



Published in final edited form as:

*Cell Mol Life Sci.* 2020 September ; 77(18): 3611–3626. doi:10.1007/s00018-019-03386-w.

## Molecular mechanism of olesoxime-mediated neuroprotection through targeting $\alpha$ -synuclein interaction with mitochondrial VDAC

Amandine Rovini<sup>1,#</sup>, Philip A. Gurnev<sup>1</sup>, Alexandra Beilina<sup>2</sup>, María Queralto-Martín<sup>1</sup>, William Rosencrans<sup>1,3</sup>, Mark R. Cookson<sup>2</sup>, Sergey M. Bezrukov<sup>1</sup>, Tatiana K. Rostovtseva<sup>1,\*</sup>

<sup>1</sup>Section on Molecular Transport, Eunice Kennedy Shriver National Institute of Child Health and Human Development, National Institutes of Health, Bethesda, MD 20892

<sup>2</sup>Cell Biology and Gene Expression Section, Laboratory of Neurogenetics, National Institute of Aging, National Institutes of Health, Bethesda, MD 20892

<sup>3</sup>Colgate University, Hamilton, NY 13346

### Abstract

An intrinsically disordered neuronal protein  $\alpha$ -synuclein ( $\alpha$ Syn) is known to cause mitochondrial dysfunction, contributing to loss of dopaminergic neurons in Parkinson's disease. Through yet poorly defined mechanisms,  $\alpha$ Syn crosses mitochondrial outer membrane and targets respiratory complexes leading to bioenergetics defects. Here, using neuronally differentiated human cells overexpressing wild-type  $\alpha$ Syn, we show that the major metabolite channel of the outer membrane, the voltage-dependent anion channel (VDAC), is a pathway for  $\alpha$ Syn translocation into the mitochondria. Importantly, the neuroprotective cholesterol-like synthetic compound olesoxime inhibits this translocation. By applying complementary electrophysiological and biophysical approaches, we provide mechanistic insights into the interplay between  $\alpha$ Syn, VDAC, and olesoxime. Our data suggest that olesoxime interacts with VDAC  $\beta$ -barrel at the lipid-protein interface thus hindering  $\alpha$ Syn translocation through the VDAC pore and affecting VDAC voltage gating. We propose that targeting  $\alpha$ Syn translocation through VDAC could represent a key mechanism for the development of new neuroprotective strategies.

### Keywords

Voltage-Dependent Anion Channel; VDAC-facilitated protein translocation; mitochondria; SH-SY5Y cells; Proximity Ligation Assay; planar lipid membrane; channel reconstitution; voltage gating; Fluorescent Correlation Spectroscopy

\*To whom correspondence should be addressed: Tatiana K. Rostovtseva, Section on Molecular Transport, Eunice Kennedy Shriver National Institute of Child Health and Human Development, National Institutes of Health, 9000 Rockville Pike, Bldg. 29B, Room 1G09, Bethesda, MD 20892-0924. Phone: (301) 402-4702, rostovtt@mail.nih.gov.

#Current address: Department of Drug Discovery & Biomedical Sciences, Medical University of South Carolina, Charleston, SC 29425.

## INTRODUCTION

One of the pathological hallmarks of Parkinson's disease (PD) is abnormal accumulation of alpha-synuclein ( $\alpha$ Syn), a 14.4-kDa, natively unfolded, neuronal protein into Lewy bodies and Lewy neurites and degeneration of dopaminergic neurons within the substantia nigra [1,2]. Under normal physiological conditions,  $\alpha$ Syn is abundant throughout the brain and mainly in presynaptic terminals of neurons. While  $\alpha$ Syn is essentially associated with the central nervous system, various reports detect smaller amounts in peripheral tissues (salivary glands, heart, muscle) and body fluids [3]. A strong body of genetic evidence including point mutations, genomic duplications, triplications, and polymorphisms in regulatory elements of the encoding *SNCA* gene, has nominated  $\alpha$ Syn as a causal gene for both familial and sporadic forms of PD [4]. The alterations in  $\alpha$ Syn expression levels, associated with its accumulation into misfolded oligomers and larger aggregates, are proposed to be the main reason for its toxicity towards dopaminergic neurons that are lost in PD. However, the detailed mechanisms by which  $\alpha$ Syn is involved in neurodegeneration remain enigmatic. Known pathways are diverse and involve interplay with multiple organelles including synaptic vesicles, ER, Golgi apparatus, lysosomes, and mitochondria [5]. A specific relationship between  $\alpha$ Syn toxicity and mitochondrial dysfunction is supported by studies showing intra-mitochondrial accumulation of the protein and impairment of the mitochondrial respiratory complexes I [6-9], IV [10] and the ATP synthase [11]. However, several important aspects of these functional interactions remain undefined, including identification of the mechanism(s) by which the normally cytosolic  $\alpha$ Syn protein gains access to mitochondria.

To date, neuroprotective strategies aimed at slowing progression of PD constitute an unmet clinical need. Recently, olesoxime, an investigational neuroprotective compound, was shown to be protective in a neuronally differentiated human cell model of  $\alpha$ Syn-mediated toxicity [12]. This low molecular weight, cholesterol-like compound was originally identified in a medium-throughput screen for rescuing motor neurons from neurotrophic factor deprivation or Fas-induced cell death [13-15]. Gouarne et al., showed that olesoxime-mediated neuroprotection was characterized by reduction of  $\alpha$ Syn-induced cell death while preserving mitochondrial integrity and function [12]. The authors associated the pro-survival effects of olesoxime with a decrease in mitochondrial cytochrome c release and inhibition of caspase-9 activation. Olesoxime was found associated with two major proteins of the mitochondrial outer membrane (MOM): translocator protein 18 kDa (TSPO), the mitochondrial cholesterol transporter, and the voltage-dependent anion channel (VDAC) [13,14]. Our interest in the role of VDAC in the olesoxime-promoted neuroprotection is motivated by our *in vitro* data showing an interaction of VDAC with monomeric  $\alpha$ Syn [16]. We found that VDAC, reconstituted into planar lipid membranes, can be reversibly blocked by recombinant monomeric  $\alpha$ Syn with nanomolar efficiency. Using electrophysiological experiments in combination with modeling, we demonstrated that  $\alpha$ Syn molecules translocate through the VDAC pore [16-18]. These results lead us to conclude that  $\alpha$ Syn-membrane binding is the required first step of  $\alpha$ Syn-VDAC interaction controlling both VDAC blockage by  $\alpha$ Syn and its translocation through the VDAC pore. We proposed a model where monomeric  $\alpha$ Syn bound to the MOM from the cytosolic side by its N-terminus domain could disrupt

ATP/ADP fluxes through VDAC by dynamically blocking the pore with its anionic C-terminal domain. Thus, in normal cells  $\alpha$ Syn could be a functional regulator of mitochondrial bioenergetics. In pathology,  $\alpha$ Syn can cross the MOM barrier by translocating through VDAC and target the electron transport chain (ETC) complexes in the mitochondrial inner membrane (MIM) causing their impairment and leading to mitochondrial dysfunction [16]. Experiments on live cells with a yeast strain deficient in VDAC1 (por1<sup>-</sup>) supported our *in vitro* findings by demonstrating that  $\alpha$ Syn toxicity in yeast depends on VDAC. However, whether or not  $\alpha$ Syn-VDAC interaction is a physiological means to regulate cellular respiration and energy production in neuronal cells, remained unknown.

Here, using neuronally differentiated SH-SY5Y cells overexpressing wild type (wt)  $\alpha$ Syn as a cell model of PD, we studied the mechanism of olesoxime-induced neuroprotection focusing on the potential role of VDAC as one of the olesoxime mitochondrial targets [13,14]. Using a combination of confocal imaging and molecular biology approaches, we gathered *in situ* evidence for  $\alpha$ Syn translocation through VDAC into the mitochondria and inhibition of this translocation in the presence of olesoxime. Using complementary *in vitro* electrophysiology method on VDAC reconstituted in planar lipid membranes, we showed that olesoxime hinders  $\alpha$ Syn translocation through the VDAC pore and affects VDAC voltage-gating properties. Our data suggest that olesoxime directly interacts with VDAC  $\beta$ -barrel at the lipid-protein interface. These results identify VDAC as a key player in olesoxime neuroprotection against  $\alpha$ Syn-induced mitochondria toxicity.

## MATERIALS AND METHODS

### Cell culture conditions and transfection

Human neuroblastoma SH-SY5Y cells were bought from the American Tissue Culture Collection (Manassas, VA) and grown in RPMI 1640 (12-702F, Gibco) medium supplemented with 10% fetal bovine serum (ThermoFisher Scientific). Cultures were maintained at 37 °C in a humidified atmosphere with 5% CO<sub>2</sub>. Constructs for transient expression of  $\alpha$ Syn were built up by insertion of wt human  $\alpha$ Syn coding sequence into the pcDNA<sup>TM</sup> 3.1 vector. SH-SY5Y cells were transiently transfected with either an empty plasmid or plasmid containing wt  $\alpha$ Syn using Lipofectamine<sup>TM</sup> 3000 transfection reagent (Invitrogen) following manufacturer's instruction and a ratio of 0.8  $\mu$ g of DNA per 100,000 cells. At the same time, differentiation was induced for all conditions by addition of 20  $\mu$ M all-trans retinoic acid (RA; Sigma-Aldrich). Cells were treated with 3  $\mu$ M olesoxime (TRO 19622, Tocris Bioscience) in 0.5% final DMSO (Sigma-Aldrich) (control cultures received DMSO only) was initiated as well. For all experiments, the entire medium was changed every 2 days and all reagents were replaced at the same concentration.

### siRNA transfection

Cells were transfected with either the VDAC1 On-Target Plus SMART pool siRNA or siGenome Non-Targeting siRNA as a control (Dharmacon, Waltham, MA) at a final concentration of 5 nM, using DharmaFECT<sup>TM</sup> 1 as per the manufacturer's instructions (GE Healthcare). Media was renewed 24 hours post transfection, with cells typically incubated for a further 24-48 hours before harvesting/analyzing.

### Trypan blue exclusion test of cell viability

To evaluate  $\alpha$ Syn cytotoxicity following transient transfection, cell viability was measured by the trypan blue exclusion assay 96 h after transfection. After the addition of 0.4% trypan blue for 2 min (Sigma), the percentage of dead cells (trypan blue-stained) was counted manually with a hemocytometer, within 5 min.

### Live cell imaging of mitochondrial membrane potential

For all confocal imaging, cells were grown on 35 mm MatTek dishes (MatTek corporation). Mitochondrial membrane potential ( $\Psi_m$ ) was assessed by loading cells with 100 nM MitoTracker™ Red CMXRos (Cell Signaling Technology) for 25 min. After the incubation, the fluorescent probe was washed out with Hank's balanced salt solution. The cells were placed in a thermostated chamber on the stage of a Zeiss LSM 710 confocal microscope and images were taken with a  $63 \times 1.4$  N.A. Plan-Apochromat. Image analysis and quantification of the fluorescence intensity was performed with ImageJ (NIH) as described in [19], and averaged for five independent experiments.

### Proximity ligation assay (PLA)

Cells were grown, transfected and differentiated on Nunc™ Lab-Tek™ II CC2™ Chamber Slide System (ThermoFisher scientific). PLA was carried out with Duolink® In Situ Detection Reagents Orange (Sigma Aldrich) with little modifications from the manufacturer's instructions, as reported in [20]. The cells were treated with primary antibodies for anti- $\alpha$ -synuclein (Cell Signaling Technology, 1:100, rabbit) and anti-VDAC1 (Abcam, 1:100, mouse) or anti-COXIV (Cell Signaling Technology, 1:100, mouse). Slides were mounted using a minimal volume of Duolink in situ Mounting Medium containing DAPI and images were taken with a Zeiss LSM 710 confocal microscope at  $63 \times$  magnification. The microscope settings were kept constant for all images to enable direct comparison. Quantification of signals (number of dots per cell) was obtained from thresholded images using the "analyze particles" feature of ImageJ, which detects isolated continuous objects in the image. Neither PLA puncta size nor circularity was constrained for detection.

### Western Blotting

Proteins were extracted in cell lysis buffer (Cell Signaling Technology) with additional phosphatase and protease inhibitors (Sigma). Proteins were separated on 4–20% Criterion TGX pre-cast gels (Biorad) in SDS/Tris-glycine running buffer and transferred to PVDF membranes by semi-dry trans-Blot Turbo transfer system (Biorad). The membranes were blocked for 1h with Odyssey Blocking Buffer (Licor) and then incubated for 1h at room temperature (RT) with antibodies: anti  $\alpha$ -syn (Cell Signaling Technologies, 1:1000, rabbit), anti-VDAC1 (Abeam, 1:2000, mouse), anti-GAPDH (Sigma, 1:2000, rabbit), and anti- $\beta$ -actin (Santa Cruz, 1:1000, mouse). The membranes were washed in TBST (3 $\times$ 5 min) at RT followed by incubation for 1h at RT with fluorescently conjugated goat anti-mouse or rabbit IR Dye 680 or 800 antibodies (Licor). The blots were washed in TBST (3 $\times$ 5 min) at RT and scanned on an ODYSSEY® CLx (Licor). Quantitation of western blots was performed using

Image Studio (Licor), the intensity of target proteins was standardized with the loading control.

### Materials and protein purification

1,2-dioleoyl-sn-glycero-3-phosphocholine (DOPC), 1,2-dioleoyl-sn-glycero-3-phosphoethanolamine (DOPE), cholesterol (Chol), and a soybean Polar Lipid Extract (PLE) were purchased from Avanti Polar Lipids (Alabaster, AL). Olesoxime was purchased from Tocris. All other chemicals were obtained from Sigma-Aldrich unless noted otherwise.

Recombinant wt  $\alpha$ Syn and Alexa-488-labeled  $\alpha$ Syn were the generous gifts of Dr. Jennifer Lee (NHLBI, NIH).  $\alpha$ Syn was expressed, purified, and characterized as described previously [21] and stored at  $-80^{\circ}$  C.  $\alpha$ Syn Y136C mutant was labeled with fluorophore Alexa488 at position Y136C as described previously [22]. VDAC was isolated from frozen mitochondrial fractions of rat liver that were a kind gift of Dr. Marco Colombini (University of Maryland, College Park, MD) following the standard method [23] and purified on a hydroxyapatite/celite (2:1) column as described previously [24]. VDAC purified from mitochondrial fraction of rat liver contains all three isoforms with VDAC1 being the predominant one (~ 80% of total VDACS) [25].

### Channel reconstitution and conductance measurements

Planar lipid membranes for single-channel experiments were formed from DOPC/DOPE (1:4 w/w) mixtures with 5% (w/w) of olesoxime or cholesterol. For multichannel experiments planar membranes were prepared from PLE with or without 20% olesoxime (w/w) or from PLE with 4% cholesterol (w/w) and with or without 16% olesoxime (w/w). The mixtures of lipids were prepared from 10 mg/ml aliquots of two lipids, cholesterol or olesoxime solutions in chloroform, followed by drying with nitrogen and then re-dissolving them in pentane to a total lipid concentration of 5 mg/ml. Planar bilayer membranes were formed from two opposing lipid monolayers across  $\sim 70$   $\mu$ m aperture in the 15- $\mu$ m-thick Teflon partition separating two  $\sim 1.2$ -mL compartments as previously described [26]. VDAC insertion was achieved by adding VDAC isolated from the mitochondria of rat liver in 2.5 % triton X-100 buffer to the aqueous phase of 1 M KCl buffered with 5 mM Hepes at pH 7.4 in the *cis* compartment. Potential is defined as positive when it is greater at the side of VDAC addition (*cis*).  $\alpha$ Syn at a final concentration of 50 nM was added symmetrically to both sides of the membrane. Records for analysis were obtained no less than 15 min after  $\alpha$ Syn addition to ensure a steady state. Single-channel conductance measurements were performed as described previously [16] using an Axopatch 200B amplifier (Axon Instruments, Inc., Foster City, CA) in the voltage clamp mode. Data were filtered by a low pass 8-pole Butterworth filter (Model 900, Frequency Devices, Inc., Haverhill, MA) at 15 kHz and a low pass Bessel filter at 10 kHz, and directly saved into computer memory with a sampling frequency of 50 kHz. For single-channel data analysis by Clampfit 10.3, a digital 8-pole Bessel low pass filter set at 5 kHz or 2 kHz was applied to current recordings and then individual events of current blockages were discriminated and kinetic parameters were acquired. Individual events of current blockages were discriminated and kinetic parameters were acquired by fitting single exponentials to logarithmically binned histograms [27] as described previously [16,28]. Four different logarithmic probability fits were generated

using different fitting algorithms and the mean of the fitted time constants was used as the mean for the characteristic open and blockage times. Each channel experiment was repeated at least three times on different membranes.

VDAC voltage-dependent properties were assessed on multichannel membranes following the protocol previously devised [29-31] in which gating is inferred from the channels response to a slow symmetrical 5 mHz triangular voltage wave of  $\pm 60$  mV amplitude from an Arbitrary Waveform Generator 33220A (Agilent). Data were acquired at a sampling frequency of 2 Hz and analyzed as described previously [31,30] using pClamp 10.3 software. In each experiment, current records were collected from membranes containing 20-850 channels in response to 5-10 periods of voltage waves. Only the part of the wave during which the channels were reopening was used for the subsequent analysis.

### Statistics

Differences between groups were analyzed by a one-way ANOVA using  $p < 0.05$  as the criterion of significance. Data points are the means  $\pm$  S.D. of 3–6 independent experiments or presented as box-plots. Images in figures are representative of three or more independent experiments.

## RESULTS

### Olesoxime attenuates $\alpha$ Syn-induced loss of mitochondrial membrane potential

Olesoxime neuroprotective capacity against  $\alpha$ Syn-induced mitochondrial toxicity was studied using retinoic acid differentiated SH-SY5Y cells overexpressing  $\alpha$ Syn. Overexpression was confirmed by western blot and immunostaining which also demonstrated the lack of effect of olesoxime on  $\alpha$ Syn expression (Supplemental Fig. S1). As seen in Supplemental Fig. S1C, enhanced level of  $\alpha$ Syn resulted in a punctate staining of the protein expanded all over the cytoplasm. Under these conditions, no aggregates were detected. The toxicity of  $\alpha$ Syn and the neuroprotection of olesoxime were confirmed in this cell model by assessing morphological features and cell survival parameters. Fig 1A shows morphological alterations associated with overexpression of wt  $\alpha$ Syn while olesoxime treatment maintained a normal phenotype, as previously described [12]. By counting the number of Trypan-blue excluded cells, we determined that  $\alpha$ Syn overexpression resulted in an average of  $21.2 \pm 4.3\%$  dead cells (Fig. 1B,  $\alpha$ Syn), a significant increase of cell death ( $p < 0.01$ , here and elsewhere, by one-way ANOVA) when compared with the  $1.5 \pm 0.7\%$  of dead cells in basal conditions (Fig. 1B, pEV). Addition of  $3\mu\text{M}$  olesoxime to  $\alpha$ Syn overexpressing cells caused a significant decrease, down to  $6.9 \pm 5.8\%$  ( $p < 0.01$ ) of dead cells (Fig 1B,  $\alpha$ Syn+Oxime), thus confirming olesoxime's protective effect. Importantly,  $3\mu\text{M}$  olesoxime alone is not toxic to the cells (Fig. 1A, B, Oxime).

To assess the effect of olesoxime on mitochondria function in  $\alpha$ Syn overexpressing cells, we evaluated mitochondrial membrane potential using the cell permeant MitoTracker™ probe which passively diffuses across the cell membrane and accumulates in energized mitochondria [32]. Analysis of the MitoTracker™ fluorescence showed that the basal fluorescence intensity of  $\alpha$ Syn overexpressing cells is significantly lower ( $46 \pm 15\%$ ,  $p <$

0.05) than in control cells (Figure 1 C, D, pEV, normalized to 100%). Olesoxime treatment of  $\alpha$ Syn overexpressing cells significantly rescued the basal fluorescence intensity ( $88 \pm 5\%$  vs  $46 \pm 15\%$ ,  $p < < < 0.05$ ), while cells treated with olesoxime alone had basal fluorescence intensity similar to control cells ( $103 \pm 18\%$ ,  $p > 0.5$ ) (Fig. 1D). These results confirm a key role of mitochondria in olesoxime neuroprotection against  $\alpha$ Syn-mediated toxicity.

### In cells, olesoxime reduces the $\alpha$ Syn association with mitochondrial membrane proteins

$\alpha$ Syn has been found to associate with both outer and inner mitochondrial membranes [33,34,7,35]. However, the pathway by which  $\alpha$ Syn enters mitochondria has not been identified. Recent *in vitro* data suggest that VDAC could be such a pathway for  $\alpha$ Syn to cross the MOM [16]. Although  $\alpha$ Syn binding to and translocation through VDAC have been extensively studied biophysically [16-18], a visual assessment of this model *in situ* in cells has never been done. The use of conventional confocal microscopy approaches for such a task is challenging due to high expression levels of both  $\alpha$ Syn in the cytosol and VDAC in the MOM. To overcome these limitations, we performed an *in situ* Proximity Ligation Assay (PLA), a method that allows to visualize the presence of two proteins at close distances with a resolution up to 30 nm [36].

To evaluate the effect of olesoxime on  $\alpha$ Syn association with mitochondrial membranes, we used PLA to assess proximities between  $\alpha$ Syn and VDAC and between  $\alpha$ Syn and complex IV (COX IV), as markers of the MOM and MIM respectively. Analysis of confocal images of cells after PLA showed that overexpression of  $\alpha$ Syn correlates with a significant enrichment of  $\alpha$ Syn found in close proximity to VDAC at the MOM (Fig 2A, upper row). Quantification of PLA signals for pairwise  $\alpha$ Syn /VDAC antibodies revealed a  $\sim 3.5$  times increase of the average number of  $\alpha$ Syn in close proximity to VDAC relative to the basal conditions (pEV) (Fig. 2 B). To provide further evidence of  $\alpha$ Syn accumulation at the MOM, we performed an additional PLA for  $\alpha$ Syn and the translocase of the outer membrane 20 (TOM20). Using this other marker of the MOM, we found a significant increase ( $p = 0.01$ ) in PLA signals for  $\alpha$ Syn and TOM20 (Supplemental Fig. S2). Olesoxime treatment of cells overexpressing  $\alpha$ Syn significantly decreased the  $\alpha$ Syn and VDAC co-localization (Fig. 2A, B). However, olesoxime treatment did not affect association of endogenous  $\alpha$ Syn with VDAC in non-transfected cells (NS;  $p > 0.5$ ) (Fig. 2A, B). These results indicate that olesoxime can diminish the co-localization of  $\alpha$ Syn and VDAC. Next, using a pair of  $\alpha$ Syn /COX IV antibodies, we found a  $\sim 3$  times increase of the average number of  $\alpha$ Syn molecules in a close proximity to COX IV at the MIM (Fig 2A, lower panels and C) in  $\alpha$ Syn overexpressing cells. Olesoxime treatment of cells overexpressing  $\alpha$ Syn resulted in a significant reduction of  $\alpha$ Syn co-localization with COX IV ( $p < 0.05$ ). Olesoxime *per se* did not affect association of endogenous  $\alpha$ Syn with COX IV (Fig. 2C). Collectively, these results show that olesoxime significantly reduces the association of  $\alpha$ Syn with both inner and outer mitochondrial membranes proteins.

### VDAC1 is involved in $\alpha$ Syn translocation into mitochondria

The finding that  $\alpha$ Syn localizes in a close proximity ( $\sim 30$ - $40$  nm) to VDAC and COX IV is suggestive of a direct interaction with these two proteins but does not provide definitive

evidence in either case (Fig. 2) and does not show whether VDAC forms a pathway for  $\alpha$ Syn translocation across MOM. As the current knowledge indicates VDAC1 as the major docking site at the MOM for misfolded proteins involved in neurodegenerative diseases [37], to further evaluate the role of VDAC in  $\alpha$ Syn mitochondrial localization, VDAC1 expression was downregulated in differentiated SH-SY5Y cells using siRNA. Western blot in Fig. 3A, B shows 85% decrease of VDAC1 protein after siRNA treatment. PLA was then used to monitor the ability of  $\alpha$ Syn to cross MOM and target COX IV in VDAC1 depleted cells. Under these conditions, we found that overexpression of  $\alpha$ Syn did not result in a significant difference ( $1.3 \pm 1.2$ ;  $p > 0.4$ ) in PLA signal for  $\alpha$ Syn/COX IV co-localization (Fig 3C, D) in contrast with 3-fold increase of  $\alpha$ Syn/COX IV co-localization in cells with endogenous VDAC1 (Fig. 2C, D). Olesoxime treatment of these cells did not affect PLA signals (Fig. 3D). Altogether, *in situ* results show that in cells overexpressing  $\alpha$ Syn, VDAC1 is important for  $\alpha$ Syn translocation into mitochondria.

### Olesoxime prevents $\alpha$ Syn translocation through VDAC

The decrease of  $\alpha$ Syn co-localization with VDAC and COX IV in cells treated with olesoxime could be explained by various mechanisms. One explanation is based on the previously shown direct interaction between olesoxime and VDAC using competition between olesoxime and photolabeled analog of pregnanolone for VDAC [14]. Therefore, we hypothesized that olesoxime binding to VDAC could interfere with  $\alpha$ Syn-VDAC interaction and  $\alpha$ Syn translocation through the channel. To assess the hypothesis, we employed electrophysiological measurements on VDAC channels reconstituted into planar lipid membranes. This method enables highly sensitive measurements of  $\alpha$ Syn's dynamic interaction with VDAC at the *single-molecule* level [16]. Using VDAC electrophysiology, we recently reported that  $\alpha$ Syn is able to reversibly block channel conductance in a concentration- and highly voltage-dependent manner. By measuring ionic selectivity of  $\alpha$ Syn-blocked state in salt gradient conditions, we showed that the presence of the anionic C-terminal domain of  $\alpha$ Syn in the pore reduces VDAC's selectivity, while the uncharged N-terminal region leaves it relatively unaffected [17]. By using an experimental approach based on this observation, we established a method for direct, model-free, real-time monitoring of the translocation of  $\alpha$ Syn polypeptide through the VDAC pore [18]. Therefore, our data imply that  $\alpha$ Syn is able to translocate through the channel [16-18]. We capitalized on this earlier established system and compared the effects of olesoxime on VDAC basic channel properties and on the kinetic parameters of  $\alpha$ Syn-induced VDAC blockages with the effect of cholesterol as a relevant control for the cholesterol-like olesoxime molecule.

Two zwitterionic lipids, DOPC and DOPE, were chosen to mimic MOM phospholipid composition where PC and PE make up to 44 and 35% of the total lipid content of the rat liver MOM, respectively [38]. A natural content of cholesterol in mammalian MOM is estimated as ~ 5 - 7% (w/w) of total phospholipid [39,40]. VDAC was reconstituted into planar membranes formed from DOPC/DOPE (1:4 w/w) mixture with 5% (w/w) of olesoxime or cholesterol or without them. Fig. 4A shows representative current traces through a single VDAC reconstituted into planar membranes formed from PC/PE with olesoxime (*a*), with the equal amounts of cholesterol (*b*), or without either of them (*c*). Before addition of  $\alpha$ Syn, a typical ion current through VDAC's high conducting or "open"

state is steady and corresponds to 4 nS conductance in all three lipid compositions (in 1 M KCl solution) [29,41] (Fig. 4A, left traces) showing that neither cholesterol nor olesoxime affect VDAC open state conductance. Addition of 50 nM of  $\alpha$ Syn to the membrane bathing solution resulted in rapid and well-resolved millisecond range current fluctuations between the open state and the lower conducting  $\alpha$ Syn-blocked state (Fig. 4A, right traces), which is a characteristic effect of  $\alpha$ Syn on reconstituted VDAC [16]. Importantly, the current corresponding to the open state did not change in the presence of  $\alpha$ Syn indicating the absence of any pore-forming activity of  $\alpha$ Syn at this concentration [16]. Each individual blockage event corresponds to a steric blockage of the VDAC pore by individual membrane-bound  $\alpha$ Syn molecule [17]. Addition of olesoxime to the lipid membrane did not visibly affect the general pattern of the characteristic current blockage behavior induced by  $\alpha$ Syn (b). Therefore, olesoxime does not prevent  $\alpha$ Syn from blocking VDAC pore. However, the kinetic analysis of blockage times, the intervals of time when channel is blocked by  $\alpha$ Syn, revealed a strikingly different behavior in the presence of olesoxime in comparison with (PC/PE/Chol) or without cholesterol (PC/PE). The blockage times obtained in the presence of olesoxime (PC/PE/Oxime) increased exponentially with the amplitude of the applied voltage up to  $-47.5$  mV (Fig. 4B). In contrast, the voltage dependences obtained in PC/PE and PC/PE/Chol membranes showed a clear biphasic behavior: blockage time increased exponentially up to  $\sim -35$  mV (indicated by the vertical dashed line in Fig. 4B) and then markedly decreased at  $|V| > 35$  mV (highlighted in yellow in Fig. 4B).

As shown previously [17,16,18], the increase of blockage time with the applied voltage corresponds to the regime of reversible VDAC blockage when the acidic C-terminal of  $\alpha$ Syn is captured and retracted from the pore while the  $\alpha$ Syn molecule remains at the same side of the channel (Fig. 6). The decrease of dwell time at higher potentials observed in PC/PE or PC/PE/Chol membranes is indicative of the translocation regime when the applied voltage drives the entire  $\alpha$ Syn molecule through the pore. This kind of behavior has been demonstrated for polypeptides and DNA translocation through various nanopores (e.g, [42,43]). Importantly, both voltage dependences obtained on PC/PE membranes with or without cholesterol are overlapping, confirming that cholesterol does not affect either blockage or translocation regimes of  $\alpha$ Syn through VDAC. This is drastically different from the exponential increase of blockage times with the voltage amplitude in the olesoxime-containing membranes (Fig. 4B).

Complementary experiments were performed to confirm the effect of olesoxime on the translocation regimes of  $\alpha$ Syn through VDAC. To that end, olesoxime was added to the membrane-bathing solution after recording of  $\alpha$ Syn-induced blockages of VDAC inserted into PC/PE/Chol membranes (Supplemental Fig. S3 A, traces *a* and *b*). Successive additions of olesoxime (from 10  $\mu$ M up to a final concentration of 100  $\mu$ M; Supplemental Fig. S3 A, traces *c* and *d*) resulted in a significant increase of blockage events duration (at high applied voltages) following addition of olesoxime, which is indicative of olesoxime interference with  $\alpha$ Syn translocation (Supplemental Fig S3B).

Measurable blockage of the VDAC pore occurs when negative potential is applied from the side of  $\alpha$ Syn addition [16].  $\alpha$ Syn blocks VDAC in qualitatively similar way from both sides of the channel [16]. Hence, for the sake of clarity, we present results for the blockage from

the presumably cytosolic side of the channel which corresponds to the “*cis*” side of our experimental chamber or to the side of VDAC addition [44]. Considering  $\alpha$ Syn’s cytosolic localization in cells, its interaction with cytosolic side of the channel is physiologically most relevant.

### Olesoxime affects VDAC voltage gating

These electrophysiological results suggest that olesoxime delays  $\alpha$ Syn translocation through the VDAC pore and consequently, prevents  $\alpha$ Syn from targeting the ETC at the MIM. The absence of cholesterol effect on  $\alpha$ Syn-VDAC interaction points toward a specific interaction of olesoxime with VDAC that modifies the channel or its immediate lipid environment in a way that prevents  $\alpha$ Syn translocation at  $|V| < 50$  mV. To further test this hypothesis using reconstituted VDAC, we examined the effect of olesoxime on VDAC voltage gating, an important VDAC’s property that allows assessing VDAC’s functions [23,29,31,30]. Quantification of VDAC gating requires a different experimental approach that involves membranes with multiple reconstituted channels [29,30,45].

Voltage gating, which is the voltage-triggered switching between high and low conducting states, the so-called “open” and “closed” states [29], is a characteristic property of VDAC and its namesake. VDAC voltage gating is known to be sensitive to the surrounding lipid environment [30,46,47], pH [31,48], and osmotic stress [49]. To achieve the reliable statistical data on VDAC gating which is an inherently stochastic process, we used the well-established experimental approach where gating is inferred by monitoring the response of multiple channels to triangular voltage waves [29,30]. The periodic voltage wave is chosen slow enough (5 mHz, corresponding to a period of 200 s) to ensure that the channel reopening is much faster (milliseconds range) so that the obtained results do not depend on the time parameters of the voltage protocol used [45].

Fig. 5A shows the normalized conductance plots ( $G/G_{max}$ , where  $G$  is the conductance at a given voltage and  $G_{max}$  is the maximum conductance at voltages close to 0 mV) versus the applied voltage obtained for the system of multiple VDAC channels reconstituted into the membranes formed from soybean Polar Lipid Extract (PLE), with cholesterol (PLE/Chol), and with olesoxime (PLE/Oxime). PLE is a natural lipid mixture which was chosen for these experiments as it ensures a pronounced VDAC gating in electrophysiological experiments [47] and also closely mimics MOM phospholipid composition in rat liver mitochondria, including the content of charged lipids and acyl chains [38]. This condition is important because VDAC gating was shown to be especially sensitive to the lipid headgroup charges [47]. All  $G/G_{max}$  plots display the characteristic bell-shaped voltage dependences of the normalized conductance (Fig. 5A) with some asymmetry with respect to the polarity of the applied voltage: the minimum conductance  $G_{min}$  calculated at  $|V| > 50$  mV, is slightly lower at negative than at positive voltages. Addition of olesoxime (20% w/w in the PLE) enhanced VDAC gating by decreasing  $G_{min}$ . This effect was more pronounced at the negative applied voltages (~8% decrease of  $G_{min}$  in comparison with control) (Fig. 5A). The effect of olesoxime on VDAC gating was more pronounced when olesoxime was added to the PLE membranes with 4% of cholesterol (w/w) (Fig. 5B, showing the data with or without olesoxime). The olesoxime content in experiments with PLE/Chol/Oxime membranes was

reduced to 16% w/w to keep the total amount of cholesterol and olesoxime at the same level as in experiments shown in Fig. 5A (20%). In PLE/Chol/Oxime membranes  $G_{min}$  decreased by ~15% at negative voltages in comparison with  $G_{min}$  obtained in experiments without olesoxime. Importantly, cholesterol, in contrast to olesoxime, does not affect VDAC gating (Fig. 5A) [47].

Next, we tested the effect of olesoxime on VDAC gating in the membranes made of the same lipid compositions as those used in single-channel experiments in Fig. 4. In these experiments, addition of olesoxime (5% w/w) to the PC/PE (1:4 w/w) mixture significantly decreased closed state conductance (\*\*:  $p < 0.01$ ; one-way ANOVA) at negative potentials, while cholesterol did not affect gating (Fig. 5C), confirming the results in Fig. 5A, B.

### Olesoxime does not measurably change mechanical properties of lipid membranes

The results with reconstituted VDAC suggest that olesoxime modulates VDAC gating either by directly interacting with VDAC or, alternatively, indirectly by changing integral properties of lipid bilayer. Considering a large amount of olesoxime (5 to 20%) used in the gating experiments one could expect a change of bilayer properties based on the previously reported data that olesoxime concentrates on mitochondrial membranes and modulates mitochondrial membrane fluidity [50,51]. Therefore, we tested the possibility that olesoxime changes the integral properties of the lipid bilayer by using two independent biophysical approaches. We first employed a channel-forming polypeptide, gramicidin A (gA), as a well-recognized highly sensitive molecular probe of planar lipid bilayer mechanics [52,30,53,54,28]. The advantage of using gA for such a task is that it allows to use exactly the same system of planar lipid membranes as in the experiments with reconstituted VDAC. Several studies have demonstrated that the lifetime of a gA conducting dimer is exquisitely sensitive to the bilayer thickness, lipid packing stress, bilayer curvature, etc. [55,54,52,30]. In particular, it was demonstrated that cholesterol causes a ~10 times reduction of gA lifetime due to the increase in the lipid packing stress [52] or membrane stiffness [55]. In contrast with drastic cholesterol effect, we did not find a significant effect of olesoxime addition (20% w/w) to the DOPE/DOPC membranes on gA lifetime (Supplemental Fig. S4), which suggests that olesoxime does not change appreciably the lipid bilayer mechanical properties under our experimental conditions. Therefore, we conclude that olesoxime most likely affects VDAC gating by directly interacting with the channel.

We then tested a possibility of olesoxime to influence  $\alpha$ Syn membrane binding using Fluorescence Correlation Spectroscopy (FCS). The interaction of  $\alpha$ Syn with liposome membranes was assessed using Alexa488-labeled  $\alpha$ Syn and large unilamellar vesicles (LUVs) (Supplemental Methods). We found that FCS correlation functions obtained with the same concentration of total lipid in LUVs prepared from PC/PE and PC/PE with 10% of olesoxime or cholesterol overlap with each other, indicating that neither olesoxime nor cholesterol affect membrane binding of  $\alpha$ Syn (Supplemental Fig. S5 A). In order to quantify  $\alpha$ Syn binding to the membranes with different lipid compositions, we followed the approach described in [22]. The analysis of FCS correlation functions showed that there is no significant difference between LUVs containing olesoxime or the same amount of

cholesterol (Supplemental Fig. S5 B). This suggests that olesoxime does not affect  $\alpha$ Syn membrane binding.

## DISCUSSION

The relationship between mitochondrial dysfunction and  $\alpha$ Syn is central to the understanding of PD pathogenesis and, therefore, could be instructive for the development of neuroprotective compounds. Based on evidence of mitochondrial function protection [13,56,57,50,51,58,59], the experimental compound olesoxime has recently been assessed for its potential neuroprotective effect in a human cell model of  $\alpha$ Syn toxicity [12]. In this model, olesoxime favored cell survival leading to the hypothesis that its protective effect is exerted through preservation of mitochondrial function. In the current work we have chosen olesoxime for its protective properties but also as a tool to investigate the mechanism of  $\alpha$ Syn-induced mitochondrial toxicity.

Here, we first confirmed olesoxime neuroprotective potential in our cell model and then assayed its effect on mitochondrial function. We found a ~ 60% loss of the MitoTracker™ fluorescence intensity in differentiated SH-SY5Y cells overexpressing  $\alpha$ Syn (Fig. 1C, D). This in line with previous studies conducted on cell lines or transgenic mice overexpressing  $\alpha$ Syn [60-64,9], as well as in recent models of dopaminergic neurons derived from PD patients [65]. We found that olesoxime significantly protects against  $\alpha$ Syn overexpression-induced dissipation of the mitochondrial membrane potential (Fig. 1C, D), thus confirming olesoxime's protective effects in our cell model. Mechanisms underlying a disturbance of mitochondrial membrane potential include insufficient ETC activity or uncoupled oxidative phosphorylation. As a matter of fact, detrimental effect of  $\alpha$ Syn on ETC components, in particular complex I [7,6,66] and complex IV [10,67], and lately ATP synthase [11,65], have been consistently reported (most recently reviewed in [68]).

Recently, we characterized the interaction between  $\alpha$ Syn and VDAC by using VDAC reconstituted into planar lipid membranes and proposed a three-step model in which  $\alpha$ Syn binding to the membrane is a necessary pre-requisite to interaction with VDAC. According to this model [16-18], VDAC can serve as a gateway for  $\alpha$ Syn translocation across the MOM into the mitochondrial intermembrane space. Interaction of  $\alpha$ Syn with VDAC has been reported in various cell and animal models of PD (reviewed in [37]) using co-localization or co-immunoprecipitation approaches. In the present work, using VDAC and COX IV as markers for MOM and MIM, respectively, we found that overexpression of  $\alpha$ Syn significantly increases  $\alpha$ Syn accumulation in close proximity to VDAC in the MOM and to COX IV in the MIM (Fig 2). These data confirm, first,  $\alpha$ Syn association with both mitochondrial membranes as shown previously [33,7,35], and, second,  $\alpha$ Syn ability to translocate across MOM, in accordance with our model. We further verified the role of VDAC as a gateway for  $\alpha$ Syn by detecting a significant loss of  $\alpha$ Syn localization at the MIM in VDAC1 downregulated cells (Fig. 3). The ~ 2 times decrease in normalized  $\alpha$ Syn /COX IV PLA signal in VDAC1 SiRNA cells (Fig. 3D) compared to cells with endogenous VDAC1 (Fig. 2C) after olesoxime treatment supports our hypothesis that VDAC is involved in  $\alpha$ Syn ability to reach MIM. However, these data play rather supportive than conclusive role in verifying this model. These results support the proposed model and are in

line with our previous study with yeast strain deficient in VDAC1 (*por1*) showing that  $\alpha$ Syn toxicity in yeast depends on VDAC expression [16].

Our findings are of particular interest because of the current lack of consensus regarding the molecular identity of the MOM pathway for  $\alpha$ Syn.  $\alpha$ Syn has been shown to bind to different subunits of the translocases of the outer membrane complex (TOM20, TOM22, TIM23, TOM40), among which TOM40 has been reported to participate in  $\alpha$ Syn import to the matrix [7]. In order to test this possibility, we performed experiments with the recombinant TOM40 channel reconstituted into planar membranes and found that  $\alpha$ Syn, up to 100 nM, does not affect TOM40 conductance (Supplemental Fig. S6). This result is in striking contrast with established  $\alpha$ Syn-VDAC interaction [16,22], but is not surprising because TOM40 is a highly cation selective channel [69-71]. Besides, TOM complex recognizes specific mitochondrial sequences, that are not present on  $\alpha$ Syn. Therefore, VDAC remains as the most plausible candidate for MOM translocator of  $\alpha$ Syn.

Since its identification, olesoxime has been described as a mitochondrial-targeting compound because it selectively binds to two important components of the MOM, TSPO and VDAC. While phenotypical observation of mitochondrial protection was confirmed in various models of neurodegeneration [51,50,56], a VDAC-related molecular mechanism has not been explored. We took advantage of VDAC being a potential target to both  $\alpha$ Syn and olesoxime to understand the role of VDAC in olesoxime neuroprotection. The PLA experiments showed a significant decrease of  $\alpha$ Syn co-localization with VDAC and COX IV in cells treated with olesoxime (Fig. 2). Whether  $\alpha$ Syn specifically binds to both VDAC and COX IV, nonspecifically targets mitochondrial membranes, interacts with other membrane proteins, or all of the above, is not certain considering the crowded protein environment in both mitochondrial membranes and the limitation of PLA resolution to 30 nm between a pair of proteins of interest. Based on the earlier report that olesoxime could modulate mitochondrial membrane properties, such as membrane fluidity [50,51], and the well-described effects of membrane lipid composition on  $\alpha$ Syn membrane-binding efficiency [72-74], we considered the possibility for olesoxime to affect  $\alpha$ Syn affinity to MOM. Olesoxime, upon partitioning into MOM and thus modifying membrane properties, could cause a depletion of  $\alpha$ Syn from MOM, and consequently, abolish its intramitochondrial targeting (Fig. 2). Under this scenario, olesoxime should nonspecifically modulate all cellular organelle membranes affecting mitochondrial and cell integrity. On the contrary, our data show that olesoxime is not toxic for cells and mitochondria (Fig. 1) and does not significantly affect association of endogenous  $\alpha$ Syn with VDAC or COX IV (Fig. 2) or  $\alpha$ Syn expression level (Supplemental Fig. S1A, B). The results of FCS experiments support this conclusion because 10% of olesoxime in liposome membranes does not measurably affect  $\alpha$ Syn binding to liposomes (Supplemental Fig. S5). Considering that extracellular addition of 3  $\mu$ M olesoxime should result in a significantly lower than 10% doping of the mitochondrial membrane with olesoxime, we thus concluded that olesoxime does not induce  $\alpha$ Syn detachment from mitochondrial membranes *in situ*. The complementary experiments on planar membranes, using a gA channel as a sensitive molecular probe of lipid bilayer mechanics, further confirmed that olesoxime does not change integral properties of lipid membranes even at a high content of the compound in the membrane (20% w/w of the total lipid) that is not physiologically achievable (Supplemental Fig. S4). However, olesoxime

affects VDAC voltage gating (Fig. 5) pointing to direct interaction between olesoxime and VDAC in *in vitro* experiments. The effect on gating is even stronger in the presence of a physiologically relevant amount of cholesterol in planar membranes.

To address a question of how olesoxime could impair  $\alpha$ Syn ability to reach MIM complexes by translocating through VDAC, we used single channel experiments with VDAC reconstituted into planar lipid membranes. Notably, while olesoxime does not qualitatively change the pattern of the reversible VDAC blockage by  $\alpha$ Syn, it hinders  $\alpha$ Syn translocation through the channel (Fig. 4B and Supplemental Fig. S3).

The cartoon in Fig. 6 illustrates a proposed mechanism of olesoxime action, by affecting  $\alpha$ Syn translocation through and most likely binding to VDAC. The absence of the olesoxime effect on VDAC conductance (Fig. 4A) indicates that highly hydrophobic olesoxime does not enter the water-filled pore and therefore, most likely interacts with VDAC at the protein-lipid interface. In the proposed model (Fig. 6) olesoxime interacts with hydrophobic exterior of the VDAC's  $\beta$ -barrel, targeting it from the lipid medium and leaving the hydrophilic interior of the pore –and consequently the channel conductance –unaffected. It should be noted that if olesoxime were to interact with channel interior, it inevitably should change the dynamics of  $\alpha$ Syn molecule capture by the VDAC pore, i.e. the blockage parameters at all applied voltages. The results (Fig. 4B) show that this is not the case and that olesoxime does not affect the regime of VDAC blockage at low voltages (at  $|V| < 35$  mV), but prevents or delays  $\alpha$ Syn translocation through the pore (VDAC blockage at high voltages,  $|V| > 35$  mV). Olesoxime could interact with VDAC at lipid-pore interface between the lipid shell consisting of one or two lipid layers [75] and the  $\beta$ -barrel, while leaving membrane integral properties unchanged. The ability of VDAC to modify its surrounding lipid shell to compensate for a hydrophobic mismatching between its  $\beta$ -barrel and an adjusted lipid has been proposed recently for the VDAC2 isoform [76] and should be applicable for other VDAC isoforms. We further speculate that olesoxime may also induce subtle structural changes in the flexible loops of VDAC, which connect  $\beta$ -strands and form channel entrances, by interfering with the specific interactions between VDAC loops and lipid headgroups as proposed by Mlayeh et al. for plant VDAC [46]. A similar mechanism was recently suggested to explain interaction between VDAC2 and MOM-located pro-apoptotic Bak and Bax proteins [77,78]. Even subtle changes in channel structure could cause the increase of the energy barrier for  $\alpha$ Syn translocation through the pore [17,18] without altering the VDAC  $\beta$ -barrel integrity. Taken together, our *in vitro* data suggest that olesoxime could affect  $\alpha$ Syn translocation through VDAC by interacting with the lipid-pore interface. To the best of our knowledge, such a mechanism, where a neuroprotective compound inhibits transport of a neurodegenerative-related protein into mitochondria through interaction with another mitochondrial membrane protein, has never been previously described.

We previously proposed that under physiological conditions  $\alpha$ Syn may play a role as a regulator of VDAC (and, consequently, the MOM) permeability by modulating ATP/ADP and other respiratory substrates fluxes through it (Fig. 6). Under stress conditions, such as  $\alpha$ Syn overexpression,  $\alpha$ Syn translocates through VDAC, targeting proteins in the MIM and potentially impairing ETC. This leads to the loss of mitochondrial membrane potential and

finally to mitochondrial dysfunction. This model is in accord with the current trend positing VDAC as the main mitochondrial docking site for various neurodegenerative disease-related misfolded proteins [37]. In particular, similar to  $\alpha$ Syn, some of these proteins – i.e. amyloid  $\beta$  associated with Alzheimer disease – could have the ability to translocate through the VDAC pore to reach and impair the ETC [37,79]. We speculate that by partitioning to the MOM, olesoxime interferes with  $\alpha$ Syn-VDAC interaction, thus hindering  $\alpha$ Syn translocation into mitochondria where it targets ETC at the MIM (Fig. 6). As a result, olesoxime maintains mitochondria integrity and eventually promotes neuronal cell survival.

To conclude, the findings presented here suggest  $\alpha$ Syn interaction with VDAC as a new target for the development of anti- $\alpha$ Syn toxicity drugs for treating PD. In particular, the use of molecules interacting with the  $\alpha$ Syn-VDAC complex could be a promising and effective pharmacological treatment of neurodegenerative diseases, aimed to decrease mitochondrial deficiencies in affected neurons.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## ACKNOWLEDGEMENTS

This work was supported by the Intramural Research Program of the National Institutes of Health (NIH), *Eunice Kennedy Shriver* National Institute of Child Health Human Development (NICHD) and National Institute of Aging. Authors are grateful to Megha Rajendran (NICHD, NIH) for fruitful discussion.

## References

1. Kim WS, Kagedal K, Halliday GM (2014) Alpha-synuclein biology in Lewy body diseases. *Alzheimers Res Ther* 6 (5):73. doi:10.1186/s13195-014-0073-2 [PubMed: 25580161]
2. Goedert M, Jakes R, Spillantini MG (2017) The Synucleinopathies: Twenty Years On. *J Parkinsons Dis* 7 (s1):S53–S71. doi:10.3233/JPD-179005
3. Ma LY, Liu GL, Wang DX, Zhang MM, Kou WY, Feng T (2019) Alpha-Synuclein in Peripheral Tissues in Parkinson's Disease. *ACS Chem Neurosci* 10 (2):812–823. doi:10.1021/acscchemneuro.8b00383 [PubMed: 30714719]
4. Nussbaum RL (2018) Genetics of Synucleinopathies. *Cold Spring Harb Perspect Med* 8 (6). doi:10.1101/cshperspect.a024109
5. Wong YC, Krainc D (2017) alpha-synuclein toxicity in neurodegeneration: mechanism and therapeutic strategies. *Nat Med* 23 (2):1–13. doi:10.1038/nm.4269
6. Chinta SJ, Mallajosyula JK, Rane A, Andersen JK (2010) Mitochondrial alpha-synuclein accumulation impairs complex I function in dopaminergic neurons and results in increased mitophagy in vivo. *Neurosci Lett* 486 (3):235–239. doi:10.1016/j.neulet.2010.09.061 [PubMed: 20887775]
7. Devi L, Raghavendran V, Prabhu BM, Avadhani NG, Anandatheerthavarada HK (2008) Mitochondrial import and accumulation of alpha-synuclein impair complex I in human dopaminergic neuronal cultures and Parkinson disease brain. *J Biol Chem* 283 (14):9089–9100. doi:10.1074/jbc.M710012200 [PubMed: 18245082]
8. Loeb V, Yakunin E, Saada A, Sharon R (2010) The transgenic overexpression of alpha-synuclein and not its related pathology associates with complex I inhibition. *J Biol Chem* 285 (10):7334–7343. doi:10.1074/jbc.M109.061051 [PubMed: 20053987]
9. Reeve AK, Ludtmann MH, Angelova PR, Simcox EM, Horrocks MH, Klenerman D, Gandhi S, Turnbull DM, Abramov AY (2015) Aggregated alpha-synuclein and complex I deficiency:

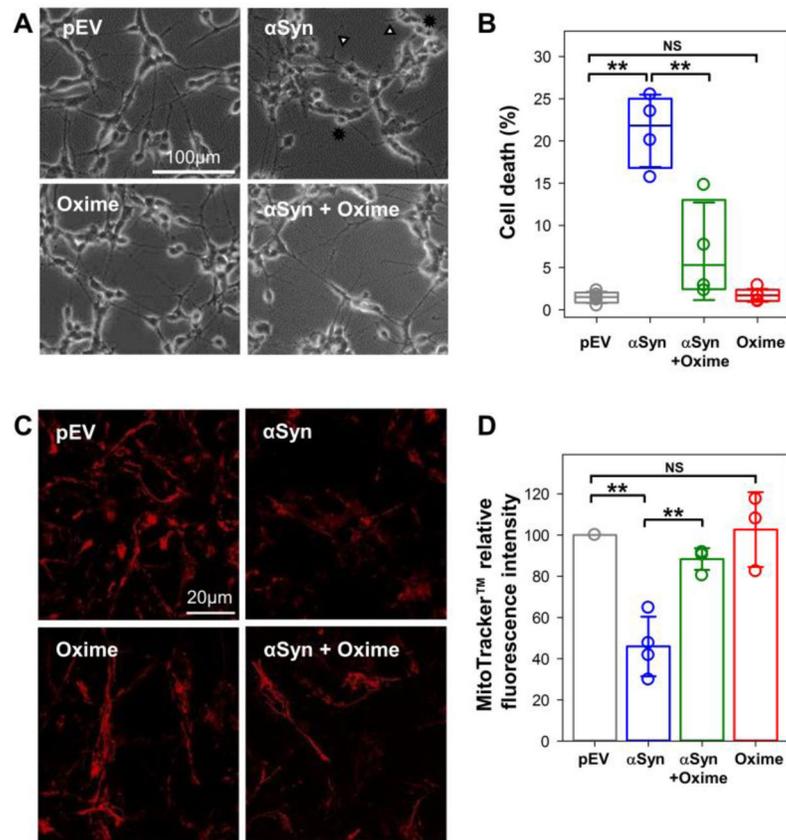
- exploration of their relationship in differentiated neurons. *Cell Death Dis* 6:e1820. doi:10.1038/cddis.2015.166 [PubMed: 26181201]
10. Elkon H, Don J, Melamed E, Ziv I, Shirvan A, Offen D (2002) Mutant and wild-type alpha-synuclein interact with mitochondrial cytochrome C oxidase. *J Mol Neurosci* 18 (3):229–238. doi:10.1385/JMN:18:3:229 [PubMed: 12059041]
  11. Ludtmann MH, Angelova PR, Ninkina NN, Gandhi S, Buchman VL, Abramov AY (2016) Monomeric Alpha-Synuclein Exerts a Physiological Role on Brain ATP Synthase. *J Neurosci* 36 (41):10510–10521. doi:10.1523/JNEUROSCI.1659-16.2016 [PubMed: 27733604]
  12. Gouarne C, Tracz J, Paoli MG, Deluca V, Seimandi M, Tardif G, Xilouri M, Stefanis L, Bordet T, Pruss RM (2015) Protective role of olesoxime against wild-type alpha-synuclein-induced toxicity in human neuronally differentiated SHSY-5Y cells. *Br J Pharmacol* 172 (1):235–245. doi:10.1111/bph.12939 [PubMed: 25220617]
  13. Bordet T, Berna P, Abitbol JL, Pruss RM (2010) Olesoxime (TRO19622): A Novel Mitochondrial-Targeted Neuroprotective Compound. *Pharmaceuticals (Basel)* 3 (2):345–368. doi:10.3390/ph3020345 [PubMed: 27713255]
  14. Bordet T, Buisson B, Michaud M, Drouot C, Galea P, Delaage P, Akentieva NP, Evers AS, Covey DF, Ostuni MA, Lacapere JJ, Massaad C, Schumacher M, Steidl EM, Maux D, Delaage M, Henderson CE, Pruss RM (2007) Identification and characterization of cholest-4-en-3-one, oxime (TRO19622), a novel drug candidate for amyotrophic lateral sclerosis. *J Pharmacol Exp Ther* 322 (2):709–720. doi:10.1124/jpet.107.123000 [PubMed: 17496168]
  15. Sunyach C, Michaud M, Arnoux T, Bernard-Marissal N, Aebischer J, Latyszenok V, Gouarne C, Raoul C, Pruss RM, Bordet T, Pettmann B (2012) Olesoxime delays muscle denervation, astrogliosis, microglial activation and motoneuron death in an ALS mouse model. *Neuropharmacology* 62 (7):2346–2352. doi:10.1016/j.neuropharm.2012.02.013 [PubMed: 22369784]
  16. Rostovtseva TK, Gurnev PA, Protchenko O, Hoogerheide DP, Yap TL, Philpott CC, Lee JC, Bezrukov SM (2015) alpha-Synuclein Shows High Affinity Interaction with Voltage-dependent Anion Channel, Suggesting Mechanisms of Mitochondrial Regulation and Toxicity in Parkinson Disease. *J Biol Chem* 290 (30): 18467–18477. doi:10.1074/jbc.M115.641746 [PubMed: 26055708]
  17. Hoogerheide DP, Gurnev PA, Rostovtseva TK, Bezrukov SM (2017) Mechanism of alpha-synuclein translocation through a VDAC nanopore revealed by energy landscape modeling of escape time distributions. *Nanoscale* 9 (1):183–192. doi:10.1039/c6nr08145b [PubMed: 27905618]
  18. Hoogerheide DP, Gurnev PA, Rostovtseva TK, Bezrukov SM (2018) Real-Time Nanopore-Based Recognition of Protein Translocation Success. *Biophysical journal* 114 (4):772–776. doi:10.1016/j.bpj.2017.12.019 [PubMed: 29338842]
  19. Maldonado EN, Patnaik J, Mullins MR, Lemasters JJ (2010) Free tubulin modulates mitochondrial membrane potential in cancer cells. *Cancer Res* 70 (24):10192–10201. doi:10.1158/0008-5472.CAN-10-2429 [PubMed: 21159641]
  20. Rostovtseva TK, Gurnev PA, Hoogerheide DP, Rovini A, Sirajuddin M, Bezrukov SM (2018) Sequence diversity of tubulin isoforms in regulation of the mitochondrial voltage-dependent anion channel. *J Biol Chem*. doi: 10.1074/jbc.RA117.001569
  21. Pfefferkorn CM, Lee JC (2010) Tryptophan probes at the alpha-synuclein and membrane interface. *The journal of physical chemistry B* 114 (13):4615–4622. doi:10.1021/jp908092e [PubMed: 20229987]
  22. Jacobs D, Hoogerheide DP, Rovini A, Zhiping J, Lee JC, Rostovtseva TK, Bezrukov SM (2019) Probing Membrane Association of  $\alpha$ -Synuclein Domains with VDAC Nanopore Reveals Unexpected Binding Pattern. *Scientific Reports* 9 (1):4580. doi:10.1038/s41598-019-40979-8 [PubMed: 30872688]
  23. Blachly-Dyson E, Peng SZ, Colombini M, Forte M (1990) Selectivity changes in site-directed mutants of the VDAC ion channel - structural implications. *Science* 247 (4947):1233–1236. doi:10.1126/science.1690454 [PubMed: 1690454]
  24. Palmieri F, De Pinto V (1989) Purification and properties of the voltage-dependent anion channel of the outer mitochondrial membrane. *J Bioenerg Biomembr* 21 (4):417–425 [PubMed: 2478528]

25. Yamamoto T, Yamada A, Watanabe M, Yoshimura Y, Yamazaki N, Yoshimura Y, Yamauchi T, Kataoka M, Nagata T, Terada H, Shinohara Y (2006) VDAC1, having a shorter N-terminus than VDAC2 but showing the same migration in an SDS-polyacrylamide gel, is the predominant form expressed in mitochondria of various tissues. *J Proteome Res* 5 (12):3336–3344. doi:10.1021/pr060291w [PubMed: 17137335]
26. Rostovtseva TK, Gurnev PA, Chen MY, Bezrukov SM (2012) Membrane lipid composition regulates tubulin interaction with mitochondrial voltage-dependent anion channel. *J Biol Chem* 287 (35):29589–29598. doi:10.1074/jbc.M112.378778 [PubMed: 22763701]
27. Sigworth FJ, Sine SM (1987) Data transformations for improved display and fitting of single-channel dwell time histograms. *Biophysical journal* 52 (6):1047–1054. doi:10.1016/S0006-3495(87)83298-8 [PubMed: 2447968]
28. Weinrich M, Worcester DL, Bezrukov SM (2017) Lipid nanodomains change ion channel function. *Nanoscale* 9 (35):13291–13297. doi:10.1039/c7nr03926c [PubMed: 28858358]
29. Colombini M (1989) Voltage gating in the mitochondrial channel, VDAC. *J Membr Biol* 111 (2):103–111 [PubMed: 2482359]
30. Rostovtseva TK, Kazemi N, Weinrich M, Bezrukov SM (2006) Voltage gating of VDAC is regulated by nonlamellar lipids of mitochondrial membranes. *J Biol Chem* 281 (49):37496–37506. doi:10.1074/jbc.M602548200 [PubMed: 16990283]
31. Teijido O, Rappaport SM, Chamberlin A, Noskov SY, Aguilera VM, Rostovtseva TK, Bezrukov SM (2014) Acidification asymmetrically affects voltage-dependent anion channel implicating the involvement of salt bridges. *J Biol Chem* 289 (34):23670–23682. doi:10.1074/jbc.M114.576314 [PubMed: 24962576]
32. Davidson SM, Yellon D, Duchon MR (2007) Assessing mitochondrial potential, calcium, and redox state in isolated mammalian cells using confocal microscopy. *Methods Mol Biol* 372:421–430. doi:10.1007/978-1-59745-365-3\_30 [PubMed: 18314743]
33. Li WW, Yang R, Guo JC, Ren HM, Zha XL, Cheng JS, Cai DF (2007) Localization of alpha-synuclein to mitochondria within midbrain of mice. *Neuroreport* 18 (15):1543–1546. doi:10.1097/WNR.0b013e3282f03db4 [PubMed: 17885598]
34. Cole NB, Dieuliis D, Leo P, Mitchell DC, Nussbaum RL (2008) Mitochondrial translocation of alpha-synuclein is promoted by intracellular acidification. *Experimental cell research* 314 (10):2076–2089. doi:10.1016/j.yexcr.2008.03.012 [PubMed: 18440504]
35. Robotta M, Gerding HR, Vogel A, Hauser K, Schildknecht S, Karreman C, Leist M, Subramaniam V, Drescher M (2014) Alpha-synuclein binds to the inner membrane of mitochondria in an alpha-helical conformation. *Chembiochem* 15 (17):2499–2502. doi:10.1002/cbic.201402281 [PubMed: 25209675]
36. Soderberg O, Gullberg M, Jarvius M, Ridderstrale K, Leuchowius KJ, Jarvius J, Wester K, Hydbring P, Bahram F, Larsson LG, Landegren U (2006) Direct observation of individual endogenous protein complexes in situ by proximity ligation. *Nat Methods* 3 (12):995–1000. doi:10.1038/nmeth947 [PubMed: 17072308]
37. Magri A, Messina A (2017) Interactions of VDAC with Proteins Involved in Neurodegenerative Aggregation: An Opportunity for Advancement on Therapeutic Molecules. *Curr Med Chem* 24 (40):4470–4487. doi:10.2174/0929867324666170601073920 [PubMed: 28571556]
38. de Kroon AI, Dolis D, Mayer A, Lill R, de Kruijff B (1997) Phospholipid composition of highly purified mitochondrial outer membranes of rat liver and *Neurospora crassa*. Is cardiolipin present in the mitochondrial outer membrane? *Biochimica et biophysica acta* 1325 (1):108–116 [PubMed: 9106488]
39. Ardail D, Lerme F, Louisot P (1990) Further characterization of mitochondrial contact sites: effect of short-chain alcohols on membrane fluidity and activity. *Biochem Biophys Res Commun* 173 (3):878–885 [PubMed: 2176492]
40. Horvath SE, Daum G (2013) Lipids of mitochondria. *Prog Lipid Res* 52 (4):590–614. doi:10.1016/j.plipres.2013.07.002 [PubMed: 24007978]
41. Rostovtseva TK, Bezrukov SM (2015) Function and Regulation of Mitochondrial Voltage-Dependent Anion Channel Electrophysiology of Unconventional Channels and Pores, vol 18 Springer, Switzerland

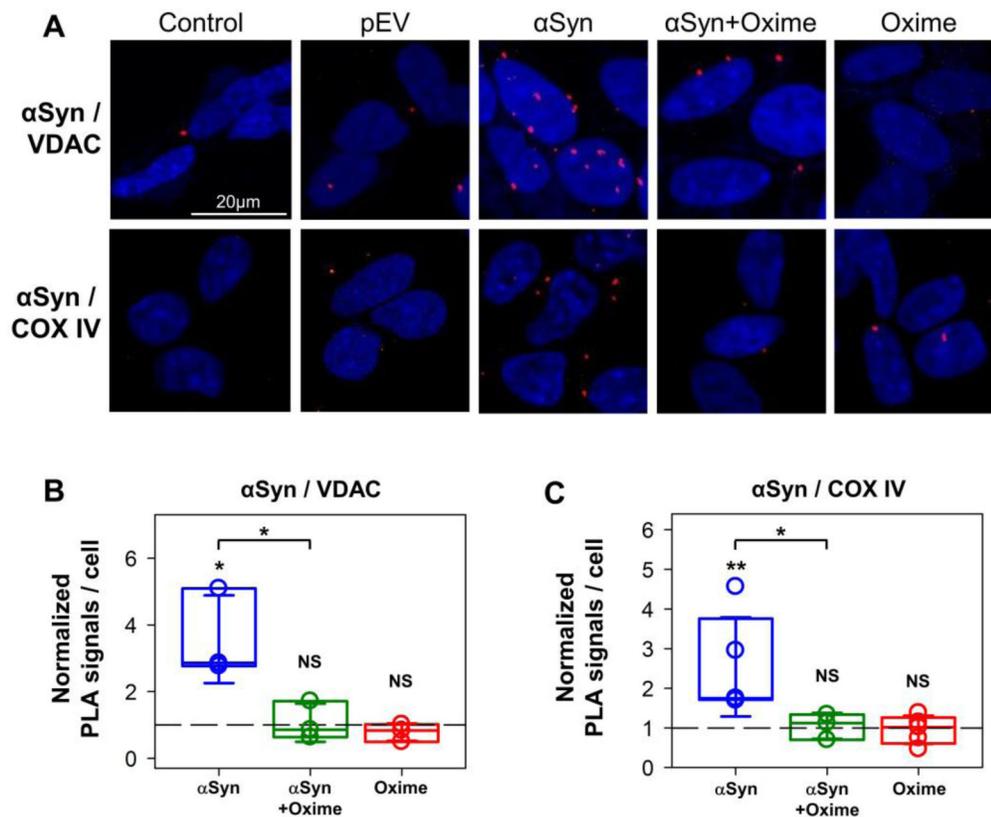
42. Kasianowicz JJ, Brandin E, Branton D, Deamer DW (1996) Characterization of individual polynucleotide molecules using a membrane channel. *Proc Natl Acad Sci U S A* 93 (24):13770–13773. doi:10.1073/pnas.93.24.13770 [PubMed: 8943010]
43. Movileanu L, Schmittschmitt JP, Scholtz JM, Bayley H (2005) Interactions of peptides with a protein pore. *Biophysical journal* 89 (2):1030–1045. doi:10.1529/biophysj.104.057406 [PubMed: 15923222]
44. Rostovtseva TK, Bezrukov SM (2012) VDAC inhibition by tubulin and its physiological implications. *Biochim Biophys Acta* 1818 (6):1526–1535. doi:S0005-2736(11)00391-9 [pii] 10.1016/j.bbame.2011.11.004 [PubMed: 22100746]
45. Rappaport SM, Teijido O, Hoogerheide DP, Rostovtseva TK, Berezhkovskii AM, Bezrukov SM (2015) Conductance hysteresis in the voltage-dependent anion channel. *Eur Biophys J* 44 (6):465–472. doi:10.1007/s00249-015-1049-2 [PubMed: 26094068]
46. Mlayeh L, Krammer EM, Leonetti M, Prevost M, Homble F (2017) The mitochondrial VDAC of bean seeds recruits phosphatidylethanolamine lipids for its proper functioning. *Biochimica et biophysica acta* 1858 (9):786–794. doi:10.1016/j.bbabi.2017.06.005 [PubMed: 28666835]
47. Queralt-Martin M, Bergdoll L, Jacobs D, Bezrukov SM, Abramson J, Rostovtseva TK (2019) Assessing the role of residue E73 and lipid headgroup charge in VDAC1 voltage gating. *Biochim Biophys Acta Bioenerg* 1860 (1):22–29. doi:10.1016/j.bbabi.2018.11.001 [PubMed: 30412693]
48. Bowen KA, Tam K, Colombini M (1985) Evidence for Titratable Gating Charges Controlling the Voltage Dependence of the Outer Mitochondrial-Membrane Channel, Vdac. *J Membrane Biol* 86 (1):51–59. doi:10.1007/Bf01871610 [PubMed: 2413210]
49. Zimmerberg J, Parsegian VA (1986) Polymer inaccessible volume changes during opening and closing of a voltage-dependent ionic channel. *Nature* 323 (6083):36–39 [PubMed: 2427958]
50. Eckmann J, Clemens LE, Eckert SH, Hagl S, Yu-Taeger L, Bordet T, Pruss RM, Muller WE, Leuner K, Nguyen HP, Eckert GP (2014) Mitochondrial membrane fluidity is consistently increased in different models of Huntington disease: restorative effects of olesoxime. *Mol Neurobiol* 50 (1):107–118. doi:10.1007/s12035-014-8663-3 [PubMed: 24633813]
51. Magalon K, Le Grand M, El Waly B, Moulis M, Pruss R, Bordet T, Cayre M, Belenguer P, Carre M, Durbec P (2016) Olesoxime favors oligodendrocyte differentiation through a functional interplay between mitochondria and microtubules. *Neuropharmacology* 111:293–303. doi:10.1016/j.neuropharm.2016.09.009 [PubMed: 27618742]
52. Weinrich M, Rostovtseva TK, Bezrukov SM (2009) Lipid-dependent effects of halothane on gramicidin channel kinetics: a new role for lipid packing stress. *Biochemistry* 48 (24):5501–5503. doi:10.1021/bi900494y [PubMed: 19405539]
53. Andersen OS, Nielsen C, Maer AM, Lundbaek JA, Goulian M, Koeppe RE 2nd (1999) Ion channels as tools to monitor lipid bilayer-membrane protein interactions: gramicidin channels as molecular force transducers. *Methods Enzymol* 294:208–224 [PubMed: 9916229]
54. Lundbaek JA, Andersen OS (1999) Spring constants for channel-induced lipid bilayer deformations. Estimates using gramicidin channels. *Biophysical journal* 76 (2):889–895. doi:10.1016/S0006-3495(99)77252-8 [PubMed: 9929490]
55. Lundbaek JA, Birn P, Girshman J, Hansen AJ, Andersen OS (1996) Membrane stiffness and channel function. *Biochemistry* 35 (12):3825–3830. doi:10.1021/bi952250b [PubMed: 8620005]
56. Rovini A, Carre M, Bordet T, Pruss RM, Braguer D (2010) Olesoxime prevents microtubule-targeting drug neurotoxicity: selective preservation of EB comets in differentiated neuronal cells. *Biochem Pharmacol* 80 (6):884–894. doi:10.1016/j.bcp.2010.04.018 [PubMed: 20417191]
57. Martin LJ (2010) Olesoxime, a cholesterol-like neuroprotectant for the potential treatment of amyotrophic lateral sclerosis. *IDrugs* 13 (8):568–580 [PubMed: 20721828]
58. Gonzalez S, Berthelot J, Jiner J, Perrin-Tricaud C, Fernando R, Chrast R, Lenaers G, Tricaud N (2016) Blocking mitochondrial calcium release in Schwann cells prevents demyelinating neuropathies. *J Clin Invest* 126 (7):2773. doi:10.1172/JCI88179
59. Xiao WH, Zheng FY, Bennett GJ, Bordet T, Pruss RM (2009) Olesoxime (cholest-4-en-3-one, oxime): analgesic and neuroprotective effects in a rat model of painful peripheral neuropathy produced by the chemotherapeutic agent, paclitaxel. *Pain* 147 (1-3):202–209. doi:10.1016/j.pain.2009.09.006 [PubMed: 19833436]

60. Parihar MS, Parihar A, Fujita M, Hashimoto M, Ghafourifar P (2009) Alpha-synuclein overexpression and aggregation exacerbates impairment of mitochondrial functions by augmenting oxidative stress in human neuroblastoma cells. *Int J Biochem Cell Biol* 41 (10):2015–2024. doi:10.1016/j.biocel.2009.05.008 [PubMed: 19460457]
61. Vekrellis K, Xilouri M, Emmanouilidou E, Stefanis L (2009) Inducible over-expression of wild type alpha-synuclein in human neuronal cells leads to caspase-dependent non-apoptotic death. *J Neurochem* 109 (5):1348–1362. doi:10.1111/j.1471-4159.2009.06054.x [PubMed: 19476547]
62. Banerjee K, Sinha M, Pham Cle L, Jana S, Chanda D, Cappai R, Chakrabarti S (2010) Alpha-synuclein induced membrane depolarization and loss of phosphorylation capacity of isolated rat brain mitochondria: implications in Parkinson's disease. *FEBS Lett* 584 (8):1571–1576. doi:10.1016/j.febslet.2010.03.012 [PubMed: 20226185]
63. Bir A, Sen O, Anand S, Khemka VK, Banerjee P, Cappai R, Sahoo A, Chakrabarti S (2014) alpha-Synuclein-induced mitochondrial dysfunction in isolated preparation and intact cells: implications in the pathogenesis of Parkinson's disease. *J Neurochem* 131 (6):868–877. doi:10.1111/jnc.12966 [PubMed: 25319443]
64. Esteves AR, Gozes I, Cardoso SM (2014) The rescue of microtubule-dependent traffic recovers mitochondrial function in Parkinson's disease. *Biochimica et biophysica acta* 1842 (1):7–21. doi:10.1016/j.bbadis.2013.10.003 [PubMed: 24120997]
65. Ludtmann MHR, Angelova PR, Horrocks MH, Choi ML, Rodrigues M, Baev AY, Berezhnov AV, Yao Z, Little D, Banushi B, Al-Menhali AS, Ranasinghe RT, Whiten DR, Yapom R, Dolt KS, Devine MJ, Gissen P, Kunath T, Jaganjac M, Pavlov EV, Klenerman D, Abramov AY, Gandhi S (2018) alpha-synuclein oligomers interact with ATP synthase and open the permeability transition pore in Parkinson's disease. *Nat Commun* 9 (1):2293. doi:10.1038/s41467-018-04422-2 [PubMed: 29895861]
66. Liu J, Wang X, Lu Y, Duan C, Gao G, Lu L, Yang H (2017) Pink1 interacts with alpha-synuclein and abrogates alpha-synuclein-induced neurotoxicity by activating autophagy. *Cell Death Dis* 8 (9):e3056. doi:10.1038/cddis.2017.427 [PubMed: 28933786]
67. Martin LJ, Pan Y, Price AC, Sterling W, Copeland NG, Jenkins NA, Price DL, Lee MK (2006) Parkinson's disease alpha-synuclein transgenic mice develop neuronal mitochondrial degeneration and cell death. *J Neurosci* 26 (1):41–50. doi:10.1523/JNEUROSCI.4308-05.2006 [PubMed: 16399671]
68. Vicario M, Cieri D, Brini M, Cali T (2018) The Close Encounter Between Alpha-Synuclein and Mitochondria. *Front Neurosci* 12:388. doi:10.3389/fnins.2018.00388 [PubMed: 29930495]
69. Hill K, Model K, Ryan MT, Dietmeier K, Martin F, Wagner R, Pfanner N (1998) Tom40 forms the hydrophilic channel of the mitochondrial import pore for preproteins [see comment]. *Nature* 395 (6701):516–521. doi:10.1038/26780 [PubMed: 9774109]
70. Suzuki H, Kadowaki T, Maeda M, Sasaki H, Nabekura J, Sakaguchi M, Mihara K (2004) Membrane-embedded C-terminal segment of rat mitochondrial TOM40 constitutes protein-conducting pore with enriched beta-structure. *J Biol Chem* 279 (48):50619–50629 [PubMed: 15347672]
71. Kuszak AJ, Jacobs D, Gurnev PA, Shiota T, Louis JM, Lithgow T, Bezrukov SM, Rostovtseva TK, Buchanan SK (2015) Evidence of Distinct Channel Conformations and Substrate Binding Affinities for the Mitochondrial Outer Membrane Protein Translocase Pore Tom40. *J Biol Chem* 290 (43):26204–26217. doi:10.1074/jbc.M115.642173 [PubMed: 26336107]
72. O'Leary EI, Jiang Z, Strub MP, Lee JC (2018) Effects of phosphatidylcholine membrane fluidity on the conformation and aggregation of N-terminally acetylated alpha-synuclein. *J Biol Chem* 293 (28):11195–11205. doi:10.1074/jbc.RA118.002780 [PubMed: 29853639]
73. Pfefferkorn CM, Jiang Z, Lee JC (2012) Biophysics of alpha-synuclein membrane interactions. *Biochimica et biophysica acta* 1818 (2):162–171. doi:10.1016/j.bbame.2011.07.032 [PubMed: 21819966]
74. Rhoades E, Ramlall TF, Webb WW, Eliezer D (2006) Quantification of alpha-synuclein binding to lipid vesicles using fluorescence correlation spectroscopy. *Biophysical journal* 90 (12):4692–4700. doi:10.1529/biophysj.105.079251 [PubMed: 16581836]
75. Eddy MT, Ong TC, Clark L, Tejjido O, van der Wel PC, Garces R, Wagner G, Rostovtseva TK, Griffin RG (2012) Lipid dynamics and protein-lipid interactions in 2D crystals formed with the

- beta-barrel integral membrane protein VDAC1. *J Am Chem Soc* 134 (14):6375–6387. doi:10.1021/ja300347v [PubMed: 22435461]
76. Srivastava SR, Zadafiya P, Mahalakshmi R (2018) Hydrophobic Mismatch Modulates Stability and Plasticity of Human Mitochondrial VDAC2. *Biophys J*. doi:10.1016/j.bpj.2018.11.001
77. Naghdi S, Varnai P, Hajnoczky G (2015) Motifs of VDAC2 required for mitochondrial Bak import and tBid-induced apoptosis. *Proc Natl Acad Sci U S A* 112 (41):E5590–5599. doi:10.1073/pnas.1510574112 [PubMed: 26417093]
78. Chin HS, Li MX, Tan IKL, Ninnis RL, Reljic B, Scicluna K, Dagley LF, Sandow JJ, Kelly GL, Samson AL, Chappaz S, Khaw SL, Chang C, Morokoff A, Brinkmann K, Webb A, Hockings C, Hall CM, Kueh AJ, Ryan MT, Kluck RM, Bouillet P, Herold MJ, Gray DHD, Huang DCS, van Delft MF, Dewson G (2018) VDAC2 enables BAX to mediate apoptosis and limit tumor development. *Nat Commun* 9 (1):4976. doi:10.1038/s41467-018-07309-4 [PubMed: 30478310]
79. Smilansky A, Dangoor L, Nakdimon I, Ben-Hail D, Mizrahi D, Shoshan-Barmatz V (2015) The Voltage-dependent Anion Channel 1 Mediates Amyloid beta Toxicity and Represents a Potential Target for Alzheimer Disease Therapy. *J Biol Chem* 290 (52):30670–30683. doi:10.1074/jbc.M115.691493 [PubMed: 26542804]

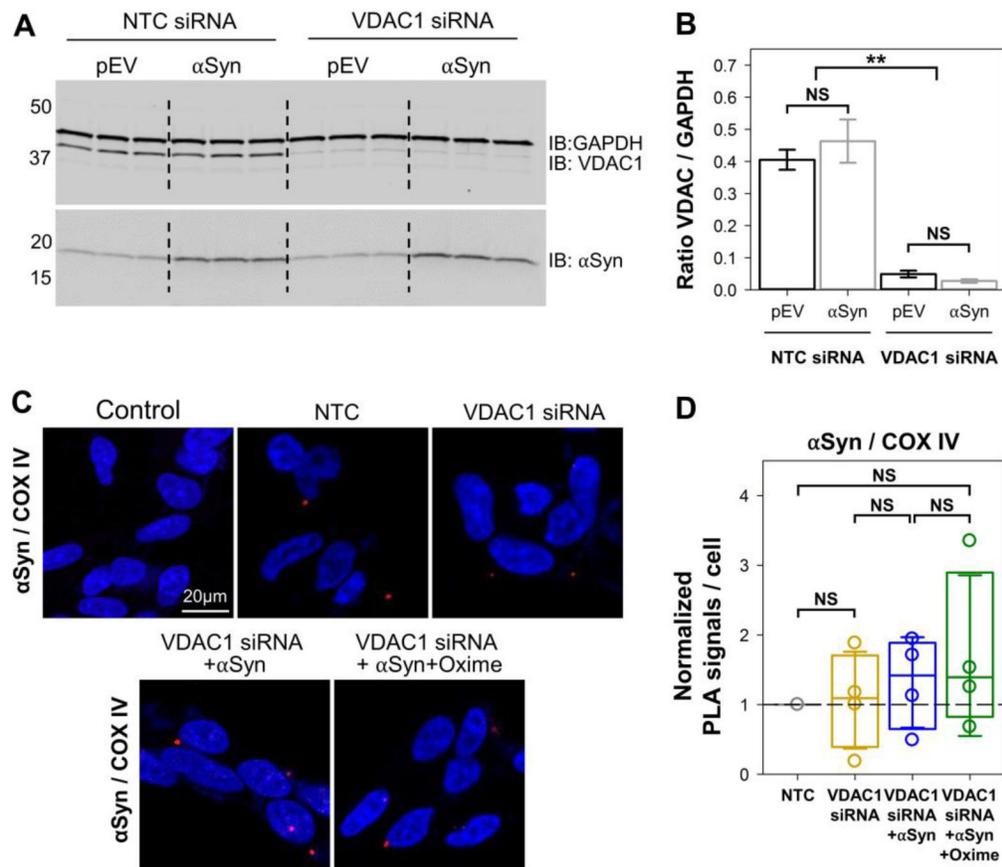


**Fig. 1. Olesoxime preserves mitochondrial integrity in a cell model overexpressing  $\alpha$ Syn.** (A) Phase contrast representative images (20x objective, scale bar 100 $\mu$ m) show morphological alterations such as reduced processes (arrow heads) and rounded floating cells (stars), following transient transfection of wt  $\alpha$ Syn in differentiated SH-SY5Y cells. Olesoxime treatment restores cell morphology. (B) Box plot showing quantification of cell death assessed 96 h after transfection by Trypan blue exclusion assay. Here and in Figs. 2 B,C and 3D the boundary of the box closest to zero indicates the 25<sup>th</sup> percentile, a line within the box marks the median, and the boundary of the box farthest from zero indicates the 75<sup>th</sup> percentile. Error bars indicate S.D. (n=4). (C) Representative images of cells loaded with MitoTracker™ CMXRos to evaluate mitochondrial membrane potential for which the relative fluorescence intensity changes are quantified as ratios over pEV control in (D) (63x objective, scale bar 20 $\mu$ m). In D data is presented as mean  $\pm$  S.D. (n = 4). In B and D, NS (not significant):  $p > 0.5$ ; \*\*:  $p < 0.01$ ; one-way ANOVA



**Fig. 2. Effect of olesoxime on  $\alpha$ Syn localization at the mitochondrial membranes.**

(A) Representative confocal images of PLA showing  $\alpha$ Syn in close proximity ( $\sim 30$  nm) to VDAC at the MOM and with Complex IV (COX IV) at the MIM in cells overexpressing  $\alpha$ Syn before and after olesoxime treatment. The nuclei are stained in blue (DAPI) (63x objective, scale bar 20 $\mu$ m). Controls omitting the primary antibodies are shown for each pairwise combination. (B, C) Box plots represent the normalized distribution of signals per cell, obtained from an average of 160 cells per experiment with a minimum of 60. Error bars indicate S.D. The PLA signals were normalized to the average PLA values in pEV control cells of the corresponding experiment. The dashed lines indicate pEV controls. At least 3 independent experiments were performed for each pairwise of antibodies. Unless indicated by brackets, significance was tested against the corresponding pEV controls (\*\*:  $p < 0.01$ , \*:  $p < 0.05$ , NS (not significant):  $p > 0.5$ ; one-way ANOVA)



**Fig. 3.  $\alpha$ Syn colocalization with COX IV at the inner membrane is lost upon downregulation of VDAC1 expression.**

(A) Western blot of VDAC1 protein levels in differentiated SH-SY5Y cells transfected with a non-targeting sequence (NTC) siRNA or VDAC1-siRNA. GAPDH was used as a loading control and maintenance of  $\alpha$ Syn overexpression under these conditions has been validated. (B) Quantification of the Western blot analysis as intensity ratios of VDAC1/GAPDH. Data is presented as mean  $\pm$  S.D. (n = 3). In VDAC1-siRNA cells, the VDAC1/GAPDH ratio is significantly less than in NTC control. (C) Representative confocal images of PLA of VDAC1 and COX IV colocalization in controls and cells depleted of VDAC1, overexpressing  $\alpha$ Syn alone or treated with olesoxime. The nuclei are stained in blue (DAPI) (63x objective, scale bar 20 $\mu$ m). (D) Box plot represent the normalized distribution of PLA signals per cell, obtained from an average of 180 cells per experiment with a minimum of 60. The PLA signals were normalized to averaged PLA value measured in the NTC control cells of the corresponding experiment. The dashed line indicates a normalized PLA signal in NTC cells. Error bars indicate S.D. At least 3 independent experiments were performed for each pairwise of antibodies. In B and D, \*\*:  $p < 0.01$ ; NS (not significant):  $p \geq 0.2$ ; one-way ANOVA.



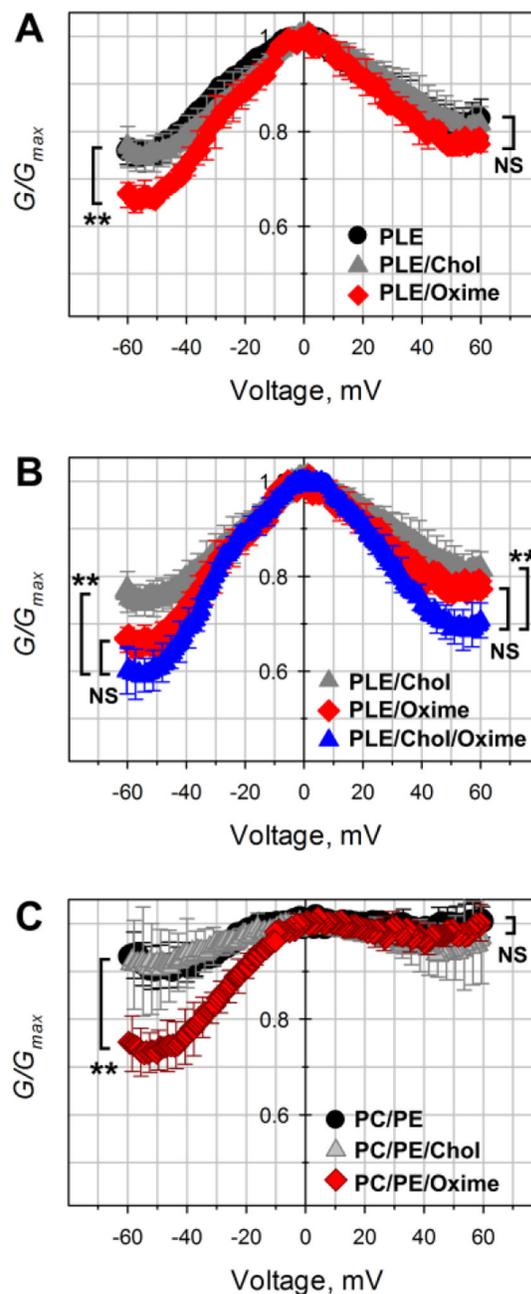
membranes) corresponds to regime of  $\alpha$ Syn translocation through the pore. In the presence of olesoxime, translocation of  $\alpha$ Syn is not observed at applied voltages up to  $-48$  mV, while in PC/PE membranes with or without cholesterol, the translocation regime starts at  $-35$  mV. Data points and error bars represent the mean and S.D. for 4 - 6 independent experiments.

Author Manuscript

Author Manuscript

Author Manuscript

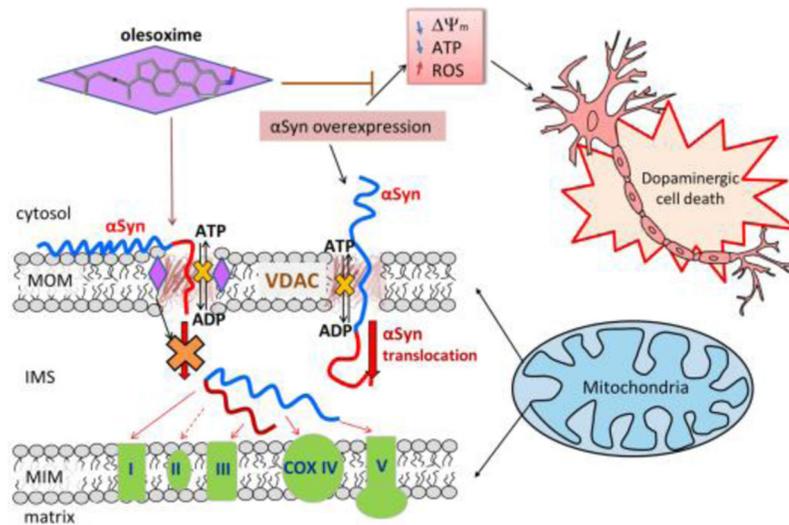
Author Manuscript



**Fig. 5. Olesoxime increases VDAC voltage gating.**

Characteristic bell-shaped plots of normalized average VDAC conductance as function of the applied voltage obtained on planar bilayers of different lipid compositions. (A) PLE with olesoxime (20% w/w) (PLE/Oxime) or cholesterol (4% w/w) (PLE/Chol) or without either of them (PLE). (B) PLE with cholesterol (4% w/w) (PLE/Chol) or olesoxime (20% w/w) (PLE/Oxime) or both (4% w/w and 16% w/w, respectively) (PLE/Chol/Oxime). (C) PC/PE (1:4) with cholesterol (10% w/w) (PC/PE/Chol) or olesoxime (10% w/w) (PC/PE/Oxime) or without either of them (PC/PE). Data were obtained on multichannel membranes with 10-200 reconstituted VDAC channels. Normalized conductance is defined as  $G/G_{max}$ , where

$G_{max}$  is the maximum conductance at voltages close to 0 mV. Membrane-bathing solutions consisted of 1 M KCl buffered with 5 mM HEPES at pH 7.4. Data are means of 3-5 experiments  $\pm$  S.D. (error bars shown every 5 points for clarity). To check significance (\*\*:  $p < 0.01$ , NS (not significant):  $p > 0.05$ ; one-way ANOVA), the minimum  $G/G_{max}$  value of each data set was used, corresponding to  $|V| \sim 52$  mV. Each voltage polarity was tested independently.



**Fig. 6. The proposed model of olesoxime neuroprotective effect.**

When  $\alpha$ Syn is captured by the VDAC pore it disrupts ATP/ADP fluxes through VDAC. Under normal conditions endogenous  $\alpha$ Syn regulates these fluxes by reversibly and dynamically blocking VDAC. Under  $\alpha$ Syn overexpression-induced stress  $\alpha$ Syn translocates across MOM through VDAC and targets ETC in the MIM causing their impairment, mitochondrial dysfunction and eventually neuronal cell death. Olesoxime partitions into the MOM and hinders  $\alpha$ Syn translocation through VDAC by interacting with the pore-lipid interface. This model suggests a tentative mechanism of olesoxime protection of mitochondria integrity and promotion of neuronal cell survival.