in **a** is magnified in **b**. Immunoreactivity of pS129- α Syn is observed as intracellular fibrous aggregates or inclusion bodies. GAD-positive neurons indicated by # are free of α -synuclein aggregate formation. GAD signals are not colocalized with pS129- α Syn. **c** In the absence of PFF treatment, exogenous human α -synuclein (Exo- α Syn) was distributed diffusely in the cell body of GAD neurons (*control*). After PFF treatment, intracellular inclusions positive for α Syn are induced in the GAD-positive cells expressing Exo- α Syn. Cell bodies are enclosed by *white dotted lines*. *N* indicates the location of the nucleus. *Bars* 10 μ m (Taguchi et al. 2014)



reported that parvalbumin-containing cortical neurons are free of LBs and spared from neurodegeneration, although the basal expression level of α -synuclein was not determined (Gomez-Tortosa et al. 2001). Formation of LBs composed of α -synuclein might be related to the endogenous expression levels of α -synuclein, which is regulated in a neuronal cell type-dependent manner.

Presynaptic α -synuclein aggregates in the cortex of DLB brain correlate with reduced dendritic spines, suggesting that these aggregates contribute to synapse loss and cognitive dysfunction (Kramer and Schulz-Schaeffer 2007). It was recently demonstrated that exposure of wild-type neurons to PFFs causes a significant reduction in mushroom-like stable spine density (Froula et al. 2018). Interestingly, this reduction of spine density is observed only in wild-type neurons expressing endogenous α -synuclein, but not in α -synuclein knockout neurons. The authors hypothesized that these changes in spine morphology result from PFF-induced corruption of endogenous α -synuclein expressed in the hippocampal neurons. This latter study focused on the morphology and function of glutamatergic excitatory synapses at early pathological stages before neuronal cell death induced by PFF-treatment, and further indicated the reduced frequency and amplitudes of spontaneous Ca^{2+} transients. Thus, endogenous expression levels of α -synuclein might be a critical factor for synapse impairment at early pathological stages during the progression of neurodegeneration.

Brain region-dependent differential expression of α -synuclein

In the pathological brain, Braak and colleagues proposed a caudorostral process associated with sporadic PD progression from the lower brain stem through the basal midbrain and forebrain into the cerebral cortex (Braak and Del Tredici 2009; Braak et al. 2003). Their studies indicated specifically affected brain regions, such as the olfactory bulb, dorsal motor nucleus of the vagus (DMN), and substantia nigra at early stages of PD, and also the amygdala, hippocampus, and neocortex at later stages. As discussed above, endogenous α -synuclein expression is required for LB-like aggregate formation (Volpicelli-Daley et al. 2011; Taguchi et al. 2014). Therefore, we further investigated the precise expression profile of α -synuclein in the wild-type adult mouse brain, particularly in the vulnerable regions affected during the progression of idiopathic PD (Taguchi et al. 2016).

α -Synuclein is broadly expressed in the mouse brain (Fig. 4). There is a similar distribution pattern between vGluT-1 and α -synuclein, except for the lateral and medial globus pallidus (LGP and MGP) and substantia nigra pars reticulata (SNR) (Fig. 4a). In contrast, GAD shows a complementary weak expression in the cerebral cortex, hippocampus, thalamus, and striatum (Str), but shows a strong expression in the LGP, MGP, and SNR, where vGluT-1 expression is very weak (Fig. 4b).

As summarized in Table 1, α -synuclein is highly expressed in the neuronal cell bodies of some early PD-affected regions, such as the olfactory bulb, DMN, substantia nigra pars compacta (SNC; Fig. 5a, b), and lateral and

Fig. 4a,b Sagittal plane brain distribution of α -synuclein compared with vGluT-1 or GAD. **a** There is a similar distribution pattern between α -synuclein (α Syn) and vGluT-1 except for some regions, including the lateral globus pallidus (LGP), medial globus pallidus (MGP), and substantia nigra pars reticulata (SNR). In these regions, α Syn is colocalized with GAD (**b**). Str striatum. Bars 1 mm (Taguchi et al. 2016)

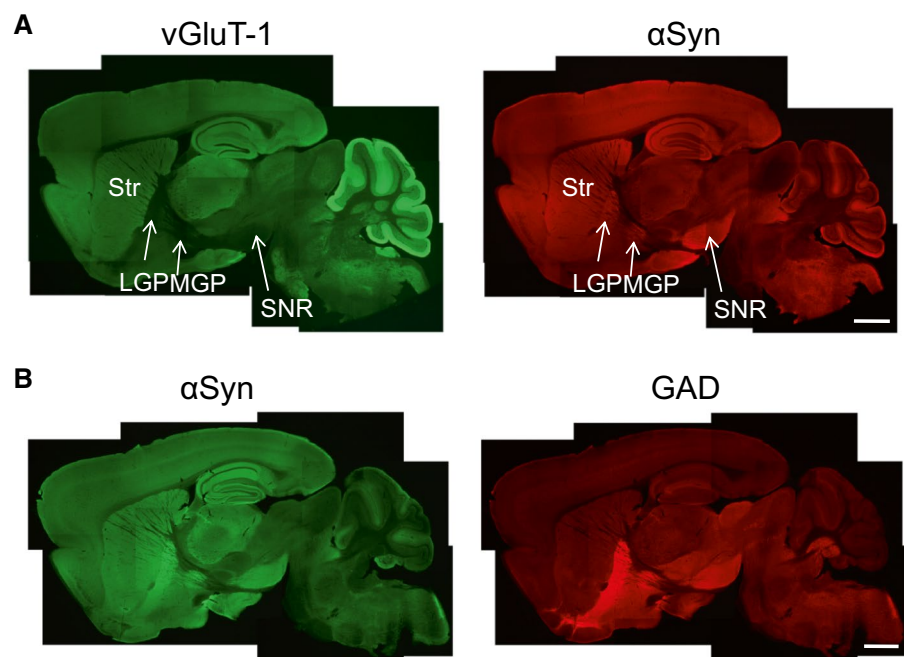


Table 1 Expression profile of α -synuclein in wild-type mouse brain. *GL* Glomerular layer, *EPL* external plexiform layer, *IML* inframitral layer, *Ant* anterior, *N* nucleus, *DMN* dorsal motor nucleus of the vagus, *GRN* gigantocellular reticular nucleus, *SNC* substantia nigra pars compacta, *SNR* substantia nigra pars reticulata, *GP* globus pallidus, *LMN* lateral mammillary nucleus, *MMN* medial mammillary nucleus. This table was modified from Taguchi et al. (2016)

	Presynaptic IR ^a	Somatic IR ^b	vGluT-1/ α Syn ^b	GAD/ α Syn ^b
Olfactory bulb ^c				
GL	++	++	+	+
EPL	+++	(-)	(-)	++
IML	++	(-)	++	(-)
Ant olfactory N	++	(-)	++	(-)
Medulla oblongata				
DMN ^c	++	++	+	+
GRN ^c	+	(-)	+	+
Raphe pallidus N ^c	+	(-)	+	+
Raphe obscurus N ^c	+	(-)	+	+
Inferior olive	+	(-)	+	+
Pons				
Locus coeruleus ^c	++	(-)	+	+
Parabrachial N ^c	+++	(-)	+	+
Midbrain				
SNC ^c	+	++	+	+
SNR	+++	(-)	+	++
Cerebrum				
Amygdala				
Central ^c	+++	(-)	+	+
Basolateral ^c	++	(-)	++	+
Cerebral cortex	++	(-)	++	(-)
Piriform cortex ^c	++	(-)	++	+
Entorhinal cortex	++	(-)	++	+
Hippocampus (CA1)	+++	(-)	++	(-)
Striatum	++	(-)	++	(-)
Lateral GP	+++	(-)	+	++
Medial GP	+++	(-)	+	++
Thalamus	++	(-)	++	(-)
Subthalamic N	+	(-)	++	(-)
LMN/MMN	++	++	+	+
Cerebellum				
Molecular layer	+++	(-)	+	(-)
Granule cell layer	+	(-)	++	(-)

^aIntensity of α -synuclein immunoreactivity graded as absent (-), weak (+), moderate (++), and strong (+++)

^b+++: clearly colocalized or positive, +: partially colocalized, (-): not colocalized or negative

^cEarly-affected PD brain regions

medial mammillary nucleus (LMN and MMN). Synaptic expression of α -synuclein is mostly accompanied by the expression of vGluT-1. In contrast, expression profiles of α -synuclein in inhibitory synapses are different among brain regions. α -Synuclein is clearly expressed in GAD-positive inhibitory synapses in the external plexiform layer of the olfactory bulb, LGP, MGP, and SNR, but not in the cerebral cortex, hippocampus, subthalamic nucleus, or thalamus. In the SNC, α -synuclein partially colocalizes with vGluT-1; there are α -synuclein-positive and -negative excitatory synapses (Fig. 5c). Regarding GAD expression, there are

also α -synuclein-positive and -negative inhibitory synapses (Fig. 5d). In this region, therefore, α -synuclein-positive synapses comprise both excitatory and inhibitory synapses. In the SNR, dense α -synuclein and GAD-immunoreactive terminals are clearly colocalized, whereas excitatory terminals are only sparsely distributed (Fig. 5c,d). Although α -synuclein is partially colocalized with vGluT-1, those excitatory terminals co-express α -synuclein. Therefore, in the SNR, α -synuclein is expressed mainly in inhibitory synapses. Conversely, in the cerebral cortex, α -synuclein is

Fig. 5a–d Localization of α -synuclein in the substantia nigra. **a** Tyrosine hydroxylase (*TH*)-positive neurons are distributed in the substantia nigra pars compacta (*SNC*) and the ventral tegmental area (*VTA*). These cells are α *Syn*-positive. **b** In *TH*-positive neurons located in the *SNC*, α *Syn* is enriched within the cell body. **c** α *Syn* is partially colocalized with vGluT-1 in the *SNC* and the substantia nigra pars reticulata (*SNR*). Although vGluT-1-positive synapses without the expression of α *Syn* are also observed in the *SNC* (*arrow*), scattered punctate signals of vGluT-1 in the *SNR* are clearly colocalized with α *Syn* (*arrowhead*). **d** α *Syn* is partially colocalized with GAD in the *SNC* (*arrowhead*). In the *SNR*, α *Syn* is clearly colocalized with GAD. *N* nucleus. Bars **a** 500 μ m, **b–d** 20 μ m (Taguchi et al. 2016)

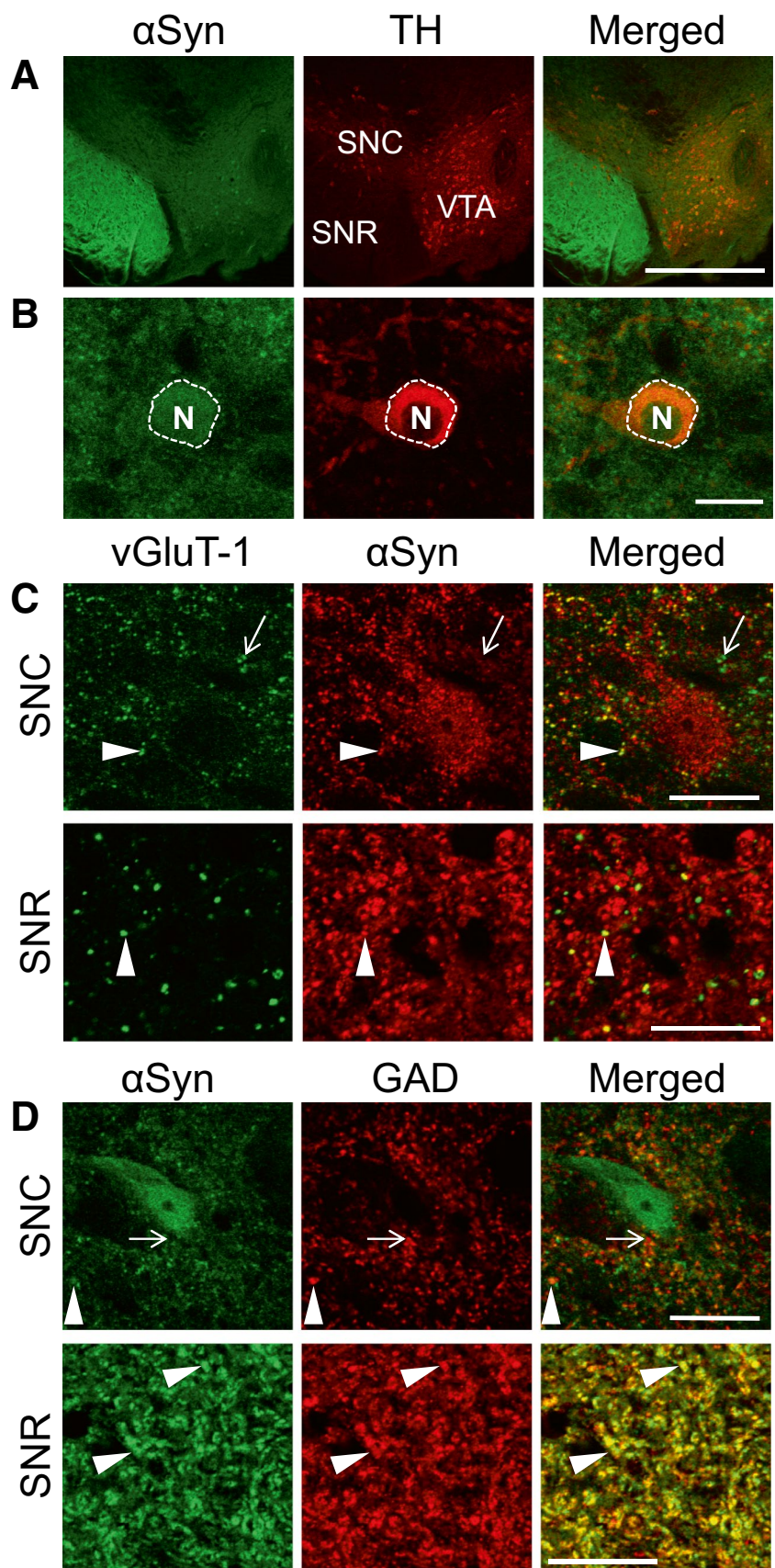
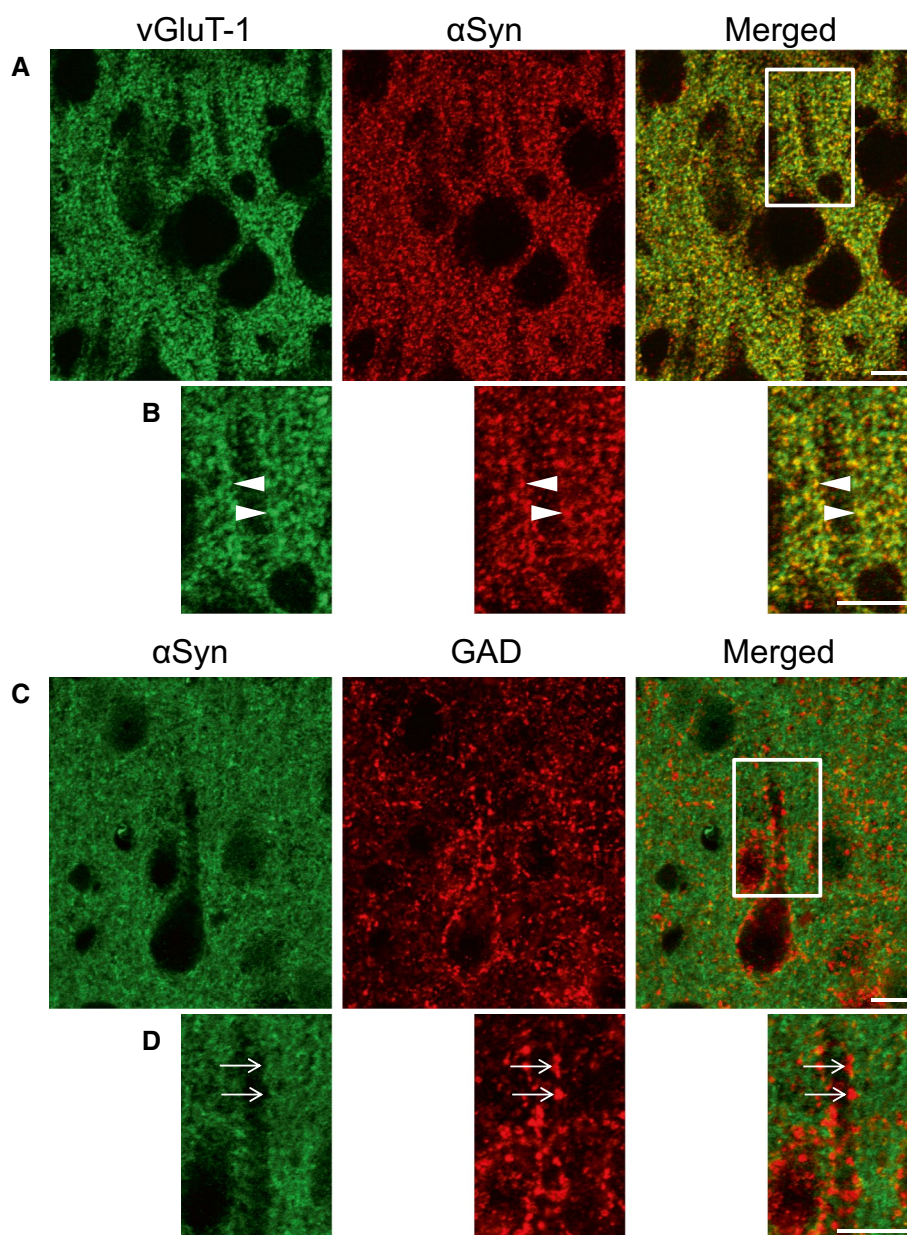


Fig. 6a–d Localization of α -synuclein in the cerebral cortex. **a, b** α Syn is clearly colocalized with vGluT-1 (*arrowheads*). **c, d** α Syn is not colocalized with GAD (*arrows*). The regions marked by *white squares* in **a** and **c** are magnified in **b** and **d**, respectively. Bars 10 μ m (Taguchi et al. 2016)



clearly expressed in excitatory synapses (Fig. 6a, b), but not in inhibitory synapses (Fig. 6c, d).

Collectively, some neurons in the early PD-affected regions express high levels of perikaryal α -synuclein, as happens in the mouse brain (Table 1). Synaptic expression of α -synuclein is different in various brain regions. Interestingly, early PD-affected regions tend to have both excitatory and inhibitory synapses expressing α -synuclein, as indicated by superscript c in Table 1. However, α -synuclein is expressed only in excitatory synapses of the affected regions, such as the cerebral cortex, hippocampus, thalamus, and subthalamic nucleus at later PD stages, although

there are also inhibitory synapses in those regions. Because α -synuclein is involved in synaptic function, expression of α -synuclein in inhibitory synapses of the early PD-affected regions might influence the local milieu and the fragility of vulnerable neurons expressing high levels of α -synuclein over time.

In the LGP, MGP and SNR, α -synuclein is clearly expressed in inhibitory synapses. These regions receive GABAergic afferents from striatal medium spiny neurons (MSNs) (Smith and Bolam 1990; Utter and Basso 2008). Therefore, it is suggested that striatal MSNs express α -synuclein (Fig. 7). However, other inhibitory neurons

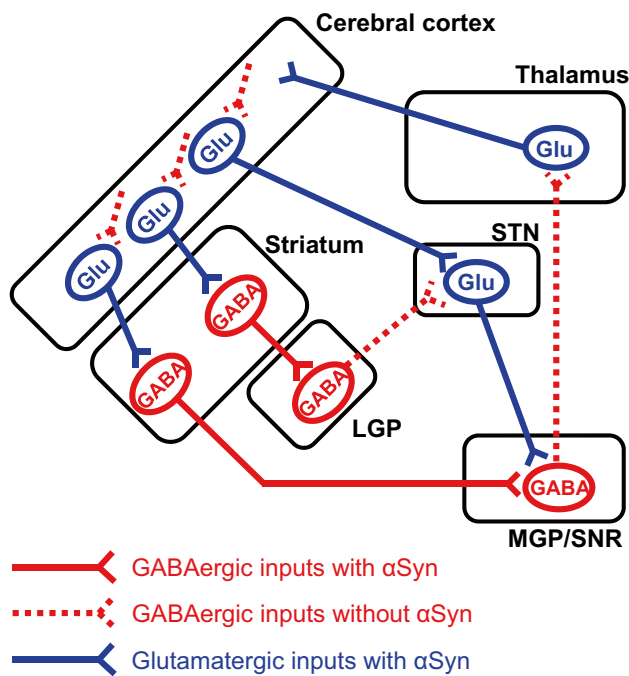


Fig. 7 Expression profile of α -synuclein in the basal ganglia circuit. α Syn-positive inhibitory afferents are located in the lateral globus pallidus (LGP), medial globus pallidus (MGP), and substantia nigra pars reticulata (SNR). These regions receive the inhibitory inputs from striatal GABAergic (medium spiny) neurons (Taguchi et al. 2016)

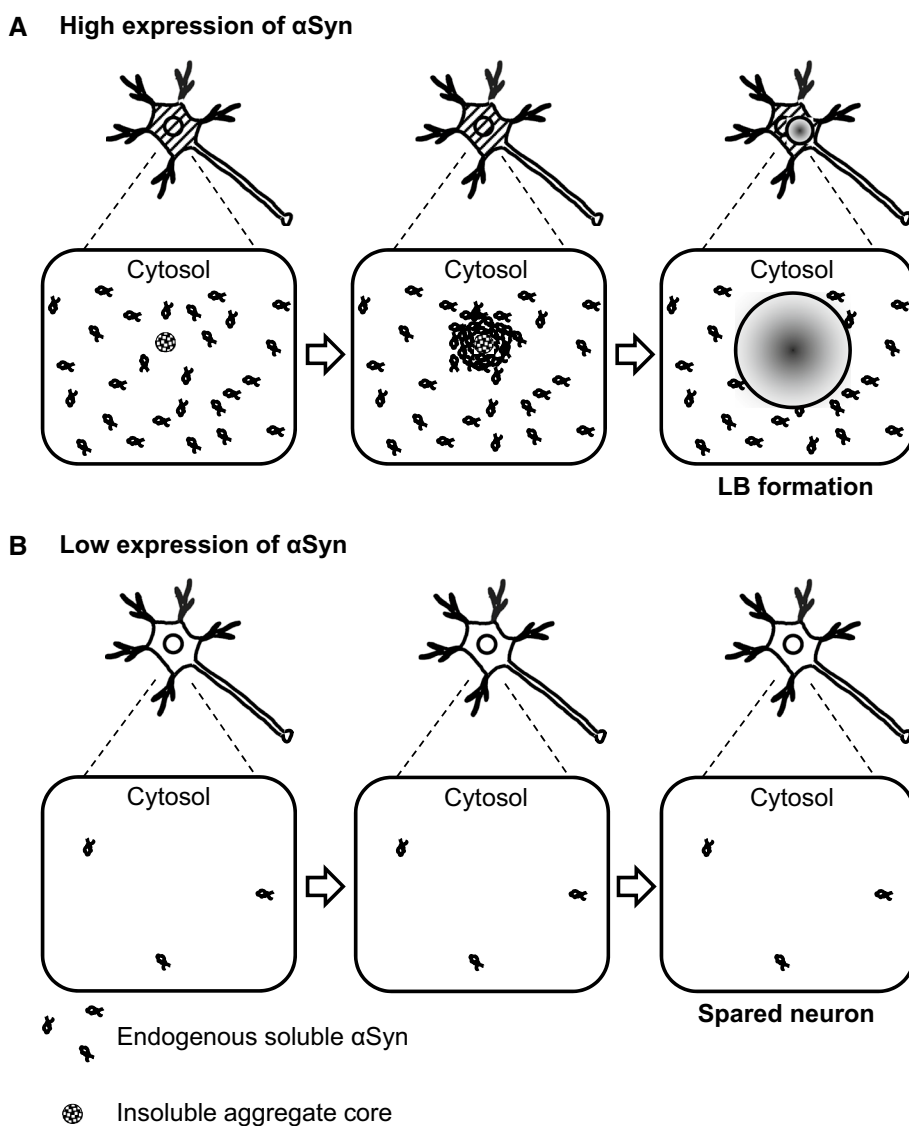
located in the LGP, MGP, and SNR project their axons to the thalamus and subthalamic nuclei, but they do not express α -synuclein. This difference might be a result of the unique neurochemical and electrophysiological properties of MSNs. Besides GABA, MSNs express substance P or enkephalin-like peptides as their neurotransmitters (Smith and Bolam 1990; McGinty 2007; Utter and Basso 2008) and fire at low rates in irregular bursts (Wilson 1993). In addition, MSN axons are unmyelinated (Miyazaki et al. 2014). Braak and Del Tredici described that neurons prone to LB pathology are projection neurons with thin, long axons that are unmyelinated or only sparsely myelinated (Braak and Del Tredici 2004, 2009). Actually, LBs are found in MSNs at stage III of the PD brain (Mori et al. 2002). In the cerebral and cerebellar cortices, GABAergic interneurons with short axons do not express α -synuclein (Taguchi et al. 2016). These results suggest that neuronal expression of α -synuclein is dependent on the morphological characteristics of axons, such as whether they are myelinated or whether they are long or short.

Selective vulnerability of α -synuclein-enriched neurons

Recently, it was reported that differential α -synuclein expression contributes to selective vulnerability of hippocampal neuron subpopulations to fibril-induced toxicity (Luna et al. 2018). Previously, Luk and colleagues reported that, in non-transgenic wild-type mice, a single intra-striatal inoculation of PFFs of α -synuclein leads to the propagation of LB-like pathology in anatomically interconnected brain regions (Luk et al. 2012). Using this experimental model, they demonstrated that Math2-expressing hippocampal *Cornu ammonis* (CA) neurons are highly susceptible to pathological seeding with the PFFs, in contrast to the dentate gyrus (DG) neurons expressing Prox1 (Luna et al. 2018). Glutamatergic excitatory neurons derived from CA regions express the transcription factor Math2, whereas DG neurons express another transcription factor, Prox1 (Bagri et al. 2002; Sugiyama et al. 2014). The hippocampal CA is a region significantly affected by α -synuclein pathology in advanced PD and DLB compared with the DG (Armstrong et al. 2014; Hall et al. 2014). Interestingly, Math2-expressing CA neurons show higher levels of α -synuclein expression, whereas Prox1-expressing DG neurons express low levels of α -synuclein and are resistant relative to the Math2-expressing subpopulations (Luna et al. 2018). These results indicate that PFF-induced cell susceptibility is closely related to the endogenous expression levels of α -synuclein, which are regulated differentially in each neuronal cell-type.

In postmortem human brains, regional levels of physiological α -synuclein are directly associated with LB pathology (Erskine et al. 2018). Quantitative imaging and western blotting analysis demonstrated that brain regions less relevant to LB pathology, such as primary visual cortex and cerebellar cortex, show strikingly lower expression levels of α -synuclein. Recruitment of soluble α -synuclein into the intracellular aggregates is required for the process of pathological LB propagation in brain, as described above. Therefore, these results are consistent with previous studies using various in vitro and in vivo models. However, brain regions with the greatest proclivity to LB pathology did not have the highest levels of endogenous α -synuclein expression, and it was suggested that expression levels of α -synuclein are not the sole determinants of cell vulnerability. Vulnerability to LB pathology is the product of anatomical connectivity and region autonomous factors, with a baseline level of physiological α -synuclein expression necessary for pathology to develop (Erskine et al. 2018). For instance, higher α -synuclein expression is observed within the cell bodies of dopamine neurons in both the SNC and ventral tegmental area (VTA). However, dopaminergic neurons in the VTA are much less affected in PD. This difference might

Fig. 8 Models of the relationship between α -synuclein expression levels and Lewy body (LB) formation. **a** LB formed by recruitment of endogenous soluble α Syn into the insoluble aggregate core. **b** Neurons with low expression of α Syn are spared from LB pathology



not be attributable to the expression levels of α -synuclein, however, because the difference in pacemaking mechanisms and engagement of L-type calcium channels in neurons of the SNC and VTA has been reported to cause a difference in their vulnerability (Guzman et al. 2010; Khaliq and Bean 2010).

Conclusion

High expression of α -synuclein is a critical risk factor for aggregate formation and neuronal loss (Fig. 8). Accumulated evidence suggests the close relationship between differential expression of α -synuclein and selective vulnerability of certain neuronal populations. Factors other than α -synuclein expression level may also be involved

in vulnerability. Further investigation of the regulation of α -synuclein expression will help understand the mechanism of LB pathology and provide an innovative therapeutic strategy to prevent PD and DLB onset.

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Compliance with ethical standards

Conflict of interest The authors declare no competing interests exist.

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