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# Advances in modelling alpha-synuclein-induced Parkinson's diseases in rodents: Virus-based models versus inoculation of exogenous preformed toxic species

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## ABSTRACT

Aggregates of alpha-synuclein ( $\alpha$ Syn) have been described in Parkinson's disease (PD) patients, and recent evidence has suggested that the most toxic  $\alpha$ Syn species in PD are small soluble aggregates including oligomers, prefibrils, protofibrils. The physiological function of  $\alpha$ Syn is still highly debated, with a possible role in synaptic vesicle trafficking and release at the presynaptic compartment, and in the regulation of gene expression in the nucleus. Emerging evidence indicate that most of  $\alpha$ Syn functions are related with the crucial ability to bind biological membranes, which is associated with structural conversion from a disordered monomer to an  $\alpha$ -helical enriched structure. Conformational properties of  $\alpha$ Syn can be modulated by a number of factors including post-translational modifications, gene duplication and triplication-driven overexpression, single point mutations, environmental changes, which affect membrane binding and the protein propensity to aggregate in toxic species. The recognized toxic role of  $\alpha$ Syn in PD has laid the rational for purposing of  $\alpha$ Syn-based, neuropathologically relevant preclinical models of PD. Different approaches have led to the establishment of transgenic models, viral vector-based models, and more recently models based on the intracerebral inoculation of exogenous  $\alpha$ Syn preformed fibrils/oligomers. Here, we overview and compare viral vector-based models of  $\alpha$ Syn overexpression and models obtained by direct intracerebral infusion of *in vitro* preformed  $\alpha$ Syn species. The advantages and pitfalls associated with these different approaches are discussed.

#### 1. Background

Parkinson's Disease (PD) is the second most common neurodegenerative disease after Alzheimer's Disease. Nowadays, PD is considered a synucleinopathy, a family of neurodegenerative diseases shearing the abnormal accumulation of misfolded, often largely phosphorylated,  $\alpha$ -synuclein ( $\alpha$ Syn) in neuronal and non-neuronal cells in the brain. Synucleinopathies including clinically different diseases such as PD, dementia with Lewy bodies, multiple system atrophy and pure autonomic failure, differ for the sites where the main protein deposition occur, however the factors leading to different clinical phenotypes remain elusive. The discovery that a Syn may form different aggregated structures in vitro, has suggested that differences in strains, but also different post-translational modifications, may account for a variety of clinical traits of synucleinopathies. In PD, neuropathological hallmarks are the progressive loss of dopaminergic neurons in the Substantia Nigra pars compacta (SNpc) (Lees et al., 2009) and the presence of deposits mainly containing fibrillar aSyn, namely Lewy Bodies (LB) (Spillantini

Aggregates of  $\alpha$ Syn have been described in sporadic PD patients and have been genetically associated with inherited forms of PD, including point mutations in the SNCA gene, namely A53 T, A30 P, E46 K, H50Q, G51D and A53E (Appel-Cresswell et al., 2013; Kruger et al., 1998; Lesage et al., 2013; Pasanen et al., 2014; Petrucci et al., 2016; Proukakis et al., 2013; Singleton et al., 2003). Although the underlying mechanisms of  $\alpha$ Syn neurotoxicity associated with PD are still largely unknown, its central neuropathological role is widely acknowledged, and has laid the rational for purposing of  $\alpha$ Syn-based preclinical models of PD. These models have been developed through different approaches, leading to the establishment of transgenic, viral vector-based, and exogenous  $\alpha$ Syn injection models (Koprich et al., 2017; Lo

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et al., 1997; Tofaris et al., 2003). Cardinal symptoms of the disease are motor impairment including bradykinesia, resting tremors, rigidity and postural instability, and a plethora of non-motor symptoms such as anxiety, depression, sleep disorders and cognitive impairments related to damage in non- dopaminergic areas (Braak et al., 2003; Chaudhuri and Odin, 2010).

Bianco et al., 2002). Modelling the aSyn-related neuropathology of the disease clearly offers a superior tool to investigate new therapeutic targets and novel neuroprotective therapies if compared with classical toxin-based models, namely the 6-OHDA rat model, and the MPTP mouse model. Ideally, experimental models of PD should display: (i) face validity, reproducing most features of the disease; (ii) predictive validity, with progressive development of the neuropathology as in PD, and therapeutic responses comparable to the human disease; (iii) construct validity, reproducing the molecular aspects of the disease. Toxin-based models only partly meet these requirements at least in rodents, where the main caveat relates to the limited predictive and construct validity, while owning face validity. Toxin-based models are based on the ability of toxins to kill dopaminergic neurons with molecular mechanisms not necessarily relevant for PD (construct validity). The neurotoxins 6-OHDA acts acutely to kill neurons, failing to model the progressive pathology of PD and aSyn-related neuropathology. For these aspects, 6-OHDA models have been extremely successful in recognizing symptomatic targets and to develop treatments of PD symptoms and drug-induced motor complications (Cenci et al., 2011), but are not recommended to investigate neurodegenerative targets and to develop disease-modifying therapies. The other widely used neurotoxin MPTP offers consistent advantages over 6-OHDA in terms of construct validity and is highly used to test neuroprotective treatments, but fails to reproduce αSyn pathology. Moreover in mice, repeated MPTP administrations are required to guarantee the chronic presence of the neurotoxin in the brain, an essential condition to increment the neurodegeneration (Schintu et al., 2009). Of note, none of the treatments showing neuroprotection in toxin models has been translated to positive results in PD patients (Athauda and Foltynie, 2015). aSyn-based models, by recapitulating relevant pathological mechanisms of the human disease, may hold a higher construct and predictive validity, and may offer a more reliable tool to investigate the neuropathology and to develop novel neuroprotective therapies.

The first αSyn-based models have been obtained by genetic manipulation, and either overexpressed human WT aSyn, modelling SNCA multiplications, or human A53 T, A30 P mutant αSyn, modelling SNCA missense mutations (Hansen et al., 2013; Kurz et al., 2010; Richfield et al., 2002; Rieker et al., 2011; van der Putten et al., 2000). Few studies generated transgenic mice overexpressing mutant  $\alpha$ Syn or its truncated form limited to dopaminergic neurons, under the control of the thyrosine hydroxylase (TH) promoter (Koprich et al., 2017; Richfield et al., 2002; Tofaris et al., 2006; Wegrzynowicz et al., 2019). Besides rodents, invertebrates such as the drosophila and the C.elegans have been used to develop a Syn-based models of PD. Although the drosophila does not have an aSyn gene ortholog, several models have been developed by overexpressing human WT or mutant aSyn in the fly (Feany and Bender, 2000; Ordonez et al., 2018). Likewise, a number of C.elegans models of PD that express a Syn have been generated (Cooper and Van Raamsdonk, 2018; Kuwahara et al., 2006; Lakso et al., 2003). In general, αSyn-based transgenic models in rodents have shown abnormalities at the striatal level, such as reduced TH-immunoreactivity and dopamine levels, without evident degeneration of dopaminergic cell bodies in the SNpc. Notably, in a recent study the transgenic overexpression of truncated aSyn in dopaminergic neurons resulted in several pathological features of PD, such as reduced dopamine levels and death of dopaminergic neurons, motor impairment, aSyn aggregation in the SNpc (Wegrzynowicz et al., 2019). Transgenic models have largely contributed to current knowledge on PD neuropathology, and have been successfully used to investigate the early stages of the disease and cognitive deficits associated with it, as extensively discussed elsewhere (Hansen et al., 2013; Hatami and Chesselet, 2015; Kurz et al., 2010; Richfield et al., 2002; Tofaris et al., 2006).

Thereafter, new models based on the overexpression of  $\alpha$ Syn have been settled by using viral vectors as an alternative approach to reproduce PD-like pathology in rats and mice (Decressac et al., 2012). Based on the recent findings pointing to oligomers, protofibrils and fibrils as the most toxic forms of  $\alpha$ Syn, PD models based on the exogenous inoculation of toxic  $\alpha$ Syn species are currently under investigation (Barrett and Timothy Greenamyre, 2015).

Here, we will discuss viral vector-based models of  $\alpha$ Syn overexpression and models obtained by direct intracerebral infusion of synthetic preformed oligomers and fibrils of  $\alpha$ Syn. We will highlight the advantages and pitfalls that in our opinion are associated with the different approaches pursued to date.

#### 2. Physiological role and conformational properties of aSyn

αSyn is a 14 kDa protein encoded by the gene SNCA, widely distributed throughout the body but highly expressed in neurons (Jakes et al., 1994; Maroteaux et al., 1988; Shibasaki et al., 1995). The physiological functions of aSyn are still highly debated. The localization in the presynaptic compartment and the association with synaptic vesicles has suggested a role in synaptic vesicle trafficking and release (Diao et al., 2013; Larsen et al., 2006; Maroteaux et al., 1988; Nemani et al., 2010), via a control of vesicle pool size, mobilization and endocytosis (Bendor et al., 2013; Faustini et al., 2018; Fusco et al., 2016; Lautenschlager et al., 2018; Vargas et al., 2014). Accordingly, aSyn interacts with presynaptic proteins such as VAMP-2 from the (SNARE)-complex (soluble N-ethylmaleimide- sensitive factor attachment protein receptor) promoting its assembly and synaptic release of neurotransmitters (Burre et al., 2010). In addition, the interaction of  $\alpha$ Syn with the vesicular transporter of monoamines (VMAT2) has been described at the presynaptic compartment (Butler et al., 2015; Guo et al., 2008; Swant et al., 2011; Wersinger et al., 2006). In line with its presynaptic role, a Syn depletion in vivo decreased the reuptake of striatal dopamine (Chadchankar et al., 2011) and caused functional deficits of the nigrostriatal dopaminergic system (Abeliovich et al., 2000). Finally, a negative role of  $\alpha$ Syn in the dopamine synthesis has been suggested (Baptista et al., 2003; Perez et al., 2002; Yu et al., 2004). All these evidences support the hypothesis that  $\alpha$ Syn plays a key role in the regulation of dopaminergic transmission at presynaptic level.

Besides the presynaptic terminal, aSyn has been identified in other cellular compartments, such as endoplasmic reticulum and Golgi apparatus (Cooper et al., 2006; Thayanidhi et al., 2010), mitochondria (Devi et al., 2008; Li et al., 2007; Nakamura, 2013) and nucleus (Goncalves and Outeiro, 2013; Kontopoulos et al., 2006; Maroteaux et al., 1988; Mori et al., 2002). In mitochondria, aSyn has been proposed to improve ATP synthase efficiency and mitochondrial function (Ludtmann et al., 2016). Within the nucleus, aSyn expression levels were shown to increase under oxidative stress conditions and appear to regulate the expression of genes related to DNA repair and mitochondrial biogenesis (Paiva et al., 2017; Siddiqui et al., 2012). Other studies have suggested that aSyn may have chaperone-like activity, sharing some homology and binding a family of cytoplasmic chaperones to induce oxidative stress and neuronal death (Chandra et al., 2005; da Costa et al., 2000; Kanda et al., 2000; Ostrerova et al., 1999). aSyn has also been associated with apoptosis, by affecting the MAPK pathway (Iwata et al., 2001; Menges et al., 2017).

Most of these functions are associated with the ability of  $\alpha$ Syn to bind biological membranes, which involves a dramatic structural conversion from an intrinsically disordered monomer in solution to a membrane-bound state that is enriched in  $\alpha$ -helical structures (Bodner et al., 2009; Chandra et al., 2003; Davidson et al., 1998; Fauvet et al., 2012; Fusco et al., 2014, 2016; Fusco et al., 2018; Maltsev et al., 2013; McLean et al., 2000; Nuber et al., 2018; Snead and Eliezer, 2014; Weinreb et al., 1996). This structural conversion is driven by 7 imperfect repeats of 11 residues located in the region 1–90 of the  $\alpha$ Syn sequence and encoding for amphipathic class A2 lipid-binding  $\alpha$ -helical segments. Other characteristic sequence properties of  $\alpha$ Syn include the N-terminal 25 residues, having the role of membrane anchor (Bodner et al., 2009; Fusco et al., 2014), the hydrophobic non-amyloid beta component (NAC) segment (residues 61–95), which is believed to play a role in  $\alpha$ Syn aggregation (Breydo et al., 2012), and the C-terminal acidic domain (96–140), containing several phosphorylation sites on Tyr-125, 133, 136 and Ser-129, crucial for protein-protein interactions (Burre et al., 2012, 2010; Woods et al., 2007), as well as for switching the conformational properties of the protein upon Ca<sup>2+</sup> binding (Lautenschlager et al., 2018).

Conformational properties of  $\alpha$ Syn and membrane binding can be modulated by post-translational modifications (PTM), including serine/ tyrosine phosphorylation and tyrosine nitration. These PTMs have been observed in both the physiological and pathological contexts involving  $\alpha$ Syn (Barrett and Timothy Greenamyre, 2015; Cariulo et al., 2019; Tenreiro et al., 2014). While phosphorylation may occur physiologically, only 4% of monomeric  $\alpha$ Syn is phosphorylated under physiological condition, while extensively phosphorylated  $\alpha$ Syn was found in synucleinopathy lesions, suggesting a pivotal role of this process in disease neuropathology (Anderson et al., 2006; Fujiwara et al., 2002).

#### 3. Aggregation of αSyn

aSyn exhibits highly complex aggregation, with primary and secondary nucleation mechanisms observed depending on the experimental conditions (Buell et al., 2014). Membrane binding has also key relevance in the aggregation process of aSyn. It has indeed been shown that the interaction with lipid bilayers can modulate the kinetics and the mechanisms of aSyn aggregation, with effects ranging from inhibition to enhancement (Necula et al., 2003; Perrin et al., 2001; Zhu and Fink, 2003). The structural details of amyloid aggregates of  $\alpha$ Syn are now known, including the characteristic cross- $\beta$  spine motif and a unique orthogonal Greek key topology (Tuttle et al., 2016). Cryo-EM structures of aSyn fibrils have also revealed a crucial interface between protofilaments spanning the region 50–57 of the  $\alpha$ Syn sequence (Guerrero-Ferreira et al., 2018). This region, which appears to stabilize the fibrillar structure of the protein, is highly relevant for hosting three pathological mutations of  $\alpha$ Syn that are associated with early forms of PD (H50Q, G51D, A53 T), suggesting possible alterations of the properties of the fibrillar form in these  $\alpha$ Syn variants.

Several factors may trigger the aggregation of  $\alpha$ Syn *in vivo*, including overexpression due to gene duplication and triplication, single point mutations, post-translational modifications and environmental changes (Barrett and Timothy Greenamyre, 2015; Fujiwara et al., 2002; Kruger et al., 1998; Lesage et al., 2013; Polymeropoulos et al., 1997; Proukakis et al., 2013; Singleton et al., 2003; Uversky et al., 2001; Zarranz et al., 2004). Intense phosphorylation was found in transgenic mice over-expressing human mutant  $\alpha$ Syn (Freichel et al., 2007; Kahle et al., 2002; Wakamatsu et al., 2007). Phosphorylation of Ser-129 has been associated with the accumulation of  $\alpha$ Syn oligomers and exacerbated the deposition of inclusions (Anderson et al., 2006; Sugeno et al., 2008), with variable lifespan depending on intramolecular interactions of these species (Alam et al., 2019).

Although LBs contain  $\alpha$ Syn in its fibrillar aggregates, several evidences suggested that the most toxic species are small aggregates that maintain a degree of solubility, including oligomers, prefibrils, protofibrils. These diffusible aggregates have been observed in degenerating areas in the PD brain (Karpinar et al., 2009; Sharon et al., 2003; Winner et al., 2011) and in biological fluids of PD patients (Majbour et al., 2016; Tokuda et al., 2010), suggesting that LBs

may represent a protective mechanism that sequesters the toxic species from the cytoplasm (Bengoa-Vergniory et al., 2017; Bucciantini et al., 2002; Espa et al., 2019; Muchowski, 2002; Soto and Estrada, 2008).  $\alpha$ Syn mutations that increase the propensity to form oligomeric species over fibrils displayed the most toxic effect in cell lines, primary neurons and in dopaminergic neurons of *C. -elegans* and Drosophila (Karpinar et al., 2009). This concept was further demonstrated *in vivo* by Winner and colleagues, showing that inoculation of  $\alpha$ Syn variants that form oligomers caused the most severe dopaminergic loss in the rat SN, whereas the  $\alpha$ Syn variants that rapidly formed fibrils were less toxic (Winner et al., 2011).

#### 4. Toxic mechanisms of αSyn

#### 4.1. $\alpha$ Syn oligomers/prefibrils interaction with neurons

Several mechanisms have been described to mediate the toxicity of  $\alpha$ Syn oligomers against neurons (Villar-Pique et al., 2016). Toxic mechanisms may involve a direct interaction with neuronal membranes or subcellular organelles, as well as the interaction with glial cells, such as microglia and astroglia.

A widely accepted primary mechanism for the neurotoxicity of aSyn oligomers is the disruption of the membrane integrity (Stockl et al., 2012; van Rooijen et al., 2010), causing calcium influx into the cytoplasm (Angelova et al., 2016). Recent studies have identified the key elements for this mechanism (Fusco et al., 2017). In particular, it was shown that the disruption of biological membranes by these  $\alpha$ Syn oligomers requires a highly lipophilic element, the N-terminal region that effectively anchors the oligomers onto the membrane surface, and a fibrillar core spanning segments of the NAC, which effectively inserts into the membrane interior and disrupts its integrity. The permeabilization of the cellular membrane has also been proposed to occur via annular pore-like oligomers in studies of mutational variants of αSyn (Danzer et al., 2007; Lashuel et al., 2002; Tosatto et al., 2012; van Rooijen et al., 2010; Volles et al., 2001). Indeed, the significant heterogeneity of a Syn oligomers is likely to be associated with a variety of multiple pathways of membrane binding and disruption, including the stabilization of pre-existing membrane defects, the alteration of membrane electrical properties, and the reduction of neuronal excitability (Chaudhary et al., 2016; Kaufmann et al., 2016). Notably, targeting the interaction of  $\alpha$ Syn oligomers with neuronal membrane protected from αSyn neurotoxicity (Cascella et al., 2019; Ysselstein et al., 2017).

Intracellularly, aSyn oligomers may induce toxicity via the inhibition of tubulin polymerization and cytoskeletal damage (Chen et al., 2007), oxidative stress into the endoplasmic reticulum (Colla et al., 2012), mitochondrial dysfunction (Di Maio et al., 2016; Lindstrom et al., 2017; Nakamura, 2013; Plotegher et al., 2014; Prots et al., 2018) and impairment of the protein degradation systems (Emmanouilidou et al., 2010b; Tanik et al., 2013; Xilouri et al., 2009). Mitochondria are a main target of a Syn toxicity, as suggested by the presence of aSyn aggregates within mitochondria in dopaminergic neurons of PD brains and their correlation with complex I dysfunction (Devi et al., 2008). Transgenic mice overexpressing human αSyn A53 T display damaged mitochondria preceding neurodegeneration, that are defectively removed by the autophagic system (Chen et al., 2015). aSyn interact with mitochondria with the N-terminal region, resulting in the impairment of mitochondrial membrane potential and permeability, and increase of ROS levels (Sarafian et al., 2013; Shen et al., 2014). When the damage was investigated in relation with different aSyn species, oligomers specifically inhibited the mitochondrial protein influx (Di Maio et al., 2016) and disrupted the mitochondrial axonal transport (Prots et al., 2018), consistent with the observation of a structure-related toxicity against multiple cellular compartments.

The neuronal clearance mechanisms, including the autophagy-lysosomal pathway (ALP) and the ubiquitin-proteasome system, are targeted by  $\alpha$ Syn (Stefanis et al., 2019; Xilouri et al., 2013). The PD brain displays a depletion of lysosomes and lysosomal markers in nigral neurons positive for  $\alpha$ Syn inclusions (Chu et al., 2009; Dehay et al., 2010). Several studies have shown that ALP dysfunction increases the accumulation of aggregated  $\alpha$ Syn, which in turn impairs the clearance machinery, suggesting a reciprocal deleterious relationship (Cuervo et al., 2004; Tanik et al., 2013). Moreover, mutant  $\alpha$ Syn may inhibit the proteasome activity and induce the accumulation of ubiquitin-positive deposits (Stefanis et al., 2019; Tanaka et al., 2001). Notably, soluble  $\alpha$ Syn oligomers specifically bind to proteasome particles and inhibit their function (Emmanouilidou et al., 2010b; Lindersson et al., 2004; Snyder et al., 2003).

Several studies have demonstrated that oligomers are the species mostly responsible for aSyn synaptic toxicity. In dopaminergic terminals, aSyn plays a key functional role in the presynaptic SNARE complex, cooperating with the protein synapsin III to regulate dopamine release (Longhena et al., 2018). In contrast, aSyn oligomers are synaptotoxic and negatively impact the neuronal signaling and excitability by targeting specific synaptic components (Choi et al., 2013; Diogenes et al., 2012; Kaufmann et al., 2016). In vitro studies demonstrated that αSyn oligomers displayed a negative impact on the development of long-term potentiation in hippocampal slices (Diogenes et al., 2012) and decreased the excitability of pyramidal neurons (Kaufmann et al., 2016). Moreover, the expression in vivo of the oligomer-prone variant E57 K induced synaptic dysfunction via the disruption of presynaptic vesicles (Rockenstein et al., 2014). Interestingly, synapsin III seems to be directly involved in the  $\alpha$ Syn aggregation process, since  $\alpha$ Syn overexpression in synapsin III KO mice neither led to deposition of aggregated or phosphorylated a Syn, nor to synaptic damage and degeneration (Faustini et al., 2018). Accordingly, synapsin III was found to colocalize with a Syn aggregates within LB of PD patients (Longhena et al., 2018). Recently, Spillantini and coworkers showed that, in transgenic mice expressing aggregation-prone truncated  $\alpha$ -Syn, the progressive aSyn aggregation in striatal terminals was associated with the reduction of dopamine release, that was restored by aggregation inhibitors (Wegrzynowicz et al., 2019).

#### 4.2. $\alpha$ Syn oligomers interaction with the glia

Importantly,  $\alpha$ Syn oligomers released from neurons may spread in the brain and establish contacts with microglia and astroglia, which are the cells in charge of clearing extracellular protein aggregates (Emmanouilidou and Vekrellis, 2016; Stefanis et al., 2019; Yamada and Iwatsubo, 2018). Deficits in these clearing mechanisms may result in an increased concentration of the extracellular protein and consequent neurotoxicity. Moreover, the interaction with microglia induces an unbalanced activation of these cells ultimately leading to microglia-mediated neuronal toxicity. In addition, uptake of  $\alpha$ Syn by astrocytes induces the production of pro-inflammatory mediators by these cells, likely contributing to the inflammatory milieu (Lee et al., 2010).

It has been proposed that  $\alpha$ Syn, both as WT or mutational variants, behaves as a chemoattractant for microglia (Kim et al., 2009) and post-mortem studies found activated microglia in close contact with neurons showing  $\alpha$ Syn deposits (Croisier et al., 2005). A number of *in vitro* studies have demonstrated that aggregated and mutated  $\alpha$ Syn stimulate pro-inflammatory responses in microglia, and the release of pro-inflammatory mediators (Klegeris et al., 2008; Lee et al., 2010). Nevertheless, physiological levels of  $\alpha$ Syn are essential to modulate microglia function and promote phagocytosis, since microglia from mice lacking  $\alpha$ Syn show an exaggerated response to LPS, displaying a pro-inflammatory profile and impaired phagocytic function

(Austin et al., 2006). Moreover, oligomer/protofibril forms of the protein hold a greater inflammatory potential when interacting with microglia than the native monomeric protein (Wilms et al., 2009; Zhang et al., 2005). Specific interaction of αSyn with microglia occurs mainly via the Toll-like receptors (TLRs), involving both the TLR2 and TLR4 subtypes, leading to NF-kB nuclear translocation and affecting the phagocytic function of these cells (Fellner et al., 2013; Stefanova et al., 2011). Accordingly, high levels of both the TLR2 and TLR4 were found in proximity of a Syn deposits in peripheral immune cells and in microglia (Doorn et al., 2014; Drouin-Ouellet et al., 2014). Although data are not univocal, several studies have demonstrated that the interaction with TLRs is also conformation-specific (Fellner et al., 2013; Kim et al., 2013). Oligomeric αSyn but not the monomer binds to microglia via TLR2, triggering the inflammatory cascade via NF-kB activation (Daniele et al., 2015; Kim et al., 2013). TLR4 are involved in microglial phagocytosis of extracellular aSyn (Stefanova et al., 2011). The phagocytic process seems to depend upon levels and type of  $\alpha$ Syn although it has been proposed that  $\alpha$ Syn, both as WT or mutational variants, behaves as a chemoattractant for microglia (Lee et al., 2008; Park et al., 2008). Park et al. (2008) showed that αSyn monomers stimulated while αSyn oligomers inhibited both basal and LPS-induced phagocytosis (Park et al., 2008). Microglia incubated with A53 T mutant a Syn displayed a pro-inflammatory profile which was associated with impaired phagocytic function (Rojanathammanee et al., 2011). In contrast, Roodveldt et al. (2010) showed that both WT and A53 T aSyn similarly promoted the phagocytosis in microglia, while the A30 P and E46 K asyn induced the opposite effect (Roodveldt et al., 2010). Finally, the exposure to either soluble monomers or fibrillar or truncated aSyn increased the phagocytic activity of microglia, but fibrils induced a milder inflammatory response than the other forms (Fellner et al., 2013). Despite the discrepancies, these studies clearly highlight that the microglia phenotype and phagocytic function are differently affected by a Syn structural variants.

#### 4.3. Seeding activity of $\alpha$ Syn oligomers

Increasing evidence suggests that oligomers of aSyn can induce misfolding and aggregation of protein monomers with a seeding effect (Kordower et al., 2008; Li et al., 2008, Olanow and Prusiner 2009). Several studies in vitro and in vivo suggested that the oligomeric forms of  $\alpha$ Syn may induce the intracellular  $\alpha$ Syn aggregation and in turn may promote the spreading of protein aggregates (Danzer et al., 2009; Hansen et al., 2011). Extending this concept, aberrant  $\alpha$ Syn was able to induce misfolding in several endogenous proteins (Breydo et al., 2012; Guo et al., 2013; Peelaerts et al., 2015). Besides intracellular spreading, a further concern is the capability of a damaged cell to entangle nearby cells. Several mechanisms have been proposed to explain the cell-to-cell propagation, including passive diffusion (Ahn et al., 2006; Chandra et al., 2003; Grozdanov and Danzer, 2018), membrane pores (Stockl et al., 2013), exosomal transport (Emmanouilidou et al., 2010a), tunneling nanotubes (Abounit et al., 2016; Dieriks et al., 2017), and the transport through carrier proteins (Sung et al., 2001; Yang et al., 2017).

Interestingly the seeding activity of oligomers seems to be structure-specific. Long and stable  $\alpha$ Syn oligomers promoted the aggregation of soluble  $\alpha$ Syn more powerfully than low-molecular weight  $\alpha$ Syn oligomers (Pieri et al., 2016). Accordingly, small annular oligomers displayed low seeding activity, despite the toxic activity *in vitro*, while larger oligomers induced protein seeding despite a low *in vitro* toxicity (Danzer et al., 2007). Of note, following phagocytosis of  $\alpha$ Syn oligomers, failure in the degradation process within the phagolysosome may lead to their release into the cytoplasm, where oligomers may seed the aggregation of  $\alpha$ Syn from the physiological pool (Flavin et al., 2017; Freeman et al., 2013; Grozdanov and Danzer, 2018).

#### 5. αSyn-based rodent models of Parkinson's disease

The transgenic models developed to overexpress either WT or pathological aSyn still represent an excellent tool to investigate pre-motor and cognitive symptoms associated with idiopathic or genetic PD (Hatami and Chesselet, 2015). In the last decade, the need of specifically investigate the seeding mechanism triggered by pathological  $\alpha$ Syn and the consequent spreading pathology, and the need to investigate the role of  $\alpha$ Syn in the later and more severe phases of PD, has motivated the researchers to develop alternative a Syn-based models of PD. Currently two approaches are being pursued, one of which involves the local overexpression of WT or mutant aSyn in specific brain regions via the inoculation of viral vectors to drive the protein expression, the other one involving the direct inoculation of exogenous aggregated aSyn, structured in different species such as fibrils, protofibrils and oligomers. Critical parameters when comparing the validity of the models are the extent and progression of the cell damage/loss at the inoculation site or distant sites and the occurrence of non-specific damage, together with the presence of the associated motor impairment and neuroinflammation, and their reproducibility by independent studies.

#### 5.1. Virus-based models

Viral vector-based models of PD have rapidly become a widely used model to mimic PD neuropathology and represent a useful tool to investigate neuroprotective strategies. The choice of virus, animal and site of inoculation can differ from model to model. Both lentivirus and adenovirus have been used to overexpress  $\alpha$ Syn into dopaminergic areas (Chung et al., 2009; Kirik et al., 2002; Lo Bianco et al., 2002; Winner et al., 2011), however the adeno-associated virus (AAV) is most commonly used to overexpress either WT or A53 T  $\alpha$ Syn most

Table 1

Key Features of Virus-based models.

likely because it allows to target neurons more specifically (Decressac et al., 2012; Kirik et al., 2002; Low and Aebischer, 2012; St Martin et al., 2007; Ulusoy et al., 2010). Among rodents, rats are mostly used for virus-based models, however an AAV-aSyn mouse model has been proposed as well (St Martin et al., 2007). As far as the site of virus inoculation the neuropathological effects of both intranigral and intrastriatal a Syn overexpression have been investigated. In particular the first pioneering viral vector-based studies date back to 2002, when at least three independent articles appeared, followed by similar short after, that used either lentiviral or first generation adenoviral vectors (LV and rAAV2/2, see also Table 1) to overexpress WT, A53 T or A30 P human αSyn within the SNpc (Chung et al., 2009; Kirik et al., 2002; Klein et al., 2002; Lo Bianco et al., 2002; Yamada et al., 2004) and the striatum (Kirik et al., 2002). By using a specific antibody against human  $\alpha$ Syn, these studies have observed an extensive immunoreactivity (IR) in the injection area, either the SNpc (Chung et al., 2009; Kirik et al. 2002, Lo Bianco et al., 2002) or the striatum (Kirik et al., 2002). Moreover, aSyn IR extended from the injected SNpc to the SN reticulata, the ventral tegmental area and some adjacent nuclei as early as 3 weeks after the virus infusion, and to terminal fields including the striatum, accumbens and cingulate cortex, suggesting that a Syn was intra-axonally transported from the infusion site to terminals (Kirik et al., 2002). This was associated with neuropathological signs such as granular cytoplasmic inclusions positive for human αSyn that firstly appeared within SNpc neurons followed by striatal terminals. Importantly, these studies reproduced a progressive and specific loss of dopamine neurons in the SNpc and of striatal fibers, suggesting that the virus-based approach may mimic advanced stages of PD (Kirik et al., 2002; Lo Bianco et al., 2002).

However, these studies also highlighted how the nigral cell loss was extremely variable, ranging from 30 % to 80 %, and did not differ between WT and mutant  $\alpha$ Syn-infused animals (Kirik et al., 2002). Moreover, despite the sometime great loss of DA neurons, rats did not

Model	cell loss in SNpc (%)*	loss of striatal DA and/or fibers	P-αSyn pathology	neuroinflammation	motor impairment
Virus- based	20. 00 (2. 27 weeks) (Virile et al. 2002; Churg et al. 2000)	Vec (Visile	ND	Inflormations.	NC (Winile
Generation	50-60 (3-27 weeks) (kirik et al, 2002; Cilding et al, 2009)	et al, 2002; Chung et al, 2009)	INK	citokines and activated microglia (Chung et al, 2009)	NS (RITIK et al, 2002)
rAAV2/2 LV	15–35 (3–20 weeks) (Lo Bianco et al, 2002)	Yes (Lo Bianco et al. 2002)	NR	NR	NR
2 <sup>nd</sup> Generation (rAAV1/2, rAAV2/5 and others)	28–80 (4–26 weeks) (Gorbatyuk et al, 2008; Koprich et al, 2011)	Yes (Gorbatyuk et al, 2008; Koprich et al, 2011)	NR	NR	Yes (Koprich et al, 2011)
3 <sup>rd</sup> Generation (rAAV2/6 or rAAV2/7)	20–80 (3–44 weeks) (Decressac et al, 2012; Van der Perren et al, 2015)	Yes (Decressac et al, 2012; Van der Perren et al, 2015)	Yes (Van der Perren et al, 2015)	NR	Yes (Decressac et al, 2012; Van der Perren et al, 2015)

\*Progression of cell loss in SNpc (percentage of contralateral side). The time lapse (in weeks) for the neurodegeneration to occur is specified.

 $AAV = Adeno-associated virus; DA = Dopamine; LV = Lentivirus; NR = not reported; NS = not significant; P-\alpha Syn = Phospho \alpha-Synuclein; SNpc = Substantia nigra pars compacta.$ 

show a significant motor impairment, which may reflect the high variability reported in neurodegeneration among animals (Kirik et al., 2002).

These studies have clearly demonstrated that the extent of  $\alpha$ Syn-induced neuropathology depends on the level of protein expression, and most importantly that high levels of expression are necessary to induce a significant pathology with motor impairment (Koprich et al., 2011). Thereafter, a second generation of AAV vectors carrying a higher expression potential such as rAAV1/2, rAAV2/5, have been developed and tested by both intranigral and intrastriatal infusions (Table 1) (Gorbatyuk et al., 2008; Koprich et al., 2011). However, high titers of GFP-carrying vectors used as a control were toxic themselves, introducing a factor of non-specific toxicity (Koprich et al., 2017).

Nevertheless, this aspect offers the appealing opportunity to modulate the vector titer in order to reach a cell loss of various extent and to investigate different stages and aspects of the disease. To overcome this pitfall, subsequent studies have infused into the rat SNpc third generation AAV serotypes with higher transduction efficiency and transgene expression (rAAV2/6 and rAAV2/7, Table 1) (Van der Perren et al., 2011), with a modified vector construct, which included an enhancer element in order to potentiate  $\alpha$ Syn expression (Decressac et al., 2012). These studies reported a significant loss of dopamine neurons in the SNpc, drop of striatal dopamine, and a frank motor impairment, within 4 months from infusion. Moreover, histopathological features, such as phosphorylated  $\alpha$ Syn, were observed both in the SN and striatum of AAV- $\alpha$ Syn infused rats (Decressac et al., 2012; Van der Perren et al., 2015).

An intense and persistent neuroinflammatory milieu in mesencephalic areas is an important recognized hallmark in PD neuropathology. However, reports of the inflammatory response in the striatum and SN following AAV-αSyn infusion are still sporadic. One study showed that intranigral AAV-A53 T aSyn infusion induced an increase of markers of microglia reactivity and of inflammatory cytokines in the Str but not in the SN, that preceded the neurodegenerative event (Chung et al., 2009). Other studies reported that the intranigral infusion of AAV-WT  $\alpha$ Syn induced a transient microglia activation in the SNpc in the absence of cell loss, that was slightly more persistent when dopamine neurons degenerated (Sanchez-Guajardo et al., 2010; Theodore et al., 2008). Interestingly, the injection of AAV-GFP as a control vector also induced some degree of microglia activation (Sanchez-Guajardo et al., 2010) suggesting that the high sensibility of microglia may strengthen the toxic effect of non-selective or weaker toxic tools. Altogether these studies concordantly suggest that microglial pathology precedes neuronal degeneration in these models. Intriguingly, the neuroinflammatory response may start in the projection area with respect to the AAV- $\alpha$ Syn infusion site, *i.e.* in the striatum when infusion is made within the SNpc (Chung et al., 2009). Several in vitro studies have shown that the inflammatory response to aSyn is structure-related as aggregated  $\alpha$ Syn produced a stronger response than monomeric  $\alpha$ Syn (Klegeris et al., 2008; Lee et al., 2010). Although seeding and spreading of toxic  $\alpha Syn$  species from the infusion site have been poorly characterized in the AAV-aSyn models, it is conceivable that toxic aggregates of human aSyn are formed in the infusion site and are axonally transported to striatal terminals, were they are released extracellularly and interact with local microglia to trigger the inflammatory phenotype. These findings pose a question on the validity of AAV-aSyn models to reproduce the chronic inflammatory milieu typical of PD, and to investigate inflammatory-related neuropathological processes.

In synopsis, most recent AAV- $\alpha$ Syn models recapitulate several hallmarks of PD neuropathology such as the progressive loss of dopaminergic cells, the increasing motor impairment and the deposits of pathological  $\alpha$ Syn within neurons, suggesting that they are reliable models in relation to face and construct validity, however a high variability of results has been considered. Notably, great advances have been made in recent years in modelling AAV- $\alpha$ Syn-induced neuropathology in monkeys (Eslamboli et al., 2007; Kirik et al., 2003; Koprich et al., 2016). These studies demonstrated the neurotoxic effect of AAV-WT  $\alpha$ Syn and to a greater extent of AAV-A53 T  $\alpha$ Syn inoculated into the monkey SN, showing a slow loss of dopaminergic neurons associated with the development of motor impairment, suggesting that the AAV- $\alpha$ Syn models may hold a pivotal translatability value and predictive validity.

As compared to transgenic models, viral-based aSyn overexpression offers some concrete advantages. The use of high titer viral vectors or highly efficient vectors yields higher gene expression and a more intense neurodegeneration, that enables to model the advanced disease stages better than transgenesis techniques; viral-based models can be established in rats, which allow a more accurate behavioral and histological evaluation than mice; local injection of viral vectors allows to target specific brain regions sparing non-dopaminergic areas; finally, infusion can be performed unilaterally in order to use the contralateral side as a control. As mentioned above, limitations of some AAV-aSyn models are represented by the inadequacy in reproducing some aspects of PD such as the chronic intense neuroinflammatory response, particularly in mesencephalic areas. The other methodological limitation is the need to express levels of the protein much higher than in human disease or physiological condition in order to induce a dopaminergic cell loss, and consequently a high variability in the degree of neurodegeneration can be observed. Moreover, this model requires to titer the virus for each new batch, that is to measure the number of viral particles per ml. In addition, level of expression does not seem to depend solely upon the number of particles injected, but also on the viral infectivity potential, suggesting that vector batches holding similar titer may result in different infectivity and potentially different expression levels (Sanchez-Guajardo et al., 2010).

An interesting finding that came out from studies in the virus-based model is that, differently from the human form, the overexpression of rat  $\alpha$ Syn did not induce any cell damage despite the presence of  $\alpha$ Syn deposits within neurons (Lo Bianco et al., 2002). *In vitro* studies have demonstrated that the rodent  $\alpha$ Syn aggregates in amyloid fibrils more rapidly than the human WT or mutated A53 T/A30 P protein, which first aggregates in nonfibrillar soluble oligomers instead (Rochet et al., 2000). These findings are an evidence that soluble protein oligomers are the most toxic form of  $\alpha$ Syn. Moreover, they suggest that the expression of a large amount of  $\alpha$ Syn is needed in virus-based models in order to produce toxic oligomers responsible for the dopaminergic neurodegeneration.

#### 5.2. Exogenous $\alpha$ Syn injection

These models are based on the unilateral or bilateral infusion of soluble oligomers of aSyn, or preformed fibrils (PFFs) of various length, or human brain tissue homogenate containing LB, into the SNpc (Abdelmotilib et al., 2017; Masuda-Suzukake et al., 2013; Peelaerts et al., 2015; Recasens et al., 2014; Thakur et al., 2017) striatum (Abdelmotilib et al., 2017; Blumenstock et al., 2017; Espa et al., 2019; Luk et al., 2012a; Paumier et al., 2015; Recasens et al., 2014), olfactory bulb (Rey et al., 2016) or more diffusely into the brain via intracerebroventricular (ICV) injection (Fortuna et al., 2017). Most of the studies addressing the properties of aSyn oligomers have been based on the generation of toxic oligomers in vitro, as the isolation of pure samples of oligomers ex vivo has revealed highly challenging. Some of the methods include the formation of highly pure oligomers using lyophilization steps in the incubation protocol. The resulting oligomers provide the advantage to be relatively homogeneous and induce neurotoxic effects to neuronal cells that are similar to those observed in cells carrying a triplication in the SNCA gene (Angelova et al., 2016; Deas et al., 2016; Devine et al., 2011). Injections of oligomers/PFFs are typically made by stereotaxic surgery using a glass capillary or a steel needle connected to a Hamilton syringe. The main aim of these models has been to investigate the toxicity of  $\alpha$ Syn strains in terms of seeding and spreading properties, and the results obtained have largely increased our knowledge of the mechanisms involved in these processes. Moreover, the oligomers or PFFs inoculation into dopaminergic areas has been performed with the specific aim of pursuing a valuable PD model that might reproduce the cardinal features of the disease, including cell loss, motor impairment, pathological deposits and neuroinflammatory response, or prodromal/non motor symptoms (Abdelmotilib et al., 2017; Espa et al., 2019; Rey et al., 2016; Thakur et al., 2017) (Table 2).

Upon neuron internalization, aSyn oligomers or fibrils may interact with endogenously expressed a Syn to seed new inclusions, and propagate within the cellular compartments in a retrograde or anterograde direction (Volpicelli-Daley et al., 2016). Fibrils may be transmitted to neighboring neurons or released from degenerating neurons triggering the spreading process. Results from these studies strongly suggest that structural and length differences in a Syn species are critical for determining the seeding potential and the spreading of pathology within the brain. Most soluble species, i.e. oligomers and short fibrils, hold the highest spreading property, however fibrils seem mostly responsible for the formation of intraneuronal p-aSyn deposits, which is the seeding hallmark (Abdelmotilib et al., 2017; Luk et al., 2012b; Masuda-Suzukake et al., 2013; Peelaerts et al., 2015). On the other hand, studies using preformed toxic species to model PD neuropathology have consistently shown that monomeric aSyn did not induce any αSyn pathology in any area and at any time (Luk et al., 2012b; Masuda-Suzukake et al., 2013). These studies have clearly demonstrated that the propagation of the  $\alpha$ Syn pathology occurs in functionally interconnected areas and is time-dependent. Moreover, the spreading pattern was observed after the inoculation of  $\alpha$ Syn PFFs in different brain areas, including the SNpc and striatum, confirming in vivo the concept that the propagation of toxic species may occur with a retrograde or anterograde direction, as previously demonstrated by in vitro studies

#### Table 2

Key Features of Exogenous αSyn inoculation-based models.

(Luk et al., 2012b; Masuda-Suzukake et al., 2013; Volpicelli-Daley et al., 2016).

In SN-injected mice, the p-aSyn pathology was observed in the striatum and amygdala, both projection areas of the SN, and in the stria terminalis, which receives amygdala projections, suggesting that the pathology spreads unidirectionally through the neural circuit (Masuda-Suzukake et al., 2013). Of note, when the toxic species used for these models were of human origin, a mouse Ab was used to detect p-αSyn deposits (Masuda-Suzukake et al., 2013). In addition, while the human protein disappeared briefly after inoculation, the mouse pathological protein was detected up to 15 months after inoculation (Masuda-Suzukake et al., 2013). This suggests that the inoculated αSyn species converted endogenous αSyn into toxic forms, and supports the concept that the transmission of pathological αSyn, recruitment and conversion of the endogenous protein is critical for the progression of PD neuropathology in these models. The inoculation of LB-enriched fractions obtained from the SN of PD brains into the mice SN induced a slow spreading of  $p-\alpha Syn$  in anatomically interconnected brain regions such as the striatum and cingulate, motor, somatosensory cortices (Recasens et al., 2014).

In striatum-injected mice or rats, PFFs were unilaterally inoculated by a single injection within the dorsal striatum at variable coordinates in mice (AP + 0.2–1.0 mm; ML + 1.85–2.0 mm; DV -2.6-3.0 mm from skull) (Abdelmotilib et al., 2017; Luk et al., 2012b),or rats (AP + 0.7–1.6 mm; ML + 2.4–3.0 mm; DV -4.2–5.5 mm from skull or beneath the dura) (Abdelmotilib et al., 2017; Paumier et al., 2015) or by two injections within the dorsomedial and dorsolateral striatum (Paumier et al., 2015). In this model, the spreading of p- $\alpha$ Syn was observed with sequential involvement of interconnected areas, including pre-frontal, insular, cingulate and motor cortical regions (layers IV and V), olfactory bulb and amygdala, followed by ventral striatum, thalamus, occipital cortex and SNpc.

WT mouse  $\alpha$ -syn PFFs or WT human  $\alpha$ -syn PFFs have been infused directly into the mouse olfactory bulb in order to investigate the relationship between  $\alpha$ Syn deposits and prodromal symptoms of PD, namely olfactory dysfunction, and to characterize the spreading of p- $\alpha$ Syn pathology in the early disease stages, based on Braak hypothesis

Model	cell loss in SNpc (%)*	loss of striatal DA and/or fibers	P-αSyn pathology	neuroinflammation	motor impairment
Exogenous α- synuclein - based	(				
SNpc infusion	25–50 (3–68 weeks) (Abdelmotilib et al, 2017; Recasens et al, 2014; Thackur et al., 2017)	Yes (Abdelmotilib et al, 2017; Recasens et al, 2014; Thackur et al., 2017)	Yes (Abdelmotilib et al, 2017; Masuda-Suzukake et al, 2013; Peelaerts et al, 2015; Recasens et al, 2014; Thackur et al., 2017)	Transient microglial activation (Thakur et al, 2017) astrogliosis (Recasens et al, 2014)	Yes (Recasens et al, 2014)
Str infusion	15–35 (8–24 weeks) (Luk et al, 2012a; Paumier et al, 2015)	Yes (Luk et al, 2012a; Paumier et al, 2015)	Yes (Abdelmotilib et al, 2017; Blumenstock et al, 2017; Luk et al, 2012a, 2012b; Paumier et al, 2015)	NS (Blumenstock et al, 2017)	Yes (Luk et al, 2012a)
<i>OB infusion</i> (Rey et al, 2016)	NO	NO	Yes	NS	NS
ICV infusion (Fortuna et al, 2017)	45 (6 weeks)	Yes	NR	NR	Yes
Combined Models	41-70 (3–24 weeks) (Espa et al, 2019; Peelaert et al, 2015; Thakur et al, 2017)	Yes (Peelaert et al, 2015; Thakur et al, 2017)	Yes (Espa et al, 2019; Peelaert et al, 2015; Thakur et al, 2017)	Transient microglial activation. p- $\alpha$ Syn in Iba <sup>+</sup> cells (Thakur et al., 2017)	Yes (Peelaert et al, 2015; Thakur et al, 2017)

\*Progression of cell loss in SNpc (percentage of contralateral side). The time lapse (in weeks) for the neurodegeneration to occur is specified.

AAV = Adeno-associated virus; DA = Dopamine; ICV; Intracerebroventricular; LV = Lentivirus; NR = not reported; NS = not significant; OB = Olfactory Bulb; P- $\alpha$ Syn = Phospho  $\alpha$ -Synuclein; SNpc = Substantia nigra pars compacta; Str = Striatum.

(Braak et al., 2003; Rey et al., 2019, 2016). In this model  $p-\alpha$ Syn pathology appeared first in olfactory regions directly connected with the olfactory bulb (1 month after infusion). At later time points pathology spread to non-olfactory regions located 1 synapse (3 months after infusion) and 2 synapses (6 months after infusion) away from the infusion nucleus, such as the central amygdala and the dentate gyrus, and to more distant widespread regions such as the SN and locus coeruleus at later time points (12 months after infusion) (Rey et al., 2016). While mice PFFs-injected into the olfactory bulb did not display any motor abnormality, they exhibited a progressive alteration of odor detection, detectable as early as 1-month post injection.

Few studies have started to use the PFFs model to address cognitive symptoms of PD, by investigating the correlation between the accumulation of p- $\alpha$ Syn aggregates in cortical areas and dendritic spines abnormalities in the same area or cognitive impairment. One study addressed this issue by infusing PFFs into the dorsal striatum (coordinates relative to the bregma: +0.2 mm anterior, +2.0 mm from midline, +2.6 mm beneath the dura), reporting that spine density in cortical layers I, IV, V was lowered in presence of  $\alpha$ Syn aggregates (Blumenstock et al., 2017). Another study infused both the viral vector coding for human wild-type  $\alpha$ Syn bilaterally *via* a double injection into the prelimbic and infralimbic cortex, and PFFs bilaterally into the rostromedial striatum (Espa et al., 2019). This study reported a significant deficit in working memory, attention, and inhibitory control, demonstrating a role of pathological  $\alpha$ Syn in non-motor symptoms (Espa et al., 2019).

Although the seeding and spreading properties of oligomers and PFFs were consistent across different works, a recent study reported the different seeding and spreading propensity of 5 aSyn strains when injected in vivo in the olfactory tubercle, confirming the structure-dependent toxicity (Rey et al., 2019). Moreover, contrasting results have been observed in relation to the neurodegenerative and proinflammatory properties of aSyn species (Table 2). The intracerebroventricular infusion of αSyn oligomers induced non-motor symptoms such as olfactory deficits and motor symptoms accompanied by dopaminergic loss in mice (Fortuna et al., 2017). In some instances, intrastriatal infusion of toxic αSyn, either oligomers or PFFs, did not induce the degeneration of nigrostriatal neurons, despite the presence of PD pathology epitomized by p-aSyn deposits (Masuda-Suzukake et al., 2013, Peelaerts et al., 2015). In other studies, the infusion of PFFs into the dorsal striatum lead to a gradual and partial degeneration of nigral dopamine neurons projecting to the dorsal striatum, detectable 6 months post-infusion and associated with a deterioration of motor performance in the rotarod test (Luk et al., 2012b). The intranigral infusion of PFFs or oligomers was sufficient itself to induce a significant loss of dopamine neurons of the SNpc 3-6 months later in some studies (Abdelmotilib et al., 2017, Thakur et al., 2017) but not in others (Peelaerts et al., 2015). Both oligomeric and PFFs species increased or accelerated the cell loss when infused into AAV-aSyn treated rats (Peelaerts et al., 2015, Thakur et al., 2017). Interestingly, similarly to deposits formation the extent of cell loss was strictly dependent upon the length of PFFs infused into the SN, with shorter 49-29 nm PFFs displaying the most toxic outcome (Abdelmotilib et al., 2017).

On the last years, the AAV-induced human  $\alpha$ Syn expression was combined with the injection of PFFs in the SNpc (Thakur et al., 2017). PFFs were inoculated into the SNpc 4 weeks after the infusion of low dose AAV within the same area but in a different site, leading to the formation of  $\alpha$ Syn aggregates and to dopaminergic degeneration associated with motor deficits (Thakur et al., 2017). Despite the long and complex methodological procedure that involves a double stereotaxic surgery, such a combined AAV/PFF-based approach has been successful in reproducing both neuropathological and symptomatic features of PD (Table 2). It is worth to note that in this case the nigral cell loss appeared as early as 3 weeks after PFFs inoculation, which may represent an advantage to faster results but a limit when reproducing the slow disease progression (Thakur et al., 2017).

The neuroinflammatory response to oligomers or PFFs inoculation has been poorly investigated and various results have been reported (Table 1). In general, PFFs-induced PD pathology without neurodegeneration failed to induce any inflammatory response in the infused area, although microglia and astroglia proliferation was only investigated at 15 months post-infusion (Masuda-Suzukake et al., 2013). Moreover,  $\alpha$ Syn aggregates were not found within microglia cells after PFFs inoculation in this study (Masuda-Suzukake et al., 2013). In contrast, when the inoculation of PFFs into the SNpc induced PD pathology and neurodegeneration, it was associated with a transient microglial activation, which was exacerbated by previous inoculation of AAV- aSyn, and which tended to remit 3 months post-infusion (Thakur et al., 2017, Wilms et al., 2009). Microglia positive for  $p-\alpha$ Syn were detected in the SN of AAV + PFFs inoculated rats for a few weeks after infusion (Thakur et al., 2017). Interestingly, the inflammatory response induced by the inoculation of short oligomers into the SNpc was intense, unbalanced and lasted for several months (Boi et al., in preparation). Although more studies are necessary to clarify the interrelationship between  $\alpha$ Syn species and microglia *in vivo*, it is plausible that short oligomers are the most proinflammatory species, in accord with in vitro observations (Park et al., 2008). Moreover, the poor pathological microgliosis observed after PFFs inoculation may account for the absence (Masuda-Suzukake et al., 2013) or scarce (Thakur et al., 2017) nigral cell loss observed after PFFs inoculation.

All together these studies strongly suggest that the toxicity of aSyn is structure-dependent both in terms of seeding/spreading properties and neurogenerative properties. Exogenous  $\alpha$ Syn inoculation models are a great tool to investigate a Syn seeding and spreading pathology and are promising models to evaluate approaches interfering with these processes. Evidence accumulated so far indicates that these models display face validity, recapitulating motor and non-motor symptoms of PD although with some variability, and construct validity by modeling cellular mechanisms of PD. Importantly, the intranigral or intrastriatal inoculation of PD-derived LB extracts, and the intrastriatal injection of αSyn PFFs in monkeys, induced a progressive nigrostriatal neurodegeneration, providing evidence for translatability and potential predictive validity of aSyn PFFs-based models (Chu et al., 2019; Recasens et al., 2014). A practical advantage of this model is the easiness of manipulation of oligomers/PFFs as compared to the virus-based technique, permitting an approach to the model by laboratories worldwide. Moreover, the exogenous preparations of the toxic species, i.e. oligomers and PFFs, offers the advantage of a more rigorous control of the amount injected and intracerebral concentration of these species, offering a better starting point for the development of a highly standardized PD model. Alike virus-based models, exogenous  $\alpha$ Syn inoculation can be made unilaterally, although unilateral injection of aSyn oligomers/PFFs may produce bilateral pathology, and therefore caution should be made in using the contralateral side as a control. The inoculation of oligomers/ PFFs can be made in specific areas and thus it is useful to investigate pathological properties and effects of toxic aSyn species, in relation to both cardinal symptoms and non-motor symptoms of PD. The long time required to develop the neuropathology and the motor symptomatology after oligomers/PFFs infusion reproduces the slow progression of human neuropathology, useful when testing disease-modifying agents, although this may be considered a disadvantage in the practical use of this model. A limitation of oligomers and PFFs is their poor stability and the tendency to dissociate, therefore requiring particular care in the storage and handling, and the need to program the intracerebral infusion within 10 days from preparation (Polinski et al., 2018). Finally, as discussed above, not all oligomeric/PFFs species hold the same toxicity, and the mechanism of structure-related toxicity is still largely unclear. In synthesis, further investigations are required to definitely establish the exogenous  $\alpha$ Syn inoculation as a valid and standardized preclinical model of PD; however, it is predictable that it will become a widely used approach because the many advantages over the other available models.

#### **Declaration of Competing Interest**

None.

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