Lanosterol induces mitochondrial uncoupling and protects dopaminergic neurons from cell death in a model for Parkinson’s disease

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Parkinson’s disease (PD) is a neurodegenerative disorder marked by the selective degeneration of dopaminergic neurons in the nigrostriatal pathway. Several lines of evidence indicate that mitochondrial dysfunction contributes to its etiology. Other studies have suggested that alterations in sterol homeostasis correlate with increased risk for PD. Whether these observations are functionally related is, however, unknown. In this study, we used a toxin-induced mouse model of PD and measured levels of nine sterol intermediates. We found that lanosterol is significantly (~50%) and specifically reduced in the nigrostriatal regions of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine-treated mice, indicative of altered lanosterol metabolism during PD pathogenesis. Remarkably, exogenous addition of lanosterol rescued dopaminergic neurons from 1-methyl-4-phenylpyridinium (MPP+) induced cell death in culture. Furthermore, we observed a marked redistribution of lanosterol synthase from the endoplasmic reticulum to mitochondria in dopaminergic neurons exposed to MPP+, suggesting that lanosterol might exert its survival effect by regulating mitochondrial function. Consistent with this model, we find that lanosterol induces mild depolarization of mitochondria and promotes autophagy. Collectively, our results highlight a novel sterol-based neuroprotective mechanism with direct relevance to PD.

Received 01.3.11; revised 30.6.11; accepted 04.7.11; Edited by N Bazan; published online 05.8.11

Lanosterol induces mitochondrial uncoupling and protects dopaminergic neurons from cell death in a model for Parkinson’s disease.

Parkinson’s disease (PD) is a movement disorder marked by the selective degeneration of dopaminergic neurons in the nigrostriatal pathway. About 5–10% of PD are genetically linked with mutations in genes such as leucine-rich repeat kinase 2 (LRRK2), alpha-synuclein, PTEN-induced putative kinase 1 (PINK1), Parkin and DJ-1.1 These genetic PD cases, while rare, have provided insights into mechanisms of PD pathogenesis and many of which point toward mitochondrial dysfunction.1–3 For example, PINK1, Parkin and DJ-1 control clearance of mitochondria by mitophagy in response to cellular stress.3 Mitochondrial defects are also seen in idiopathic PD, whereby the catalytic activity of brain mitochondrial complex I is compromised.4 Finally, environmental toxins that affect complex I, such as 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP),5 can also induce Parkinsonism. The active metabolite of MPTP, 1-methyl-4-phenylpyridinium (MPP+), selectively enters dopaminergic neurons via the dopamine transporter, inhibits mitochondrial complex I and eventually induces clinical symptoms reminiscent of PD.

As MPTP/MPP+ toxicity emulate PD symptoms, it has been widely used in animal and cellular models to study neuronal cell death and to screen for neuroprotective agents. Of the neuroprotective metabolites identified, many are found in mitochondria, including L-carnitine, creatine and coenzyme Q (CoQ)-10.6 Although their precise protective mechanisms are still poorly understood, some evidence suggests that they act as mitochondrial uncouplers.7 Consistent with this model, uncoupling proteins (UCPs) are protective in the MPTP model of PD.8,9 and their expression is downregulated in DJ-1 knockout mice.10 Neuronal cell death induced by glutamate toxicity can also be rescued by mitochondrial uncouplers.11 More recently, misregulation of sterol metabolism has also been implicated in PD. In clinical studies, elevation of

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Keywords: lanosterol; Parkinson’s disease; mitochondrial membrane potential; lipids; dopaminergic neurons

Abbreviations: ATP, adenosine triphosphate; AV, autophagosome vacuole; GCPP, m-chlorophenylhydrazone; CCD, charge-coupled device; cdk5, cyclin-dependent kinase 5; CoQ, coenzyme Q; ubiquinone; DIV, days in vitro; DJ-1, (PARK7) Parkinson disease 7; ER, endoplasmic reticulum; GC–MS, gas chromatography–mass spectrometry; GSK-3β, glycogen synthase kinase 3 beta; HPLC, high-pressure liquid chromatography; JCI, 5,5′,6,6′-tetrachloro-1,1′-3,3′-tetrachlorobenzimidazo-locarbocyanine iodide; KDEL, ER retention sequence (lys-asp-glu-leu); LC3, microtubule-associated protein light chain 3; LDL-C, low-density lipoprotein cholesterol; LRRK2, leucine-rich repeat kinase 2; LSS, lanosterol synthase; MPP+, 1-methyl-4-phenylpyridinium; MPTP, 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine; MMN, multiple reaction monitoring; p35, cdk5 activator protein with 35 kDa; PBS, phosphate-buffered saline; PC, phosphatidylcholine; PD, Parkinson’s disease; PINK1, PTEN-induced putative kinase 1; SREBP2, sterol regulatory element-binding protein 2; TH, tyrosine hydroxylase; TOMM20, translocase of outer mitochondrial membrane 20; TUJ1, neuron-specific class III beta tubulin; UCP, uncoupling protein; 1/9, mitochondrial membrane potential

Received 01.3.11; revised 30.6.11; accepted 04.7.11; Edited by N Bazan; published online 05.8.11
low-density lipoprotein cholesterol (LDL-C) in serum correlates with higher prevalence of PD, whereas higher serum levels of total cholesterol are associated with a decreased risk of PD. Biochemical studies showed that alpha-synuclein, the major component of Lewy bodies found in PD brains, binds to cholesterol with high affinity and its aggregation is accelerated in presence of oxidized cholesterol. In a separate line of investigation, oxidative stress has been shown to increase levels of lanosterol, a cholesterol precursor, in mitochondria and several other intracellular compartments of macrophages, suggesting that this sterol metabolite may be part of a global cellular response to stress. Yet, how sterol metabolism is altered in the brains of PD patients, or which sterol metabolites, if any, have an impact on the survival of dopaminergic neurons is unknown.

Our study investigated the role of sterol metabolism in the MPTP/MPP+ model of PD. Our results provide evidence for the specific role of lanosterol as a neuroprotective agent in dopaminergic neurons. We also show that lanosterol induces mild uncoupling of mitochondria and promotes autophagy, two events that have been previously linked to neuroprotection in various models of PD. To our knowledge, our work provides the first link between sterol metabolism and mitochondrial function, and identifies lanosterol as a potential therapeutic agent for PD.

Results

Lanosterol levels are decreased in the striatum and ventral midbrain of MPTP-injected mice. To determine if sterol metabolism is altered in a rodent model of PD, we measured levels of sterol metabolites in the striatum and ventral midbrain of control and MPTP-treated mice by gas chromatography-mass spectrometry (GC–MS). This acute MPTP injection regime results in about 35% loss of dopaminergic neurons after 48 h. Of the nine metabolites analyzed, we found that lanosterol, the first cyclic sterol, was reduced by ~50% in affected areas (Figure 1). Owing to the specific and highly significant reduction in the levels of lanosterol in the affected areas of MPTP-treated animals, we reasoned that lanosterol might be important for dopaminergic neuronal survival.

Lanosterol protects dopaminergic neurons from MPP+-induced cell death. We asked whether exogenous addition of lanosterol protects dopaminergic neurons against MPP+-induced cell death. Primary postnatal neuron cultures from the ventral midbrain were treated with MPP+, and the survival of dopaminergic neurons was determined in the absence or presence of exogenously added lanosterol. In all, 48% of dopaminergic neurons survived on treatment with MPP+ under these culture conditions (Figures 2a and c). Co-incubation of MPP+ with phosphatidylcholine (PC, vehicle control) or cholesterol did not improve survival of dopaminergic neurons (Figures 2b and c). In contrast, co-treatment of the cultures with MPP+ and lanosterol increased survival of dopaminergic neurons to 76% (Figures 2b and c). Thus, lanosterol, but not cholesterol or PC, rescues dopaminergic neurons from MPP+-induced cell death.

To determine if addition of lanosterol in culture affects other sterol intermediates, we analyzed an array of metabolites (squalene, lanosterol, lathosterol, 7-dehydrocholesterol, desmosterol, cholesterol, 7-beta hydroxycholesterol, 7-ketocholesterol and 24-hydroxycholesterol) in extracts from ventral midbrain cultures by GC–MS. In cells treated with lanosterol, we observed a 12-fold increase in lanosterol levels relative to control (no treatment, Figures 2d and e) and no detectable effect on other sterol metabolites tested (Figure 2e), suggesting that cells have a high capacity to accumulate lanosterol, consistent with previous findings in macrophages. This is not the case for cholesterol, as we did not detect any changes in cholesterol levels on exogenous addition of cholesterol, perhaps because neuronal cholesterol homeostasis is tightly regulated. In addition to the metabolites indicated above, we also tested the impact of lanosterol and cholesterol addition on levels of ubiquinones, isoprenoid-derived electron carriers in mitochondria, and found no significant differences (Figures 3a and b).

In these experiments, postnatal dopaminergic neurons were grown in direct contact with a glia feeder. It is therefore possible that neuroprotection is indirectly mediated by a modification of astrocyte physiology. To address whether lanosterol addition leads to a specific increase in lanosterol concentration in neurons, we repeated these experiments using hippocampal neurons cultured according to Banker’s method, which allows physical separation of neurons and astrocytes. We found that lanosterol levels increased in both neurons and astrocytes to a similar extent on treatment with exogenous lanosterol (data not shown). Although we cannot rule out the possible contribution of astrocytes in lanosterol-mediated neuroprotection, our results indicate that lanosterol levels are significantly elevated in neurons, arguing for a direct role of lanosterol in promoting survival of dopaminergic neurons.
To explore the mechanisms underlying lanosterol-mediated neuroprotection, we next investigated the effects of MPP⁺ and lanosterol on various signaling pathways previously implicated in cellular metabolism and neurodegeneration. First, we examined the expression level of sterol response element-binding protein (SREBP2, Figure 4a), a transcription factor reported to exacerbate neuronal degeneration. 20 We found that MPP⁺ induced a modest increase in SREBP2 levels, which was not rescued by lanosterol addition (Figure 4a, top panel). We also checked for levels of cdk5 activator protein with 35 kDa (p35), the activator of cyclin-dependent kinase 5 (cdk5), because the genetic ablation of p35/cdk5 confers neuroprotection in the MPTP model. 21 Although we observed a decrease in p35 levels in MPP⁺-treated cells, addition of lanosterol did not restore p35 expression to control levels (Figure 4a, second panel).
from top). Consistent with previous findings implicating glycogen synthase kinase 3 beta (Gsk-3/β) (Ser9) phosphorylation in PD pathogenesis, we found increased phospho-Gsk-3/β in MPP+ treated cells. However, this increase was not affected by the addition of lanosterol (Figure 4a, third panel from top).

Finally, we examined the levels of lanosterol synthase (LSS), the enzyme that catalyzes cyclization of oxidosqualene, the rate limiting step in lanosterol synthesis. The levels of LSS did not markedly change on MPP+ addition (Figure 4a, fourth panel from top). Interestingly, however, we observed a different localization pattern of LSS in MPP+ -treated dopaminergic neurons. LSS immunostaining appeared much more punctate after MPP+ treatment (Figure 4b), suggesting drug-induced redistribution of LSS to a different intracellular compartment. We thus proceeded to examine the subcellular localization of LSS in control and MPP+ -treated dopaminergic neurons.

**Redistribution of LSS in dopaminergic neurons on MPP+ insult.** LSS is a membrane-associated enzyme, which is targeted to the cytoplasmic leaflet of the endoplasmic reticulum (ER). Biochemical analysis of purified fractions from mouse liver showed that the bulk of LSS is indeed associated with the microsomal/ER fractions and that minor amounts are found in purified mitochondria (Figure 4c).

Consistent with our biochemical fractionation data, we found that LSS colocalized with an ER marker (ER retention sequence (lys-asp-glu-leu) (KDEL)) in both dopaminergic and non-dopaminergic neurons (Figure 5a). Extensive overlap between LSS and the ER marker was observed for both dopaminergic and non-dopaminergic neurons ($R^2 = 0.83 \pm 0.01$ and $0.84 \pm 0.01$, respectively). Less overlap was found between LSS and translocase of outer mitochondrial membrane 20 (TOMM20) (a mitochondrial marker) for both dopaminergic and non-dopaminergic neurons ($R^2 = 0.70 \pm 0.02$ and $0.69 \pm 0.02$, respectively, Figure 5a, second panel from the top). This pattern changed noticeably in dopaminergic neurons treated with MPP+ . The colocalization of LSS with TOMM20 was markedly increased ($R^2 = 0.856 \pm 0.01$, $0.70 \pm 0.02$, and $0.69 \pm 0.02$, respectively, Figure 5a, second panel from the bottom). Translocation of LSS from the ER to mitochondria was also observed in dopaminergic neurons co-treated with MPP+ and lanosterol (Figure 5b). MPP+ did not, however, affect LSS localization in non-dopaminergic neurons (Figure 5b), consistent with the inability of these cells to take up MPP+. These data indicate that a significant fraction of LSS redistributes from the ER to mitochondria in dopaminergic neurons exposed to MPP+, suggesting that LSS (and its product lanosterol) may have a role in regulating mitochondrial function.

**Lanosterol uncouples mitochondria in dopaminergic neurons.** An important aspect of mitochondrial physiology can be assessed by measuring the mitochondrial membrane potential ($\Delta \Psi$), which is the electrical and chemical gradient that drives protons across the inner membrane during electron transport and oxidative phosphorylation. This process is never completely ‘coupled’, allowing a fraction of electrons and protons to be transported without concurrent production of adenosine triphosphate (ATP). We first examined the impact of lanosterol on $\Delta \Psi$ in hippocampal neurons using the ratiometric voltage-sensitive fluorescent dye, 5,5′,6,6′-tetrachloro-1,1′,3′,3′-tetraethylbenzimidazolocarbocyanine iodide (JC-1). Using time-lapse, multi-positioning imaging to capture about 3–4 fields in 30-s intervals, we measured changes of $\Delta \Psi$ on lipids addition by the ratiometric analysis of JC-1 red to green emission. As expected, treatment of neurons with 200 nM m-chlorophenylhydrazine (CCCP), a known uncoupler of oxidative phosphorylation, induced an immediate and sharp reduction of the mitochondrial membrane potential ($\sim 25\%$ within 30 s; Figure 6a). Under these same experimental conditions, exogenous addition of lanosterol reduced the membrane potential by $\sim 20\%$ over 15 min, whereas PC and cholesterol had no significant effect (Figure 6b).

We repeated these experiments in ventral midbrain cultures (Figures 6c and d). As these cultures were grown at a lower density than the hippocampal cultures, we needed to capture 10–15 fields in time-lapse multi-positioning imaging at lower

**Figure 3** Measurements of ubiquinone on addition of lipids. (a) Simplified pathway to show the cross-talk between sterol and ubiquinone biosynthesis in mammalian cells. Both pathways use isoprenyl units as precursors. (b) Plot of ubiquinone/CoQ levels normalized to total cholesterol levels in neuronal cultures from various treatment conditions. Error bars represent S.E.M. over four independent experiments.
magnification to gain enough statistical power. As a consequence, this limited time intervals to 2 min. We observed a ~20% of reduction in membrane potential on lanosterol addition (Figures 6c and d), consistent with the results obtained in hippocampal neurons.

As MPP⁺ uptake depends on ATP levels, it is possible that the addition of lanosterol causes mitochondrial depolarization, and subsequently ATP depletion. This would, in turn, inhibit MPP⁺ uptake in the mitochondria. To determine if lanosterol alters ATP levels, we measured levels of ATP in

**Figure 4** Analyses of SREBP2, Gsk-3β, p35/ck5 and LSS in ventral midbrain treated with lipids and MPP⁺. (a) Immunoblot analyses of several factors previously linked to neuroprotection in PD and/or MPTP treatment. (b) Immunofluorescence images of ventral midbrain neurons stained with LSS and TH treated with or without MPP⁺. LSS appears more punctate in TH⁺ neurons on MPP⁺ treatment. (c) Subcellular fractionation of liver tissue. M, mitochondrial, ER, microsome/ER, C, cytosolic, WL, fractions were purified from whole liver. Immunoblot of calnexin (ER marker) and VDAC/porin (mitochondrial marker) were used to assess the purity of each fraction. LSS showed a clean single band at the expected weight of 78 kDa and is enriched in the ER fraction. A small amount of LSS is also detected in the mitochondrial fraction.

**Figure 5** LSS is redistributed from ER to mitochondria in dopaminergic neurons on addition of MPP⁺. (a) Confocal images of ventral midbrain neurons stained with TH (white), LSS (green), either KDEL (red, first and third panels from the top) or TOMM20 (red, second and fourth panels from the top) of ventral midbrain neurons in control and MPP⁺ treated conditions. White boxes show dopaminergic neurons as determined by TH⁺ staining. Right panels show pixel intensity correlation plots of LSS with either KDEL or TOMM20 in dopaminergic neurons. Scale bar represents 10 μm. (b) Average R² values (a measure of colocalization) of two classes of neurons in control and MPP⁺-treated conditions. In control, R² values from co-staining of KDEL-LSS, n = 26 and n = 27, TOMM20-LSS, n = 20 and n = 21, were assessed for dopaminergic and non-dopaminergic neurons respectively. In MPP⁺-treated cells, R² values from co-staining of KDEL-LSS, n = 20 and n = 20, for TOMM20-LSS, n = 18 and n = 17, were assessed for dopaminergic and non-dopaminergic neurons respectively. In MPP⁺-lanosterol co-treated cells, R² values from co-staining of KDEL-LSS, n = 18 and n = 23, for TOMM20-LSS, n = 19 and n = 18, were assessed for dopaminergic and non-dopaminergic neurons respectively. ****P<0.00001
neurons treated with PC, cholesterol or lanosterol. We did not see any significant changes in the levels of ATP across all treatment conditions (Supplementary Figure S1), indicating that lanosterol is unlikely to inhibit MPP⁺ uptake. Thus, one mechanism by which lanosterol could mediate neuroprotection is through the uncoupling of mitochondria.

**Figure a**

- **TH**
- **LSS**
- **KDEL**
- **LSS-KDEL-DAPI**

**Figure b**

- **TOMM20-LSS**
- **KDEL-LSS**

**Graph**

- **TH⁺ neuron**
- **TH⁺ neuron**
- **TH⁺ neuron**
- **TH⁺ neuron**

**Legend**

- **Control**
- **MPP⁺**
- **Lanosterol & MPP⁺**

**Images**

- **Intensity in TOMM20 channel (red)**
- **Intensity in KDEL channel (red)**
- **Intensity in LSS channel (green)**

**R² values**

- 0.89
- 0.72
- 0.76
- 0.89
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Cell Death and Differentiation
Lanosterol increases autophagy in dopaminergic neurons. Previous studies have shown that the loss of mitochondrial membrane potential can initiate the autophagic degradation of damaged mitochondria.26,27 Gene linked to familial forms of PD, such as PINK1 and Parkin, are thought to regulate this process, and PD-associated Parkin mutations cause a decrease in mitophagy in mammalian cell lines.28 We thus asked if lanosterol mediates autophagosome formation in dopaminergic neurons. Using microtubule-associated protein light chain 3 (LC3) as a marker for autophagy, we quantified both the size and number of autophagic vacuoles (AVs) in primary ventral midbrain neurons on MPP\(^+\)/lanosterol treatment. Similar to other studies,29 we found that addition of MPP\(^+\) in primary dopaminergic neurons increased the number of AVs by about 2.5-fold (Figure 7). There is also about a 75% increase in the average size of AVs with MPP\(^+\) treatment. Remarkably, when neurons were exposed to lanosterol alone, we observed a similar increase in AV size and number. Co-treatment of MPP\(^+\) and lanosterol led to an additive effect on autophagy, with significant increases in both the number and size of AVs, compared with MPP\(^+\) or lanosterol alone (Figure 7).

Next, we addressed whether lanosterol has a specific effect on axonal mitophagy. In PD, axons of dopaminergic neurons progressively degenerate and ‘die back’, in a process that may be accelerated by mitochondrial dysfunction and involve mitophagy. For this, we grew hippocampal neurons in microfluidic chambers to segregate axons from neuronal cell bodies and dendrites.30 Mitochondria were visualized by expression of the fluorescent reporter MitoRed and neurons were immunostained for endogenous LC3. We found a small but significant increase in colocalization of MitoRed with LC3 (Figure 8) on lanosterol treatment, suggesting an increase in axonal mitophagy. Taken together, these results suggest that the protective effects of lanosterol are mediated by mitochondria uncoupling, and subsequent clearance of damaged mitochondria.

Discussion

The etiology of PD implicates several factors, including mitochondrial dysfunction and misregulation of sterol metabolism. Evidence implicating impaired mitochondrial function in PD is substantial. This evidence is based on (1) the identification of rare PD-associated mutations in genes that affect mitochondrial function such as the putative kinase, PINK1 (PARK6), the E3 ligase Parkin (PARK2) and DJ-1 (PARK7); (2) similarities between PD and clinical symptoms that arise on exposure to the neurotoxin MPTP, a complex I inhibitor; and (3) a significant decrease in complex I/II activity in the platelets of patients with PD.31 On the other hand, evidence for a role of aberrant sterol metabolism in PD is rather controversial. For example, lower levels of serum LDL-C are considered a risk factor for PD,\(^{12,32}\) but it is unclear how serum LDL-C relates to levels of cholesterol in the brain. Statins, which lower serum cholesterol levels by inhibiting 3-hydroxy-3-methyl-glutaryl-coenzyme A reductase, have been used to treat PD, but there are many unresolved questions regarding their benefits (see review in Becker and Meier\(^{33}\)).

In this study, we measured a selection of sterol metabolites in the ventral midbrain and striatum of MPTP-treated mice, two brain regions characterized by cell death and/or axonal loss in both the MPTP model as well as in PD. Our results show a highly selective reduction of lanosterol levels in these affected areas (Figure 1) and point to an alteration of lanosterol metabolism in MPTP-treated mice. We cannot rule out the possibility that lanosterol is oxidized or metabolized to di-hydrolanosterol or other products in MPTP-treated animals because we are unable to measure these oxidized metabolites.

Consistent with a role of lanosterol in PD pathogenesis, we observed an improved survival of MPP\(^+\)-treated dopaminergic neurons on exogenous addition of lanosterol (Figure 2). We also demonstrated that LSS relocalized from the ER to mitochondria in dopaminergic neurons on MPP\(^+\) treatment (Figures 4b and 5), suggesting an increase in lanosterol synthesis in mitochondria. Interestingly, recent lipidomic analysis of macrophages during stimulation with Lipid A (a condition that leads to oxidative bursts) showed a pronounced increase in lanosterol levels in several intracellular compartments, including mitochondria,\(^{16}\) implying that modulation of lanosterol metabolism may be part of a global cellular response to stress. If upregulation of lanosterol synthesis is part of a cellular defense mechanism, it is not clear why lanosterol levels drop in response to MPTP treatment. In this regard, it is interesting to note that two recent papers reported lowered lanosterol levels in the serum of patients with Alzheimer’s disease,\(^{34}\) and in fibroblasts challenged by virus infection.35 One possible explanation for these seemingly contradicting results is that distinct types of stress differentially impact lanosterol metabolism. Perhaps, in some cases, the substantial decrease in lanosterol levels cannot be compensated by upregulation of lanosterol synthesis as part of the cell’s protective response. Alternatively, we cannot exclude the possibility that translocation of LSS to mitochondria in response to MPP\(^+\) is an epiphenomenon, which is not indicative of a cellular response to stress.

Our results, however, show that exogenous addition of lanosterol leads to mild uncoupling of mitochondria in dopaminergic and glutamatergic neurons, with no detectable impact on ATP levels (Figure 6 and Supplementary Figure...
The mitochondria uncoupling and protective effects of lanosterol are strikingly similar to those observed with low dose of the uncoupler, FCCP, which improves cellular survival in ischemic preconditioning but has no significant impact on ATP levels. In the context of PD, the mitochondrial uncoupling effect of lanosterol has important implications. For example, Parkin is recruited to mitochondria via PINK1 on membrane depolarization, and regulates the clearance of damaged mitochondria by mitophagy in mammalian cell lines. In addition, the translocation of Parkin to mitochondria has etiological significance, as a number of disease-associated Parkin mutant proteins fail to translocate. Together, these data point to a role of mitochondrial uncoupling and autophagy in PD pathogenesis. In line with this model, our results reveal that lanosterol induces mitochondrial uncoupling and promotes autophagy.

Figure 7  Lanosterol and MPP+ increase the number of AVs in dopaminergic neurons. (a) Confocal images of ventral midbrain neurons stained with TH (top panels), LC3 (middle panels) and AV quantification (ImageJ software output, bottom panels) with various treatment conditions indicated on the top. Scale bar represents 10 μm. (b) Graphical plot showing the average number of AV identified per TH+ cells and the average size of AV in pixel area during treatment with various lipids and MPP+ co-treatment. For each condition, n>40 TH+ cells from three independent experiments. Error bars represent S.E.M. ***P<0.001

S1). The mitochondria uncoupling and protective effects of lanosterol are strikingly similar to those observed with low dose of the uncoupler, FCCP, which improves cellular survival in ischemic preconditioning but has no significant impact on ATP levels. In the context of PD, the mitochondrial uncoupling effect of lanosterol has important implications. For example, Parkin is recruited to mitochondria via PINK1 on membrane depolarization, and regulates the clearance of damaged mitochondria by mitophagy in mammalian cell lines. In addition, the translocation of Parkin to mitochondria has etiological significance, as a number of disease-associated Parkin mutant proteins fail to translocate. Together, these data point to a role of mitochondrial uncoupling and autophagy in PD pathogenesis. In line with this model, our results reveal that lanosterol induces mitochondrial uncoupling and promotes autophagy.

Although mitochondrial uncoupling has been shown to be neuroprotective in various models, including MPTP-induced neurodegeneration, the mechanisms involved are still unclear. Some studies have shown that uncoupling reduces superoxide species, offering an explanation for improved
neuronal survival in the MPTP model, because oxidative stress is thought to be the primary cause of cell death.8,9 In other studies, transient mitochondrial uncoupling is neuroprotective in glutamate-induced neurotoxicity, as it prevents uptake of calcium from the cytosol to mitochondria.11 Finally, a recent study showed that DJ-1, a gene involved in early onset PD, regulates the expression of two UCPs (UCP4 and UCP5) and controls oxidative stress in mitochondria of dopaminergic neurons in the substantia nigra.10 Although these studies cited above have identified different modulators by which a cell/neuron alters mitochondrial membrane potential, they are in good agreement with our findings, whereby uncoupling mechanism proves to be a central regulator of cellular response to stress.

In conclusion, we report that in addition to its role as a precursor for cholesterol biosynthesis, lanosterol acts as a survival factor for dopaminergic neurons. Furthermore, our findings point to an unexpected role of this sterol metabolite in regulating mitochondrial function and autophagy, and bring sterol metabolism to the forefront of neurobiological disease.

Materials and Methods

PD animal model: MPTP injections. All procedures performed in rodents were in accordance with IACUC guidelines. MPTP injections were performed according to previously published methods, following the acute schedule.17 Briefly, C57B6 mice were given four i.p. doses of either 18 mg/kg of MPTP (Sigma-Aldrich, St. Louis, MO, USA) or saline (control) every 2 h. Mice were decapitated 48 h after the last dose, and the ventral midbrain and striatum were dissected and snap-frozen according to previously published methods, following the acute schedule.17 Briefly, dried lipid extracts were resuspended in culture medium on the day of treatment.

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Lanosterol, cholesterol and PC were dissolved in chloroform/methanol (1:1). Cholesterol or lanosterol was mixed in equimolar proportion with PC and dried by vacuum in a Speedvac (Thermosavant, Waltham, MA, USA). The lanosterol/PC or cholesterol/PC mixture was resuspended in culture medium on the day of treatment to make a 0.5 mM stock liposome. Each type of stock liposome was used in the neuronal cultures within a day of preparation.

Lipid extraction. For tissues and cells, extraction of lipids was performed using Bligh and Dyer method.29 Briefly, cells were washed 3–5 times with phosphate-buffered saline (PBS), scraped in 400 μl ice-cold methanol and transferred to a 1.5-ml Eppendorf tube. Chloroform (200 μl) was added and samples were vortexed for 1 min. Next, 300 μl of 1 M KCL was added, and the homogenates were microcentrifuged at 14 000 r.p.m., for 5 min at 4 °C to separate phases. The lower organic phase was carefully transferred to a new Eppendorf tube. The aqueous phase was re-extracted twice with 300 μl chloroform. All organic phases were pooled and dried under vacuum using a Speedvac and stored at ~80 °C until derivatization and subsequent GC–MS analysis.

Ventral midbrain (~20–25 mg) and striatum (~15–20 mg) were dissected and tissues were homogenized using a Dounce homogenizer in 600 μl ice-cold chloroform/methanol (1:2). Another 300 μl chloroform was added to the homogenate followed by the addition of 450 μl of 1 M KCL. The homogenates were microcentrifuged at 14 000 r.p.m., for 5 min at 4 °C. The lower organic phase was carefully transferred to a new Eppendorf tube, and the aqueous phase was re-extracted twice with 300 μl chloroform. All organic phases were pooled and dried under vacuum in a Speedvac. Dried samples were then stored at ~80 °C until derivatization and subsequent analyses.

Sample preparation for GC–MS. Briefly, dried lipid extracts were resuspended in chloroform/methanol (1:1) to a concentration of 0.1 mg tissue/μl solvent. A 20-μl sample of lipid extract was removed and completely dried in a glass vial. For each sample, we added a mixture of heavy isotopes: 40 ng of 7α-hydroxycholesterol-d-4, 40 ng of 7β-hydroxycholesterol-d-4, 40 ng of 26(27)-hydroxycholesterol-d3, 80 ng of 7-ketocholesterol-d7, 0.2 μg of 5α-cholestanol, 0.2 μg of desmosterol-d7, 0.2 μg of lathosterol-d7, 0.2 μg of campsterol-d7, and 0.2 μg of β-sitosterol-d7 in 25 μl of ethanol. Standards and sample mixtures were dried under a stream of N2 before adding the derivatizing agent (15 μl acetonitrile and 15 μl BSTFA – TMCS; Pierce Thermoscientific, Waltham, MA, USA). The derivatized samples were analyzed with an Agilent 5975 inert XL mass selective detector (Santa Clara, CA, USA). Selective ion monitoring was performed using the electron ionization mode at 70 eV (with the ion source maintained at 230 °C and the quadrupole at 150 °C) to monitor one target ion. Two qualifier ions were selected for the mass spectrum of each compound to optimize for sensitivity and specificity.

Ventral midbrain cultures. Ventral midbrains from 20 postnatal day 0 to day 2 rodents were dissected and digested in papan solution and plated on a glia layer. Cells were cultured in serum-free neurobasal/B27 medium (Invitrogen, Carlsbad, CA, USA) supplemented with superoxide dismutase 1 (5 μg/ml), apo-transferrin (95 μg/ml) and insulin (21 μg/ml) (all from Sigma). In contrast to Rayport et al.,16 our...
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culture medium contained no serum because the cultures were subjected to lipid addition in experimental conditions. One hour after plating the cells, 10 ng/ml of glial-derived neurotrophic factor (Millipore, Billerica, MA, USA) was added. To inhibit glia growth, a solution of 16.5 mg/ml uridine and 6.7 mg/ml 5-fluorodeoxyuridine was added 1 day after plating. Cells were then cultured for 7 days (days in vitro (DIV7)) before treatment with MPP+ and various types of liposome.

Hippocampal neuron cultures. Hippocampal neurons from rats E16.5 embryos were cultured in neurobasal/B27 as described. Briefly, neurons were plated on a wax-dotted coverslip coated with 1 mg/ml of poly-c-lysine (Sigma). Two hours after plating, when neurons had attached, the coverslips were flipped into a six-well plate containing a glia feeder layer. PC, cholesterol or lanosterol liposome (5 μM each) was added after DIV7.

Western and antibodies. For western blots, cultured cells were washed three times with 1× PBS. Cells were lysed in 100 μl of RIPA buffer (50 mM Tris-HCl, pH 7.4, 1% NP-40, 0.25% Na-deoxycholate, 150 mM NaCl and 1 mM EDTA), supplemented with a cocktail of protease inhibitors (Complete Mini-EDTA inhibitors, Roche, Diagnostics, Indianapolis, IN, USA). In all, 20 μg of cell protein lysate was loaded in each well of a 10% polyacrylamide gel containing 0.1% SDS. After electrophoresis, proteins were transferred to nitrocellulose membrane and probed with the following antibodies: (i) rabbit anti-p38 (C91; 1:1000, Santa Cruz Biotechnology, Santa Cruz, CA, USA), (ii) rabbit anti-β-actin (1:1000, AVIA, Systems Biology, San Diego, CA, USA), (iii) rabbit anti-tyrosine hydroxylase (TH) (1:20 000, Covance, Princeton, NJ, USA), (iv) rabbit anti-SREBP2 (1:1000, Abcam, Cambridge, MA, USA), (v) rabbit anti-calnexin (1:2000, Abcam), (vi) rabbit anti-VDAC/porin (1:2000, Abcam) and (vii) anti-rabbit-pDSDK-3/β(ε)2/3 (1:1000, Cell Signaling, Danvers, MA, USA). Peroxidase-conjugated anti-rabbit or anti-mouse secondary antibodies (1:10000) were purchased from Bio-Rad (Hercules, CA, USA). Immunoblots were visualized with enhanced chemiluminescence reagent from Pierce Thermoscientific.

Fluorescence microscopy and quantification of dopaminergic neuronal survival. Ventral midbrain cultures plated on 12-mm coverslips were treated for 24 h with 10 μM MPP+ with or without 5 μM PC, 5 μM cholesterol or 5 μM lanosterol liposome. Cells were washed three times with PBS, then fixed with 4% paraformaldehyde for 20 min, followed by permeabilization and blocking in 5% bovine serum saline (0.1% TritonX-100 for 30 μl. Cells were then stained with anti-TH (secondary: Alexa-fluor 488, green) and neuron-specific class III beta tubulin (TUJ1) (secondary: Alexa-fluor 555, red). Anti-mouse and anti-rabbit Alexa-fluor 488 and 555 (1:1000) were purchased from Molecular Probes (Eugene, OR, USA). Invitrogen. TUJ1-positive and TH-positive cells were counted on an Olympus fluorescence microscope (Tokyo, Japan) with FITC and TRITC filter sets. Every 5 μM PC, 5 μM cholesterol or 5 μM lanosterol liposome was plated at a higher density (300 cells/mm²), and multi-position time-lapse imaging of 3–4 fields was performed at 30-s intervals for 20 min.

For both types of cultures, the intensity of the JC-1 red-to-green ratio was measured in each frame, and the change in mitochondrial membrane potential was plotted as ΔF/Δt, where F is the average JC-1 red-to-green ratio over the first 10 frames before treatment. Decay curves were fitted to a mono-exponential function, y = xe⁻ᵏᵗ, using IGOR Pro 6.1 (WaveMetrics Inc., Lake Oswego, OR, USA).

Measurement of CoQ. For each condition, 3 to 4 million hippocampal neurons (DIV7) were treated with various lipids and incubated for 24 h. Cells were washed twice with cold PBS and scraped in 0.5 ml of cold PBS. Cells were centrifuged at 1000 × g to pellet the cells, and resuspended in 100 μl fresh PBS. Next, 750 μl of hexane/ethanol (5:2 v/v) was added, and samples were vigorously vortexed for 1 min. To extract CoQ, 400 μl of the organic phase was collected and completely dried under a stream of N₂, followed by LC–MS analysis.

An Agilent HPLC 1200 system coupled with an Applied Biosystems 3200 QTRAP mass spectrometer (Foster City, CA, USA) was used for measuring CoQ6, Q9, and Q10 and free cholesterol. Chromatographic separation was carried out using an Agilent Zorbax Eclipse XDB-C18 column (i.d. 4.6 × 150 mm). High-pressure liquid chromatography (HPLC) conditions were as the following: mobile phase: chloroform/methanol 1:1 (v/v); flow rate: 0.5 ml/min; column temperature: 30 °C; injection volume: 10 μl. The LC–MS instrument was operated in positive atmospheric pressure chemical ionization mode with a vaporizer temperature of 500 °C and a corona current of 3 μA. CoQ6 was used as an internal standard and monitored with an multiple reaction monitoring (MRM) transition of 591.4→197.0. MRM transitions of 727.5→197.0, 795.5→197.0 and 863.6→197.0 were set up for analysis of CoQ6, Q9 and Q10, respectively.

Quantification of AVs. Ventral midbrain cultures plated on 12-mm coverslips were treated for 24 h with 10 μM MPP+ with or without 5 μM PC, 5 μM cholesterol or 5 μM lanosterol liposome. Cells were washed three times with PBS, then fixed with 4% paraformaldehyde for 20 min and permeabilized and blocking in 5% fetal bovine serum in 0.1% TritonX-100 for 30 μl. Cells were then stained with anti-TH (secondary: Alexa-fluor 488, green) and neuron-specific class III beta tubulin (TUJ1) (secondary: Alexa-fluor 555, red). Anti-mouse and anti-rabbit Alexa-fluor 488 and 555 (1:1000) were purchased from Molecular Probes (Eugene, OR, USA). Invitrogen. TUJ1-positive and TH-positive cells were counted on an Olympus fluorescence microscope (Tokyo, Japan) with FITC and TRITC filter sets. Every neuron on the 12-mm coverslip was counted. The percentage of dopaminergic neurons in each group was determined by the number of TH+/TUJ1+ cells. Typically, in each control coverslip, there were 2000–3000 TUJ1+ cells, of which 400–1200 were TH+ cells. For each treatment, we assessed 4–6 coverslips per independent experiment. The averages of 4–5 independent experiments are shown in the figures.

Confocal microscopy and colocalization studies. Ventral midbrain cultures in control and MPP+–treated cells were stained with rabbit anti-LSL (1:100), monoclonal mouse anti-TGMR20 (1:500) (Abcam) or monoclonal mouse anti-KDEL (1:500) (Abcam), and shear anti-TH (1:500) (Abcam). The secondary antibodies, goat anti-mouse Alexa-fluor 555, goat anti-rabbit Alexa-fluor 488 and donkey anti-sheep Alexa-fluor 633, were obtained from Invitrogen. Cells were imaged with a laser-scanning confocal microscope (LSM510, Carl Zeiss, Oberkochen, Germany) with excitation and emission filters filtering the secondary Alexa-fluor antibody dye specifications. Images were taken using a 63X objective with the same laser power and gain. The 12-bit images were quantified using ImageJ (analyze particle drop-in, National Institute of Health, Bethesda, MD, USA). For each image, detected LC3 puncta were intensity thresholded (<1000) and gated for size (<15 pixel). For each condition, 40–60 TH+ cells were assessed from three independent experiments.

As a positive control for our method of evaluating AV, mouse embryonic fibroblast grown in serum or serum deprived were stained with LC3 and quantified according to the same parameters. As expected and shown in supplementary data (Supplementary Figure S2), there is approximately a 25-fold increase in AV on serum starvation.

Quantification of mitophagy in axons. E18.5 hippocampal neurons were electroporated using the Amaza poration system (Lonza, Basel, Switzerland) with MitoRed construct (Clenetech, Mountain View, CA, USA, cat. No.: PT-3635-3). Cells were then plated in microfluidic chambers (Xona Microfluidics, Temecula, CA, USA), which allow for physical separation of axons and cell somas. At DIV7, automated stage and built in autofocusing system (PFS, Nikon, Tokyo, Japan) and driven by Metamorph 7.6 (Universal Imaging, Ypsilanti, MI, USA). JC-1 is excited at 488-nm, and its fluorescence emission was collected at 530 ± 10 nm (green) and 590 ± 17 nm (red), corresponding to peak fluorescence from the monomer and aggregate signals, respectively. Mitochondrial membrane potential was measured by taking the red to green emission ratio.
neurons were treated with 5 μM PC, 5 μM cholesterol, 5 μM lanosterol or 200 nM of rapamycin (positive control) for 24 h. Cells were then fixed with 4% paraformaldehyde for 20 min and permeabilized with 100 μg/ml of digitonin for 10 min. Following permeabilization, cells were washed three times with PBS, and stained with anti-LC3 (secondary: Alexa-fluor 488, green). To quantify mitophagy, the percentage of MitoRed and LC3-positive mitochondria were plotted as a percentage to total Mitotracker-positive mitochondria using the Metamorph 7.6 colocalization drop-in. For each condition, images of 40–50 axons were taken from three independent experiments.

**Statistical analyses.** Error bars represent S.E.M. For fold changes in sterol intermediates, P-values were calculated with two-tailed Mann–Whitney U-test. For in vitro assessment of dopaminergic neuron survival, CoQ levels, ATP levels, AV 3D number and size, percentage of mitophagy, P-values were calculated with two-tailed Student’s t-test.

**Conflict of Interest.** The authors declare no conflict of interest.

**Acknowledgements.** We thank Thilo Hagen for providing us with an aliquot of CCCP. This study was supported by the National University of Singapore (NUS), Department of Biological Sciences fellowship, the Singapore Biomedical Research Council (BMRC) Grant ID 08/1/21/19/558, the Singapore National Research Foundation grant R-123000-165-571, the Department of Biological Sciences fellowship, the Singapore Biomedical Research Council (BMRC) Grant ID 08/1/21/19/558, the Singapore National Research Foundation grant R-123000-165-571, and the Singapore National Research Foundation grant R-123000-165-571.

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Supplementary Information accompanies the paper on Cell Death and Differentiation website (http://www.nature.com/cdd)