| 1 | Pyrroloquinoline Quinone (PQQ) Influences Intracellular Alpha-Synuclein |
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| 2 | Aggregates |
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20 Abstract

21 Parkinson's disease (PD) is an irreversible neurodegenerative disorder clinically manifesting in 22 uncontrolled motor symptoms. There are two primary hallmark features of Parkinson's disease - an 23 irreversible loss of dopaminergic neurons of the substantia nigra pars compacta and formation of 24 intracellular insoluble aggregates called Lewy bodies mostly composed of alpha-synuclein. Using a 25 clinical improvements-first approach, we identified several clinical trials involving consumption of a 26 specific diet or nutritional supplementation improved motor and non-motor functions. Here, we aimed 27 to investigate if and how pyrroloquinoline quinone (PQQ) compound disrupts pre-formed alpha-28 synuclein deposits using SH-SY5Y cells, widely used Parkinson's Disease cellular model. 29 SH-SY5Y neuroblastoma cells, incubated in presence of potassium chloride (KCl) to induce alpha-30 synuclein protein aggregation, were treated with PQQ for up to 48 hours. Resulting aggregates were

examined and quantified using confocal microscopy. Overall, nutritional compound PQQ reduced the
 average number and overall size of intracellular cytoplasmic alpha-synuclein aggregates in a PD cellular
 model.

34

35 Keywords

36 alpha-synuclein, α-synuclein, aggregation, Lewy Body, nutrition, nutritional supplements, Parkinson's

37 disease, Pyrroloquinoline quinone; PQQ

38 Background

Parkinson's disease is a physically restricting, incurable, and progressive neurodegenerative disease.
Patients with Parkinson's disease suffer from increasing and irreversible loss of dopamine due to
dopaminergic neuron cell death leading to loss of motor function, manifesting as resting tremors, muscle
rigidity and bradykinesia (1). As Parkinson's disease worsens, everyday activities become progressively
compromised and increasing care level is needed.

44 The aggregation of alpha-synuclein has been identified in both familial and sporadic forms of 45 Parkinson's disease and is the main protein found in Lewy body (LB) pathology (2). Molecular 46 mechanisms of alpha-synuclein toxicity in the cells include an increase in oxidative stress (2) and 47 damage to the cell membrane structure through the formation of fibrils and pore-like structures (3), 48 leading to disturbance in cellular homeostasis and increase in cellular stress. Recent reviews have 49 highlighted that specific cellular pathways may trigger changes in cellular homeostasis leading to 50 dopaminergic cell death, including cellular oxidative damage, mitochondrial dysfunction, inflammation, 51 the breakdown of cellular mechanisms such as an impairment of protein degradation via chaperone-52 mediated autophagy and the aggregation of the alpha-synuclein protein (4-6). Overall, there appears to 53 be an interplay of alpha-synuclein misfolding and/or aggregation and the cellular pathways leading to 54 the death of dopaminergic neurons. Therefore, identifying a compound that targets and/or fragments the 55 aggregation may be key in future treatment strategies for Parkinson's disease.

56 The gold standard treatment for Parkinson's disease is pharmacological and based on dopamine 57 replacement therapy using a dopamine precursor agent such as L-Dopa. Unlike dopamine, L-Dopa can 58 cross the blood-brain barrier and metabolise to dopamine in the central nervous system and the 59 peripheral tissues. L-Dopa is often administrated with peripheral decarboxylase inhibitors such as 60 carbidopa to prevent L-Dopa's peripheral conversion to dopamine, allowing more L-Dopa to cross over 61 the blood-brain barrier for dopamine conversion (7). However, pharmacological treatments carry 62 significant side effects, risk and diminished response rates as the disease progresses (8) leaving limited 63 treatment options available for Parkinson's disease patients.

64 Currently, there are limited targeted dietary and/or nutritional interventions integrated into clinical 65 practice to manage Parkinson's disease symptoms in patients. Dietary interventions focus on managing 66 the symptoms of comorbidities such as constipation, weight loss and malnourishment (9). However, 67 several clinical trials and in vitro studies identified specific diets and nutrient compounds may assist in 68 improving both motor and non-motor functions. For instance, ketogenic diet trials have shown an 69 improvement in both motor and non-motor symptoms (10,11), supplementation with potent intracellular 70 antioxidants CoQ10 and glutathione, often depleted in Parkinson's disease patients, showed minor 71 improvement in motor symptoms (12-14) and food containing certain nutrient compounds such as 72 carotenoids, vitamin E, sulforaphane, Omega-3, resveratrol, epigallocatechin-3-gallate and caffeine 73 might offer important neuroprotective properties, including the potential to alter the underlying 74 pathological cellular and molecular changes identified in Parkinson's disease (15). Nutrients including 75 selenium, vitamin A, C and E may serve as potent antioxidants protecting the dopaminergic cell from 76 oxidative damage to preserve dopamine producing neurons (8).

Interestingly, in a dietary study where participants were given broad beans to consume as part of their daily diet, improvements in motor function and increased levels of endogenous levodopa were observed similar to levels achieved with L-Dopa therapy (16). On further investigation, it was also found that broad beans contain high levels of a nutritional compound Pyrroloquinoline Quinone (PQQ) (17), that has shown to modify alpha-synuclein aggregation *in vitro* (18).

82 PQQ is a water-soluble organic molecule found in varying quantities in vegetables, fruits, and beverages 83 such as tea. POQ is a vitamin-like compound with combined chemical properties similar to ascorbic 84 acid, vitamin B_2 and vitamin B_6 , but with a nutritional requirement similar to folate and biotin (19). In 85 animal studies, PQQ has been demonstrated to be highly absorbable, predominantly in the lower 86 intestine, and non-toxic in both animals and humans, indicating low or no side effects (20). Recent 87 studies have demonstrated that PQQ permeability to the blood-brain barrier is enhanced using 88 esterification processes (21). Importantly for molecular events leading to Parkinson's disease, PQQ was 89 shown to have an anti-aggregation effect on the alpha-synuclein protein *in vitro* (18) and inhibits fibril

90 formation in both a full length and C-terminal deleted mutant alpha-synuclein *in vitro* (22). However, 91 none of the other alpha-synuclein studies had been conducted in the PD-relevant cellular context. 92 Our study demonstrated that the addition of PQQ to pre-formed alpha-synuclein aggregates can reduce 93 the number and size of these aggregates in a Parkinson's disease SH-SY5Y cellular model. This study 94 is the first to show PQQ aggregation-modifying properties in a Parkinson's disease-specific cellular 95 model. It confirms previous in vitro and clinical trial reports establishing this nutritional compound and 96 food containing PQQ as an important player in the regulation of Parkinson's disease symptoms and 97 possible novel treatment approach to Parkinson's disease. 98 Methods

99 Materials

100 SH-SY5Y neuroblastoma cells (94030304-1VL; Sigma). Hams F-12 nutrient mixture (N48880), Eagles 101 Minimum Essential Medium (EMEM; M2279), Glutamine (GT513), Non-essential amino acids 102 (M7145), Pyrroloquinoline Quinone (D7783-1MG) were purchased from Sigma-Aldrich. Monoclonal 103 antibody to alpha-synuclein (ab27766) was purchased from Abcam, secondary goat anti-mouse Alexa 104 Fluor 488 IgG antibody (A-11001) was from ThermoFisher Scientific and Foetal Bovine Serum (FBS, 105 SFBSF8) from Bovogen. Pyrroloquinoline Quinone (PQQ) was dissolved in 1M sodium hydroxide 106 (NaOH) solution to prepare a stock concentration of 50 mM, and 50µM PQQ was used as the final 107 concentration.

108 Methods

109 Cell Culture

- 110 The SH-SY5Y neuroblastoma cells were maintained in the following cell growth media mix EMEM,
- 111 Ham's F12 nutrient mixture, 1% glutamine, 1% non-essential amino acids and 15% FBS. Cells were
- 112 maintained in a humidified incubator at 37°C supplied with 5% CO₂.

113 Experimental Procedures

114 Cell seeding: 24 hours (h) prior to the experiment, cells were cultured on glass coverslips at 20,000 cells

- per well of a 12-well multi-well dish. Each experiment contained the following groups in triplicates: No
- treatment, 50mM KCl treatment, 50µM PQQ treatment, 50mM KCl followed by 50µM PQQ treatment.
- 117 PQQ and KCl treatment: Cells were treated with 50µM PQQ for 24 h, followed by treatment with
- 118 50mM KCl to the appropriate samples for 60 minutes (min). After 60 min, the PQQ/KCl containing
- 119 media was decanted and replaced with growth media and incubated further for 24 or 48 h.

Immunofluorescence Assay: At the 24- and 48-h time points, cells were fixed using 4% paraformaldehyde (prepared in 1 X PBS; PFA/PBS) and permeabilised with 0.1% Triton X-100. Fixed coverslips were stained with primary mouse anti-alpha-synuclein antibody (optimised at 1:50 dilution) and secondary goat anti-mouse Alexa 488 IgG antibody. Hoechst DNA stain diluted in 1X PBS was used for nuclear staining. The coverslips were mounted onto microscopy slides using Fluorescence mounting medium (Dako) and analysed using confocal microscopy.

126 **Confocal Microscopy**: Digitised fluorescent cellular images were collected using a Nikon Eclipse Ti 127 confocal laser-scanning microscope, with a Nikon 60x/1.40 oil immersion objective. Images were 128 captured at an optical thickness of 0.68 µm and an optical resolution of 0.12 µm. An average of 10 - 15129 images for each treatment group from two independent experiments were captured, resulting in 130 approximately 250 images.

131 **Image Analysis:** Image analysis was performed in ImageJ v1.51 - images were opened as a hyperstack, 132 split channel colour composite, 16-bit image. Each colour composite image was separated into two 133 multi-level channels (alpha-synuclein and nucleus), and both channels were then merged into one 134 composite image while preserving the original multi-level composite channels. To determine the 135 aggregates' approximate size, a scale was set in ImageJ at 1 pixel to 0.21 µm, and an outline of the 136 aggregates was traced. The ImageJ pre-set function was used to calculate an approximate perimeter size 137 of the aggregate. Aggregate size data and the total number of aggregates was captured for both the 24 138 and 48-h per treatment group and entered into Prism GraphPad v9 to produce a scatter plot and determine 139 differences between groups.

140 Images were discounted if there was either insufficient immunofluorescence staining of the alpha-141 synuclein or the nucleus, if cells had not shown sufficient growth and proliferation, or if cells were out 142 of focus. Cells were discounted from analyses if the DNA staining did not define a distinct nucleus or

143 there were morphological abnormalities, such as condensing or blebbing of the cell nucleus.

144 Results

145 Confocal microscopy revealed two general patterns of alpha-synuclein localisation at both time points
146 investigated - generalised diffusion surrounding the cell nucleus and aggregates predominantly located
147 near the nucleus of the cell (arrows, Figure 1A and 1B).

148 To objectively determine the number of alpha-synuclein aggregates per cell, images were analysed using 149 ImageJ by counting the total number of aggregates and normalising this count to the total number of 150 cells in the image for each sample or by measuring the size of the individual aggregate (Figure 2A and 151 2B, respectively). The highest percentage of aggregates were observed in 50mM KCl only treated 152 samples at both time points, 25.71% and 6.48% compared to 1.47% and 0.66% in no treatment samples 153 at 24h and 48h, respectively. A three-fold decrease in the number of aggregates was observed in cells 154 treated with PQQ only at both time points, 8.43% and 1.65% at 24h and 48h, respectively. This decrease 155 was two-fold in cells treated with PQQ and stimulated with 50mM KCl compared to control, 11.40% 156 and 3.49% at 24h and 48h, respectively (Figure 2).

157 The same images were used to determine the average size of all aggregates (Figure 2). A measurable 158 difference in mean aggregate size between the 24 and 48h control and experimental samples were 159 observed. Across all samples, 24h samples contained on average smaller aggregates (Figure 2A). The 160 mean aggregate size in the presence of PQQ only or PQQ + KCl were 1.84 µm and 2.51 µm, respectively, 161 compared to 3.05 µm in the presence of KCl only at 24h. The same trend was evident at 48h with a 162 decrease in mean aggregate size in the presence of POQ (1.94 µm; POQ only) and 2.85 µm (POQ + 163 KCl), compared to KCl only (3.88 µm). In comparison to their controls, in samples incubated with 164 50mM KCl, the mean aggregate size in the presence of PQQ + KCl decreased to 17.7% and 26.5% at 165 24h and 48h, respectively (Figure 2A and 2B).

Overall, the cells treated with PQQ without or with KCl consistently demonstrated an overall reduction
in the number and mean aggregate size of alpha-synuclein at both 24- and 48-h compared to cells that
were not treated with PQQ.

169 Discussion

In the current study, we used an established Parkinson's disease cellular model, neuroblastoma SHSY5Y cell line, to demonstrate the reduction in number and size of pre-formed intracellular alphasynuclein aggregates at 24 and 48h post treatment with PQQ.

PQQ was first identified in 1979 as a redox cofactor for bacterial dehydrogenase. Although PQQ is not synthesised in humans, it is found in trace amounts in tissues due to absorption from everyday food, such as broad beans, green peppers, spinach, fermented soybeans, oolong tea, green tea, papaya and kiwi fruit (17). The nutritional requirement for PQQ from a dietary perspective is not defined, but animal studies that modified diets to omit PQQ resulted in impaired growth, immune dysfunction and fertility issues (23).

In humans, the nutritional importance of PQQ and the possible clinical applications are increasingly emerging with a primary focus in PQQ supplementation is its potential to have a pharmacological-like role in human disease. In human and animal studies, PQQ is observed to have positive physiological benefits for various diseases including dementia, neurodegenerative disorders, Parkinson's disease,

183 cardiovascular disease, chronic inflammation, insulin resistance, mood disorders and accelerated aging 184 (24). Animal research models have identified the benefits of PQQ treatments to include overall 185 improvement and even reversal of cognitive impairment, stimulation of nerve growth factors for cell 186 proliferation, protecting cells from oxidative stress and neurotoxicity caused by cerebral ischemia, 187 reduction in neurotoxin-induced injury to the brain (25-27) and protecting nerve cells from beta-amyloid 188 damage in Alzheimer's disease (28). Molecular mechanisms underpinning these observed phenotypes 189 include an increase in mitochondrial energy production, promotion of mitochondrial biogenesis, and 190 AMP kinase activity which serve as a master switch for energy production (29).

191 Human clinical studies using PQQ as a supplement showed an increased presence of the compound in 192 blood and reduced levels of inflammatory markers C-reactive protein (CRP) and interleukin 6 (IL-6) 193 indicating enhanced mitochondria-related function (30). Further studies support the fact that PQQ 194 enhances mitochondrial function and improves cognitive performance, increases cerebral blood flow 195 and oxygen use in the prefrontal cortex of the brain, and improves indicators for stress, fatigue and sleep 196 (19). Interestingly, human trials investigating the consumption of food high in PQQ levels identified 197 improvement in motor skills in Parkinson's disease patients (16). While these studies show PQQ 198 absorption into the blood and improvements in the measured outcomes, they do not address specific 199 tissue, cellular or molecular pathways influenced by the PQQ or PQQ-rich food.

200 The normal biological function of the alpha-synuclein protein is poorly understood. However, in 201 Parkinson's disease, LB pathology indicates that abnormal aggregation of the protein is a crucial factor 202 in progressive and irreversible loss of dopaminergic cells in both familial and sporadic forms of 203 Parkinson's disease (31). Previous in vitro studies have shown PQQ has an anti-aggregation effect on 204 the alpha-synuclein protein in both truncated and full-length forms by reducing the protein's ability to 205 form fibrils and aggregate in addition to reducing its overall toxic influence. However, these studies 206 were not conducted in a Parkinson's disease relevant cellular context (18,32). In line with these studies, 207 our study demonstrates that the addition of PQQ to pre-formed aggregates in SH-SY5Y cells not only 208 reduced the number of the aggregates but also indicated the dispersion or reduced size of the aggregates.

209 Identifying underlying cellular and molecular mechanisms that can clear alpha-synuclein aggregates 210 will allow for greater understanding of Parkinson's disease and allow targeted research. Using in vitro 211 and in vivo studies, a model of these mechanisms has been suggested for PQQ to act as a ligand-like 212 compound for the cell surface receptors that influence key pathway(s) involved in alpha-synuclein post-213 translational modification, including ubiquitination, cross-linking, truncations, oxidation (like nitration) 214 and phosphorylation (18,32,33). Extensive research has been conducted in phosphorylated alpha-215 synuclein and its role in Parkinson's disease with some research in the context of PQQ. Increase of 216 phosphorylated Serine 129 (pSyn-129) alpha-synuclein levels have been noted in Parkinson's disease 217 patients (34), leading to increased aggregation, changes in cell membrane receptor distributions (35) 218 and cytotoxicity (36). In vitro studies have indicated that PQQ-addition reduces the oligomerisation of 219 the alpha-synuclein aggregates to the levels seen with removal of the Ser129 residue (32), indicating 220 that mechanism of action for PQQ may be via interference in phosphorylation of the alpha-synuclein. 221 While the exact cellular pathway is still unclear, it is clear that PQQ can act on the aggregate itself 222 (18,32) and when enclosed within a cellular context as shown by our study.

223 To date, Parkinson's disease research has predominately focused on the brain and risks factors that may 224 induce defects in the cellular systems triggering the gradual degeneration and death of dopamine neurons 225 including the potential causal role of alpha-synuclein protein aggregations in cell death. Interestingly, 226 recent evidence points to the gut-brain axis as an essential pathway in Parkinson's disease manifestation, 227 including consistent constipation and gastrointestinal issues that have recently been identified as one of 228 the first symptoms to appear before a patient is diagnosed with Parkinson's disease (37). Further, recent 229 insights indicate that alpha-synuclein protein may have prion-like characteristics and propagate between 230 cells infecting different regions of the brain (38) as well as evidence of alpha-synuclein aggregation in 231 enteroendocrine cells (EECs) located in the enteric nervous system of the gastrointestinal tract before 232 they are found in the brain. This suggests that the alpha-synuclein protein's pathological origin may stem 233 from the gastrointestinal tract and spread to the central nervous system via a cell to cell propagation 234 using the vagus nerve (39). Considering the high levels of PQQ in particular food and its use as a 235 supplement, one unexplored avenue of Parkinson's disease research is the influence of food on the gut-

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brain axis as linked to health and disease. While different models would need to be developed (e.g. enteric neuron cellular model, induction of alpha-synuclein aggregates in the gut cell of an animal model) to further explore the possibility of disease origin it is important to consider in light of food/diet/supplement as preventative and/or therapeutic intervention.

240 Extending further research into the PQQ mode of action across multiple models is essential to uncover 241 how PQQ impacts the interplay of the development of LBs and the death of dopaminergic neurons. 242 Understanding further the mechanism of action of PQQ on a cellular and molecular level strengthens 243 the clinical trials of diets or supplementation high in this compound, and our findings provide a basis 244 for further cellular, animal and clinical studies as is specifically shows modifications of intracellular 245 alpha-synuclein aggregates. Finally, targeting of the LB formation or therapeutic intervention of formed 246 aggregates using a naturally derived product found in a variety of food while undergoing L-Dopa 247 substitutions may be considered an integrated, evidence-based combinatorial therapeutic strategy for 248 both sporadic and familial Parkinson's disease patients.

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- 257 Data availability: The data that support the findings of this study are available from the
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- 259
- 260
- 261

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- 369 Figure 1: Images of alpha-synuclein aggregates show morphology change in the presence of
- 370 PQQ.

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| 372 | SH-SY5Y cells were incubated in the presence or absence of 50mM PQQ for 1 hour and then for a |
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| 373 | further hour with 50mM KCl treatment as appropriate for the sample to induce alpha-synuclein |
| 374 | aggregation. After KCl treatment the cells were washed into full cell media containing 50 mM PQQ |
| 375 | ("PQQ treatment "and "PQQ & KCl treatment") or full cell media only ("no treatment" and "KCl |
| 376 | treatment") for 24 h (A) or 48 h (B). The cells were fixed and then stained using anti-alpha-synuclein |
| 377 | antibody and secondary goat anti-mouse Alexa Fluor 488 IgG antibody. Hoechst 3342 diluted in 1X |
| 378 | PBS was used for nuclear staining (shown in red). The images were taken with a Nikon Ti Eclipse |
| 379 | confocal laser-scanning microscope and NIS Elements AR software. Scale bar: 16 µm. |





Images for both 24 h (A) and 48 h (B) were analysed for inclusion based on sufficient immunofluorescence staining of the alpha-synuclein protein or nucleus and sufficient visibility of cells under confocal microscopy. On identification of images for inclusion, total cells were counted only if DNA staining showed a distinct nucleus, and there was no condensing or blebbing of the cell nucleus. The total number of identified alpha-synuclein aggregates were counted for each sample and divided by the total number of cells in that sample for each cell treatment group per time point. All the identified aggregates of alpha-synuclein proteins were measured in ImageJ v1.51 by tracing around the aggregate's perimeter with a set scale of 1 pixel = $0.21 \,\mu$ m. The mean size of aggregates for each cell treatment group was calculated by totalling the mean size of each aggregate and then dividing by the total number of aggregates. Plots were obtained using Prism GraphPad v9. The line represents the median size of aggregates.