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PINK1-induced mitophagy promotes neuroprotection in Huntington's disease

B Khalil¹, N El Fissi², A Aouane², M-J Cabirol-Pol¹, T Rival^{2,3} and J-C Liévens*^{1,3}

Huntington's disease (HD) is a fatal neurodegenerative disorder caused by aberrant expansion of CAG repeat in the huntingtin gene. Mutant Huntingtin (mHtt) alters multiple cellular processes, leading to neuronal dysfunction and death. Among those alterations, impaired mitochondrial metabolism seems to have a major role in HD pathogenesis. In this study, we used the Drosophila model system to further investigate the role of mitochondrial damages in HD. We first analyzed the impact of mHtt on mitochondrial morphology, and surprisingly, we revealed the formation of abnormal ring-shaped mitochondria in photoreceptor neurons. Because such mitochondrial spheroids were previously detected in cells where mitophagy is blocked, we analyzed the effect of PTEN-induced putative kinase 1 (PINK1), which controls Parkin-mediated mitophagy. Consistently, we found that PINK1 overexpression alleviated mitochondrial spheroid formation in HD flies. More importantly, PINK1 ameliorated ATP levels, neuronal integrity and adult fly survival, demonstrating that PINK1 counteracts the neurotoxicity of mHtt. This neuroprotection was Parkin-dependent and required mitochondrial outer membrane proteins, mitofusin and the voltage-dependent anion channel. Consistent with our observations in flies, we demonstrated that the removal of defective mitochondria was impaired in HD striatal cells derived from HdhQ111 knock-in mice, and that overexpressing PINK1 in these cells partially restored mitophagy. The presence of mHtt did not affect Parkin-mediated mitochondrial ubiguitination but decreased the targeting of mitochondria to autophagosomes. Altogether, our findings suggest that mitophagy is altered in the presence of mHtt and that increasing PINK1/Parkin mitochondrial guality control pathway may improve mitochondrial integrity and neuroprotection in HD. Cell Death and Disease (2015) 6, e1617; doi:10.1038/cddis.2014.581; published online 22 January 2015

Huntington's pathology is a devastating inherited neurodegenerative disease clinically characterized by involuntary movements, cognitive decline, emotional changes and dementia. Huntington's disease (HD) results from an increased length of a polyglutamine stretch in the N terminus of Huntingtin protein, due to expanded CAG repeat in the first exon of the *IT15/huntingtin* gene. Symptoms are observed when patients carry more than 36 CAG repeats and there is still no cure to halt the progression, which leads to death after 10 to 20 years. Despite ubiquitous expression of mutant Huntingtin (mHtt) throughout the brain, only specific neuronal sub-populations degenerate mostly in the striatum and, to a lesser extent, in the cerebral cortex.

The presence of mHtt leads to multiple cellular dysfunctions, including alterations in intracellular signaling pathways, defects in cellular trafficking, transcription deregulation, abnormal synaptic transmission, proteasomal dysfunction and mitochondrial alterations. One challenging question is to determine to what extent each of those defects contributes to the disease progression.

Growing evidence suggest that mHtt disrupts mitochondrial functions, resulting in energetic defect, reactive oxygen

species overload and release of proapoptotic molecules. More specifically, the respiratory chain is affected with impairment of the mitochondrial complex II/III activity in the caudate and putamen samples of HD patients.^{1,2} The maintenance of functional mitochondria requires biogenesis as well as mitochondrial fusion/fission dynamics to replenish stores of damaged components. Peroxisome proliferatoractivated receptor gamma coactivator-1-alpha (PGC-1 α) is a key transcriptional coactivator that controls mitochondrial biogenesis and energy metabolism. PGC-1 α is downregulated by mHtt through interference with the CREB/TAF4-dependent transcriptional pathway.³ More recently, fragmented mitochondria were linked to HD,4 due to increased dynaminrelated protein 1 (DRP1) activity.5,6 The accumulation of mitochondrial damages in postmitotic neurons is therefore considered as a key process in HD pathogenesis.

The ultimate mechanism that cells use when they are overwhelmed by dysfunctional mitochondria is to remove the damaged organelles by selective autophagy, the so-called mitophagy. PTEN-induced putative kinase 1 (PINK1) is a molecular sensor that accumulates at the outer membrane of damaged mitochondria to recruit the E3 ubiquitin ligase Parkin

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Abbreviations: CCCP, carbonyl cyanide *m*-chlorophenyl hydrazone; DRP1, dynamin-related protein 1; HD, Huntington's disease; Mfn, mitofusin; mHtt, mutant Huntingtin; PD, Parkinson's disease; PGC-1*α*, peroxisome proliferator-activated receptor gamma coactivator-1-alpha; PINK1, PTEN-induced putative kinase 1; TRAP1, TNF receptor-associated protein 1; VDAC, voltage-dependent anion channel

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and initiates the mitophagic process.^{7–9} So far, the impact of PINK1 and Parkin has been investigated mainly in Parkinson's disease (PD). Indeed, mutations in both genes are directly linked to familial parkinsonism.

In the present study, we first analyzed the impact of mHtt on mitochondrial morphology in a Drosophila model of HD. Surprisingly, we revealed the formation of mitochondrial spheroids in non-apoptotic photoreceptor neurons. Because those atypical mitochondria were recently described in cells where mitophagy was blocked.^{10,11} it suggests that mHtt may cause an imbalance between the accumulation of damaged mitochondria and their clearance through mitophagy. To test this hypothesis, we analyzed the effect of PINK1 overexpression in HD flies. We found that PINK1 reduces mitochondrial spheroid formation and, more importantly, confers substantial protection against mHtt-induced neuronal phenotype in adult flies. We further showed that mHtt disrupted colocalization between mitochondria and autophagosomes and, thereby, the elimination of defective mitochondria in striatal cells derived from knock-in HdhQ111 mice. Consistent with our obsevations made in HD flies, PINK1 overexpression partially restored mitophagy in HdhQ111 cells. Altogether, our results highlight the value of PINK1/Parkin pathway to alleviate mitochondrial defects in HD.

Results

Mitochondrial accumulation and spheroid formation precede mHtt-induced neurodegeneration and are suppressed by PINK1. To determine whether neuronal expression of mHtt (Httex1p Q93) in *Drosophila* alters mitochondrial morphology, we performed transmission electron microscopy analysis of the retina, a CNS structure that shows widespread neuronal loss in HD flies.¹² We analyzed 1-day-old flies expressing mHtt specifically in neurons because the neurodegeneration is moderate at this age, although some photoreceptors start to display rhabdomere atrophy (Figure 1a, panel 2, black asterisk) or cytoplasmic condensation typical of apoptosis (Figure 1a, panel 2, white asterisk). We found that non-apoptotic photoreceptors contained two

types of mitochondria: typical round-shaped mitochondria and, unexpectedly, mitochondrial tubules that seem to bend and often fuse at their extremities to adopt a ring-shaped structure (Figure 1b, panels 2-5). Sometimes those bended mitochondria engulf cytoplasmic components such as pigment granules (Figure 1b, panel 6). Those abnormal mitochondria were strikingly reminiscent to the mitochondrial spheroids recently described in cells where mitophagy was blocked.^{10,11} With the aim of boosting mitophagy, we decided to overexpress PINK1. Drosophila PINK1 overexpression (PINK1^{OE} line) significantly reduced mitochondrial spheroid formation in photoreceptor neurons (Figure 1a, panel 3 and Figure 1c, histograms 1 and 2). We also performed guantitative analyses on the mitochondria that were not forming spheroids. In photoreceptors, these organelles were generally round shaped, or sometimes appeared as short tubules. Drosophila expressing mHtt showed a decrease in the average size (Figure 1c, histogram 3) and an increase in the average number of non-spheroid mitochondria (Figure 1c, histogram 4), suggesting that mHtt leads to mitochondrial fragmentation. More importantly, the total area covered by the mitochondria in the cytoplasm was significantly higher (Figure 1c, histogram 5). This indicates that the mitochondria tend to accumulate in HD flies. Overexpression of PINK1 in HD flies restored mitochondrial area. number and density to control values (Figure 1c, histograms 3-5).

PINK1 overexpression rescues HD neuronal pathology. Next, we determined whether PINK1 mitigates photoreceptor neurodegeneration in the retina. Pseudopupil analysis revealed that PINK1 overexpression (*PINK1^{OE}*) markedly reduced photoreceptor loss in 1-day-old flies (Figures 2a and b). Whereas only 23% of ommatidia remained intact with seven visible photoreceptors in HD flies, this percentage reached 44% in flies coexpressing mHtt and PINK1. Similar neuroprotection was found in 6-day-old flies (Figure 2b). PINK1 overexpression in control flies had no impact on ommatidia even at 1 month of age. We also explored the effect of PINK1 on postmitotic neurons in the central brain. Although HD flies at 12 days displayed

Figure 1 Mitochondrial spheroid formation precedes Httex1p Q93-induced neurodegeneration and is suppressed by Drosophila PINK1 overexpression. (a) Transmission electron microscopy (TEM) pictures of eve sections from 1-day-old adult Drosophila. Panel 1: Ommatidia of control flies normally contain seven visible photoreceptor neurons (PhRs), each photoreceptor harboring a rhabdomere membrane subdomain (Rha) that detects light. Black arrowheads show typical round-shaped mitochondria. Panel 2: Targeted neuronal expression of httex1p Q93 triggers rhabdomere degeneration (black asterisk) and photoreceptor apoptosis (white asterisk). Circular electron-dense structures corresponding to mitochondrial spheroids (white arrows) are observed in the cytoplasm of non-apoptotic PhRs. Black arrowheads point to round-shaped mitochondria. Panel 3: TEM picture of an ommatidium from 1-day-old Drosophila coexpressing httex1p Q93 and Drosophila PINK1 (PINK1^{OE}) in neurons. No mitochondrial spheroid is detected in this ommatidium. Black arrows correspond to round-shaped mitochondria. (b) High magnification TEM images show that circular electron-dense structures observed in Httex1p Q93 photoreceptors are mitochondrial spheroids. Panels 1 and 2, respectively, show control and Httex1p Q93-expressing photoreceptors with no sign of degeneration. Black arrowheads point to mitochondria with normal round-shaped morphology, whereas white arrows correspond to spheroid mitochondria. Panel 3 represents an httex1p Q93expressing photoreceptor containing a massive accumulation of mitochondria within its cytoplasm (white arrows: mitochondrial spheroids; black arrows: round-shaped mitochondria). Panels 4-6 show the ultrastructure of representative mitochondrial spheroids from httex1p Q93-expressing photoreceptors. Mitochondrial spheroids often form ring-like structures harboring an enlarged region. Mitochondrial spheroids can also appear as bended barbell-like structures (panels 5 and 6) sometimes involving multiple mitochondria (panel 5) or even enwrapping cytoplasmic components such as pigment granules (panel 6). (c) Quantitative analyses from 1-day-old flies expressing no transgene (control), httex1p Q93 alone or together with PINK1 (PINK1^{OE}) in neurons. Histograms show the number of spheroids per ommatidium (histogram 1), the number of photoreceptors containing mitochondrial spheroids (histogram 2), the average area of individual mitochondria (µm²) (histogram 3), the number of mitochondria per 10 µm² of the cytoplasm (histogram 4) and the total area occupied by the mitochondria within the photoreceptor cytoplasm (ratio of mitochondrial area/cytoplasmic area in %) (histogram 5). Measurements in histograms 3-5 do not include spheroids. Observations were made on eye sections from at least three independent flies. Spheroid quantifications were performed on 20 ommatidia and 120 non-apoptotic photoreceptors for httex1p Q93, and 32 ommatidia and 190 non-apoptotic photoreceptors for httex1p Q93+PINK1^{OE}. Quantification of mitochondrial area, number and density were performed on 51 photoreceptors and 278 mitochondria for control, 55 photoreceptors and 875 mitochondria for httex1p Q93, and 58 photoreceptors and 374 mitochondria for httex1p Q93+PINK1^{OE}. Statistics were performed by using the Mann-Whitney U-test (***P<0.0001 versus control; ^{ØØØ}P<0.0001 versus httex1p Q93)

about 7% of degenerative neurons showing cytoplasmic vacuolization in the olfactive lobes (Figure 2c), <2% of vacuolized neurons were detected in the presence of PINK1 (Figure 2d).

It is well documented that mHtt neuronal expression triggers the premature death of adult flies. We found that PINK1 significantly increases both mean and median survival (+40%) of HD flies (Figure 3a). By contrast, PINK1 had no impact on the longevity of control flies (Figure 3b). On the other hand, reducing PINK1 endogenous expression in HD flies by the mean of an heterozygous PINK1 loss-of-function allele ($PINK1^{B9}$) did not decrease HD fly lifespan (Figure 3c). The impact of $PINK1^{B9}$ homozygosity was not tested, as *Drosophila* carrying two $PINK1^{B9}$ mutant alleles already show decreased lifespan.¹³ We next monitored ATP levels to evaluate mitochondrial function. We found that 7-day-old





Figure 2 Drosophila PINK1 overexpression rescues neurodegeneration in flies expressing httex1p Q93 in neurons. (a) Pseudopupil images from 1-day-old flies expressing no transgene (control), httex1p Q93 alone, httex1p Q93 together with Drosophila PINK1 (PINK1^{OE}) or PINK1^{OE} alone under the pan-neuronal driver elav-GAL4. In control flies, seven photoreceptors are visible, whereas the expression of httex1p Q93 induces photoreceptor neuron loss. $PINK1^{OE}$ counteracts the httex1p Q93-induced photoreceptor degeneration. (b) Number of photoreceptors per ommatidium in flies expressing no transgene (control), httex1p Q93, httex1p Q93, httex1p Q93, httex1p Q93+PINK1^{OE} or PINK1^{OE} alone at day 1 or 6 after eclosion. Significant rescue of neurodegeneration was observed at both ages when PINK1 was overexpressed in HD flies ($n \ge 10$ flies; P < 0.0001, Mann–Whitney U-test). (c) Transmission electron microscopy images showing highly vacuolized neurons in 12-day-old flies expressing no transgene (control), httex1p Q93 alone or together with PINK1^{OE}. PINK1^{OE} significantly reduces mHtt-induced neurodegeneration (Student's t-test, **P < 0.01, ***P < 0.001 versus control; ⁶⁰⁰⁰P < 0.001 versus httex1p Q93). Quantifications were performed as followed: n > 150 neurons in six flies per condition

flies expressing mHtt in neurons exhibited a 24% reduction of ATP levels in the brain (Figure 3d). In contrast, HD flies overexpressing PINK1 did not show decreased ATP levels at the same age (Figure 3d).

PINK1 neuroprotection against mHtt toxicity is dependent on Parkin. To determine if PINK1-mediated neuroprotective effect involves Parkin, we examined whether Drosophila parkin overexpression (*parkin^{OE}*) alleviates HD phenotype in *Drosophila*. Unexpectedly, overexpressing Parkin failed to modify the life expectancy or the photo-receptor loss of flies expressing Httex1p Q93 under the control of the neuronal driver *elav-GAL4* (Supplementary Figure S1). Coexpression of Parkin with PINK1 was deleterious as no adult flies emerged from pupae. This observation is in agreement with previous data showing that

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Figure 3 Drosophila PINK1 overexpression ameliorates survival of flies expressing httex1p Q93 in neurons. (a) Survival rates of flies expressing httex1p Q93 alone or together with $PINK1^{OE}$ in neurons. Statistical significance was determined by the log-rank test (n > 80 flies per condition; ***P < 0.0001). (b) Survival rates of flies expressing no transgene or $PINK1^{OE}$ under the control of the neuronal driver *elav-GAL4*. No significant change in survival is detected (n > 60 flies per genotype). (c) Survival rates of flies expressing *httex1p Q93* alone or in the presence of one mutant allele of $PINK1 (PINK1^{B9})$ in neurons. Lifespan of HD flies expressing $PINK1^{B9}$ was not significantly different from those expressing *httex1p Q93* alone. Data from at least five independent experiments were averaged and were presented as mean \pm S.E.M. Statistical analysis was performed using a Student's *t*-test (***P < 0.001 versus control; $^{OO}P < 0.01$ versus httex1p Q93)

coexpression of PINK1 and Parkin under the heat-shock-GAL4 driver leads to pupal lethality.¹⁴ Next, we determined if partial loss of function of Parkin can prevent rescue by PINK1. We generated HD flies overexpressing PINK1 and carrying a heterozygous Parkin loss-of-function allele $(park^{1})$. In those flies, PINK1^{OE} failed to increase lifespan (Figure 4a). We also demonstrated that PINK1 neuroprotection of photoreceptors was dependent on Parkin (Figure 4b). At 4 days, flies expressing mHtt alone exhibited 86% of ommatidia with 5-6 photoreceptors and only 9% of intact ommatidia (Figure 4b). As shown previously, PINK1 markedly reduced photoreceptor degeneration as 32% of the ommatidia remained intact. However, when PINK1^{OE} was combined with the $park^1$ mutation, the neuroprotection was abolished as 88% of ommatidia contained 5-6 photoreceptors and only 9% of ommatidia were intact. No significant change on both survival and photoreceptor number was observed in control *park*¹ heterozygous flies. Therefore, we demonstrate that overexpressing PINK1 provides neuroprotection against mHtt through the Parkin

pathway. However, our data also show that Parkin by itself is inefficient against mHtt toxicity.

PINK1 neuroprotective effect requires mitofusin. The mechanisms by which PINK1/Parkin pathway mediates phagophore formation and mitophagy are still elusive. Mitofusin (Mfn) is a target of Parkin and its degradation might promote mitochondrial fission and facilitate mitochondrial removal.¹⁵ We then analyzed whether or not increasing mitochondrial fission could influence mHtt toxicity. We found that RNAi silencing of the Drosophila mfn homolog, marf (marf^{IR}), or overexpressing Drosophila fission effector, DRP1 (DRP1^{OE}), failed to increase the lifespan of flies expressing mHtt in neurons (Figure 5a). The survival rate of HD flies was even reduced by about 23% in the presence of DRP1^{OE}. We also determined whether Mfn/Marf inactivation could impact the neuroprotective effect of PINK1. Whereas overexpressing PINK1 increased the life expectancy of HD flies, no significant effect was detected when marther was coexpressed with PINK1 (Figure 5a). We then analyzed the impact of Mfn/Marf



Figure 4 Loss of function of Parkin suppresses *Drosophila* PINK1 neuroprotective effect in HD flies. (a) Survival rates of flies expressing *httex1p Q93* alone, *httex1p Q93* with *PINK1^{OE}*, *httex1p Q93* in the presence of one mutant allele of *parkin (park¹)* or *httex1p Q93* together with *PINK1^{OE}* and *park¹*. Loss of function of Parkin reduces the PINK1 neuroprotective effect on the survival of HD flies. Statistical significance was determined by the log-rank test (n > 80 flies per genotype; ***P < 0.0001). (b) Number of photoreceptors per ommatidium in 4-day-old flies expressing no transgene (control), *httex1p Q93* alone, *httex1P Q93* with *PINK1^{OE}*, *httex1p Q93* together with *PINK1^{CE}* and *park¹* in neurons. Loss of function of Parkin abolishes the PINK1-mediated rescue of photoreceptor loss in HD flies (n = 10-12 flies per genotype; P < 0.0001, Mann–Whitney *U*-test)

inactivation on the loss of photoreceptors (Figure 5b). Interestingly, while silencing Mfn/Marf did not affect neuronal loss in HD fly retina, we observed a significant reduction of the PINK1-mediated neuronal rescue. Our data suggest that Mfn/Marf is involved in PINK1-mediated neuroprotection.

VDAC/Porin is necessary for PINK1-mediated rescue. It has been proposed that voltage-dependent anion channel (VDAC) could be part of the PINK1/Parkin pathway.^{16,17} As shown in Figure 5c, the presence of one mutant allele of porin (porink05123), the Drosophila homologous of VDAC, was sufficient to counteract PINK1-mediated rescue on the survival of HD flies. Whereas flies coexpressing mHtt and PINK1 in neurons showed a 35% increase in life expectancy versus flies expressing mHtt alone, PINK1 was uneffective on HD flies carrying one copy of *porin^{k05123}*. The loss of function of VDAC/ Porin led by itself to a very slight amelioration of the survival of HD flies, which could be related to the direct role of VDAC in the apoptotic release of mitochondrial cytochrome C. We also tested the impact of VDAC/Porin on the loss of photoreceptor neurons induced by mHtt (Figure 5d). Whereas the presence of one copy of porink05123 allele did not affect neuronal loss, porink05123 significantly reduced the neuronal rescue by PINK1. Thus, we propose that PINK1 neuroprotection against mHtt toxicity also involves VDAC/Porin.



Figure 5 Decreasing Mfn/Marf and VDAC/Porin reduce PINK1 neuroprotective effect. (a) Survival rates of flies expressing httex1p Q93 alone, httex1p Q93 with RNA interference (RNAi) against *mfn/marf* (*marf*^{*R*}), *httex1p* Q93 with a DRP1 transgene (DRP1^{OE}), *httex1p* Q93 with *PINK1^{OE}* or *httex1p* Q93 together with *PINK1^{OE}* and marf^R in neurons. Silencing Mfn/Marf reduces the PINK1 neuroprotective effect on the survival of HD flies. Statistical significance was determined by the log-rank test (n>60 flies per genotype; ***P<0.0001). (b) Number of photoreceptors per ommatidium in 4-day-old flies expressing no transgene (control), httex1p Q93 alone, httex1p Q93 with PINK1^{OE}, httex1p Q93 with the RNAi transgene mart^R or httex1p Q93 together with PINK1^{OE} and mart^{IR} in neurons (n = 7-10 flies per genotype; P<0.0001, Mann–Whitney U-test). (c) Survival rates of flies expressing httex1p Q93 alone. httex1p Q93 with PINK1^{OE}, httex1p Q93 in the presence of one mutant allele of VDAC/porin (porin^{k05123}) or httex1p Q93 together with PINK1^{OE} and porin^{k05123}. Loss of function of Porin prevents the PINK1 neuroprotective effect on the survival of HD flies. Statistical significance was determined by the log-rank test (n > 60 flies per genotype; ***P < 0.0001). (d) Number of photoreceptors per ommatidium in 4-day-old flies expressing no transgene (control), httex1p Q93 alone, httex1p Q93 with PINK1^{OE}, httex1p Q93 with the mutant allele porin^{k05123} or httex1p Q93 together with PINK1^{OE} and porin^{k05123} in neurons. Loss of function of Porin reduces the PINK1mediated rescue of photoreceptor loss in HD flies (n=6 flies per genotype; P<0.0001, Mann-Whitney U-test)

TRAP1 is not required for PINK1 neuroprotective effect. In parallel of PINK1/Parkin-mediated mitophagy, PINK1 can also regulate mitochondrial integrity by direct activation of mitochondrial chaperones such as TNF receptor-associated protein 1 (TRAP1).^{18,19} We found that increasing TRAP1 expression failed to reduce the organismal death of HD flies (Supplementary Figure S2). Therefore, PINK1 neuroprotection is likely not dependent on the activation of TRAP1 in the HD context.

The presence of mHtt compromises the degradation of defective mitochondria in mouse striatal cells. One important question raised by our data is whether or not mHtt may directly impact mitophagy. To investigate this issue, we used mouse striatal HdhQ7 and HdhQ111 cell lines that were immortalized from knock-in mice carrying 7 and 111 CAG, respectively, in the mouse htt gene.²⁰ As those cells did not strongly express Parkin (data not shown), they were transfected with Parkin plasmids. Parkin is specifically recruited to the mitochondria in response to dissipation of mitochondrial membrane potential. Accordingly, treatment with the mitochondrial uncoupler, carbonyl cyanide *m*-chlorophenyl hydrazone (CCCP; $10 \mu M$ for 6 h) led to the translocation of transfected mCherry-Parkin from the cvtoplasm to Tom20-immunostained mitochondria, whereas striatal cells treated with DMSO exhibited diffuse localization of mCherry-Parkin in the cytoplasm (Figure 6a). We found that 55% of HdhQ7 striatal cells showed Parkin translocation after CCCP treatment (Figure 6b). Importantly, mHtt in HdhQ111 did not modify the proportion of cells showing Parkin translocation to the mitochondria. As previously observed in HeLa cells,^{16,21} striatal cells treated with CCCP for a longer period (10 μ M for 38 h) exhibited a clustering of their mitochondria around the nucleus, followed by subsequent clearance of mitochondria as revealed by the loss of Tom20 staining (Figure 6c). However, strikingly, whereas 17% of HdhQ7 cells were devoid of Tom20-immunostained mitochondria, this amount fell down to only 2% for HdhQ111 cells (Figure 6d). At the same time, no change was detected between HdhQ7 and HdhQ111 cells regarding the capacity of Parkin to translocate to the mitochondria (Figure 6e) or perinuclear mitochondrial clustering (Figure 6f). We also confirmed these observations by using another mitochondrial marker, Tim23, which stains the inner membrane (Supplementary Figure S3). Altogether, our data indicate that mHtt could perturb the mitophagy process, likely downstream of Parkin translocation and mitochondrial perinuclear clustering.

PINK1 overexpression is able to ameliorate mitophagy process in mouse striatal cells. As mHtt compromises mitophagy, we determined whether PINK1 overexpression was still able to boost mitophagy in HdhQ111 cells. After 38 h CCCP treatment, PINK1 transfection increased the proportion of HdhQ7 and HdhQ111 cells showing Parkin mitochondrial translocation (Figure 6e) or mitochondrial perinuclear clustering (Figure 6f). More importantly, PINK1 overexpression increased the number of HdhQ7 and HdhQ111 cells with no detectable mitochondria (Figure 6g). However, the number of cells with no Tom20-immunostained mitochondria remained reduced for HdhQ111 *versus* HdhQ7 cells, further confirming that mHtt impacts mitophagy. Similar results were obtained when Tim23 immunostaining was used to visualize mitochondria (Supplementary Figure S3). Thus, our data show that overexpressing PINK1 is still able to increase partially the removal of defective mitochondria even in the presence of mHtt.

The targeting of mitochondria to autophagosomes is decreased by mHtt. Next, we further investigated at which step mitophagy process may be compromised by mHtt. Parkin promotes mitophagy by triggering mitochondrial ubiquitination, which in turn recruits ubiquitin-binding autophagic components. As shown in Figure 7a, CCCP treatment (10 µM for 38 h) strongly increased the ubiquitination of Tim23-immunostained mitochondria as compared with DMSO. Quantitative analysis revealed no change in the ubiquitination rate of CCCP-treated mitochondria between HdhQ111 and HdhQ7 cells (Figure 7b; Mander's coefficients: 0.88 ± 0.01 and 0.89 ± 0.01 for HdhQ7 and HdhQ111 cells, respectively). Treatment with CCCP also induced the accumulation of LC3-positive autophagic vesicles in striatal cells (Figure 7c). Interestingly, colocalization between GFP-tagged LC3 and mitochondria was significantly reduced in the HdhQ111 versus HdhQ7 cells (Figure 7d: Mander's coefficients: 0.45 ± 0.02 and 0.23 ± 0.01 for HdhQ7 and HdhQ111 cells, respectively). This suggests that mHtt impairs mitophagy, likely by perturbing the targeting of defective mitochondria to autophagosomes.

Discussion

Our data demonstrate that enhancing mitochondrial quality control by overexpressing PINK1 confers protection against mHtt toxicity. Indeed, PINK1 overexpression rescues abnormal mitochondrial spheroid formation, neuronal loss, ATP levels and, more importantly, it increases the survival of adult flies expressing mHtt in neurons. Additionally, we have dissected the mechanisms underlying PINK1-mediated neuroprotection. We showed that it requires the E3 ubiquitin ligase Parkin and the outer membrane mitochondrial proteins Mfn/Marf and VDAC/Porin. Furthermore, we provided the first evidence that HD mutation affects mitophagy in HD striatal cells derived from HdhQ111 knock-in mice. Thus, our findings highlight the potential role played by damaged mitochondria accumulation in HD pathogenesis, and support the PINK1/Parkin pathway as a valuable therapeutic target.

Many studies have described abnormal mitochondrial fragmentation and cristae disruption in HD models and patients.^{4–6,22,23} It was proposed that increased fission might have a direct role in HD pathogenesis. In agreement with this, reducing DRP1 activity protected cultured neurons expressing mHtt from axonal trafficking defects and cell death,⁶ and rescued motility defects in a *Caenorhabditis elegans* model of HD.²² More recently, a selective inhibitor of DRP1, P110-TAT, was shown to inhibit the pathology of HD mouse models.²⁴ It was also reported that mHtt directly interacts with DRP1 and, thereby, increases DRP1 GTPase activity and its association with mitochondria.^{6,23,24} This may explain why we found that



Figure 6 Parkin-directed mitophagy is altered by mHtt in HdhQ111 striatal cells but is partially restored by PINK1. (a) Striatal cells derived from HdhQ7 and HdhQ111 knockin mice were transfected with mCherry-Parkin and treated by dimethyl sulfoxide (DMSO) or 10 μ M CCCP for 6 h. Mitochondria were immunostained by an anti-Tom20 antibody and nuclei were visualized by Hoechst stain. Arrowheads show translocation of mCherry-Parkin to the mitochondria in CCCP-treated cells. Scale bar: 10 μ m. (b) Quantitative analyses of the percentage of cells with Parkin translocated to the mitochondria. Data from at least three independent experiments (3–6 coverslips each) were averaged and were presented as mean \pm S.E.M. (c) HdhQ7 and HdhQ111 striatal cells were transfected with mCherry-Parkin and treated by DMSO or 10 μ M CCCP for 38 h. Mitochondria were immunostained by an anti-Tom20 antibody and nuclei were visualized by Hoechst stain. Arrowheads show cells with Parkin translocation and perinuclear mitochondria (lustering. White asterisk corresponds to a CCCP-treated cell devoid of Tom20-immunostained mitochondria. Scale bar: 10 μ m. (d) Quantitative analyses of the percentage of cells showing perinuclear clustering of the mitochondria (**9**) cuantitative analyses of the percentage of cells showing Parkin translocation to the mitochondria (e), the proportion of cells showing perinuclear clustering of the mitochondria (f) or cells with no mitochondria (g). HdhQ7 and HdhQ111 striatal cells were transfected with mCherry-Parkin together with mock plasmid or PINK1-HA and treated with 10 μ M CCCP for 38 h. Mitochondria were detected by Tom20 immunostaining. Data from at least three independent experiments (3–6 coverslips each) were averaged and were presented as mean \pm S.E.M. Statistical analysis was performed using a Student's *t*-test ($^{OOO}P < 0.001$ PINK1-transfected *versus* mock-transfected cells; ***P < 0.001 *versus* HdhQ7 cells).



Figure 7 The presence of mHtt alters the formation of mitochondria-containing autophagosomes. (a) HdhQ7 and HdhQ111 striatal cells were transfected with EGFP-Parkin, PINK1-HA and mRFP-ubiquitin (mRFP-Ub), and then treated by dimethyl sulfoxide (DMSO) or 10 μ M CCCP for 38 h. Mitochondria were visualized by anti-Tim23 immunostaining. Scale bar: 10 μ m. (b) Quantitative analyses showing the Mander's colocalization coefficient between mRFP-ubiquitin and mitochondria. Data from at least 30 cells from six coverslips were averaged and were presented as mean \pm S.E.M. (c) Striatal cells were transfected with mCherry-Parkin, PINK1-HA and EGFP-LC3, and then treated by DMSO or 10 μ M CCCP for 38 h. Mitochondria were visualized by anti-Tim23 immunostaining. Arrowheads show mitochondria that are not colocalized with EGFP-LC3 in CCCP-treated cells. Scale bar: 10 μ m. (d) Quantitative analyses showing the Mander's colocalization coefficient between EGFP-LC3 and mitochondria. Data from at least 30 cells from six coverslips were averaged and were presented as mean \pm S.E.M. Statistical analysis was performed using the Mann–Whitney *U*-test (****P*<0.001 *versus* HdhQ7 cells)

increasing DRP1 expression in our HD fly model exacerbates further the adult lethality.

In the present study, we detected the presence of abnormal mitochondria that adopt a ring-like form, termed mitochondrial spheroids. Such mitochondrial shape was rarely observed *in vivo* and their significance in pathological conditions remains to be fully investigated. So far, they were reported in injured livers of acetaminophen-overdosed mice and in retinal lesions of chicks reared in continuous light, suggesting that mitochondrial spheroids may appear in specific cell types npg

upon cellular stress.^{10,25} More importantly, mitochondrial spheroids were found in vitro after treating Parkin-deficient murine embryonic fibroblasts and HeLa cells with the mitochondrial uncoupler CCCP, an ionophore that usually induces mitophagy.¹⁰ In this study, Ding et al.¹⁰ showed that mitochondria engage in a self-fusion process when mitophagy is defective. Moreover, reactive oxygen species seem to be required since antioxidants suppress mitochondrial spheroid formation even in the presence of CCCP.¹⁰ Taking those data into account, our study suggests that the mitophagy pathway might be overwhelmed by the accumulation of damaged mitochondria in the presence of mHtt. leading to the formation of mitochondrial spheroids. This is consistent with mitochondrial spheroid formation being reduced when we overexpressed PINK1, which controls the activation of Parkinmediated mitophagy. Nevertheless, it is also possible that mHtt directly inhibits the mitophagy process, leading to the accumulation of defective mitochondria. In support of this hypothesis, we found that the presence of mHtt in mammalian HdhQ111 striatal cells perturbs CCCP-induced mitophagy but not Parkin translocation. Thus, HD mutation likely alters mitophagy downstream of Parkin translocation. Previous studies reported that mHtt could sequester proautophagic proteins and that there is a failure in the recognition of cargo by autophagosomes.^{26,27} Our data also provide evidence that formation of mitochondria-containing autophagosomes is disrupted by mHtt in HdhQ111 striatal cells. More recently, autophagosome transport was also found defective in primary neurons expressing mHtt, leading to inefficient degradation of engulfed mitochondria.²⁸ We show here that whereas HD mutation compromised mitophagy, PINK1 overexpression was still able to increase partially the removal of defective mitochondria in HdhQ111 cells. It is thus tempting to propose that overexpressing PINK1 might decrease the threshold of mitochondrial damage that must be crossed before mitophagy occurs. Then, PINK1 might accelerate the removal of defective mitochondria against the progressive impairment of autophagy.

Previous studies have reported that PINK1 overexpression is neuroprotective in the context of PD. PINK1 prevents cell death induced by the parkinsonian neurotoxin 1-methyl-4phenyl-1,2,3,6-tetrahydropyridine²⁹ or apoptotic inducers in cell cultures.³⁰ Similarly, overexpression of PINK1 ameliorates the lifespan and mobility of *Drosophila* expressing the PD-related gene, *a*-synuclein, in dopamine neurons.³¹ Mitochondrial stress mechanisms are not restricted to PD but are common to other neurodegenerative diseases. We provide here, for the first time, compelling evidence that PINK1 pathway mitigates neuronal loss and increases life expectancy in an animal model of HD.

PINK1 led to intensive *in vitro* studies to elucidate how it controls mitochondrial quality control. The mitochondrial chaperone TRAP1 might be a direct substrate of PINK1, allowing the refolding/removal of misfolded proteins and preventing apoptotic cytochrome *C* release.¹⁸ However, our data suggest that TRAP1 overexpression does not mimic PINK1-mediated protection in HD flies. At the organelle level, PINK1 facilitates mitophagy through the recruitment of Parkin to dysfunctional mitochondria.^{7–9} Moreover, direct phosphorylation of Parkin at Ser65 by PINK1 seems to be required to initiate mitophagy.^{32–34} Then, Parkin might tag outer membrane mitochondrial substrates for proteasomal

degradation and/or for autophagosome recognition. We here demonstrated that whereas Parkin is essential to mediate PINK1 neuroprotection, overexpressing Parkin alone was ineffective to confer protection against mHtt toxicity. This is in agreement with the crucial role of PINK1 in determining which mitochondria must be removed by mitophagy. Moreover, ubiguitination of Mfn by Parkin may account for its rapid removal from the mitochondria, therefore preventing damaged mitochondrial fragments to fuse back and corrupt healthy mitochondria.^{35,36} However, in our study, silencing Mfn failed to ameliorate the survival of HD flies and abolished the beneficial effect of PINK1 on neuronal survival. Mfn degradation is therefore not a sufficient condition to mimic PINK1 effect. On the contrary, our data suggest that normal Mfn expression is required for PINK1-mediated neuroprotection. This is consistent with recent data which demonstrate that in mammals Mfn2 is phosphorylated by PINK1 and promotes PINK1-dependent translocation of Parkin to the mitochondria.³⁷ Finally, we also showed that VDAC/Porin is involved in PINK1-mediated reduction of mHtt toxicity in Drosophila. Whether or not VDAC is necessary for PINK1/ Parkin pathway remains controversial. Mammalian VDAC1 was identified as a target for Parkin-mediated ubiguitination and to promote mitophagy in HeLa cells.¹⁶ In contrast, while ubiquitination of VDAC1 was confirmed in MEF cells with damaged mitochondria, VDAC1 appeared dispensable for mitochondrial clustering and mitophagy in those cells.²¹ The role of VDAC in mitophagy may be redundant and depend on cell types. More recently, it was hypothesized that VDAC may serve as a docking site to recruit Parkin from the cytosol to defective mitochondria.¹⁷ Finally, PINK1 could interact with the proautophagic protein Beclin1, which is also sequestered by mHtt.²⁶ Further studies are required to fully understand in which conditions and how Mfn and VDAC participate in PINK1/ Parkin pathway.

In conclusion, we propose that mitophagy may reach a saturation stage in the presence of mHtt and that increasing PINK1/Parkin pathway may improve mitochondrial integrity in HD. It is therefore conceivable that PINK1/Parkin pathway may be of therapeutic interest not only in PD but also in other mitochondrial disorders.

Materials and Methods

Drosophila strains. Flies were raised on a standard agar/cornmeal/yeast diet at 25 °C. Fly stocks carrying $PINK1^{Bg}$, $PINK1^{OE}$ and $parkin^{OE}$ were from Pr. J Chung (Taejon, Korea), flies carrying $DRP1^{OE}$ and $mart^{IR}$ from Dr. M Guo (Los Angeles, CA, USA), flies carrying $TRAP1^{OE}$ (lines 1 and 2 M) from Dr. LM Martins (Leicester, UK). All other strains were obtained from the Bloomington *Drosophila* Stock Center (Bloomington, IN, USA). In accordance with the genetic background, we used w^{1118} flies as controls.

Electron microscopy. Adult fly heads were fixed in 2% paraformaldehyde, 2.5% glutaraldehyde, 5mM CaCl₂ and 0.1mM sodium cacodylate for 24 h at 4 °C. Postfixation was performed in 2.5% glutaraldehyde, 0.8% osmium tetroxide and 0.1 mM sodium cacodylate for 2 h at 4 °C. After staining with 2% uranyl acetate and dehydratation, samples were included in Epon resin. Ultrathin sections were examined with a Philips CM10 (Eindhoven, The Netherlands) or a Leo 912 Zeiss (Zeiss Microscopy, Jena, Germany) microscope at 100 kV.

Pseudopupil analysis. Pseudopupil analysis was performed as described previously.¹² The number of photoreceptors was determined from 20 ommatidia per fly. Data from at least seven flies per condition were averaged and were presented

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as mean \pm S.E.M. Statistical analysis was performed using the non-parametric Mann–Whitney U-test.

Longevity assays. Survival analysis was performed on female flies grouped by 15–20 in independent vials. Each genotype was evaluated at least two times. Survival curves were generated by using the GraphPad Prism software (La Jolla, CA, USA) and statistical analysis was performed by log-rank test. To establish a high-confidence list of modifier genes, we considered significant only genes that modify life expectancy for more than the lifespan variance of Httex1p Q93expressing flies between tubes within each trial.

Measurement of ATP levels. ATP measurements were essentially performed according to the manufacturer's instructions (Luminescent ATP Detection Assay Kit; Abcam, Cambridge, UK). For each experiment, five adult fly brains were rapidly dissected, and the extracts were immediately deproteinized by perchloric acid precipitation. Luminescence was measured with a TriStar LB941 luminometer (Berthold Technologies, Bad Wildbad, Germany). Data from at least five independent experiments were averaged and were presented as mean \pm S.E.M. Statistical analysis was performed using a Student's *t*-test.

Striatal cell culture, transfection and CCCP treatment. Analysis of mitophagy was performed on striatal cell lines derived from wild-type and mHtt knock-in mice: HdhQ7 and HdhQ111 (from Coriell Cell Repositories, Camden, NJ, USA).²⁰ Cells were grown in Dulbecco's modified Eagle's medium (Invitrogen, Carlsbad, CA, USA) supplemented with 10% fetal bovine serum (Invitrogen), penicillin, streptomycin and G418 (Invitrogen) and maintained at 33 °C in a 5% CO₂ incubator. Striatal cells were seeded at 100 000 cells per coverslip density in 12-well plates for immunostaining. Transfection with mCherry-Parkin (generous gift from Richard J Youle, Bethesda, MD, USA), EGFP-Parkin (generous gift from Ricardo Vago, Milan, Italy), EGFP-LC3 (Addgene, Cambridge, MA, USA) or mRFP-Ubiquitin (Addgene) plasmids was performed using Lipofectamine (Invitrogen) according to the manufacturer's instructions. One day after transfection, dissipation of mitochondrial membrane potential was achieved with 10 μ M CCCP (prepared in DMSO) for 6 or 38 h.

Immunostaining. HdhQ7 and HdhQ111 striatal cells were washed with phosphate-buffered saline (PBS; Invitrogen), fixed in 4% paraformaldehyde for 10 min, permeabilized in 0.1% Triton X-100 in PBS for 10 min and incubated for 2 h at room temperature with anti-Tom20 (Santa Cruz Biotechnology, Dallas, TX, USA) or anti-Tim23 antibodies (BD Biosciences, Franklin Lakes, NJ, USA) or anti-HA antibodies for PINK1 detection (Roche, Meylan, France) diluted in 3% bovine serum albumin. Then, cells were incubated for 45 min with secondary antibodies prepared in 3% bovine serum albumin and mounted in Mowiol (Calbiochem, La Jolla, CA, USA). Quantitative analyses were achieved using an AX10 Zeiss Apotome (Zeiss Microscopy). Proportion of cells showing mitochondrial Parkin translocation, mitochondrial perinuclear clustering or with no detectable mitochondrial markers were evaluated on at least 70 cells per coverslip. Data from three independent experiments with 3-6 coverslips each were averaged and were presented as mean ± S.E.M. Statistical analysis was performed using a Student's t-test. Confocal image acquisition was performed on a Zeiss LSM780 laser scanning (Zeiss Microscopy) microscope. Colocalization between mRFP-ubiquitin or EGFP-LC3 with Tim23-immunostained mitochondria was evaluated on a minimum of 30 cells from six coverslips. Mander's colocalization coefficient was calculated with the JACoP plugin of the Image J software. Data were presented as mean ± S.E.M. and statistical analysis was performed using the non-parametric Mann-Whitney U-test.

Conflict of Interest

The authors declare no conflict of interest.

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