

Microtubule depolymerization potentiates alpha-synuclein oligomerization

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Sandra M. Cardoso, Faculdade de Medicina e Centro de Neurociências e Biologia Celular da Universidade de Coimbra, 3004 Coimbra, Portugal. e-mail: smacardoso@yahoo.com Parkinson's disease (PD) is associated with perturbed mitochondria function and alpha-synuclein fibrilization. We evaluated potential mechanistic links between mitochondrial dysfunction and alpha-synuclein aggregation. We studied a PD cytoplasmic hybrid (cybrid) cell line in which platelet mitochondria from a PD subject were transferred to NT2 neuronal cells previously depleted of endogenous mitochondrial DNA. Compared to a control cybrid cell line, the PD line showed reduced ATP levels, an increased free/polymerized tubulin ratio, and alpha-synuclein oligomer accumulation. Taxol (which stabilizes microtubules) normalized the PD tubulin ratio and reduced alpha-synuclein oligomerization. A nexus exists between mitochondrial function, cytoskeleton homeostasis, and alpha-synuclein oligomerization. In our model, mitochondrial dysfunction triggers an increased free tubulin, which destabilizes the microtubular network and promotes alpha-synuclein oligomerization.

Keywords: Parkinson disease, cybrids, ATP, alpha-synuclein, tubulin, mitochondria

INTRODUCTION

Sporadic Parkinson's disease (PD) pathology is characterized by substantia nigra pars compacta neuronal loss, systemic dysfunction of the mitochondrial enzyme complex I, and intraneuronal alphasynuclein (a-syn) aggregation (Cardoso et al., 2005). While a-syn overexpression and mutation cause familial PD, the mechanisms that mediate this are unclear (Singleton et al., 2003; Hope et al., 2004). This is not surprising, as the normal function of a-syn is itself poorly characterized.

Recent data indicate a-syn interacts with microtubules to affect membrane stability and neuronal plasticity (Lee et al., 2006). Microtubules are highly dynamic tubulin polymers that mediate organelle transport, cell motility, division, and morphology (Morris and Hollenbeck, 1995). Conditions that favor incorporation of tubulin into microtubules reduce free/polymerized tubulin ratios, while conditions that retard microtubule formation increase this ratio.

The complex I inhibitors rotenone and 1-methyl-4-phenylpyridinium (MPP⁺) alter microtubule dynamics and increase free tubulin/polymerized tubulin ratios (Cappeleti et al., 2005; Ren and Feng, 2007). Tubulin, interestingly, induces a-syn fibrillization in yeast, rat brain, and human brain (Alim et al., 2002; Kim et al., 2007).

Because complex I dysfunction occurs in PD and may contribute to a-syn aggregation (Parker et al., 1989; Schapira et al., 1989; Lee, 2003) we hypothesized a nexus between complex I dysfunction, ATP depletion, microtubule disassembly, and a-syn oligomerization might exist. We tested this hypothesis using cytoplasmic hybrid (cybrid) cells prepared via the transfer of PD subject platelet mitochondria to mitochondrial DNA (mtDNA) depleted (ρ 0) human NT2 teratocarcinoma cells. PD cybrids have been shown to model complex I dysfunction and its consequences, including increased oxidative stress, perturbed calcium homeostasis, and reduced ATP (Cassarino et al., 1997; Sheehan et al., 1997; Esteves et al., 2008, 2009).

MATERIALS AND METHODS

CHEMICALS AND CELL MEDIA

Paclitaxel (taxol), was obtained from Sigma (St. Louis, MO, USA). Optimem medium was obtained from Gibco-Invitrogen. Nondialyzed and dialyzed fetal bovine serum was obtained from Gibco-Invitrogen. NT2 ρ 0 cell growth medium consisted of Optimem supplemented with 10% non-dialyzed fetal bovine serum, 200 µg/ ml sodium pyruvate, 150 µg/ml uridine, and 100 IU/ml penicillin and 50 µg/ml streptomycin. NT2 cybrid selection medium consisted of Optimem supplemented with 10% dialyzed fetal bovine serum and penicillin–streptomycin. Cybrid expansion medium consisted of Optimem supplemented with 10% non-dialyzed fetal bovine serum and penicillin–streptomycin. Taxol medium consisted of cybrid expansion medium with 5 nM taxol, and was prepared using a DMSO-taxol stock solution. For all conditions tested, control experiments were performed using unsupplemented cybrid expansion medium.

HUMAN SUBJECTS

Subject participation was approved through the Institutional Review Board of the University Hospital of Coimbra. The two sporadic PD subjects, without any nuclear DNA mutation known to be relevant to PD, meet Gelb et al. (1999) diagnostic criteria and two healthy, age-matched control subject provided 10 ml blood samples following written informed consent. Blood was drawn directly into tubes containing acid-citrate-dextrose.

CREATION OF CYBRID CELL LINES AND CELL CULTURE

NT2 ρ 0 cells were briefly agitated in polyethylene glycol with platelets from the human subjects (Cardoso et al., 2004). Seven days after plating the resulting mixture in T75 flasks and ρ 0 growth medium, the medium was changed to cybrid selection medium. NT2 ρ 0 cells lack intact mtDNA, do not possess a functional electron transport chain (ETC), and are auxotrophic for pyruvate and uridine (Swerdlow et al., 1997; Cardoso et al., 2004). Maintaining cells in selection medium removes ρ 0 cells that have not repopulated their mtDNA with platelet mtDNA. After selection was complete, the resultant cybrid cells were switched to cybrid expansion medium.

For taxol, experiments, cybrid cell lines were placed in media containing 5 nM taxol. Flasks were maintained in these media at 37° C, 5% CO₂ for 24 h prior to harvesting.

IMMUNOCYTOCHEMISTRY

Cybrid cell lines grown on coverslips in 12 well plates were washed twice with PBS and fixed for 30 min at room temperature using 4% paraformaldehyde. The fixed cells were washed again with PBS, permeabilized with 0.2% Triton X-100, and blocked with 3% BSA. The permeabilized cells were incubated with primary antibody (1:2,000 monoclonal anti-alpha-tubulin from Sigma; 1:400 monoclonal anti-a-syn from Santa Cruz Biotechnology, Inc., Santa Cruz, SA, USA, or 1:400 Polyclonal anti-A11 from BioSource International, Inc., Flynn Road, Camarillo, CA, USA) for 1 h and then with the appropriate secondary antibody (1:250 alexa fluor 594 and 488 from Molecular probes, Eugene, OR, USA). Cells were visualized by confocal microscopy.

IMMUNOBLOTTING

Individual cybrid cell lines were scraped in buffer containing 25 mM HEPES, 1 mM EDTA, 1 mM EGTA, 2 mM MgCl,, protease inhibitors (commercial protease inhibitor cocktail from Sigma), 0.1 M PMSF (Sigma), 0.2 M DTT (Sigma), and 1% Triton X-100. Cell suspensions were frozen three times in liquid nitrogen and centrifuged at 20,000 g for 10 min. The resulting supernatants were removed and stored at -80°C. Protein concentrations were determined by the Bradford method and equal amounts of protein (30 or 60 µg) were used for immunoblotting. For the SDS-PAGE experiments samples were resolved by electrophoresis in SDSpolyacrylamide gels and transferred to PVDF membranes. For the PAGE experiments samples were not boiled and were diluted 1:2 in a sample buffer without SDS. Non-specific binding was blocked by gently agitating the membranes in 5% non-fat milk and 0.1% Tween in TBS for 1 h at room temperature. The blots were subsequently incubated with the respective primary antibodies overnight at 4°C with gentle agitation (1:1,000 Polyclonal anti-a-syn antibody from Cell Signaling Technology, Inc., Danvers, MA, USA; 1:2,500 monoclonal anti-GAPDH antibody from Chemicon International; or 1:10,000 monoclonal anti-alpha-tubulin antibody from SIGMA).

Blots were washed with TBS containing 0.1% non fat milk and 0.1% Tween three times (each time for 15 min), and then incubated with the appropriate horseradish peroxidase-conjugated secondary antibody for 2 h at room temperature with gentle agitation.

After three washes specific bands of interest were detected by developing with an alkaline phosphatase enhanced chemical fluorescence reagent (ECF, GE Healthcare, Buckinghamshire, England). Fluorescence signals were detected using a Biorad Versa-Doc Imager, and band densities were determined using Quantity One Software.

ISOLATION OF SOLUBLE AND POLYMERIC TUBULIN

To prepare soluble and polymeric tubulin fractions, cells were very gently washed twice with a microtubule stabilizing buffer (0.1 M N-morpholinoethanesulfonic acid, pH 6.75; 1 mM MgSO₄; 2 mM EGTA; 0.1 mM EDTA; 4 M glycerol). Soluble proteins were extracted for 4–6 min at 37°C in 100 μ l of microtubule stabilizing buffer containing 0.1% Triton X-100. The remaining contents of the culture dish were scraped in 100 μ l of 25 mM Tris (pH 6.8) and 0.5% SDS. This suspension was frozen three times in liquid nitrogen (Joshi and Cleveland, 1989). Protein concentrations for each sample were quantified using the Bradford method.

DATA ANALYSIS

Each experimental endpoint for each cell line was run in duplicate, and data are expressed as the mean \pm SEM from at least three different measurements. *P*-values were calculated by one-way ANOVA, followed by a *post hoc* Dunnett's or Bonferroni's *t* test when appropriate.

RESULTS

ALPHA SYNUCLEIN OLIGOMERIZATION AND PROTEIN UBIQUINATION WERE INCREASED IN THE PD CYBRID CELLS

The PD cybrid cells and the control cybrid cells analyzed in this study were originally prepared and bioenergetically characterized as part of previously reported PD cybrid studies (Esteves et al., 2008, 2009). PD cybrid lines used in this current study show a decrease of 38% in complex I activity and of 28% in ATP levels when compared to the control cybrid lines (Esteves et al., 2008). In order to confirm the presence of protein aggregates we used antibodies (A11) that detect oligomeric proteins. All antibody staining was much stronger in the PD cell cells, suggesting it contained more oligomeric protein or proteins (Figure 1A). Moreover, we performed a PAGE western blot to detect if some of these oligomers were a-syn. We observed that the a-syn oligomer/monomer ratio was higher in the PD cybrid lines (Figures 1B,C), and the PD cybrid lines showed more protein ubiquination levels (Figure 1D). We also observed an increase in a-syn monomers levels detected by SDS-PAGE in PD cybrid lines as compared to CT cybrid cells (data not shown).

THE FREE TUBULIN/POLYMERIZED TUBULIN RATIO WAS INCREASED IN THE PD CYBRID

MT assembly requires ATP, therefore the reduction of PD cybrids ATP may interfere with the efficacy of the polymerization/depolymerization process of tubulin. For that reason we immunocytochemically evaluated MT network integrity. MT networks were less well defined in the PD cybrid cells (**Figure 2A**), and the free/polymerized tubulin ratio was also elevated (**Figures 2B,C**). In the CT cybrid cells approximately one-third of the tubulin was in the free form, and in the PD cybrid approximately one-half of the tubulin was in



the free form (**Table 1**). Because of the viscosity of the polymerized tubulin all samples were corrected with GAPDH antibody in order to eliminate loading errors (**Figures 2B,C**).

TAXOL REDUCED TUBULIN AND ALPHA SYNUCLEIN ALTERATIONS IN PD CYBRID CELLS

Taxol stabilizes MT assemblies and reduces free tubulin/polymerized tubulin ratios (Cragg and Newman, 2005). Before testing its effects in our cybrid model (which consists of undifferentiated, rapidly dividing cells) we determined 5 nM taxol did not compromise cell viability (data not shown). 5 nM taxol reduced the free tubulin/polymerized tubulin ratio in the PD cybrid lines but not in the control cybrid lines (**Figures 3A,B**). Taxol lowered a-syn oligomerization in PD cybrid cells but did not altered CT cybrid cells a-syn levels (**Figures 3C,D**). These results support the idea that MT depolymerization may potentiate the formation of oligomeric a-syn.

DISCUSSION

We recently reported that mitochondrial ROS potentiates an increase in free tubulin levels, which is followed by a significant increase of a-syn oligomers (Esteves et al., 2009). In order to further clarify the relationship between mitochondrial metabolism and a-syn conformational change we now report studies performed using two carefully characterized PD and CT cybrid cell lines. Our results corroborate and extend our previous finding that PD cybrids mitochondrial dysfunction induces a-syn oligomerization, and suggest that microtubules may be a feasible therapeutic target to PD.

The cybrid approach has been used to model human disease-specific mitochondrial dysfunction in PD. It has been described that in cybrid models such as the one used in these experiments changes in cell physiology likely arise as a consequence of and in response to mitochondrial function (Swerdlow, 2007).



Numerous cybrid studies using three different nuclear back-

grounds have shown PD cybrids have a sustained reduction of com-

plex I Vmax activity (Swerdlow et al., 1996, 1998; Gu et al., 1998;

Esteves et al., 2008). In these studies biochemical data obtained

from distinct, different cybrid cell lines were averaged and group

Table 1 | Percentage of free and polymerized tubulin.

	CT cybrids	PD cybrids
Free tubulin	36.41 ± 2.49	51.45 ± 2.86***
Polymerized tubulin	63.59±2.49***	48.55 ± 2.86*##

SDS-PAGE analysis demonstrated free and polymerized tubulin percentages differed between the PD cybrid lines and the CT cybrid lines. Results are expressed as the mean \pm SE of five independent measurements. *P < 0.05, **P < 0.01, ***P < 0.001, significantly different as compared to the CT cybrid's free tubulin percentage. *P < 0.05, **P < 0.01, significantly different as compared to the CT cybrid's polymerized tubulin percentage.

We fully characterized a-syn aggregation in two PD cybrid cell lines. We demonstrated that compared to cells from two control cybrid cell lines, the PD cybrid cells had increased a-syn oligomerization, ubiquitination, and fibrillization.

If mitochondrial dysfunction does in fact promote a-syn oligomerization, then it is necessary to elucidate the mechanisms that mediate this phenomenon. Our data support the hypothesis that free tubulin directly interacts with a-syn and promotes fibril formation (Alim et al., 2002). While our data do not specifically elucidate how tubulin-a-syn interactions may physically promote oligomerization, they do provide a framework for developing further studies to address why a-syn oligomerization occurs in PD.

Taxol's ability to reduce a-syn oligomerization in our experiments suggests free tubulin can promote a-syn oligomerization. This does not preclude the possibility that a-syn oligomerization might in turn disrupt tubulin polymerization (Chen et al., 2007). It is therefore necessary to consider whether mitochondrial dysfunction supports a positive feedback loop in which mitochondrial dysfunction increases free tubulin levels, free tubulin increases a-syn oligomerization, and a-syn oligomers in turn further elevate the free/polymerized tubulin ratio. a-Syn fibrillization requires it to transition from a 'natively unfolded' random coil to a beta-sheet structure. The beta sheet form then aggregates into oligomers, protofibrils, and eventually amyloid fibrils with crossbeta structures (Shults, 2006). a-Syn polymerization is nucleationdependent and it is promoted by both tubulin and oxidative stress (Paik et al., 1999; Giasson et al., 2000; Souza et al., 2000; Paxinou et al., 2001; Sherer et al., 2002; Esteves et al., 2009).

Our data indicate for the first time that taxol prevents mitochondrial-mediated MT depolymerization and a-syn oligomerization in cells that have dysfunctional mitochondria. These *in vitro* observations were made in cell lines with a neuronal background, and it will be interesting to see whether these findings generalize to PD brains.

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means were comparable.



CT cybrid cells. (C) PAGE showing the effect of 5 nM taxol on

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