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## Oxidative and Nitritive Protein Modifications in Parkinson's Disease

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### Abstract

Parkinson's disease (PD) is a complex neurodegenerative syndrome likely involving contributions from various factors in individuals including genetic susceptibility, exposure to environmental toxins, and the aging process itself. Increased oxidative stress appears to be a common causative aspect involved in the preferential loss of dopaminergic neurons in a region of the brain prominently affected by the disorder, the substantia nigra (SN). Loss of dopaminergic SN neurons is responsible for the classic clinical motor symptoms associated with PD. Several oxidative and nitritive post-translational modifications (PTMs) have been identified on proteins pertinent to PD that may affect this or other aspects of disease progression. In this review, we discuss several examples of such PTMs to illustrate their potential consequences in terms of initiation or progression of PD neuropathophysiology.

### Keywords

Oxidation; nitration; HNE; 3NT; post-translational modification; Parkinson's disease

### Introduction

While oxygen in its many states is a requirement for life, dismutation of molecular oxygen leads to the formation of more reactive oxygen species that can lead to extensive damage to various cellular macromolecules including DNA, lipids and proteins contributing to the general process of aging as well as to various age-related diseases. Oxidative stress has indeed emerged as a common causative factor in the pathology of several neurodegenerative disorders including Parkinson's disease (PD) [1].

PD, the second most common neurodegenerative disease after Alzheimer's, is characterized by the preferential loss of dopaminergic neurons in a region of the midbrain known as the substantia nigra (SN) and the presence of  $\alpha$ -synuclein-containing inclusion bodies (referred to as Lewy bodies) in the cytoplasm of neurons in various regions of the brain including the SN, olfactory bulb, neocortex, and brain stem [2]. Whether oxidative/nitritive stress is a primary trigger for PD or merely a byproduct of cellular dysfunction is currently unknown, however several related PTMs identified on proteins with relevance to PD suggest that such modifications may indeed contribute to disease neuropathology. These oxidative and nitritive PTMs and their significance for PD are discussed in this review and summarized in Table 1.

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## Evidence for Oxidative and Nitritive Stress in PD

Extensive evidence has accumulated particularly from postmortem brain tissues which suggests that oxidative and nitritive stress is an integral factor in neurodegeneration of the SN associated with PD [3]. Decreased levels of the antioxidant glutathione (GSH) in the SN, for instance, is one of the earliest biochemical changes observed in the disease [4–6]. The PD SN also displays increased levels of lipid peroxidation as indicated by decreased levels of polyunsaturated fatty acids (substrates for lipid peroxidation) and increased levels of malondialdehyde (MDA) [7] and 4-hydroxy-2-nonenal (HNE) [8]. Another marker of oxidative stress, the nucleic acid oxidation product 8-hydroxyguanosine (8OHG), is elevated in affected neurons compared to surrounding brain regions in the PD brain as well as in comparison to the SN of age-matched controls [9,10]. A widely used assay to measure general protein oxidation involves the reaction of 2,4-dinitrophenylhydrazine with carbonyl groups on proteins which can then be quantified spectrophotometrically. This method has been used to demonstrate the presence of increased levels of oxidized protein carbonyls in the SN of PD patients [11,12]. Oxidative damage to proteins also occurs through the nitration of tyrosine residues by peroxynitrite formed from the reaction of nitric oxide with superoxide radicals or through nitrite-dependent peroxidase activity [13]. Antibodies specific for 3-nitrotyrosine (3NT) have been developed and used to demonstrate a qualitative increase in the nitration of proteins within Lewy bodies in the PD brain [14,15]. Increased 3NT levels have also been detected in baboons [16] and mice [17] following systemic administration of the PD-inducing toxin 1-methyl-4-phenyl-1,2,3,6-tetrapyridine (MPTP).

## Some Possible Sources of Oxidative/Nitritive Stress in PD

Reactive oxygen species (ROS) and reactive nitrogen species (RNS) can be generated by many biological processes. Nitric oxide (NO) can be formed by the enzyme nitric oxide synthase (NOS). Inhibition of the neural isoform NOS1 in baboons or the inducible isoform NOS2 (expressed in glial cells) in mice has been found to enhance resistance to MPTP toxicity [18–20], indicating that NO is required for MPTP-induced cell death. Iron, which can drive the formation of the reactive hydroxyl radical via Fenton chemistry, is increased in the SN of PD patients [21–23]. If this iron is present in a free state it can contribute to ROS/RNS production and subsequent neurodegeneration. Environmental toxins such as rotenone, MPTP and paraquat, all known environmental risk factors for PD, are capable of generating oxidative stress in several ways, either directly or indirectly by inhibiting complex I [24,25], activation of microglia, or through formation of  $\alpha$ -synuclein aggregates [26].

## The Possible Role of Dopamine Oxidation in PD

Scientific evidence from as early as 1895 suggested that the SN was likely involved in PD [27]. This finding was followed by the discovery of proteinaceous inclusion bodies in the brains of PD patients [28] and lesions of the SN [29]. In the 1960's it was determined that there is a severe decrease in the levels of neurotransmitter dopamine in the corpus striatum and SN of PD patients [30]. Subsequently, this dopamine deficiency was connected with the loss of dopaminergic cells in the SN [31]. This led to the use of levodopa (L-DOPA, a dopamine precursor) as a treatment for PD [32,33]. Dopamine is chemically reactive and is the most predisposed of the catecholamines towards oxidation [34]. It can, for example, be metabolized by monoamine oxidase (MAO) which also creates  $H_2O_2$  as a byproduct of the reaction [35]. Dopamine oxidation can also create dopamine quinone which can directly modify proteins [36]. Various catecholamine oxidation products including DOPAC (3,4-dihydroxyphenylacetic acid), formed via monoamine oxidase-B-mediated dopamine oxidation to 3'-4'-dihydroxyphenylaldehyde and further conversion via an aldehyde dehydrogenase, readily form adducts with GSH and cysteine. Increased levels of free 5-S-cysteinyl-

catecholamine adducts have been found in the SN of PD patients and have been reported to result in inhibition of mitochondrial complex I, a hallmark of the disease [37]. Rat nigral cells treated with 5-S-cysteinyl-catecholamines were reported to have elevated levels of DNA damage including 8OHG [38]. Another product of dopamine oxidation, 7-(2-aminoethyl)-3,4-dihydro-5-hydroxy-2H-1,4benzothiazine-3-carboxylic acid or DHTB-1, has also been reported to be an irreversible inhibitor of complex I activity, resulting in subsequent oxidative stress [39]. Dopamine oxidation and reaction with electrophiles such as cysteine and glutathione however is not necessarily neurotoxic as it constitutes the pathway for neuromelanin production which is formed as a function of the normal aging process.

## Specific Oxidative/Nitrative PTMs of Proteins Associated with PD

### $\alpha$ -Synuclein

The function of  $\alpha$ -synuclein and its role in PD has been heavily studied since the discovery of a dominant mutation in the  $\alpha$ -synuclein gene in a number of Greek and Sicilian families with a familial form of the disorder [40]. This discovery was soon followed by its identification as a major component of Lewy bodies in sporadic cases of the disease [41]. The exact function of  $\alpha$ -synuclein and the mechanism by which it aggregates to form soluble oligomers, soluble protofibrils and insoluble fibrils (the form contained in Lewy bodies) is unknown. It is also still controversial as to whether the fibrils contribute to disease pathology or whether this form of the protein acts as a sink to eliminate toxic soluble oligomers and/or protofibrils. Rotenone, paraquat, nitrating agents as well as expression of the human familial mutation A53T have all been reported to result in increased aggregation of  $\alpha$ -synuclein [42–45].

Lewy bodies appear to qualitatively contain high levels of nitrated tyrosine residues as indicated by antibodies specific for 3NT [14]. Nitrated  $\alpha$ -synuclein has also been detected in the SN and ventral midbrain of mice treated with MPTP [46]. Recombinant  $\alpha$ -synuclein treated with nitrating agents (peroxynitrite/CO<sub>2</sub> myeloperoxidase/H<sub>2</sub>O<sub>2</sub>/nitrite) generates highly stable nitrated  $\alpha$ -synuclein oligomers as a consequence of oxidation resulting in the formation of crosslinked o-o'-ditryrosine [47]. This form of aggregated  $\alpha$ -synuclein is resistant to proteolysis, perhaps due to its structural stabilization. Further work from Virginia Lee and collaborators using antibodies specific for nitrated tyrosine residues on  $\alpha$ -synuclein demonstrated that  $\alpha$ -synuclein in particular is a target for nitrating agents and that nitrated  $\alpha$ -synuclein is present in the Lewy bodies of postmortem tissues from PD patients as well as other neurodegenerative synucleinopathies [48]. Human  $\alpha$ -synuclein contains four tyrosine residues (Tyr39, Tyr125, Tyr133, Tyr136), each of which when independently mutated to phenylalanine (particularly Tyr125), play a vital role in  $\alpha$ -synuclein aggregation induced by peroxynitrite [49].  $\alpha$ -Synuclein has been suggested to function in the regulation of synaptic vesicles and its 3NT modification at Tyr39 has been demonstrated to significantly decrease the binding of  $\alpha$ -synuclein to synthetic vesicles and to decrease the rate at which it is degraded [50]. PTMs identified on  $\alpha$ -synuclein in rotenone-treated cultured PC12 cells include 3NT as well as aminotyrosines at Tyr133 and Tyr136 [51]. A recent paper from Benner *et al.* has suggested the possibility that nitrated  $\alpha$ -synuclein may escape the immune privileged area of the brain into lymphoid tissue and elicit an immune response. Mice treated with MPTP were found to generate antibodies against both un-modified and nitrated  $\alpha$ -synuclein. A strong immune response was observed when mice were immunized with the nitrated c-terminal epitope of  $\alpha$ -synuclein but not upon immunization with the un-modified protein. Nitrated  $\alpha$ -synuclein activated T cells transferred to MPTP-treated mice enhancing microglial activation and degeneration of dopaminergic neurons. Mice with low levels of B and T cells were found to have decreased MPTP-induced neurodegeneration. This suggests that oxidative/nitrative PTMs may cause an adaptive immune response that augments the neuropathology in PD [52].

In addition to tyrosine nitration, rotenone treatment of cultured cells was also found to result in the formation of a methionine sulfone at Met127 of  $\alpha$ -synuclein. Methionine residues can be readily oxidized to methionine sulfoxide through the addition of oxygen to the reactive sulfur atom. The reverse reduction reaction is catalyzed by a family of enzymes called methionine sulfoxide reductases (MSRs). In a  $\alpha$ -synuclein *Drosophila* model, overexpression of MSRA was found to prevent the decreased locomotor activity and shortened lifespan caused by elevated  $\alpha$ -synuclein [53].

Other identified PTMs of  $\alpha$ -synuclein may also be a consequence of increased oxidative stress. The most common PTM found on fibrils of  $\alpha$ -synuclein within Lewy bodies is phosphorylation of serine-129 [54]. This phosphorylation likely occurs as a consequence of oxidative kinase induction, as its levels are found to be increased after exposure of cultured SH-SY5Y cells to  $H_2O_2$  [55]; phosphorylation of this serine also appears to promote  $\alpha$ -synuclein fibril formation *in vitro* [56].

Dopamine adducts on  $\alpha$ -synuclein have been reported to stabilize protofibril formation possibly enhancing toxicity and progression of PD [57]. In addition to being degraded by the proteasome, wildtype  $\alpha$ -synuclein can also be degraded by lysosomes through chaperone-mediated autophagy (CMA). Mutations in  $\alpha$ -synuclein have been found to inhibit degradation of  $\alpha$ -synuclein by the CMA. Dopamine-modified  $\alpha$ -synuclein not only inhibited its own degradation by the CMA but also impairs the degradation of other proteins suggesting why dopaminergic cells may be particularly sensitive to various forms of stress leading to selective SN neurodegeneration in PD patients [58]. More recent data has suggested that dopamine oxidation products on alpha-synuclein may not be derived by adduct formation but rather via noncovalent interaction of oxidized catechols with the protein which may result in its inappropriate C-terminal cleavage and/or conformational changes that impact on its oligomerization [59,60].

A final PTM of  $\alpha$ -synuclein is a byproduct of lipid peroxidation malondialdehyde (MDA) which can conjugate with lysine residues of proteins (MDA-Lys).  $\alpha$ -Synuclein isolated post-mortem from the frontal cortex and SN of PD patients has been shown to contain this modification. The extent to which this is a PD specific modification is unknown as one control cortex sample and one control SN sample were found to have  $\alpha$ -synuclein containing MDA-Lys [61].

## Parkin

Parkin is a ubiquitin E3 ligase encoded by the PARK2 gene [62]. Recessive mutations in the gene are responsible for approximately half of all familial juvenile and early-onset cases of parkinsonism [2]. Overexpression of mutant forms of parkin in NT-2 or SK-NMC cell lines have been shown to increase levels of oxidative damage to proteins and lipids specifically protein carbonyl, lipid peroxidation and 3NT levels. Interestingly, increased oxidative stress did not change the anti-oxidant activity of SOD1, SOD2, catalase, glutathione peroxidase or glutathione reductase however glutathione levels did decrease [63]. Parkin  $-/-$  mice exhibited decreased serum antioxidant capacity, increased protein carbonyl levels and an age-dependent increase of HNE immunostaining in the parkin  $-/-$  brain [64]. Parkin has been suggested to be S-nitrosylated in MPTP-treated mice and in human PD and DLBD patients. S-nitrosylation results in decreased autoubiquitination of Parkin as well as decreased ubiquitination of synphilin-1 in HEK293 cells [65]. Further work will need to be done to establish what proportion of Parkin is S-nitrosylated and the functional impact of this modification on other substrates.

It has been demonstrated that when dopaminergic MES23.5 neuronal cells or cell lysates are incubated with 250  $\mu$ M  $^{14}$ C-dopamine, parkin but not DJ-1 (see below) becomes modified

by dopamine resulting in a decrease of Parkin's E3 ligase activity. Catechol-modified parkin was also found in the SN of normal human brain; whether levels are altered in the SN of PD patients is currently not clear [34].

## DJ-1

Mutations and deletions in the PARK7 gene that encodes DJ-1 [66] account for less than 1% of all familial cases of PD [2]. Several potential functions have been attributed to DJ-1 including roles in mRNA regulation, as a chaperone, a protease and antioxidant [67]. Its role as an antioxidant has garnered the most interest since it had previously been shown that DJ-1 is converted to variants with a lower pI after treatment with hydroperoxides [68]. It was subsequently reported in a cell culture model that the downregulation of DJ-1 enhanced cell death via oxidative stress [69]. DJ-1 exists as a homodimeric protein with an approximate MW of 39 kD but migrates as a monomer (~20 kD) after 1D SDS-PAGE due to denaturation by the detergent. As many as 10 different isoforms of DJ-1 may exist, 6 monomeric versions with isoelectric points ranging from 5.5 to 6.4 and 4 SDS-resistant dimeric forms with isoelectric points ranging from 7.0 to 8.4 [70]. In one study all monomeric forms of DJ-1 were found to be widely expressed in human frontal cortex, particularly in astrocytes although the most alkaline isoform was reduced in abundance in the cortex of PD brains [71]. Subsequent reports indicated that the acidic monomer isoforms of DJ-1 (pI 5.5 and 5.7) and the basic dimer isoforms (pI 8.0 and 8.4) were increased in the frontal cortex of PD and Alzheimer's disease patients as compared to controls. Overall the total expression of DJ-1 appears to be elevated in PD and AD brains [70]. Similarly Meulener *et al.* found levels of the most acidic isoforms of DJ-1 increase (~10–30%) with age in flies, mice and humans [72]. General measurements of carbonyl levels in samples only show an overall increase in oxidative stress but do not indicate if there was a specific target with higher sensitivity to oxidative stress. By first isolating DJ-1 from post mortem PD brains and then assaying for carbonyl levels, Choi *et al.* was able to determine that there was a significant increase in carbonyl levels in the one monomeric isoform (pI 6.4) and three dimeric isoforms (pI 7.4, 8.0 and 8.4) of DJ-1 [70].

Employing a mass spectrometry approach, Kinumi *et al.* analyzed the structural modifications of the acidic isoform of DJ-1 taken from whole cell extract of H<sub>2</sub>O<sub>2</sub> treated human umbilical vein endothelial cells run on 2D gels. DJ-1 contains 3 cysteines (amino acid numbers 46, 53, and 106). All 3 cysteines were found to be oxidized to cysteine sulfonic (SO<sub>3</sub>H) acid to various extents. Treatment with 100 μM H<sub>2</sub>O<sub>2</sub> for 1 hour oxidized a small percentage of Cys46 and Cys53 while Cys106 was found to be completely oxidized. This suggests that Cys106 is the most sensitive cysteine to oxidation and thus may be important for the acidification and function of DJ-1 [73]. Oxidation of Cys106 has been suggested to amongst other things predispose DJ-1 to cleavage between glycine 157 and proline 158 residues [74]. The *Drosophila* homolog (DJ-1b) of human DJ-1 also functions as an antioxidant and contains a cysteine at amino acid 104 which is analogous to Cys106 in human DJ-1. It has been determined that this cysteine (Cys104) is similarly oxidized to sulfinic (SO<sub>2</sub>H) and sulfonic acid in *Drosophila* [72]. Oxidation of Cys53 to sulfonic acid was detected in both control and PD frontal cortex brain samples from age-matched individuals (5 controls 65–87 and 5 PD 66–79 years old) indicating that some level of age-related oxidation is normal. This same study also identified the presence of an oxidized methionine Met133 to methionine sulfone in DJ-1 which was detected in PD but not control samples via mass spectrometry [70]. The relative levels of oxidized and unoxidized DJ-1 may be informative for determining if some threshold of oxidized DJ-1 levels must be overcome to induce oxidative stress.

By systematically mutating the 3 DJ-1 cysteine residues Canet-Aviles *et al.* confirmed that Cys106 is the critical residue oxidized to cysteine-sulfinic acid resulting in the acidic shift in pI of DJ-1. They also demonstrated in a separate PD model (MPP<sup>+</sup> treatment of neuroblastoma

cells) that acidification of DJ-1 occurred [75] consistent with earlier findings that acidification of DJ-1 by MPP<sup>+</sup> is caused by increased oxidation of Cys106 [76]. Interestingly, targeted mutagenesis in SH-SY5Y cells treated with 0.25 or 1 mM S-nitroglutathione identified S-nitrosylation of DJ-1 at Cys46 and Cys53, but not Cys106 suggesting these residues are differentially sensitive to nitrosylation. S-nitrosylation of Cys46 decreased the levels of homodimeric isoforms of DJ-1 [77] but the consequences of these modifications in terms of DJ-1 function as an antioxidant or any other function are unknown.

## PINK1

PTEN-induced kinase 1 (PINK1) is a widely expressed and mitochondrially-targeted protein whose function is still unclear. Mutations in PINK1 account for about 1–2% of early onset PD [2]. Homozygous expression of the PD-associated G302D mutation in PINK1 results in increased levels of lipid peroxidation and GSH levels and a mild decrease in complex I activity [78]. Inhibition of the *Drosophila* homolog of PINK1 (dPINK1) using RNAi resulted in a progressive loss of dopaminergic neurons that could be abrogated by expressing human PINK1 or through the addition of antioxidants [79]. This suggests that PINK1 may play a functional role in preventing neural loss through oxidative stress. Work from Pridgeon *et al.* identified a mitochondrial heat shock protein (Hsp75 or TNF receptor-associated protein 1, TRAP1) as a substrate for PINK1. Phosphorylation of TRAP1 by PINK1 protected cells from oxidative stress by preventing cytochrome c release from mitochondria suggesting that the serine/threonine kinase activity of PINK1 may be involved in preventing apoptotic neuronal cell death in PD [80].

## Mitochondrial complex I

MPTP can selectively inhibit mitochondrial complex I activity [81,82]. Since this discovery the role of complex I in PD has been intensely studied. Post mortem SN samples from PD patients have a complex I deficiency indicating a link between PD and reduced complex I activity [83]. The cause of the selective complex I inhibition is unknown.

Several antioxidant molecules exhibit altered expression levels in PD. Total glutathione (GSH + GSSG) depletion is the earliest known indication of oxidative stress in PD, occurring prior to decreases in mitochondrial complex I activity and dopamine levels [4,5,84]. We previously demonstrated *in vitro* that decreased glutathione levels resulted in selective inhibition of complex I activity. Complex I activity could be restored to normal levels by adding the reducing agent dithiothreitol (DTT) suggesting that the reversible oxidation of cysteine residues could be the cause [85]. Previous studies have demonstrate that oxidation of GSH to GSSG can result in glutathionylation and inhibition of complex I [86,87]. However, given that both GSH and GSSG are decreased in PD, glutathionylation of complex I is unlikely to be the cause of complex I inhibition in the disease. In order to determine the oxidative or nitrative species involved in the observed CI inhibition, we used scavengers specific for various ROS or RNS [88,89]. Data from these studies indicate that following either acute or chronic glutathione depletion, complex I inhibition appears to be dependent on nitric oxide (NO)-related mechanisms. Since complex I inhibition was found in our system to be reversible via either DTT or restoring glutathione levels to normal in our cells, this suggests an S-nitrosylation event. Direct treatment of isolated mitochondria with peroxynitrite had previously been shown to lead to significant reductions on the activities of several components of the electron transport chain including complex I, none of which were not found to be DTT-reversible [90]. By isolating complex I from peroxynitrite-treated mitochondria via sucrose gradient separation Murray *et al.* demonstrated that at least 5 subunits of complex I are modified by 3NT. Subsequent analysis using mass spectrometry identified those subunits and in several cases the specific modified residues as NDUFS2, NDUFS8, NDUFB4 (Tyr46 3NT, Tyr50 3NT, Tyr51 3NT, Tyr128 3NT), NDUFA12 (Tyr48 3NT, Tyr49 3NT, Trp61 *N*-formylkynurenine), and

NDUFA6 (Tyr122 3NT) [90]. While this demonstrates that 3NT modifications can occur in complex I as a consequence of peroxynitrite exposure, it does not prove that these modifications result in the observed complex I inhibition. Indeed, we also observe the presence of 3NT modifications in complex I subunits following glutathione depletion but as in our case enzymatic inhibition is reversible, this would argue against them playing a functional role in glutathione-mediated effects [91]. In further *in vivo* studies, we were able to demonstrate that inducible dopaminergic glutathione depletion in the adult animal results in S-nitrosylation of complex I subunits which coincides with a selective decrease in complex I activity and an age-dependent neurodegeneration of dopaminergic SN neurons [92]. Coupled with our *in vitro* data, this strongly implies that a possible mechanism of the hallmark complex I inhibition in these cells could be a consequence of early glutathione depletion.

Complex I immunocaptured from the frontal cortex of PD patients shows increased carbonyl levels as compared to age-matched controls [93]. This increase in carbonylation of complex I could be replicated by treating mitochondria from control frontal cortex samples with rotenone. Here again, the presence of such modifications does not prove whether or not they are functionally significant.

Inhibition of complex I results in two critical predicaments which can impede cell survival, decreased ATP production and increased ROS production. These two factors can augment and aggravate each other in a cyclic fashion amplifying the detrimental effect on cells and tissues. Using three models (SK-N-MC human neuroblastoma cells, rat midbrain slice cultures, and *in vivo* rat), chronic rotenone treatment was shown to cause oxidative damage (via increased total carbonyl levels) and dose-dependent cell death independent of ATP depletion [94]. ATP levels decreased only about 1/3 in SK-NMC cells treated with 100 nM rotenone, a concentration which was completely toxic after 2 days. Cells treated with 1 mM 2-deoxyglucose (2-DG) had a similar decrease in ATP levels with no corresponding toxicity or increase in carbonyl levels. Systemic rotenone infusion *in vivo* caused a slight increase in total protein carbonyl levels in the cortex and striatum but no increase in the cerebellum and hippocampus. The largest increase in carbonyl levels due to rotenone treatment occurred in the olfactory bulb and midbrain, the dopaminergic areas of the brain [94]. Cell type may be an important factor in terms of vulnerability to rotenone. Dopaminergic (MN9D) cells were found to be more sensitive than non-dopaminergic (MN9X) cells following chronic exposure to nanomolar levels of rotenone [95]. Enhanced sensitivity of dopaminergic cells to rotenone could be explained by either increased vulnerability to oxidative stress or the higher energy demand of dopaminergic cells.

## Proteasome

Increased levels of oxidatively or nitratively damaged proteins may be due to either increased oxidative/nitrative stress levels or decreased elimination of modified proteins. The proteasome is a large protein complex that degrades a wide variety of proteins in an ATP-dependent fashion, after proteins have been targeted for proteolysis by the addition of several ubiquitin molecules via E3 ubiquitin ligases such as parkin [96]. Decreased activity of the proteasome has been proposed as a mechanism involved in the neurodegeneration associated with PD due to the accumulation of damaged proteins. This theory is supported by evidence that demonstrates that proteins modified by the lipid peroxidation product HNE are more slowly degraded by the proteasome [97]. High levels of HNE can decrease proteasomal activity [98] however, such high levels of HNE may never occur physiologically [99]. Mutations in  $\alpha$ -synuclein associated with familial forms of PD also have decreased rates of elimination as compared to wildtype  $\alpha$ -synuclein [100]. Selective inhibition of the proteasome by lactacystin or epoxomicin results in increased oxidative stress through increased protein carbonyls, lipid peroxidation and DNA damage [101]. Whether decreased proteasomal activity has a causative role or is a consequence of other mechanisms in PD is still unclear.

## Tyrosine hydroxylase

One of the biochemical changes evident in PD patients is the severe decrease in dopamine levels in neurites emanating from the SN and innervating the striatum. Any detrimental effects on the activity of tyrosine hydroxylase (TH), the rate-limiting enzyme in dopamine biosynthesis, would be of obvious importance in terms of disease development. Work from Harry Ischiropoulos and colleagues has indicated that TH can be selectively nitrated by either peroxynitrite or MPP<sup>+</sup> at tyrosine residues resulting in the inhibition of TH activity [102]. Subsequent work from the laboratory has demonstrated that nitration of tyrosine residue 423 in particular is critical for the observed reduction in TH activity [103]. Conversely, work from Donald Kuhn and associates suggest that while tyrosine residues are nitrated by peroxynitrite, the reduced TH activity is instead due to the oxidation of cysteine residues [104]. Subsequently Kuhn's lab has demonstrated that tyrosine nitration is prevented when TH is treated with peroxynitrite in the presence of excess cysteine or glutathione and that TH activity is repressed by Sthiolation [105]. Work from both laboratories clearly indicate that tyrosine residues on TH are nitrated after exposure to peroxynitrite. However it is not clear to what extent nitration versus cysteine thiolation has on the activity of TH; further experiments will be needed to clarify this issue.

## Conclusions

Increased oxidative/nitrative stress has been linked with PD-associated neurodegeneration as well as several oxidative and nitrative PTMs on proteins pertinent to PD that may affect disease progression. The ever growing number of examples of such PTMs suggests that oxidative/nitrative stress is likely to play a major role in PD initiation or progression. Whether a common source exists for the various oxidative and nitrative modifications observed on the proteins discussed here is unknown. The identification of modifications that impact protein function and in doing so contribute to Parkinson's disease pathologies may prove valuable for diagnoses and preventative or therapeutic targets.

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## Glossary

### Abbreviations

<b><math>\alpha</math>-syn</b>	$\alpha$ -synuclein
<b>PD</b>	Parkinson's disease
<b>SN</b>	substantia nigra
<b>PTM</b>	post-translational modification
<b>GSH</b>	glutathione (reduced)
<b>GSSG</b>	glutathione (oxidized)
<b>MDA</b>	malondialdehyde
<b>HNE</b>	4-hydroxy-2-nonenal
<b>8OHG</b>	8-hydroxyguanosine
<b>3NT</b>	3-nitrotyrosine
<b>MPTP</b>	1-methyl-4-phenyl-1,2,3,6-tetrapyridine

<b>ROS</b>	reactive oxygen species
<b>RNS</b>	reactive nitrogen species
<b>NO</b>	nitric oxide
<b>NOS</b>	nitric oxide synthase
<b>MAO</b>	monoamine oxidase
<b>DOPAC</b>	3,4-dihydroxyphenylacetic acid
<b>DHBT-1</b>	7-(2-aminoethyl)-3,4-dihydro-5-hydroxy-2H-1, 4benzothiazine-3-carboxylic acid
<b>Tyr</b>	tyrosine
<b>Cys</b>	cysteine
<b>Met</b>	methionine
<b>Trp</b>	tryptophan
<b>Ser</b>	serine
<b>MSR</b>	methionine sulfoxide reductase
<b>CMA</b>	chaperone-mediated autophagy
<b>MDA-Lys</b>	malondialdehyde-lysine
<b>SOD</b>	superoxide dismutase
<b>DTT</b>	dithiothreitol
<b>TH</b>	tyrosine hydroxylase
<b>MPP<sup>+</sup></b>	1-methyl-4-phenylpyridinium
<b>Phos</b>	

**pI** phosphorylation  
isoelectric point



**Table 1**

<b>Protein</b>	<b>Source of Modification</b>	<b>PTM</b>	<b>Proposed Consequence</b>	<b>Reference</b>
$\alpha$ -synuclein	unknown kinase	Phosphorylated ser129	Promotes $\alpha$ -syn fibrils	[53–55]
	Lewy bodies/ nitrating agents/MPTP	Several 3NTs	Stable $\alpha$ -syn oligomers	[13,45–47]
	peroxynitrite	Tyr125 3NTs	$\alpha$ -syn dimerization	[48]
	peroxynitrite dopamine	Tyr39 3NTs Dopamine adduct formation	Decreased vesicle binding Stabilizes $\alpha$ -syn protofibrils/ inhibits degradation of proteins by CMA	[49] [56,57]
$\alpha$ -synuclein overexpression MPTP	rotenone	Aminotyrosine at Tyr133 and Tyr136	Unknown	[50]
	$\alpha$ -synuclein overexpression	Met127 sulfone formation	Prevents fibril formation	[52]
	MPTP	Nitration	Immune response	[51]
parkin	overexpression of mutant parkin	Increased lipid peroxidation, protein carbonyls and 3NTs in total protein	Decreased GSH, increased oxidative stress	[60]
	parkin $-/-$ mice	Increased HNE adducts and carbonyl levels	Decreased antioxidant capacity	[61]
	dopamine	Dopamine adduct formation	Decreased parkin ligase activity	[33]
	MPTP	S-nitrosylation	Decreased parkin ligase activity	[62]
DJ-1	DJ-1 expression decreased Hydroperoxides/H <sub>2</sub> O <sub>2</sub>	Oxidation Cys46, Cys53 and Cys106 oxidized to sulfonic acid	Cell death, oxidative stress Acidification of DJ-1, oxidative stress, DJ-1 cleavage	[66] [70,71,65]
	PD brain tissues	Cys53 sulfonic acid, increased carbonyl levels on several isoforms of DJ-1	Acidification of DJ-1, oxidative stress	[67]
	MPTP	Met133 sulfone, Cys106 sulfinic acid	Acidification of DJ-1, oxidativestress	[72,73]
	S-nitrosylation	S-nitrosylation of Cys46 and Cys53	Decreased homodimer DJ-1 formation	[74]
PINK1	G302D-PINK1 mutant	Increased lipid peroxidation	Decreased complex I activity, Increased SOD and GSH levels	[75]
	PINK1 kinase activity	Phosphorylation of Trap1	Prevention of cytochrome-c release and apoptosis	[77]
mitochondrial complex I	GSH depletion	Elevated cysteine oxidation, Decreased sulfhydryls, mitochondrial HNE adducts, and S-nitrosylation levels increased	Complex I activity inhibited	[82,85,86, 88,89]
	Peroxynitrite	Complex I subunit oxidation (several 3NTs)	Complex I activity inhibited	[87]
	PD brain, frontal cortex	Increased carbonyl levels particularly in dopaminergic areas	Complex I activity inhibited	[90]
	Rotenone	Increased carbonyl levels in dopaminergic areas of brain	Oxidative stress	[91]
proteasome	HNE Lactacystin or epoxomicin	HNE adduct formation Increased carbonyl, lipid peroxidation, and DNA damage	Proteasome Inhibition Oxidative stress	[94,95] [98]
	tyrosine hydroxylase	Tyr423 (3NT) Cysteine oxidation	TH activity inhibited TH activity inhibited	[99,100] [101,102]