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## **Modulation of cellular redox environment as a novel therapeutic strategy for Parkinson's Disease**

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## Abbreviation List

6-OHDA	6-hydroxydopamine	NADPH	nicotinamide adenine dinucleotide phosphate
AADC	aromatic l-amino acid decarboxylase	NOX	transmembrane NADPH oxidase
AD	Alzheimer's Disease;	Nrf2	nuclear factor erythroid 2-related factor 2
ARE	antioxidant responsive element	OXPPOS	oxidative phosphorylation
ATP	adenosine triphosphate	PD	Parkinson's Disease
ATP13A2	ATPase 13A2	PGC1- $\alpha$	peroxisome proliferator-activated receptor gamma coactivator 1 alpha
BSO	buthionine sulfoximine	PINK1	PTEN-induced kinase 1
CHCHD2	coiled-coil-helix-coiled-coil-helix domain containing 2	PLA2G6	phospholipase A2 group VI
CNS	central nervous system	POLG	DNA polymerase subunit gamma
EGCG	epigallocatechin-3-gallate	PPP	pentose phosphate pathway
ER	endoplasmic reticulum	Prxs	peroxiredoxins
ETC	mitochondrial electron transport chain	PUFAs	omega-3 polyunsaturated fatty acids
FBXO7	F-box protein 7	ROS	reactive oxygen species
FDA	Food and Drug Administration	RNS	reactive nitrogen species
G6PD	glucose-6-phosphate dehydrogenase	SIRT1	sirtuin 1
GIGYF	GRB10 interacting GYF protein 2	<i>SNpc</i>	substantia nigra pars compacta
GPx	glutathione peroxidase	SOD	superoxide dismutase
GR	glutathione reductase	sPD	sporadic Parkinson's Disease
GS-DAQ	glutathione to form adducts	TCA	tricarboxylic acid
GSH	glutathione	TFAM	mitochondrial transcription factor A
HCA's	hydroxycinnamic acids	TFB2m	mitochondrial transcription factor B2
HIF1- $\alpha$	hypoxia-inducible factor 1- <i>alpha</i>	TH	tyrosine hydroxylase
LB's	lewy bodies	TPP <sup>+</sup>	triphenylphosphonium cation
L-DOPA	levodopa	TPx	thioredoxin peroxidase
LRRK2	leucine-rich repeat kinase 2	Trx	thioredoxin
MAM	mitochondria-associated membranes	TrxR	thioredoxin reductase
MAO	monoamine oxidase	UCH-L1	ubiquitin C-terminal hydrolase L1
MPP <sup>+</sup>	1-methyl-4-phenylpyridinium	UPS	ubiquitin proteasome system
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine	VMAT2	vesicular monoamine transporter 2
mtDNA	mitochondrial deoxyribonucleic acid	VPS35	vacuolar protein sorting-associated protein 35
mtROS	mitochondrial reactive oxygen species		

## Abstract

Parkinson's Disease (PD) is an incurable neurodegenerative movement disorder. PD affects 2% of the population above 65 years old; however, with the growing number of senior citizens, PD prevalence is predicted to increase in the following years. Pathologically, PD is characterized by dopaminergic cell neurodegeneration in the *substantia nigra*, resulting in decreased dopamine levels in the nigrostriatal pathway, triggering motor symptoms. Although the pathological mechanisms leading to PD are still unclear, large evidence indicates that oxidative stress plays an important role, not only because it increases with age which is the most significant risk factor for PD development, but also as a result of alterations in several processes, particularly mitochondria dysfunction. The modulation of oxidative stress, especially by using dietary mitochondriotropic antioxidants, represents a promising approach to prevent or treat PD. Although most mitochondria-targeted antioxidants with beneficial effects in PD-associated models have failed to show any therapeutic benefit in clinical trials, several questions remain to be clarified. Hereby, we review the role played by oxidative stress in PD pathogenesis, emphasizing mitochondria as reactive oxygen species (ROS) producers and as targets for oxidative stress-related dysfunctional mechanisms. In addition, we also describe the importance of using dietary-based mitochondria-targeted antioxidants as a valuable strategy to counteract the deleterious effects of ROS in pre-clinical and/or clinical trials of PD, pointing out their significance to slow, and possibly halt, the progression of PD.

**Keywords (6-8):** Dietary-based antioxidants; metabolism; mitochondria; mitochondria-targeted antioxidants; oxidative stress; Parkinson's Disease.

## 1. Introduction

Parkinson's Disease (PD) is the second most common neurodegenerative disorder after Alzheimer's Disease (AD) and the most common disorder affecting motor coordination <sup>1</sup>. PD prevalence ranges from 1 to 2 per 1,000 in unselected populations, and PD affects 2% of the population above 65 years old. Before 50 years of age, PD is rare but reaches a prevalence of 4% in the oldest age groups, pointing out age as the most common risk factor for PD development. The annual prevalence in high-income countries is 14 per 100,000 people in the total population and 160 per 100,000 people over 65 years old <sup>1</sup>. The incidence of PD is 1.5 times higher in males than females <sup>2</sup>. Although the most prevalent PD cases are sporadic forms, which have an unknown cause, around 10-15% of the individuals with PD have a mutation of one or several specific genes, called familial PD form (Table 1) <sup>3</sup>. Currently, at least ten mutated genes have been linked with familial PD, including *α-synuclein* (PARK1), *Parkin* (PARK2), *ubiquitin C-terminal hydrolase L1* (UCH-L1 or PARK5), *PTEN-induced kinase 1* (*PINK1* or PARK6), *DJ-1* (PARK7), *leucine-rich repeat kinase 2* (*LRRK2* or PARK8), *ATPase 13A2* (ATP13A2 or PARK9), *phospholipase A2 group VI* (PLA2G6 or PARK14), *F-Box protein 7* (FBXO7 or PARK15), *GRB10 interacting GYF protein 2* (GIGYF or PARK11) (Table 1) <sup>4,5</sup>.

Patients with PD have severe motor symptoms, including resting tremor, muscular rigidity, slowness of movement – *bradykinesia*, postural instability, and difficulty walking and thinking <sup>6</sup>. Patients can also have symptoms not affecting motor coordination, including depression, anxiety, dementia, and sleep abnormalities <sup>6</sup>. Currently, PD is clinically diagnosed based on the motor symptoms of the patients, including the presence of *bradykinesia*, combined with resting tremors or muscular rigidity and the response to specific drugs, including levodopa (L-DOPA), which is the most widely used drug to control PD-associated motor symptoms <sup>7</sup>. However, the diagnosis validation still depends on *postmortem brain* histology <sup>8</sup>, which led to the development of several models to study PD pathogenesis and therapeutics over time.

Parkinson's Disease has been known for over 200 years, and despite intense research efforts, effective treatments and timely diagnostic markers are lacking. This review focuses on oxidative stress in PD, particularly on the role of mitochondrial dysfunction in that phenomenon. We also address the use of dietary antioxidant-based therapeutic strategies to counteract the deleterious effects of reactive oxygen species (ROS)

to prevent or treat PD. Although several antioxidants have shown minimal therapeutic benefit, we summarize the molecular and metabolic evidence that novel mitochondria-targeted antioxidants can be multifunctional compounds promoting mitochondrial health. Moreover, we also discuss why antioxidant therapies failed when used in clinical trials, thus showing new insights to achieving novel opportunities for intervention.

## **2. Oxidative Stress in Parkinson's Disease pathogenesis**

Pathologically, PD is characterized by dopaminergic neurodegeneration in *substantia nigra pars compacta (SNpc)*, culminating in decreased dopamine levels in the *dorsal striatum*<sup>9</sup>. Although the exact mechanism by which dopaminergic neuronal death occurs is unclear, several contributors to PD pathophysiology have been identified (Figure 1). The widely accepted PD disease mechanisms include accumulation of misfolded protein aggregates, alterations in dopamine metabolism, neuroinflammation, mitochondrial dysfunction, and impaired quality control mechanisms (Figure 1)<sup>10</sup>. The indefiniteness related with a specific biochemical mechanism for the progressive and complex neurodegenerative cascade makes the oxidative stress an indisputable player in that process<sup>11</sup> (Figure 1). Since oxidative stress is also aging-associated and aging is the most significant risk factor for PD development, it is unclear whether oxidative stress is cause or consequence of detrimental pathways during PD pathogenesis<sup>11</sup>. The oxidative stress theory of aging postulates that reductions in physiologic functions associated with age are caused by a slow and steady accumulation of macromolecules-induced oxidative damage, which increases with age and are associated with organisms' life expectancy<sup>12</sup>. A study based on the aging-time-control coefficient showed that 25 out of 57 molecular processes are controlled by aging<sup>13</sup>.

### **2.1. Disruption of redox circuits in PD pathogenesis**

The brain is one of the major organs that generates large amounts of ROS<sup>14</sup>. ROS are generated intracellularly as byproducts of basal metabolism,<sup>15</sup> and can exert beneficial or detrimental effects at the cellular level depending on their concentration and origin, and of antioxidant defense concentrations and activity in different sub-cellular compartments<sup>15</sup>. For instance, electrons derived from different substrates are carried into electron transport chain (ETC) in order to fuel OXPHOS. Electrons leaking from ETC can react

with oxygen, generating superoxide, which is itself a ROS and can generate other ROS <sup>16</sup>. Moreover, NADPH oxidase could product superoxide either as a by-product of ETC or from a dysfunctional variant in the conversion of xanthine dehydrogenase to xanthine oxidase, since in both mechanisms NADPH oxidase transfers electrons from NADPH to molecular oxygen through NOX catalytic subunit <sup>17</sup>. Physiologically, low ROS levels act as a second messenger, modifying oxidation-reduction (redox) states, which culminates in post-translational modifications of kinases and phosphatases and ultimately in the perturbation of processes such as cell survival and proliferation (Figure 2) <sup>15</sup>. ROS-induced modifications can also regulate enzymatic activity, drug detoxification and cellular responses to external perturbations <sup>18</sup>, all pivotal in maintaining homeostatic signaling events. In fact, mitochondrial oxidants induce an adaptive response to improve systemic defense, the so-called mitohormesis (Figure 2) <sup>18</sup>. Among the several ROS, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) acts as a second messenger due to its neutral and membrane-permeable capacity allowing its relatively free diffusion from the generation site. Furthermore, H<sub>2</sub>O<sub>2</sub> has no unpaired electrons in the last orbital making it comparatively less reactive than other ROS <sup>19</sup>.

On the other hand, ROS overproduction resulting from endogenous sources, including transmembrane NADPH oxidases (NOXs) and the mitochondrial ETC, or cumulative environmental exposure, commonly denominated as the exposome, can be detrimental to the cell's well-being. The exposome can involve nutritional cues, and exposure to drugs, toxicants, pollutants, physical (ultraviolet, X-ray, and ionizing radiation) or psychological stressors <sup>18</sup>. The steady-state cellular levels of ROS are a balance between their production and their degradation/removal. Therefore, oxidative stress can result from overproduction of reactive species or by a deficiency of enzymatic and endogenous non-enzymatic antioxidants. The impact of oxidative stress damage on the organism depends on the oxidant type, the place, duration and intensity of its production, the composition and activities of various antioxidants, and repair systems' ability <sup>20</sup>. Classically, the imbalance between ROS and endogenous antioxidants defense mechanisms results in an oxidative stress state, promoting irreversible damage to biomolecules, causing growth arrest and cell death, which are customarily associated with pathological states, including PD, and are currently referred to as oxidative distress <sup>18</sup>.

Oxidative distress that occurs in neurons can contribute to PD pathogenesis. Nigral dopaminergic neurons seem to be particularly vulnerable to oxidative stress, possibly due to long axons and millions of synapses, requiring high energy input and hence a high oxygen demand <sup>21</sup>. Moreover, compared with other organs (e.g. liver) the brain has lower antioxidant enzymes activities, namely superoxide dismutase (SOD), catalase, and glutathione peroxidase (GPx) activities. At the same time, the brain contains a high percentage of lipids with unsaturated fatty acids that are prone to lipid peroxidation. In addition, the brain glutathione (GSH) concentration is lower than in other organs, such as the liver, kidney or the small intestine <sup>14</sup>. In fact, cytosolic GSH in neurons is approximately 5 mM compared with 10-11 mM in hepatocytes <sup>22</sup>.

Moreover, oxidative distress can result from increased levels of dopamine and its metabolites and mitochondrial dysfunction. On the other hand, antioxidant defense mechanisms in the brain are inadequate, limiting its protective action against ROS <sup>23</sup>. This complex system will be described below in the context of dopaminergic neurons (Figure 3).

### **2.1.1. Disruption of redox homeostasis in dopaminergic neurons**

#### **2.1.1.1. Pro-oxidant properties of dopamine**

*SNpc* is a vital part of the brain basal ganglia, which predominantly comprises neurons that secrete dopamine, a catecholaminergic neurotransmitter <sup>24</sup>. Taking into account the selective degeneration of dopaminergic neurons of *SNpc*, this suggests dopamine itself may be an oxidative stress source. Dopamine is synthesized from tyrosine by tyrosine hydroxylase (TH) and aromatic l-amino acid decarboxylase (AADC) <sup>25</sup>. After its uptake by vesicular monoamine transporter 2 (VMAT2), dopamine is stored in synaptic vesicles <sup>26</sup>. However, in the presence of an excessive cytosolic amount of dopamine outside of the synaptic vesicle in damaged neurons, that neurotransmitter is metabolized by monoamine oxidase (MAO) generating H<sub>2</sub>O<sub>2</sub> or by auto-oxidation to cytotoxic ROS, which in the presence of free Fe<sup>2+</sup> generates hydroxyl radical (<sup>•</sup>OH), H<sub>2</sub>O<sub>2</sub>, and dopamine-quinone species (Figure 3). The byproducts of dopamine-quinone species resulting from dopamine oxidation react with GSH to form adducts (GS-DAQ) through the GSH-transferases activity <sup>27</sup>. Subsequently, mitochondrial respiration is inhibited and the permeability transition pore opening occurs in brain mitochondria promoting cell death and neuronal loss <sup>28,29</sup>. Thus, it has been proposed that dopaminergic

cell death-induced by mitochondrial inhibitors depends on dopamine oxidation <sup>27</sup>. In fact, it was demonstrated that dopamine interacts with lipid hydroperoxides produced during peroxidation events leading to the formation of adducts that could account for the selective cytotoxicity of dopaminergic neurons <sup>30</sup>. Moreover, hexanoyl dopamine, an arachidonic acid-derived adduct, causes severe cytotoxicity in human dopaminergic neuroblastoma SH-SY5Y cell line <sup>30</sup>. Since PD also affects non-dopaminergic cells, and not all dopaminergic neurons are equally affected during the disease progression <sup>31</sup>, the high toxicity of dopamine can sensitize dopaminergic neurons to oxidative damage.

#### 2.1.1.2. Superoxide dismutases (SODs) activity dysregulation

During cell metabolism, oxygen suffers a series of univalent reduction reactions leading to the production of  $\cdot\text{O}_2^-$ ,  $\text{H}_2\text{O}_2$  and  $\text{H}_2\text{O}$ . Superoxide anion is dismuted by SODs to  $\text{H}_2\text{O}_2$ , which is catalyzed to  $\text{H}_2\text{O}$  by catalase, peroxiredoxins (Prx) or GPx (Figure 3) <sup>32</sup>. SODs are the major antioxidant defense systems, which consist of three isoforms in mammals: the Cu-Zn-dependent SOD (Cu/ZnSOD or SOD1) located in the cytoplasm or the mitochondrial intermembrane space, the manganese-dependent SOD (MnSOD or SOD2) located in the mitochondrial matrix, and the extracellular Cu/ZnSOD (SOD3) <sup>32</sup>.

The protective role of SOD1 and SOD2 in both sporadic and genetic PD models has been previously demonstrated <sup>33</sup>. In fact, SOD1 and SOD2 overexpression rescue the toxic effect of products derived from dopamine in human SH-SY5Y cells <sup>34</sup>. Analysis of SODs in PD brains have shown no changes in SOD1 activity, while SOD2 activity was increased <sup>35</sup>, suggesting that SOD2 is activated in response to ROS overproduction. However, assays in human SH-SY5Y neuroblastoma cells or *drosophila melanogaster*-induced PD showed that SOD2 is important to protect against high paraquat concentrations. In contrast, at sub-lethal paraquat concentrations, the over-expression of SOD1 is enough to rescue the paraquat-associated toxicity in *drosophila melanogaster* <sup>36</sup>. The same occurs when SOD1 is specifically expressed in dopaminergic neurons. These observations indicate that other cytosolic processes inside dopaminergic neurons, namely dopamine oxidation, may amplify ROS toxicity <sup>36</sup>. Defects in SODs activity develop over time in PD, remaining unclear whether this is a later manifestation of deterioration of antioxidant activity or an epigenetic-related phenomenon <sup>37</sup>.



### 2.1.1.3. Thioredoxins and peroxiredoxins system (dys)homeostasis

The thioredoxin system consists of three proteins, namely thioredoxin (Trx), thioredoxin reductase (TrxR), thioredoxin peroxidase (TPx), and nicotinamide adenine dinucleotide phosphate (NADPH) as the electron donor <sup>38</sup>. The central core element is Trx, which has an active conserved site Cys-Pro-Gly-Cys, capable of reducing disulfide bonds in proteins <sup>39</sup>. In humans, two different isoforms of Trx are found: a cytosolic isoform (Trx1) and a mitochondrial isoform (Trx2), which can be found in different human body tissues. Trx plays essential roles in regulating redox signals and stress response, modulating several signaling pathways, transcription factors, and immunological response <sup>38</sup>.

The enzyme responsible for catalyzing the reduction of the disulfide at the Trx active site is TrxR. In humans, there are three different TrxR: TrxR1 locates in the cytosol, TrxR2 locates in mitochondria and thioredoxin-glutaredoxin reductase (TrxR3) <sup>40</sup>. While TrxR1 and TrxR2 are expressed in all mammalian cells and tissues, including the brain, TrxR3 is expressed in testis <sup>38</sup>. Trx acts as the reducing agent for peroxiredoxins and others proteins, namely p53, nuclear factor erythroid 2-related factor 2 (Nrf2), hypoxia-inducible factor 1-alpha (HIF1- $\alpha$ ), and others <sup>41</sup>.

The system dependence of Trx is also linked with TPx, which cooperatively can reduce lipid peroxides and H<sub>2</sub>O<sub>2</sub> (Figure 3). TPx is part of a family of proteins called Prx <sup>42</sup>. The Prx family are thiol-dependent peroxidases promoting the reduction of H<sub>2</sub>O<sub>2</sub>, ONOO<sup>-</sup> and alkyl hydroperoxides. Prx has a conserved active site cysteine, called the peroxidatic cysteine, which reacts with peroxide forming a cysteine sulfenic acid (R-SOH) and releases the corresponding alcohol or water <sup>43</sup>. However, some of the Prx contain also a second cysteine, called the resolving cysteine, which attacks the R-SOH to form a water molecule and a disulfide bond in the protein. In humans, there are six isoforms of Prx (Prx1-6) and their expression in central nervous system (CNS) is cell type-specific and localize in different cellular compartments <sup>21</sup>

Trx and Prx induction contributes to brain tolerance to toxic insults. The thioredoxin/ thioredoxin reductase (Trx/TrxR) system protects dopaminergic cells against toxicity induced by MPP<sup>+</sup>, 6-OHDA and paraquat <sup>27</sup>. Work performed in patients with sporadic PD (sPD) showed that oxidative forms of DJ-1 protein protect both culture cells and *substantia nigra* of mice from oxidative stress by inducing Trx1 gene expression

via the transcription factor Nrf2<sup>44</sup>. Incubation of immortalized rat dopaminergic cells with auranofin, an inhibitor of TrxR, in the presence of sub-toxic concentrations of PD-associated toxins, including 6-OHDA and paraquat resulted in H<sub>2</sub>O<sub>2</sub> increased levels, mitochondrial dysfunction, and oxidative stress-induced cell death<sup>45,46</sup>.

Overexpression of peroxiredoxin 1, 2 and 4 (Prx1/2/4) protects against dopaminergic cell death induced by 6-OHDA, while the silencing of the mitochondrial isoforms, Prx3 and Prx5 increased the sensitivity to MPP<sup>+</sup>. Dopaminergic neuronal cells treated with 6-OHDA lead to ROS generation, which causes oxidative modification of Prx1, while Prdx1 overexpression protects neuronal cells against 6-OHDA-induced cell death by means of ROS scavenging<sup>47</sup>. Moreover, brain tissues from PD patients show increased levels of Prdx2<sup>48</sup> and Prdx3<sup>49</sup>. Prdx2 overexpression attenuated 6-OHDA neurotoxicity in *in vitro* and *in vivo* models of PD through antiapoptotic effects, namely by inhibiting caspase-dependent apoptosis and suppressing of apoptosis signal-regulating kinase signaling cascade<sup>50</sup>.

<sup>51</sup>. Glutathione metabolism is regulated by the glutathione cycle, in which glutamate, the most abundant excitatory transmitter in the CNS, is added and released during this cycle<sup>52</sup>. GSH can react with  $\cdot\text{O}_2^-$ , NO, ONOO $\cdot$  and hydroxyl radical, this latter is only scavenged by GSH since there is no other known antioxidant system against hydroxyl radical (Figure 3). Moreover, GSH works as an electron donor to reduce H<sub>2</sub>O<sub>2</sub> or other peroxides by GPx<sup>53</sup>. The brain has a relatively high level of GPx when compared with catalase. However, the *substantia nigra* contains lower GSH levels than other brain parts, including *cortex*, *cerebellum*, *hippocampus*, and *striatum*<sup>54</sup>. H<sub>2</sub>O<sub>2</sub> is reduced to H<sub>2</sub>O by GPx using GSH, which is oxidized to GSH disulfide (GSSG) (Figure 3). The latter is reduced back by GSH reductase with NADPH to GSH and is reused as a GPx substrate<sup>53</sup>.

Decreased GSH levels have been considered one of the earliest biochemical alterations associated with

PD, since the loss of GSH occurs in incidental Lewy Bodies (LBs) disease, which is considered a pre-occurring PD phase in which non motor symptoms are present. A decrease in GSH levels by 40-50% has been reported in autopsied brain PD patients<sup>55</sup>. Accordingly, depletion of GSH promotes nigrostriatal degeneration in PD mice model<sup>27</sup>. Furthermore, dopaminergic neurons treated with buthionine sulfoximine (BSO), an inhibitor of GSSG reductase, showed decreased GSH levels and increased neurotoxicity induced by 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>) and 6-OHDA<sup>56,57</sup>. In addition, decreased catalase and GSH levels were demonstrated in blood and *postmortem* brain PD patients<sup>58</sup>. A clinical trials phase II using GSH to slow down the progression of PD symptoms was already completed with no results released so far (NCT02424708).

## **2.2. Mitochondria as a ROS producer and its involvement in Parkinson's Disease pathogenesis**

Mitochondria are cellular organelles that regulate important anabolic and catabolic pathways<sup>59</sup>. Although one of the main functions of mitochondria is adenosine triphosphate (ATP) production through oxidative phosphorylation (OXPHOS), that organelle also plays essential roles in calcium homeostasis, biogenesis of iron-sulfur clusters, heme and steroid synthesis, fatty acids catabolism, redox regulation of cellular signaling, and regulation of programmed cell death<sup>60</sup>. Moreover, mitochondria have essential roles in other cellular processes, including autophagy, and cellular proliferation and differentiation<sup>61</sup>. According to specific internal and external cues, including oxygen availability or oxidative stress, and because of their nature, dynamic mitochondrial distribution, structure, and function can be modified<sup>62</sup>, impacting cell physiology. Consequently, alterations in cell's redox circuits impact mitochondrial morphology and function at different levels (Figure 4), which may have pathophysiological consequences<sup>63</sup>.

Approximately 2-4% of the oxygen consumed by mitochondria is redirected to form  $\cdot\text{O}_2^-$ <sup>14</sup>. Mitochondrial complexes I and III are primary sources of mitochondrial  $\cdot\text{O}_2^-$ <sup>64</sup> and the main target of ROS-induced oxidative stress, leading to ATP production inhibition and uncontrolled ROS generation. Subsequently, the vicious cycle between the defects in the ETC and ROS production drives the uncontrolled oxidative stress that may underlie PD pathogenesis, especially in the progressive dopaminergic neuronal

degeneration<sup>65</sup>. In this sense, ROS appears as the trigger point and the one significant consequence of mitochondrial dysfunction that occurs in PD (Figure 4).

Mitochondrial dysfunction can also result from environmental toxins exposure, implicated in sPD epidemiology. This increased awareness of environmental exposure associated with the risk of developing PD later in life through mitochondrial damage and augmented oxidative stress (Figure 4)<sup>66</sup>.

MPTP-induced parkinsonism in intravenous drug users<sup>67,68</sup> was the first report showing that mitochondrial dysfunction is related to PD symptoms<sup>69</sup>. MPTP penetrates the blood brain barrier (BBB), being converted into its toxic form MPP<sup>+</sup> by glial MAO<sup>70</sup>. MPP<sup>+</sup> interferes with mitochondrial complex I of respiratory chain activity in dopaminergic neurons, causing selective neurodegeneration in human and mouse *SNpc*<sup>67,71</sup>. Rotenone is a well-studied insecticide presenting neuronal toxicity. Rotenone is highly lipophilic and can easily cross biological membranes, including BBB<sup>72</sup>, leading to dopaminergic neurodegeneration. Rotenone, a mitochondrial complex I inhibitor, results in decreased ATP levels, leading to electron leakage that can form  $\cdot\text{O}_2^-$ <sup>72</sup>. Currently, rotenone treatment is used as a PD model due to its ability to reproduce parkinsonism-like features<sup>73</sup>, including the accumulation and aggregation of specific hallmarks, such as  $\alpha$ -synuclein,  $\alpha$ -synuclein- and polyubiquitin-positive LBs and Lewy neurites, combined with apomorphine-responsive behavior deficits, early and sustained activation of microglia, oxidative modification and translocation of DJ1 into mitochondria *in vivo*, impairment of ubiquitin-proteasome system (UPS), iron accumulation in the *substantia nigra* through a mechanism involving transferrin and transferrin receptor 2,  $\alpha$ -synuclein pathology in enteric neurons and functional deficits in gastrointestinal function, including gastroparesis<sup>74</sup>. The decreased activity of mitochondrial complex I in neurons from idiopathic PD patients were associated with mitochondrial DNA (mtDNA) depletion and decreased in mitochondrial transcription factor A (TFAM) and mitochondrial transcription factor B2 (TFB2m) protein contents<sup>75,76</sup>. Furthermore, decreased complex I expression has also been shown in several brain regions of PD<sup>77-79</sup>. Complex I deficiency was observed in *substantia nigra* of patients with PD. However, this deficiency in complex I can be mtDNA damage-independent<sup>79</sup>. However, one question arises when considering the role of mitochondrial complex I dysfunction in PD etiology: why patients with mitochondrial diseases with complex I abnormalities rarely develop a PD phenotype? Currently, no mtDNA mutations were found to be associated with PD (Figure 4), and mitochondrial diseases-associated

PD is restricted to mutations affecting mtDNA maintenance genes, such as DNA polymerase subunit gamma (POLG) and twinkle mtDNA helicase<sup>80,81</sup>. In fact, the exonuclease dysfunctional POLG mutator mouse does not show a PD phenotype itself; however, when crossed with a Parkin knockout mouse, the appearance of a PD phenotype occurs. These observations suggest that other insults are required, besides accumulation of somatic mtDNA mutations to cause PD<sup>82</sup>. It is known that somatic mtDNA mutations are increased in *substantia nigra* of rotenone-treated rats<sup>80</sup>.

The mitochondrial respiratory chain in neurons from sPD patients presented defects in complex I and II and several depletions in mtDNA related to decreased mtDNA copy number in *substantia nigra*<sup>75</sup>. Aging is a risk factor for developing sPD, associated with reduced mitochondrial function, particularly an increased number of mtDNA mutations and oxidative stress, and decreasing respiratory chain activity<sup>83</sup>. Thus, mitochondrial dysfunction resulting from mtDNA and respiratory chain abnormalities contributes to PD pathogenesis by decreasing the threshold for the susceptibility to other genetic and environmental insults. Also, alterations in tricarboxylic acid (TCA) cycle fluxes occur in several PD models, including a decrease in  $\alpha$ -ketoglutarate dehydrogenase activity. Moreover, complex I inhibitors and paraquat and PINK1 mutations promoted the oxidative inactivation of aconitase<sup>27</sup>.

Several mutated genes in familial PD are directly associated with mitochondrial dysfunction, and increased oxidative stress, including  *$\alpha$ -synuclein*, *LRRK2*, *Parkin*, *PINK1* and *ATP13A2*. Similarly, several mutated genes in PD, including *DJ-1*, *LRRK2*, *PINK1*, and *Parkin*, encode several regulators of mitochondrial and ROS homeostasis<sup>83</sup>. Alterations in newly found PD-relevant genes lead to mitochondrial dysfunction, including *vacuolar protein sorting-associated protein 35 (VPS35)* and *coiled-coil-helix-coiled-coil-helix domain containing 2 (CHCHD2)*, which lead to mitochondrial fragmentation and abnormalities in mitochondrial structure, respectively<sup>83</sup>.  $\alpha$ -Synuclein has a non-canonical mitochondrial targeting sequence, being also found in mitochondria-associated membranes (MAM), affecting mitochondrial structure and function<sup>84</sup>. Moreover, increased levels of wild-type  $\alpha$ -synuclein induce mitochondrial fragmentation and production of ROS both *in vitro* and *in vivo* models<sup>85</sup>. MAM is a structure that forms an interface between the endoplasmic reticulum (ER) and mitochondria, playing important roles in calcium regulation, lipid transfer and apoptosis<sup>86</sup>. Indeed, mutant or overexpressed wild-type  $\alpha$ -synuclein leads to dissociation of ER and

mitochondria at MAM, impairing calcium exchange and decreasing mitochondrial energy production <sup>87</sup>. Mutated  $\alpha$ -synuclein in human dopaminergic neurons leads to decreased mitochondrial biogenesis through downregulation of peroxisome proliferator activated receptor gamma coactivator 1 alpha (PGC1- $\alpha$ ) <sup>80</sup>.

Several models of wild-type mutated *LRRK2* presented higher vulnerability to mitochondrial toxins, increased ROS production and defects in mitochondrial dynamics (Figure 4) <sup>80</sup>. Mutant *VPS35* triggers mitochondrial fragmentation leading to impaired mitochondrial complex I assembly and activity, promoting neurodegeneration <sup>88</sup>. In addition, loss of *ATP12A2* impairs glycolysis impacting cellular bioenergetics and aggravating mitochondrial dysfunction. Furthermore, zinc ( $Zn^{2+}$ ) homeostasis is deregulated by loss of *ATP13A2* through impairing vesicular sequestration, promoting mitochondrial and lysosomal dysfunction contributing to defective mitophagy (Figure 4) <sup>80,89,90</sup>. While Parkin-deficient models also show several defects in mitochondrial morphology and function <sup>91</sup>, defective PINK1 models impact mitochondrial function, especially degradation, morphology, and trafficking <sup>80</sup>. Moreover, *Parkin* mutations are associated with sporadic early-onset parkinsonism and *PINK1* mutations are connected with parkinsonism in pediatric patients <sup>92</sup>.

### **3. Pre-clinical and clinical studies with dietary antioxidants for Parkinson's Disease**

#### **3.1. The role of dietary polyphenols as redox modulators**

Considerable efforts have been made to search and/or develop compounds capable of reducing ROS levels and/or preventing ROS-induced damage and mitochondrial dysfunction modulating cellular redox circuits.

Exogenous antioxidants or dietary antioxidants can be beneficial for decreasing oxidative stress status as they might compensate the inability of the endogenous defense systems and enhance the overall antioxidant response emerging as attractive drug candidates for PD therapy <sup>93-95</sup>.

Dietary polyphenols are bioactive molecules with remarkable antioxidant properties since they can decrease oxidative stress by acting as primary or secondary antioxidants and/or up-regulating endogenous antioxidant defenses <sup>96</sup>. In general, polyphenols can act at the "ROS-regulating level" that include a mechanism of direct ROS- or RNS-scavenging and/or a favorable modulation of removing ROS by redox-regenerating enzyme activity <sup>97</sup>. The direct radical scavenging action of polyphenols is due to its reaction with

$\cdot\text{OH}$ ,  $\cdot\text{O}_2^-$ ,  $\cdot\text{NO}$ , alkoxyl ( $\cdot\text{RO}$ ) and peroxy radicals ( $\cdot\text{RO}_2$ ), and with non-radical species, including ONOO $\cdot$  and hypochloride acid (HClO) <sup>98</sup>. The antioxidant activity of polyphenols mediated by ROS-scavenging is attributed to the presence of phenolic groups <sup>98</sup>. Notwithstanding, polyphenols can also be considered secondary antioxidants due to their ability to up-regulate the endogenous antioxidant defense network, particularly the ROS-removing enzymes, such as SOD, catalase, and GPx enzymes <sup>99</sup>. Some redox-regenerating enzymes, including glutathione reductase (GR) and Trx can also be up-regulated by polyphenols <sup>99</sup>. Furthermore, polyphenols can target ROS-formation processes, since they can perform a direct inhibitory action on metal-dependent free radicals formation process, namely  $\cdot\text{OH}$ ,  $\cdot\text{O}_2^-$  and ROS-producing enzymes. Polyphenols are also particularly active in chelating transition metals such as iron and copper ions, rendering them unavailable to participate in free radical-generating reactions <sup>100</sup>. Under physiological conditions, polyphenols' iron-binding properties are related to their adequate capacity to modulate cellular iron homeostasis <sup>96</sup>. Moreover, the polyphenols can inhibit NADPH oxidase, possibly by interfering with the assembly or inhibiting the expression of several subunits, and the mitochondrial-bound monoamine oxidase (MAO) activity <sup>97</sup>.

### **3.2. Dietary polyphenol-based antioxidants in Parkinson's Disease**

Dietary polyphenols can overcome the defects of the antioxidant defense system and control ROS levels given their biological potential. Therefore, the neuroprotective potential of several dietary antioxidants, including resveratrol <sup>101</sup>, curcumin <sup>102</sup>, quercetin <sup>23</sup>, epigallocatechin-3-gallate (EGCG) <sup>103</sup>, apocynin and diapocynin <sup>104</sup> have been tested.

Resveratrol, a natural polyphenolic non-flavonoid compound, presents several *in vitro* and *in vivo* neuroprotective effects, protecting neurons against oxidative stress damage and toxicity and prevents apoptotic neuronal death. Resveratrol also ameliorates mitochondrial dysfunction observed in some *in vivo* PD models <sup>101</sup>. Moreover, resveratrol crosses the BBB by transmembrane diffusion <sup>101</sup>. Despite the great potential of resveratrol as a therapeutic agent, several limitations were also described. Resveratrol has low bioavailability especially due to its short biological half-life, rapid metabolism, chemical instability, clearance, high

photosensitivity, and poor solubility in water<sup>105</sup>. Low bioavailability of resveratrol hampers its accumulation at necessary concentrations for successful therapy<sup>106</sup>.

Curcumin is a polyphenol and an active turmeric component from *Curcuma longa*, a dietary spice used in Indian cuisine and medicine<sup>107</sup>. Neurological benefits of curcumin in PD models are related to its antioxidant function. In the 6-OHDA rat PD model, curcumin protected neurons in substantia nigra against ROS-induced cell death by improving striatal dopamine levels and chelate free Fe<sup>2+</sup>. The phenolic rings probably mediate its antioxidant mechanism of action, and the diketone groups curcumin serve as an electron trap preventing the generation of H<sub>2</sub>O<sub>2</sub>, •OH and •O<sub>2</sub><sup>-</sup>. Moreover, curcumin reverses the inhibition of DNA repair enzymes in an *in vivo* PD model and in SH-SY5Y cells<sup>108</sup>. Similarly, curcumin *in vitro* pre-treatment attenuated peroxynitrite (ONOO<sup>-</sup>)-mediated nitrosative stress and mitochondrial dysfunction<sup>102</sup>. Besides, curcumin can cross the BBB, while it has been regarded as a potential therapeutic factor for many nervous system diseases due to its pleiotropic therapeutic effects<sup>109</sup>. Although beneficial neuroprotective effects of curcumin have demonstrated, its clinical application has been hampered by its low aqueous solubility, reduced absorption in the gastrointestinal tract, high rate of metabolism, poor stability in the body fluids, rapid clearance, and limited bioavailability. Although a large number of clinical trials using curcumin have been demonstrated its safety, the limitations described above and the clinical trials's inconclusive results limit curcumin approval as a drug for clinical application<sup>109-112</sup>.

Quercetin, a plant flavanol from the flavonoid group, has been used as a PD supplemental therapy. Several reports showed that its neuroprotective effect is related to its anti-inflammatory and antioxidant properties<sup>113</sup>. Treatment with quercetin attenuated motor deficits, biochemical and neurotransmitter alterations against rotenone- and iron- supplemented-induced PD rats<sup>114</sup>. Moreover, quercetin's chronic administration in a 6-OHDA mice model of PD decreased ROS levels and lipid peroxidation<sup>23</sup>. Although quercetin showed beneficial effects on various *in vivo* models of neural disorders, contrasting data also exist, resulting from quercetin-induced neurotoxicity. Thus, quercetin efficacy in neurodegenerative disorders lacks evidence<sup>115</sup>. Also, the low solubility and bioavailability of quercetin limit its clinical use<sup>116</sup>.

EGCG, the most abundant green tea polyphenol, harbors free-radical scavenging, iron-chelating, and anti-inflammatory properties. Several studies reported neuroprotective effects of EGCG against MPP<sup>+</sup>- or



MPTP-induced neurodegeneration in both *in vitro* and *in vivo* PD models <sup>103</sup>. EGCG inhibited MPP<sup>+</sup>-induced oxidative stress in PC12 cells by increasing antioxidant enzymes and through sirtuin 1/ PCG1- $\alpha$  (SIRT1/PCG1- $\alpha$ ) signaling pathway. Moreover, EGCG inhibited microglial activation protecting against MPTP-induced neuronal loss. EGCG upregulates ferroportin, an iron-export protein, in *substantia nigra*, reduces oxidative stress, and rescues MPTP-induced functional and neurochemical deficits in mice <sup>103</sup>. However, EGCG bioavailability in the brain is low, limiting its use as a neuroprotective agent <sup>117</sup>. Due to the lack of evidence of EGCG activity in humans and the lack of well-controlled clinical trials, the EGCG effect in preventing oxidative stress-related disorders, namely PD, is still under evaluation.

Apocynin, a plant-derived antioxidant, also protected dopaminergic neurons against MPP<sup>+</sup>-induced oxidative stress and cell death. Apocynin is converted into diapocynin, which is an inhibitor of NADPH oxidase. In MPTP-treated mice, the administration of diapocynin protected against nigrostriatal damage and oxidative stress <sup>118</sup>. Apocynin stability is low due to the phenolic hydroxyl group in its structure <sup>119</sup>, and apocynin bioavailability is also low since it quickly metabolizes into glucuronic conjugate <sup>104</sup>. While apocynin crosses the BBB at a low rate, this effect was not observed in diapocynin <sup>104</sup>.

All the antioxidants described above, despite their demonstrated beneficial effects in several PD models, mostly *in vitro*, also presented common limitations, especially at the level of limited bioavailability, incapacity to cross BBB or unspecificity in tissue/organelle-targeted, which have hampered their clinical applications.

### 3.3. Controlling ROS production by targeting mitochondrial function

Discovering mitochondrial drug candidates is a great challenge since it requires a target-specific organelle affinity and, at the same time, a restricted drug safety window. Applying a concept pioneered by Skulachev et al. to measure  $\Delta\Psi_m$  *in vitro* <sup>120</sup>, the covalent attachment of lipophilic cations, such as triphenylphosphonium cation (TPP<sup>+</sup>), is a golden-standard method to target small bioactive molecules to mitochondria <sup>121</sup>. TPP<sup>+</sup> derivatives are rapidly and extensively accumulated *in vivo* by mitochondria driven by the large  $\Delta\Psi_m$ , negative inside the matrix <sup>122</sup>. These compounds pass through all biological membranes and accumulate in the mitochondrial matrix faster than their non-targeted molecules. The uptake mechanism is by

now well understood. TPP cations can cross the plasma and mitochondrial inner membrane due to their extensive hydrophobic surface area and the cation's large outer ionic radius, effectively lowering the activation energy needed for membrane passage <sup>123,124</sup>. The Nernst equation adequately describes the uptake dependent of membrane potential of the lipophilic cations, increasing 10-fold for every ~60 mV of  $\Delta\Psi_m$ . It is estimated that a hundred- or thousand-fold increase in the antioxidant concentration occurs inside mitochondria. The extent of TPP<sup>+</sup> derivative-uptake depends on plasma  $\Delta\Psi$  and  $\Delta\Psi_m$ , cell volume, the external media, and the number of mitochondria within a given cell. Once inside mitochondria, these derivatives can promote beneficial effects through several mechanisms, namely by scavenging reactive radicals and control mitochondrial redox signaling and/or preventing membrane lipid peroxidation <sup>121</sup>.

Considering that mitochondria are main sites for the ATP production, it is not surprising that most targets of mitochondrial drug are related and involved in energy metabolism and mostly related with mechanisms that regulate ROS, respiration and mitochondrial biogenesis. Ultimately, potential novel drugs should restore mitochondrial function as well as normalize mitochondrial ROS production <sup>125</sup>. Nowadays, the central focus on finding drugs that target mitochondrial function is related to the discovery and development of antioxidants inhibiting mitochondrial oxidative damage <sup>121</sup>.

### **3.4. Mitochondria-targeted antioxidants used in Parkinson's Disease pre-clinical studies**

By using antioxidants that target mitochondrial oxidative stress in order to decrease mitochondrial ROS (mtROS) in pathological processes may be essential to both normal cell function and in disease prevention.

The most studied mitochondria-targeted antioxidants are MitoQ and SkQ1. While MitoQ consists of a TPP<sup>+</sup> unit covalently attached to the endogenous antioxidant ubiquinone <sup>126</sup>, SkQ1 consists of a TPP<sup>+</sup> unit covalently attached to plastoquinone, a chloroplast quinone, as core moiety <sup>127</sup>. MitoQ is a promising neuroprotective compound because of its direct antioxidant action. Similar to its parent antioxidant (CoQ10), MitoQ scavenges peroxy, ONOO<sup>-</sup>, and <sup>•</sup>O<sub>2</sub><sup>-</sup>, protecting mitochondria against lipid peroxidation (Figure 4) <sup>118</sup>. After detoxifying oxidants, MitoQ is recycled by the respiratory chain complex II to the active ubiquinol antioxidant form <sup>118</sup>. MitoQ inhibited 6-OHDA-induced mitochondrial fragmentation in SH-SY5Y cells and

prevented MPP<sup>+</sup>-induced loss of neurons in a PD's dopaminergic cell culture model <sup>118</sup>. In a Parkinson's disease mouse model treated for 14 days with MitoQ, the number of dopaminergic neurons increased relative to untreated PD mouse model <sup>128</sup>. The successful of pre-clinical trials obtained with MitoQ led to clinical trials for several diseases, including PD (NCT00329056). However, the results of the trials were disappointing. Some justifications include time elapsed from PD diagnosis, in which 50% of dopaminergic neurons have already been lost or even the insufficient penetration of the BBB by MitoQ <sup>129</sup>.

Administration of SkQ1 to a MPTP-model of PD using C57BL/6 mice showed that SkQ1 increased dopamine levels (Figure 4), while the loss of dopaminergic neurons in the *substantia nigra* and signs of sensory-motor deficiency were decreased <sup>130</sup>.

Similarly, other mitochondria-targeted antioxidants have been tested with beneficial effects in several PD models, including MitoVitE (combination of the vitamin E with TPP<sup>+</sup>), MitoTEMPO (combination of the piperidine nitroxide TEMPO with TPP<sup>+</sup>), and MitoApocynin (combination of the apocynin with TPP<sup>+</sup>), MitoPBN (combination of the nitron antioxidant derived from IBTP with TPP<sup>+</sup>), MitoSOD (combination of the SOD with TPP<sup>+</sup>) <sup>118</sup>. Zhelev et al. <sup>131</sup> demonstrated that MitoTEMPO increased the activity of dopaminergic neurons in the brain of MPTP-treated mice <sup>131</sup>. Also, MitoApocynin prevented microglial activation, oxidative damage, mitochondrial dysfunction, and progressive neurodegeneration in TFAM<sup>-/-</sup> mice <sup>132</sup>. The beneficial effects of these mitochondria-targeted compounds have been extensively studied using several disease models <sup>133-137</sup>; however their effects in PD models have been poorly studied.

Although attractive, the success of these approaches has been troubled by several challenges and limitations, with none resulting yet in a Food and Drug Administration (FDA)-approved drug for PD therapy. The lack of efficacy may relate with the pre-clinical assessment of these compounds in inadequate PD animal models <sup>138</sup>. Moreover, the pre-clinical data set should define better neuroprotective dose-response relationships, pharmacokinetic-pharmacodynamic correlations, therapeutic windows, optimum doses regimens, and treatment durations <sup>138</sup>. Another possible explanation for the disastrous results of the compounds in clinical trials could be due to PD's multifactorial and heterogeneous nature, which will reduce any single therapeutic agent less beneficial than it seemed in simpler pre-clinical models. An interesting viewpoint would be not only to fine-tune the chemical-biology interactions of these compounds to eliminate

the undesirable effects, but also to identify new bioactive compounds against mitochondrial oxidative damage

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Facing the pitfalls of clinical trials, new chemical compounds directed to mitochondria must be developed. Hydroxycinnamic (HCAs) and hydroxybenzoic (HBA) acids are the main subgroup of phenolic acids present in plants. HCAs are potent antioxidant compounds primarily due to several physiological roles in biological systems<sup>93</sup>.

Their antioxidant-related mechanism of action is suggested to be through its scavenger activity, related to their ability to donate hydrogen or electrons and the stability of the resulting phenoxyl radicals. However, other mechanisms of action have been also proposed, including inhibition of enzymes generating ROS and reactive nitrogen species (RNS), gene expression modulation through the antioxidant responsive element (ARE)/Nrf2 pathway, and metal chelators, such as iron and copper<sup>139,140</sup>. New mitochondriotropic antioxidants based on HCA and HBA core have been developed<sup>141</sup>. AntiOxBEN<sub>2</sub> is a hydroxybenzoic acid derivative linked to lipophilic TPP<sup>+</sup> through an alkyl spacer, previously demonstrated to have antioxidant capacity. Treatment of SH-SY5Y cells with maximal non-lethal concentrations of AntiOxBEN<sub>2</sub> prevented H<sub>2</sub>O<sub>2</sub>-induced cytotoxicity<sup>142</sup>. Despite considered an antioxidant, AntiOxBEN<sub>2</sub> prevention of cell death was paralleled by slightly increased ROS levels, suggesting that this mitochondriotropic antioxidant can also act as a prooxidant, triggering adaptative responses<sup>142</sup>. AntiOxCIN<sub>4</sub> is a hydroxycinnamic acid derivative linked to lipophilic TPP<sup>+</sup> through an alkyl spacer and which was previously shown to have antioxidant properties (Figure 5)<sup>141,143</sup>. AntiOxCIN<sub>4</sub> expectedly accumulated within mitochondria in its reduced form<sup>144</sup> driven by the  $\Delta\Psi_m$  without affecting mitochondrial morphology and polarization<sup>145</sup>. Consequently, AntiOxCIN<sub>4</sub> showed potent antioxidant and iron-chelation properties and can inhibit oxidative damage in isolated liver mitochondria or hepatic cells<sup>145</sup>. AntiOxCIN<sub>4</sub> can indirectly influence an antioxidant by triggering a ROS-induced adaptive response<sup>146</sup>. After treatment with that compound, Nrf2 translocates to the nucleus inducing the transcription of enzymes related with the cellular antioxidant defense system, which act as homeostatic circuits to balance detoxification and production of ROS and consequently preventing cell death induced by ROS<sup>146</sup>. Furthermore, neuroprotective effects were also demonstrated in SH-SY5Y cells against 6-OHDA-induced oxidative damage<sup>142</sup>.

More recently, we demonstrated by using human skin fibroblasts from patients with sPD that AntiOxCIN<sub>4</sub> reverted metabolic and mitochondrial defects present in skin fibroblasts from sPD patients <sup>147</sup>. AntiOxCIN<sub>4</sub> works as mild pro-oxidant, which leads to the stimulation of the Nrf2 pathway (Figure 5) <sup>146,147</sup>, having an important role in the cellular response to oxidants. Nrf2 stimulates the total SOD activity, consequently facilitating the conversion of O<sub>2</sub><sup>•-</sup> into H<sub>2</sub>O<sub>2</sub> and inhibiting cell death induced by ROS. Downstream of SOD action, H<sub>2</sub>O<sub>2</sub> is converted into H<sub>2</sub>O by GPx in a glutathione GSH-dependent manner. AntiOxCIN<sub>4</sub> increased the NAD(P)H dehydrogenase (quinone) 1 (NQO1) gene expression, which is a target of NFR2. Since those fibroblasts from sPD patients have low ATP levels after treatment with AntiOxCIN<sub>4</sub>, AMPK- $\alpha$  induces cell cycle arrest in S phase, in a p53-dependent manner, in order to restore mitochondrial function, avoiding the perpetuation of cells with defective mitochondria (Figure 5). A general enhancement in mitochondrial fitness in fibroblasts from sPD patients is demonstrated by increased maximal respiration, decreased mitochondrial swelling, and increased cellular metabolic activity <sup>147</sup>. These and other findings highlight the successful development of new mitochondria-directed antioxidants based on dietary scaffolds and demonstrate their application as first-class candidates with therapeutic potential for mitochondrial oxidative stress-related diseases, such as PD.

#### **4. Conclusions**

Oxidative stress increased and mitochondrial dysfunction have been largely implicated in PD pathogenesis <sup>118</sup>. However, it is not established whether mitochondrial dysfunction is a cause, a consequence or part of a self-sustaining vicious cycle of cellular damage <sup>129</sup>.

PD-associated symptoms usually develop slowly over time and that prodromal phase may start as early 5-10 years before diagnosis <sup>148</sup>. With this in mind, this earlier phase may be an ideal time point for therapeutic interventions, in which dietary mitochondriotropic antioxidants have a huge advantage and could be a justification for the lack of success of several candidate compounds used in pre-clinical studies. The future of therapeutic strategies for PD based on mitochondriotropic antioxidant will depend on the capacity to develop better and more efficient strategies that target mitochondria with bioactive molecules or using multifunctional compounds <sup>118</sup> and consider the multifactorial contribution of PD in pre-clinical assays.

Moreover, it is essential to use sPD models in pre-clinical studies, since the etiology of sPD, the most common PD form, is not completely understood. This limitation results not only from the low number of experimental models of sPD used but also in the experimental difficulty in obtaining appropriate human tissues to investigate the pathogenesis of PD. However, a better consider of sPD models in pre-clinical studies may anticipate promising results when undergoing clinical trials.

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## 6. Conflict of interest

PJO and FB are co-founders of the start-up MitoTAG. The funding agencies had no role in the decision to publish the manuscript.

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**Table 1: List of mutated genes involved in familial PD**

<b>Gene</b>	<b>Locus name</b>	<b>Protein Product</b>	<b>Chromossome localization</b>	<b>Type of mutation</b>	<b>Mode of inheritance</b>
<i>SNCA</i>	PARK1	Alpha-synuclein	4q21.3–22	Missense, Point	Autosomal dominant
<i>LRRK2</i>	PARK8	Leucine-rich repeat kinase 2	12q12	Missense	Autosomal dominant
<i>Parkin</i>	PARK2	Parkin	6q25.2–q27	Missense, Frameshift, Splice site, Point, Nonsense	Autosomal recessive
<i>PINK1</i>	PARK6	PTEN-induced putative kinase 1	1p36.12	Missense, Frameshift, Splice site, Point, Truncating	Autosomal recessive
<i>DJ-1</i>	PARK7	Protein DJ-1	1p36.23	Point, Missense, Frameshift, Exon deletion and Splice site	Autosomal recessive
<i>ATP13A2</i>	PARK9	ATPase 13A2	1p36	Frameshift	Autosomal recessive
<i>PLA2G6</i>	PARK14	Phospholipase A2 Group VI	22q13.1	Missense	Autosomal recessive
<i>FBXO7</i>	PARK15	F-Box protein 7	22q12-q13	Missense, splice site	Autosomal recessive
<i>GIGYF2</i>	PARK11	GRB10 interacting GYF protein 2	2q36-37	Missense	Autosomal dominant
<i>UCH-L1</i>	PARK5	Ubiquitin C-Terminal Hydrolase L1	4p14	Missense	Autosomal dominant

## Figure Legends

**Figure 1: Mechanisms suggested contributing to oxidative stress in Parkinson's Disease pathophysiology.** Mechanisms suggested contributing to oxidative stress in Parkinson's Disease pathophysiology. Mitochondrial dysfunction, alterations in dopamine metabolism, quality control mechanism impairment, neuroinflammation, and protein misfolding might underlie the pathogenesis of Parkinson's Disease. Oxidative stress plays an irrefutable role in a progressive and complex cascade of neurodegeneration culminating in dopaminergic neurodegeneration. These different pathways and their alterations, resulting either from genetic mutations or environmental factors contribute to disrupted redox balance in cells. All these cellular mechanisms that threaten dopaminergic cell's function are identical, but not linked in an orderly cascade of cause and effect.

**Figure 2: The multifaceted role of ROS.** Under normal conditions, ROS can act as intracellular messengers activating protective stress-response pathways, e.g. through up-regulating antioxidant defense system. These protective mechanisms seem to be incapable of avoiding effectively the gradual accumulation of damage independent of ROS. With aging, the generation of ROS can overwhelm antioxidant systems and their toxicity will contribute in causing the very damage by stress pathways dependent of ROS. This triggers a toxic run-away process that might originate the basis of ROS involvement in Parkinson's Disease.

**Figure 3: Schematic representation of mitochondrial involvement in Parkinson's Disease pathogenesis.** Multifaceted links between changes in mitochondrial function in PD are represented. These mitochondrial alterations are related with the process of aging. The ubiquitin-proteasome system (UPS) dysfunctional is attributed to Lewy Bodies (LBs) pathology.

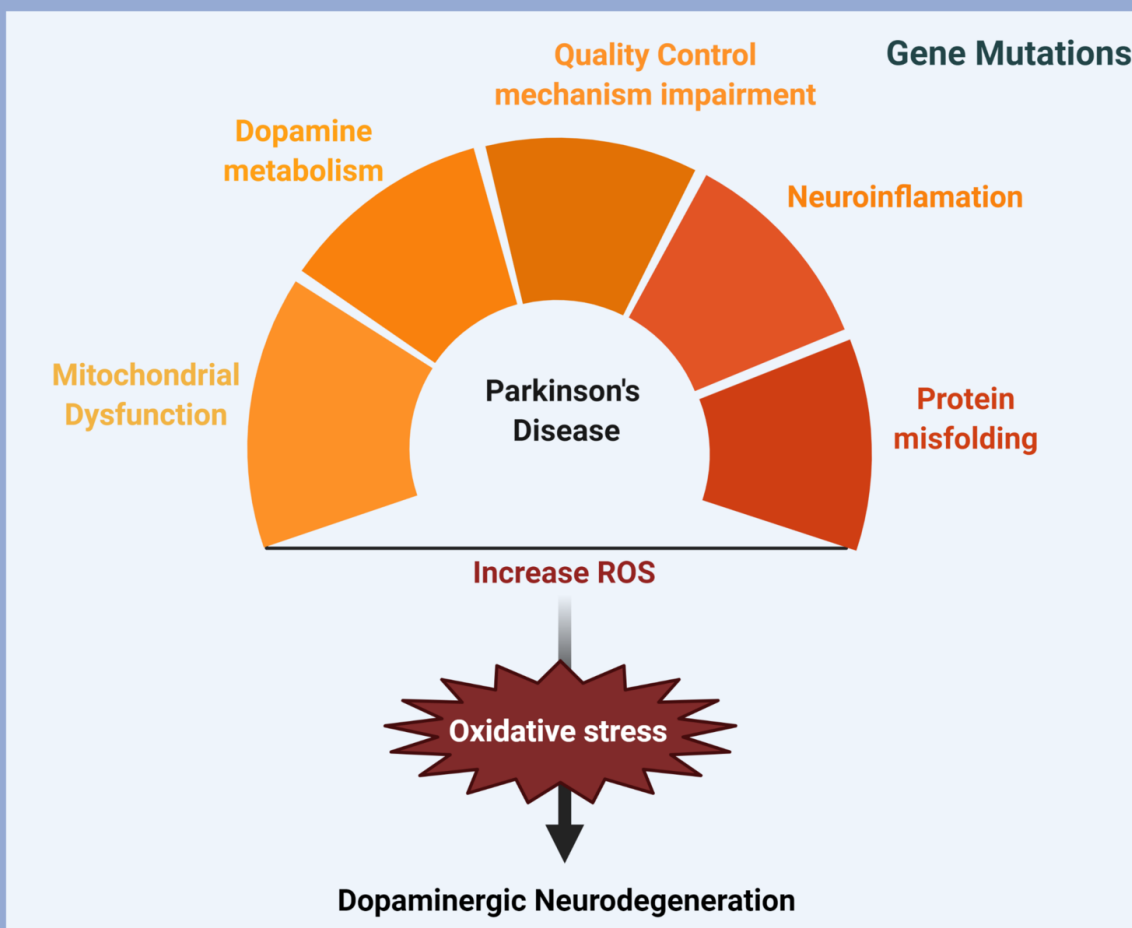
**Figure 4: Main pathways for the formation of reactive species in Parkinson's Disease pathogenesis.** Oxidative distress can result from increased dopamine levels and its metabolites, mitochondrial dysfunction, and alterations in the antioxidant defense system. Neurotoxicity induced by dopamine is due to ROS produced during dopamine metabolism by monoamine oxidase (MAO) or due to dopamine auto-oxidation. Abbreviations: DA-quinone: dopamine-quinone; DOPAC: dihydroxyphenylacetic acid; ETC: electron transport chain; GPx: glutathione peroxidase; GR: glutathione reductase; GSH: reduced glutathione; GSSG: oxidized glutathione; L-DOPA: levodopa; MAO: monoamine oxidase; NO: nitric oxide; NOX: NADPH oxidase; SOD: superoxide dismutase; PPP: pentose phosphate pathway; Trx: thioredoxin; TrxS<sub>2</sub>: oxidized thioredoxin; Trx(SH)<sub>2</sub>: reduced thioredoxin; XO: xanthine oxidase

**Figure 5: Effects of mitochondria-directed compounds in PD.** The scheme represents the main molecular targets of mitochondria-directed antioxidants tested in PD models, as discussed in the main text.

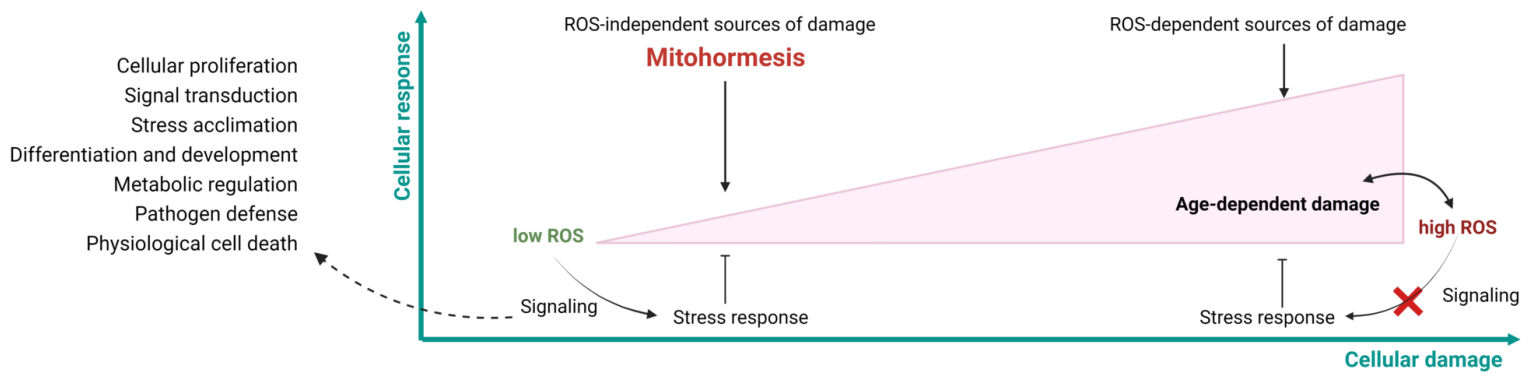
Abbreviations: DOPAC - 3,4-dihydroxyphenylacetic acid; GPx - glutathione peroxidase; GSH – reduced glutathione; GSSG – oxidized glutathione; IM – inner membrane; IMS – intermembrane space; MAO – monoamine oxidase; mtDNA – mitochondrial deoxyribonucleic acid; NQO1 – NAD(P)H quinone oxidoreductase; Nrf2 - nuclear factor erythroid 2-related factor 2; OM – outer membrane; PGC1- $\alpha$  - peroxisome proliferator-activated receptor gamma coactivator 1 alpha; SOD - superoxide dismutase

Accepted Article

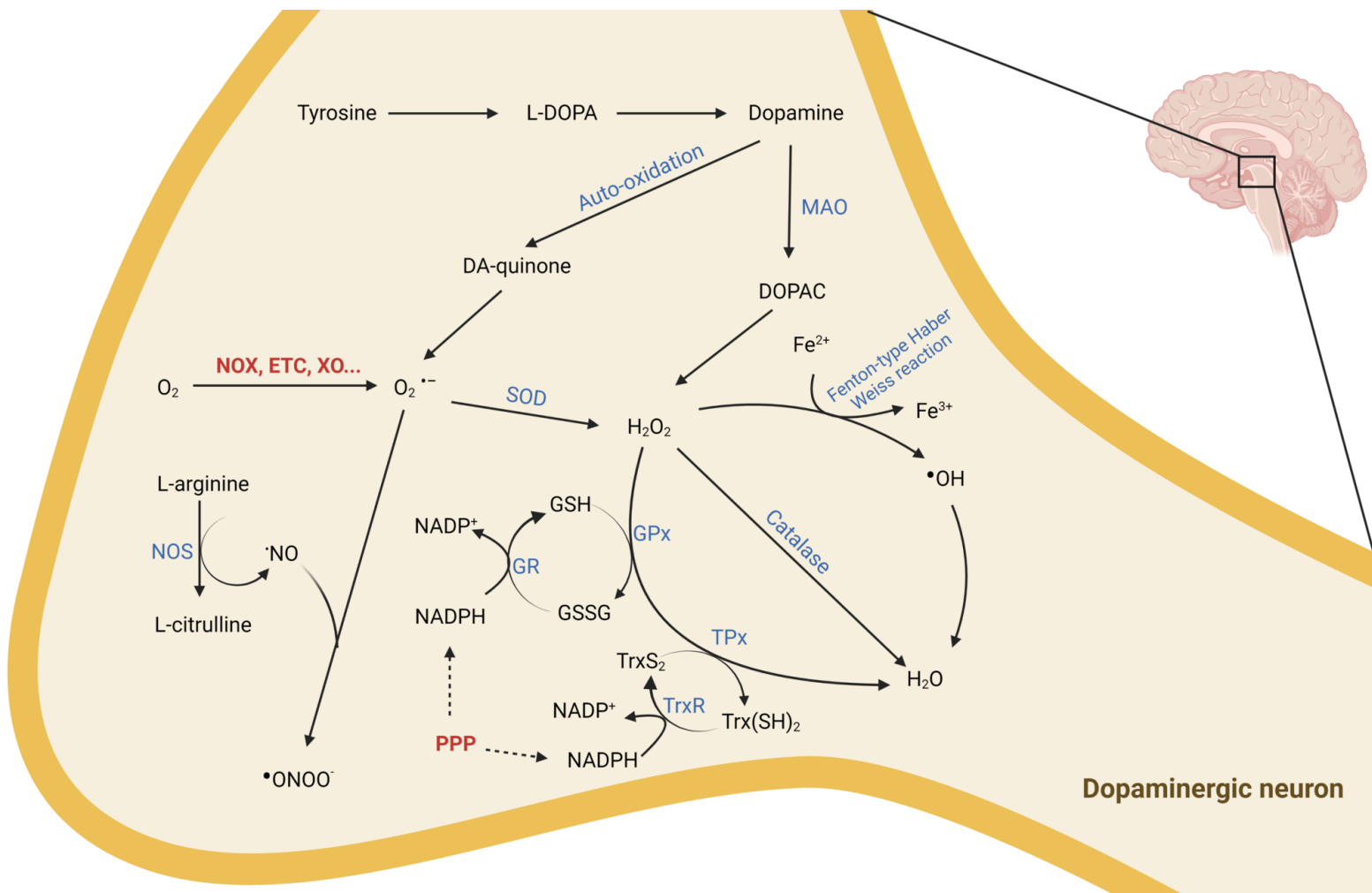
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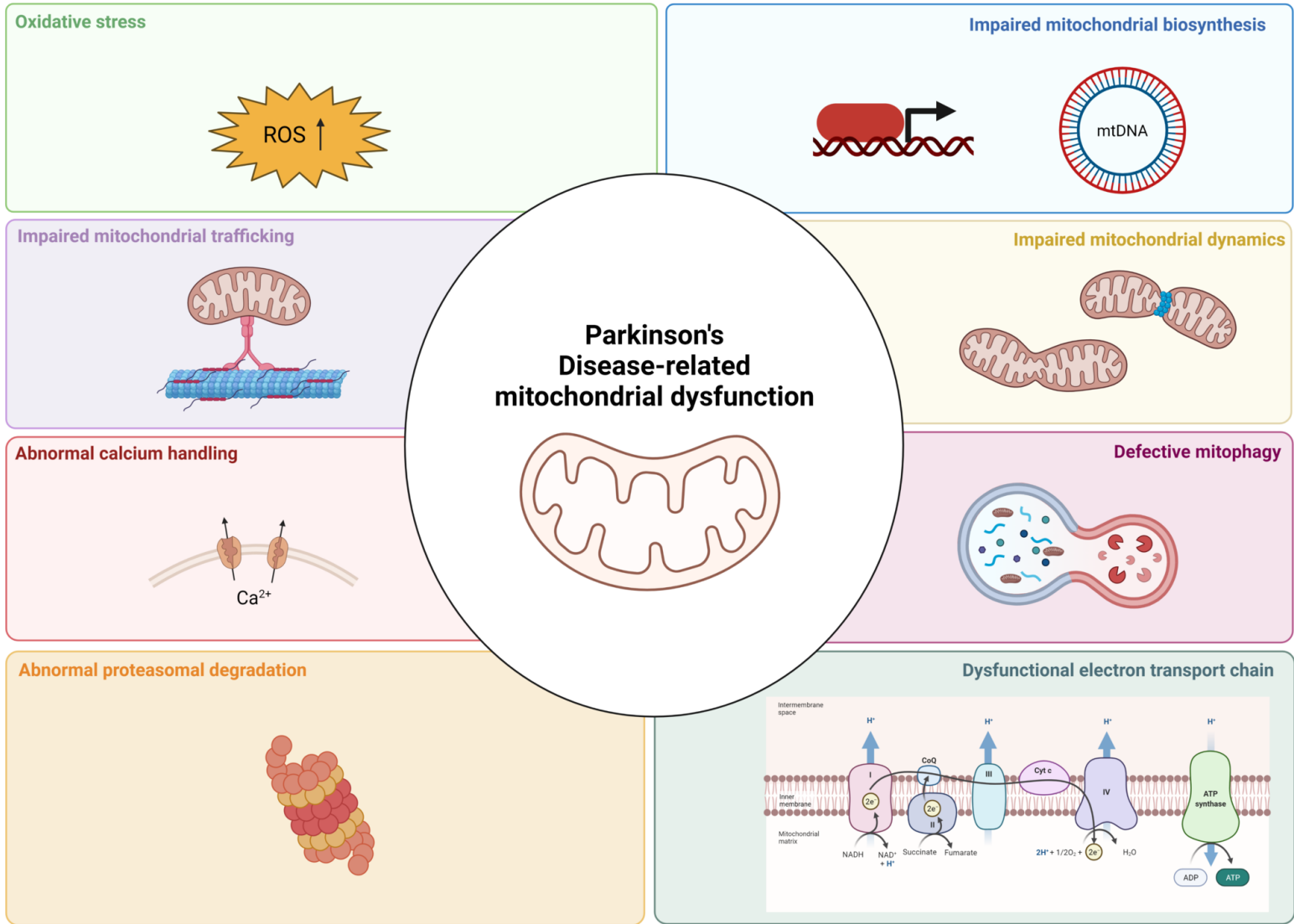
ECl\_13820\_Figure 1.png



ECI\_13820\_Figure 2.png

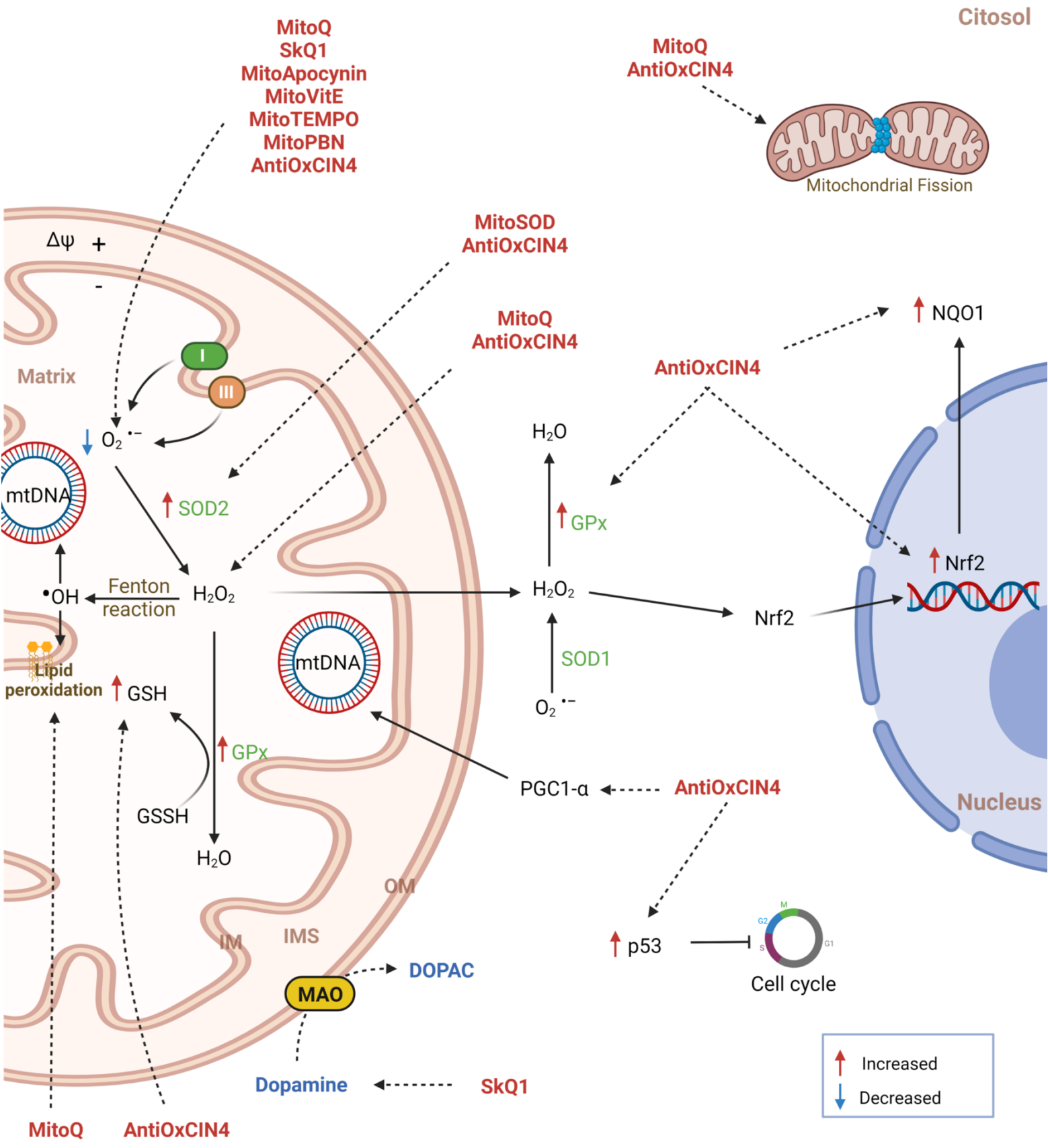


ECl\_13820\_Figure 3.png



ECI\_13820\_Figure 4.png





ECl\_13820\_Figure 5.png