

Review Article

Peroxidation and Halogenation Stress: Windows to a Better Understanding of Sporadic Parkinson's Disease

Fernández E*

Department of Medical Physiology and Biophysics, University of Seville, Spain

*Corresponding author: Emilio Fernandez, Department of Medical Physiology and Biophysics, Laboratory of Molecular Neurology and Neurophysiology (BIO127), School of Medicine, Universidad de Sevilla, Av. Sanchez Pizjuan 4, E-41009 Sevilla, Spain

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Abstract

Oxidative stress is considered as an important pathogenic mechanism in Parkinson's disease. Two types of oxidative stress, peroxidation and halogenation stress, are gaining increasing importance as biochemical windows to a better understanding of the pathogenesis of this disease. Peroxidation stress is due to excess of hydrogen peroxide, and it is related to the presence of many peroxidation-related molecular markers in Parkinsonian patients, and to misfolding of proteins. Peroxidation-induced misfolded proteins show altered functionality, such as loss of neuroprotective activity and tendency to form proteinaceous aggregates inside neurons. Peroxidation stress is also detected by a loss of activity of the main hydrogen peroxide scavengers in cerebrospinal fluid of patients. Altered hydrogen peroxide scavenging also leads to halogenation stress. Halogenation stress is characterized by the excess of halogenated molecules such as hypohalous acids, haloamines and halogenated proteins. Halogenated amines and proteins are thought to be deleterious for neurons, and they could play an important role in the etiology of Parkinson's disease.

Keywords: Oxidative stress; Peroxidation; Halogenation; Hydrogen peroxide; Haloamine; Halogenated protein; Parkinson's Disease

Abbreviations

AOPP: Advanced Oxidation Protein Products; α SYN: α -Synuclein; CNS: Central Nervous System; CSF: Cerebrospinal Fluid; EOP: Eosinophil Peroxidase; GPx: Glutathione-Peroxidase; GR: Glutathione-Reductase; GSH: Glutathione; GSSG: Glutathione Disulfide; GST: Glutathione-S-Transferase; 8-OHdG: 8-Hydroxyguanosine; H_2O_2 : Hydrogen Peroxide; HSA: Human Serum Albumin; LPO: Lactoperoxidase; MDA: Malondialdehyde; MPO: Myeloperoxidase; NADPH: Nicotinamide Adenine Dinucleotide Phosphate; NOX: Nicotinamide Adenine Dinucleotide Phosphate Oxidase; O_2 : Oxygen; $\bullet O_2^-$: Superoxide Anion; $\bullet OH$: Hydroxyl Ion; PARK2: Parkin; PD: Parkinson's Disease; PRDx: Peroxiredoxin; SULT: Sulfotransferase; SH: Thiol Group; SOD: Superoxide Dismutase; SOH: Sulfenic Acid; TPO: Thyroperoxidase

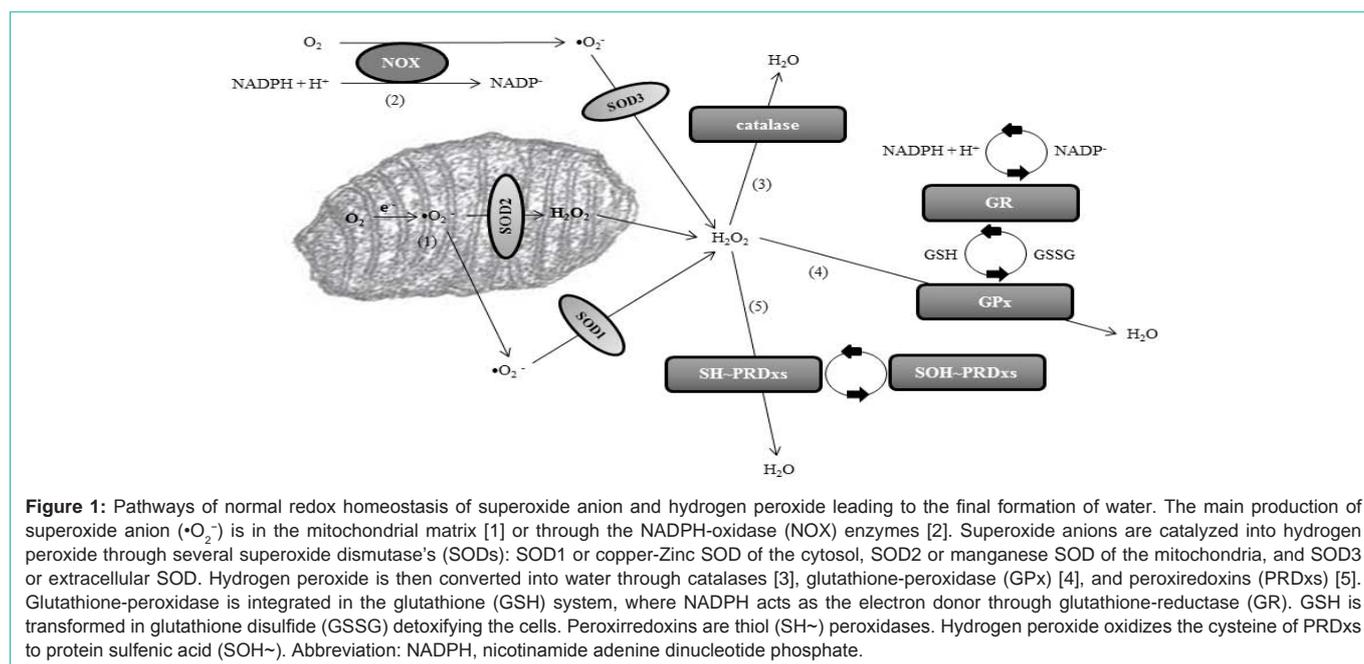
Introduction

Oxidative stress, which is defined as an imbalance between the production of reactive oxidative species and anti-oxidant mechanisms, is considered as an important pathogenic mechanism in Parkinson's disease (PD) and other neurodegenerative diseases [1-4]. Two types of oxidative stress, peroxidation and halogenation stress, are gaining increasing importance as biochemical windows to a better understanding of the pathogenesis of PD.

Peroxidation stress, defined as excess of production of hydrogen peroxide (H_2O_2) or loss of normal H_2O_2 scavenging, is an important type of oxidative stress. Peroxidation stress is related to the overproduction of free radicals and reactive oxygen species (ROS), which are otherwise products of the normal cellular metabolism. Superoxide ions ($\bullet O_2^-$) are the most important ROS, and they are

produced in the mitochondrial matrix or through the NADPH-oxidase (NOX) enzymes, mostly located extracellularly and attached to the plasmatic membrane. The first line of defense of the organism is to minimize the production of $\bullet O_2^-$ in the mitochondria, because metabolic O_2 is converted directly into water thanks to the action of cytochromes of the electronic chain. However, around 0.1-1% of electrons transferred to O_2 can yield $\bullet O_2^-$. Superoxide anions are then catalyzed into hydrogen peroxide through several superoxide dismutases (SODs): SOD1 or copper-Zinc SOD of the cytosol, SOD2 or manganese SOD of the mitochondria, and SOD3 or extracellular SOD. H_2O_2 is then converted into water after reacting with reduced glutathione (GSH), under the control of glutathione-peroxidase (GPx). The H_2O_2 produced from $\bullet O_2^-$ is also eliminated through the action of catalases and peroxiredoxins (PRDxs). Catalases carry out dismutation reactions, and most catalases are heme-containing enzymes. Peroxiredoxins are cysteine-dependent or thiol (SH~) peroxidases. Hydrogen peroxide oxidizes the cysteine of PRDxs to protein sulfenic acid (SOH~PRDx). To sum up, SODs scavenge superoxide anions, and the most important scavengers of hydrogen peroxide are GPx, catalases and PRDxs (Figure 1). It can be concluded that a failure in these anti-oxidant defenses could lead to oxidative stress, with overproduction of ROS, mainly $\bullet O_2^-$ or H_2O_2 .

Overproduction of hydrogen peroxide is also biochemically linked to halogenation stress, another type of oxidative stress that is characterized by excess of halogenated radicals or reduced scavenging of halogenated molecules. Halogenation reactions are mediated by haloperoxidases, a group of enzymes that mediate the conversion of hydrogen peroxide into hypohalous acids, such as hypochlorous or hypoiodous acids, after the incorporation of halogen elements such as chloride or iodide [5,6]. Hypohalous acids can react with primary



amines to form halogenated amines (haloamines) such as chlorinated or iodinated amines. Normally hypohalous acids and haloamines are scavenged by glutathione-S-transferase, leading to the formation of oxycacids and sulfoxides. Haloamines can also be eliminated by sulfation through the action of sulfotransferases or SULTs [7], or after the formation of nitriles and aldehydes, although these latter reactions are slow in normal physiology. However if there is peroxidation stress and excess of H_2O_2 , as it seems to be the case in PD [1-4], the enhanced formation of hypohalous acids and halogenated amines would lead to the formation of halogenated proteins, due to the incorporation of halogens at tyrosine and lysine residues of proteins. In addition, halogenated proteins could be subjected to degradation by proteasome, resulting in further release of free haloamines [8]. Haloamines in turn increase ROS production, thereby inducing a vicious cycle of oxidation and toxicity.

Peroxidation and Parkinson's disease

Peroxidation stress is detected in brain tissue, cerebrospinal fluid (CSF) and blood, among other fluids, of PD patients. Analysis of substantia nigra tissue indicates that this nucleus is subjected to oxidative stress because it shows enhancement of several peroxidative markers such as 8-hydroxyguanosine (8-OHdG, marker of oxidative stress to DNA) [9], protein carbonyls (markers of protein oxidation) [10], lipid-peroxidation products [11,12], and advanced lipoxidation end-products [13]. The cerebrospinal fluid, a fluid in close contact with neural tissue and good witness of neurodegeneration processes, also shows signs of peroxidation in PD. The enzyme DJ-1 (PARK7), reported to act as an antioxidant peroxidase, is reduced in CSF in the initial phases of PD [14]. Buhmann et al. [15] detected enhanced lipid peroxidation in the CSF, and high levels of malondialdehyde (MDA), a compound that causes toxic stress in cells, have been reported too [16,17]. Other markers for oxidative stress such as the oxidized forms of coenzyme-Q and 8-OHdG are augmented in CSF of PD patients [18]. The anti-oxidant ferroxidase activity of ceruloplasmin is also reduced in the CSF of PD [19-21]. Regarding anti-oxidant molecules

encompassing the glutathione (GSH) system, several studies have indicated that levels of total glutathione protein are diminished in CSF and brain tissue of PD [22-24], although data on glutathione-related enzymes are contradictory [25-27]. The blood of patients with PD also show signs of peroxidation stress, because it has been detected increase of 8-OHdG, MDA, and vitamin E, and reduction of the antioxidant enzyme GPx [16,28].

My research group has detected a strong peroxidation stress in CSF of patients with PD because the enzymatic activity of the three principal hydrogen peroxide scavengers is reliably reduced [29]. Thus after analyzing the activity of superoxide anion and hydrogen peroxide (H_2O_2) scavengers, the data revealed a disturbance of H_2O_2 scavenging, but not superoxide anion scavenging, as manifested through a significantly reduced activity of GPx, catalase, and PRDxs, without changes in the activity of SODs [29]. GPx are selenoproteins that reduce peroxides in water. They also reduce lipid hydroperoxides to their corresponding alcohols. Catalases are ubiquitous enzymes that mediate the decomposition of H_2O_2 in water, as explained. PRDxs are cysteine-dependent peroxidases that react with hydrogen peroxide and hydroperoxide substrates. The reliable reduction of the activity of hydrogen peroxide scavengers is expected to activate alternative pathways in order to scavenge H_2O_2 . These pathways include the Fenton reaction, the Haber-Weiss reaction, and the formation of hypohalous acids after activation of haloperoxidases. In addition, dopamine catabolism of dopaminergic neurons, cells that die in the course of PD, leads to either the formation of H_2O_2 under the action of monoamino-oxidases (MAO), or the generation of dopamine-quinones [30]. All of these pathways, even though they facilitate the elimination of H_2O_2 , give to the formation of radical species and quinones, highly oxidant molecules, which would worsen oxidative stress [30]. Figure 2 shows these alternative pathways that are activated after failure of normal H_2O_2 scavenging, and that are linked to peroxidation stress in PD. Our observation of reduced H_2O_2 scavenging in the CSF of patients leads to the hypothesis that

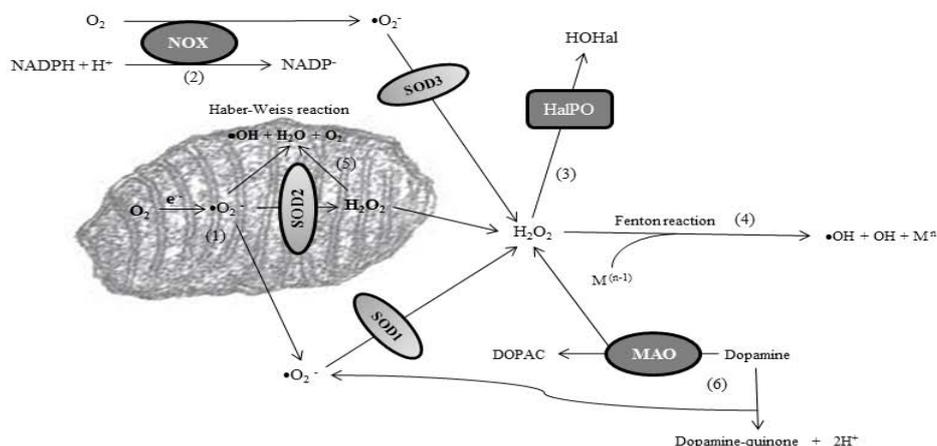


Figure 2: Alternative pathways that are activated after failure of normal H_2O_2 scavenging. As explained before, the main production of superoxide anion ($\bullet O_2^-$) is in the mitochondrial matrix [1] or through the NADPH-oxidase (NOX) enzymes [2]. Superoxide anions are catalyzed into hydrogen peroxide through superoxide dismutases (SODs). The excess of hydrogen peroxide or its inadequate scavenging leads to the action of haloperoxidases (HalPO), enzymes which catalyze the formation of hypohalous acids (HOHal) [3], as well as to the activation of the Fenton reaction if free transition metals (with [n-1] electronic structure) are also enhanced [4]. The Fenton reaction leads to the formation of reactive species such as hydroxyl ion ($\bullet OH$). In addition, excess of superoxide anion and H_2O_2 inside the mitochondria activates the Haber-Weiss reaction, another source of reactive species [5]. Finally, in dopaminergic neurons, cells that die in the course of PD, there is overproduction of H_2O_2 (under the action of monoamino-oxidases or MAO), dopamine-quinones and $\bullet O_2^-$, all worsening oxidative stress [6].

the activation of these alternative pathways is a critical event in the parkinsonian central nervous system. In this context, ferritin levels and iron storage have also been observed to be altered in CSF and central nervous system of patients, and it is well known that the enhancement of transition metals such as iron and copper facilitates the Fenton reaction to occur [30].

Misfolding of proteins is another deleterious effect of peroxidation stress, which alters normal protein function and/or facilitates the formation of proteinaceous aggregates inside neurons. Quinones, which are also formed during peroxidation stress in dopamine neurons in PD, also contribute to misfolding of proteins due to the formation of protein cross-linking [30]. Several of these misfolded proteins such as α -synuclein (α -SYN) and parkin (PARK2) are classically associated to the pathogenesis of PD. Misfolded α -SYN tends to form protofibrils which would precipitate forming fibrils which in turn constitute the core of the Lewy bodies (LBs), anatomic-pathological hallmarks of PD. Protofibrils have been shown to be the neurotoxic species of α -SYN [31]. Association of misfolded α -SYN with peroxidized lipid metabolites can lead to mitochondrial dysfunction that in turn leads to dopaminergic neuron death [31]. Peroxidation stress is also known to induce structural changes in α -SYN leading to covalent aggregation [32]. Regarding Parkin (PARK2), this protein is an E3 ubiquitin lygase involved in the ubiquitination of proteins, and this activity is inhibited by peroxidation. The inhibition of parkin function could contribute to the degenerative process by impairing the ubiquitination of parkin substrates [33]. Parkin is also a transcriptional repressor of p53 [34], and peroxidation alters this function that is also associated with PD [35]. It is worth recalling that recessive mutations in the PARK2 gene encoding Parkin are responsible of many juvenile and early-onset cases of parkinsonism [36].

Halogenation and Parkinson's disease

Halogenation stress is directly associated to peroxidation stress, because overproduction of hydrogen peroxide or its reduced

scavenging leads to the activation of halogenation pathways (Figure 2). The role of halogenation stress in PD has not been discerned, but likely it could play a significant role in the pathogenesis of PD [37,38]. Halogenation reactions are mediated by haloperoxidases, which encompass several enzymes such as white blood-cells peroxidases (myeloperoxidase or MPO, eosinophil-peroxidase or EPO), and glandular haloperoxidases (thyroperoxidase or TPO; lactoperoxidase or LPO) [5,6]. Haloperoxidases mediate the conversion of hydrogen peroxide and halogens into hypohalous acids, such as hypochlorous or hypoiodous acids. As explained, the physiological formation of hypohalous acids and haloamines is mostly scavenged by glutathione-S-transferase and sulfotransferases [7]. However if there is excess of H_2O_2 , as it seems to be the case in PD [1-4], hypohalous acids can lead to the overproduction of haloamines and halogenated proteins. Among haloamines, those derived from tyrosine are of high interest for understanding PD, because halotyrosines include molecules such as chlorotyrosines and iodotyrosines that are considered as neurotoxic for dopamine neurons [37,39]. Halogens can be incorporated at position 3 and 5 of the 6-carbon aromatic ring of tyrosine, leading to the formation of 3-monohalotyrosines or 3,5-dihalotyrosines. As for halogenated proteins, they are gaining increased importance in PD, because they can act as proinflammatory mediators. For instance, chlorinated proteins originate as a result of the action of free radicals such as chloramines and hypochlorous acid on proteins, and they act as inflammatory mediators after binding albumin [40-42]. Of note is that characterizing protein markers associated with halogenation stress is of great interest for biomedicine, because proteins or adducted protein products are good candidates for biomarkers since they show great stability, early formation, and longer lifespan [43]. The molecular pathways which are involved in the formation and degradation of haloamines and halogenated proteins are illustrated in Figure 3.

Among halogenated proteins, advanced oxidation protein products (AOPP) are considered as promising markers for

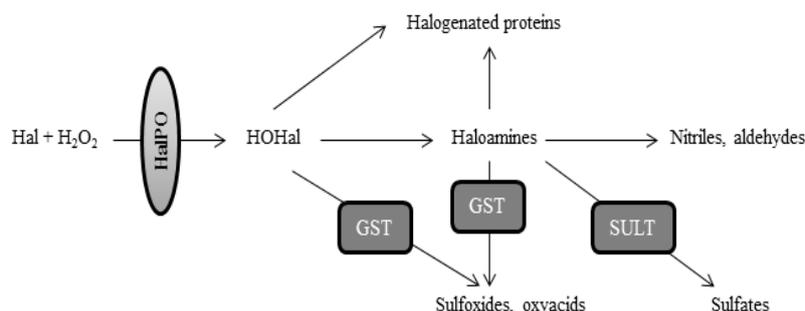


Figure 3: Molecular pathways which are involved in the formation and degradation of haloamines and halogenated proteins. Haloperoxidases (HalPO) mediate the conversion of hydrogen peroxide into hypohalous acids (HOHal), after the incorporation of halogens (Hal). Hypohalous acids can react with primary amines to form haloamines. Hypohalous acids and haloamines are scavenged by glutathione-S-transferase, leading to the formation of oxyacids and sulfoxides. Haloamines can also be eliminated by sulfation through the action of sulfotransferases (SULT), or after the formation of nitriles and aldehydes, although these latter reactions are slow in normal physiology. If there is peroxidation stress and excess of H_2O_2 , the enhanced formation of hypohalous acids and halogenated amines would lead to the formation of halogenated proteins, due to the incorporation of halogens on proteins.

halogenation stress [37,40]. These protein products originate as a result of the action of halogenation radicals on proteins, and they are known to be augmented in serum and cerebrospinal fluid of PD patients [37]. Hypohalous acids include hypochlorous (HOCl), hypobromous (HOBr) and hypoiodous acids (HOI), leading to chlorinated, brominated and iodinated AOPP, respectively [40,41,42,44,45]. As shown in Figure 3, haloperoxidases catalyze these reactions leading to hypohalous acids from halogens and hydrogen peroxide. AOPP can conjugate with human serum albumin (HSA) giving AOPP-HSA conjugates. Of note is that AOPP-HSA conjugates can act as pro-inflammatory factors, because chlorinated AOPP-HSA are known to be inflammatory mediators triggering the oxidative “ignition” of neutrophils, monocytes and T-lymphocytes [46,47]. Chlorinated AOPP are enhanced in many inflammatory diseases, and they correlate with several measures of these diseases [44,46-48]. Hence halogenated protein products could be linked to initial inflammatory processes in PD, where immunological and inflammatory responses in the vicinity of dopaminergic neurons have been detected [49,52].

Our research group has demonstrated that AOPP are enhanced in serum and CSF of parkinsonian patients [37]. Serum concentration of these protein products is progressively reduced over time, and levodopa treatment contributes to this reduction [37], although there is no a statistical relationship because only 47% of our patients received levodopa. In addition, after analyzing the correlation between duration of the disease and serum AOPP levels in patients with moderate PD, a significant correlation was found ($p < 0.003$). In other words, low serum AOPP levels can predict a longer duration of the disease and a longer duration in reaching advanced stages of the disease, and more than 80% of studied patients with a PD longer than 10 years had serum AOPP levels lower than $350 \mu\text{M}$ (normal value in control subjects is $148.4 \pm 37 \mu\text{M}$) [37].

Serum level of advanced oxidation protein products could hence be a prognostic marker of duration of PD. However, specificity of AOPP as a PD biomarker is low because these products also accumulate in other diseases such as diabetes mellitus, uremic syndrome, atherosclerosis, systemic sclerosis and acquired immune deficiency syndrome (AIDS) [53-58]. Then it is urgently needed to identify the specific molecular pathways which are involved in AOPP synthesis in PD. As explained, halogenation stress and AOPP

formation are associated to the action of haloperoxidases, and different haloperoxidases mediate different halogenation processes. MPO (haloperoxidase of white blood cells) mediates the formation of hypochlorous and hypobromide acids, with leads to the formation of AOPP with chloride and bromide radicals. The MPO-related pathway is the most activated in above cited diseases, but MPO seems not to be involved in the formation of AOPP in PD, because this enzyme is found not to be enhanced in serum or CSF of PD patients [59]. TPO, haloperoxidase of thyroid origin involved in iodide management, is found to be elevated in more than 25% of patients with PD [59]. TPO mediates the formation of hypoiodous acid, which leads to the production of iodinated AOPP. It seems plausible that serums AOPP in PD are iodinated protein products, and they could exert proinflammatory effects. In this context, the thyroid gland has been linked to PD, because some researchers have found more prevalence of some thyroid dysfunctions among PD patients or they have detected sympathetic denervation of the thyroid gland in PD patients, a hallmark of the disease because it also occurs in other organs such as heart, stomach and rectum [60-63]. Finally, the role of other human haloperoxidases such as EPO and LPO has not been studied. EPO is detected in serum and CSF like MPO, and it is involved in the formation of hypochlorous and hypobromide acids. LPO is detected in saliva and milk, and it is mostly involved in the formation of hypoiodous and hypobromide acids.

Conclusion

Oxidative stress, which is defined as an imbalance between the production of reactive oxidative species and anti-oxidant mechanisms, is considered as an important pathogenic mechanism in PD. Two types of oxidative stress, peroxidation and halogenation stress, are gaining increasing importance as biochemical windows to a better understanding of the pathogenesis of this disease. Peroxidation stress is associated with excess of hydrogen peroxide, and it induces the presence of many peroxidation-related molecular markers in Parkinsonian patients, and to misfolding of proteins. Many of these proteins, such as α -synuclein and parkin, are traditionally linked to PD. Peroxidation-induced misfolded proteins show altered functionality, such as loss of neuroprotective activity or the tendency to form proteinaceous aggregates in neurons. This latter effect could account for the formation of Lewy bodies, anatomic-pathological

hallmark of PD. Peroxidation stress is also detected by a loss of the activity of the main hydrogen peroxide scavengers, glutathione-peroxidase, catalase, and peroxiredoxins, in the cerebrospinal fluid of patients. Altered hydrogen peroxide scavenging leads to formation of oxidative radicals, alteration in iron storage, and halogenation stress. Halogenation stress is characterized by the excess of halogenated molecules such as hypohalous acids, haloamines and halogenated proteins. Among halogenated proteins, advanced oxidation protein products or AOPP are augmented in Parkinson's disease. Halogenated amines and proteins are thought to be deleterious for neurons, and the author hypothesizes that they could play a critical role in the ethiology and development of Parkinson's disease.

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