

Pink1 regulates mitochondrial dynamics through interaction with the fission/fusion machinery

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Mitochondria form dynamic tubular networks that undergo frequent morphological changes through fission and fusion, the imbalance of which can affect cell survival in general and impact synaptic transmission and plasticity in neurons in particular. Some core components of the mitochondrial fission/fusion machinery, including the dynamin-like GTPases Drp1, Mitofusin, Opa1, and the Drp1-interacting protein Fis1, have been identified. How the fission and fusion processes are regulated under normal conditions and the extent to which defects in mitochondrial fission/fusion are involved in various disease conditions are poorly understood. Mitochondrial malfunction tends to cause diseases with brain and skeletal muscle manifestations and has been implicated in neurodegenerative diseases such as Parkinson's disease (PD). Whether abnormal mitochondrial fission or fusion plays a role in PD pathogenesis has not been shown. Here, we show that Pink1, a mitochondria-targeted Ser/Thr kinase linked to familial PD, genetically interacts with the mitochondrial fission/fusion machinery and modulates mitochondrial dynamics. Genetic manipulations that promote mitochondrial fission suppress *Drosophila* Pink1 mutant phenotypes in indirect flight muscle and dopamine neurons, whereas decreased fission has opposite effects. In *Drosophila* and mammalian cells, overexpression of Pink1 promotes mitochondrial fission, whereas inhibition of Pink1 leads to excessive fusion. Our genetic interaction results suggest that Fis1 may act in-between Pink1 and Drp1 in controlling mitochondrial fission. These results reveal a cell biological role for Pink1 and establish mitochondrial fission/fusion as a paradigm for PD research. Compounds that modulate mitochondrial fission/fusion could have therapeutic value in PD intervention.

Drosophila | Parkinson's disease | muscle | neurodegeneration | mitochondrial morphogenesis

Parkinson's disease (PD) is the most common movement disorder characterized by relatively selective degeneration of dopamine (DA) neurons. Postmortem studies have identified common features associated with PD, such as mitochondrial dysfunction, oxidative stress, and aggregation of abnormal proteins (1, 2). Although most PD cases are sporadic, recent studies have highlighted the importance of genetic factors in PD pathogenesis (2, 3). At least 10 distinct loci have been associated with familial forms of PD (FPD). Understanding the molecular lesions associated with these FPD genes promises to shed light on the pathogenesis of the more common forms of disease. Interestingly, studies using animal models and cell culture models have linked mutations of FPD genes to impairments of mitochondrial structure/function. Furthermore, some of these disease-associated proteins have been shown to be present in mitochondria or interact with mitochondrial proteins (4–7), reinforcing the general involvement of mitochondrial dysfunction in PD pathogenesis.

Recent studies of Pink1 have provided direct support for a causal role of mitochondrial dysfunction in PD pathogenesis. Pink1 encodes a predicted Ser/Thr kinase with a predicted mitochondrial targeting signal (5). In *Drosophila*, inactivation of

dPink1 leads to indirect flight muscle (IFM) degeneration, DA neuron loss, photoreceptor loss, and male sterility (8–11). These effects are generally accompanied by morphological and functional defects in the mitochondria of the affected tissues. Significantly, at least in IFM, the occurrence of abnormal mitochondria precedes tissue degeneration (8), supporting a causative role for mitochondrial dysfunction in dPink1-induced muscle cell death. Furthermore, genetic studies in *Drosophila* have revealed that Parkin acts downstream of Pink1 to maintain mitochondrial function and tissue integrity (8–10).

Despite these studies, the exact mitochondrial process that is regulated by Pink1 remains elusive. Intrigued by the mitochondrial morphological phenotypes induced by dPink1 inactivation, we have examined whether Pink1 regulates mitochondrial fission/fusion, which are dynamic membrane remodeling processes that govern mitochondrial morphology in eukaryotes (12). Our results indicate that dPink1 exhibits genetic interactions with components of the mitochondrial fission/fusion machinery and it regulates mitochondrial dynamics. Our findings link mitochondrial fission/fusion to PD pathogenesis and suggest ways to understand and treat PD and related disorders.

Results

Genetic Interactions Between Pink1 and Components of the Mitochondrial Fission and Fusion Pathways. Swollen mitochondria appeared in IFM of dPink1 mutant before overt myofiber degeneration (8), suggesting that defective mitochondrial morphogenesis is a primary effect of dPink1 inactivation. To test this possibility, we manipulated the activities of the mitochondrial fission and fusion pathways. Increasing mitochondrial fission by introducing an extra copy of Drp1 (13), a key component of the fission machinery, efficiently rescued the wing posture phenotype caused by IFM degeneration in dPink1 mutant [Fig. 1 *B* and *F* and supporting information (SI) Tables S1 and S2]. Conversely, Drp1 heterozygosity significantly enhanced dPink1 mutant phenotype (Table S1). Heterozygosity of Opa1-like (14), a core component of the fusion machinery, also significantly suppressed dPink1 mutant phenotype (Fig. 1 *C* and *G* and Tables S1 and S2), although the effect subsided in older animals. Similar manipulation of Drp1 or Opa1-like did not affect a retinal degeneration phenotype caused by inhibition of DJ-1a (15), a *Drosophila* homologue of another familial PD gene affecting mitochondrial function (Fig. S1). This result demon-

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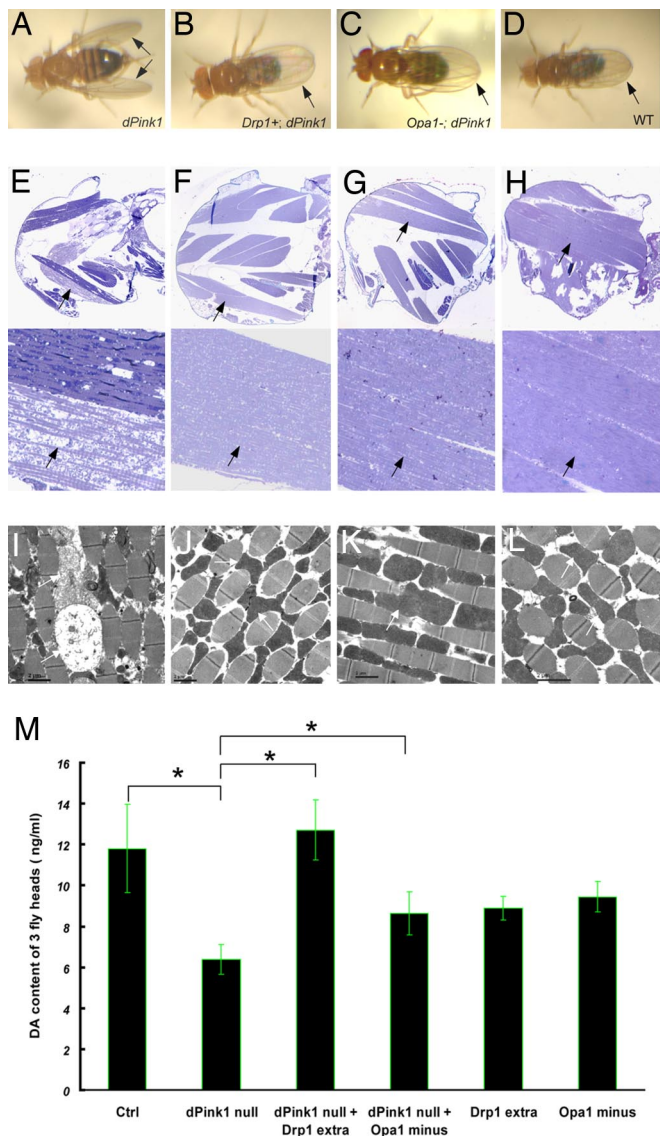


Fig. 1. Promotion of mitochondrial fission suppresses mutant phenotypes caused by *dPink1* loss of function. (A–D) Wing posture phenotypes in 5-day-old adult flies of the following genotypes: *dPink1*^{B9} (or B9) (A), *dPink1*^{B9}; *Drp1* extra copy (B), *dPink1*^{B9}; *Opa1-like*⁺ (C), and wild-type control (D). The arrows point to the abnormal wing posture in A and normal posture in B–D. (E–H) (Upper) Longitudinal sections of thoraces in 5-day-old adult flies of the following genotypes: *dPink1*^{B9} (E), *dPink1*^{B9}; *Drp1* extra copy (F), *dPink1*^{B9}; *Opa1-like*⁺ (G), and control (H). (Lower) Magnified views of corresponding IFMs. Sections from resin-embedded thoraces of 5-day-old adult flies were stained with toluidine blue to visualize tissue morphology, particularly the musculatures; anterior is to the left. The arrows point to IFMs. (I–L) Transmission electronic microscopy analysis of IFM ultrastructure in 5-day-old adult flies of the corresponding genotypes shown above. The arrows point to mitochondria. (M) Added expression of *Drp1* or loss of one copy of *Opa1-like* suppressed the reduction of head dopamine (DA) content caused by loss of *dPink1*. Two-week-old flies were subject to DA measurement. *, $P < 0.01$, one-way ANOVA test.

strates specificity of the genetic interactions between *dPink1* and the mitochondrial fission/fusion genes.

We next examined tissue integrity and mitochondrial morphology in the IFM of rescued animals. In control animals, spindle-shaped IFM fibers were aligned in an organized manner, with mitochondria packed in-between muscle fibers (Fig. 1L). These mitochondria contained electron-dense material and cris-

tae. IFM fibers were thin, atrophic, and disorganized in *dPink1* mutant (Fig. 1I). Some of the mitochondria appeared grossly enlarged, with inner membrane disorganized or disintegrated and overall electron density decreased (Fig. 1I). *Drp1* overexpression restored mitochondrial morphology and IFM morphology and organization (Fig. 1J). *Opa1*-like heterozygosity also showed effective rescue (Fig. 1K). *Drp1* overexpression or *Opa1*-like heterozygosity had little effect on IFM organization or mitochondrial morphology in an otherwise wild-type background (Fig. S2). The rescue of IFM pathology of *dPink1* mutant by the mitochondrial fission and fusion genes thus correlates with the restoration of mitochondrial morphology.

To test whether the genetic interaction between *dPink1* and mitochondrial fission/fusion genes happens in other tissues, we analyzed DA neurons. Inactivation of *dPink1* leads to dysfunction of DA neurons, as indicated by the reduction of brain DA levels (8, 9). *Drp1* overexpression restored DA levels to normal in *dPink1*-null mutant background (Fig. 1M). *Opa1*-like heterozygosity also showed rescuing effect (Fig. 1M), although not as robust as *Drp1* overexpression. *dPink1* mutants overexpressing *Drp1* or heterozygous for *Opa1*-like have a normal number of dopaminergic neurons (data not shown). It should be noted that the degree of brain DA-level reduction ($\approx 50\%$) is much greater than the minor or no-DA-neuron loss seen in *dPink1*-null mutants (9, 10), suggesting that, in addition to promoting DA neuron survival, *Pink1* may play a more prominent role in regulating dopaminergic physiology, such as DA metabolism, storage, or transmission.

Pink1 Cooperates with Mitochondrial Fission/Fusion Proteins to Regulate Mitochondrial Morphogenesis in DA Neurons. We next examined the effect of manipulating fission/fusion genes on mitochondrial morphology in the DA neurons of *dPink1* mutant by using a mito-GFP reporter that targets GFP to the mitochondrial matrix. Mito-GFP was expressed in DA neurons using the tyrosine hydroxylase (TH)-Gal4 driver. The mito-GFP reporter clearly labeled mitochondrial networks, which contained tubular and punctate units (Fig. 2A, A', and J). We focused our analysis on DA neurons in the dorsolateral protocerebral posterior (PPL1 or DL1) clusters. In *dPink1* mutant or *dPink1* RNAi animals, prominent mitochondrial aggregates formed. In addition, tubular structures were frequently observed (Fig. 2B, B', C, C', E, and J). Mito-GFP aggregates were also observed in other DA neuron clusters of *dPink1* mutant (data not shown). In *dPink1* mutant overexpressing *Drp1*, however, mitochondrial aggregates were no longer formed, and mito-GFP signals were uniformly distributed throughout the soma (Fig. 2F and J), as seen in the control (Fig. 2A, A', D, and J). Removal of one copy of *Drp1* reversed the rescuing effect of *Drp1* overexpression conferred by the extra copy (Fig. S3A). *Drp1* overexpression by introducing an extra copy in an otherwise wild-type background showed no obvious effect on mitochondrial network morphology (Fig. 2G and J). Similarly, *Opa1*-like heterozygosity also had no obvious effect in a wild-type background (Fig. 2I and J), but it suppressed mitochondria aggregation in *dPink1* mutant (Fig. 2H and J). In contrast, loss of one copy of the mitochondrial manganese superoxide dismutase (*MnSOD1*) gene had no effect on *dPink1* mutant phenotype (Fig. S3D). These results are consistent with *dPink1* specifically affecting mitochondrial fission/fusion dynamics.

Pink1 Overexpression also Leads to Abnormal Mitochondrial Morphogenesis in DA Neurons. To test whether *dPink1* is sufficient to induce changes in mitochondrial morphology, we overexpressed it in DA neurons. Surprisingly, *dPink1* overexpression also induced mitochondrial clustering (Fig. 3A and A'). However, different from that in *dPink1* mutant, the nonaggregated mitochondria appeared spherical in *dPink1* overexpression neurons,

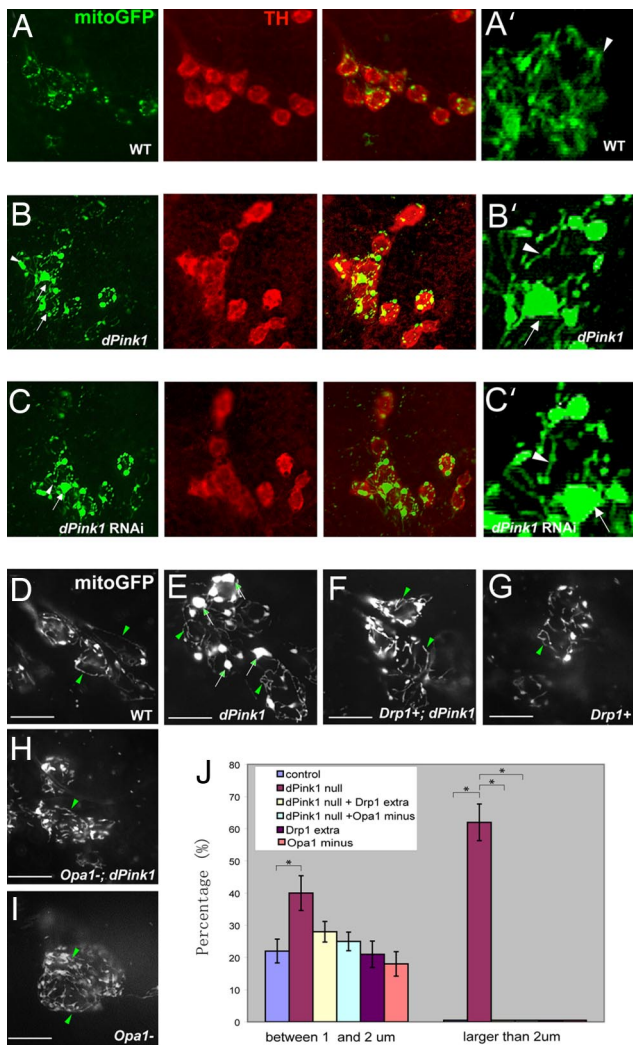


Fig. 2. Up-regulation of mitochondrial fission prevents mitochondrial clustering in DA neurons caused by loss of *dPink1*. (A–C) Whole-mount immunostaining of dorsolateral protocerebral posterior (PPL1 or DL1) cluster DA neurons in 5- to 7-day-old adult flies of the following genotypes: *TH-Gal4>UAS-mitoGFP* (A), *TH-Gal4>UAS-mitoGFP; dPink1^{B9}* (B), and *TH-Gal4>UAS-mitoGFP; UAS-dPink1 RNAi* (C). (A–C) (Left) Immunostaining of GFP to monitor mitochondria. (Center) Immunostaining of TH to mark DA neurons. (Right) Merged images. (D–I) Live imaging of mitochondria by visualizing mitoGFP within the PPL1 cluster neurons in 5- to ≈7-day-old adult flies of the following genotypes: *TH-Gal4>UAS-mitoGFP* (D), *TH-Gal4>UAS-mitoGFP; dPink1^{B9}* (E), *TH-Gal4>UAS-mitoGFP; dPink1^{B9}; Drp1 extra copy* (F), *TH-Gal4>UAS-mitoGFP; Drp1 extra copy* (G), *TH-Gal4>UAS-mitoGFP; dPink1^{B9}; Opa1-like/+* (H), and *TH-Gal4>UAS-mitoGFP; Opa1-like/+* (I). A', B', and C' show higher magnification views of the mitochondrial networks in A–C, respectively. Arrows, aggregated mitochondria; arrowheads, tubular mitochondria. (J) Quantification of the percentage of DA neurons containing mitochondrial aggregates with diameters of 1–2 or >2 μm. Note that mitochondrial aggregates >2 μm in diameter were observed almost exclusively in *dPink1* mutant. *, $P < 0.01$, one-way ANOVA test.

and tubular structures were rarely observed. Furthermore, Drp1 overexpression (Fig. 3F) or Opa1-like heterozygosity (Fig. 3H) could not modify this *dPink1* overexpression effect, suggesting that the mitochondrial aggregates in *dPink1* mutant and *dPink1* overexpression neurons are intrinsically different. To further verify this point, we tested the effect of Parkin overexpression in these backgrounds. Parkin overexpression can efficiently suppress *dPink1* mutant phenotypes (8–10). Parkin overexpression efficiently blocked the formation of mito-GFP aggregates in

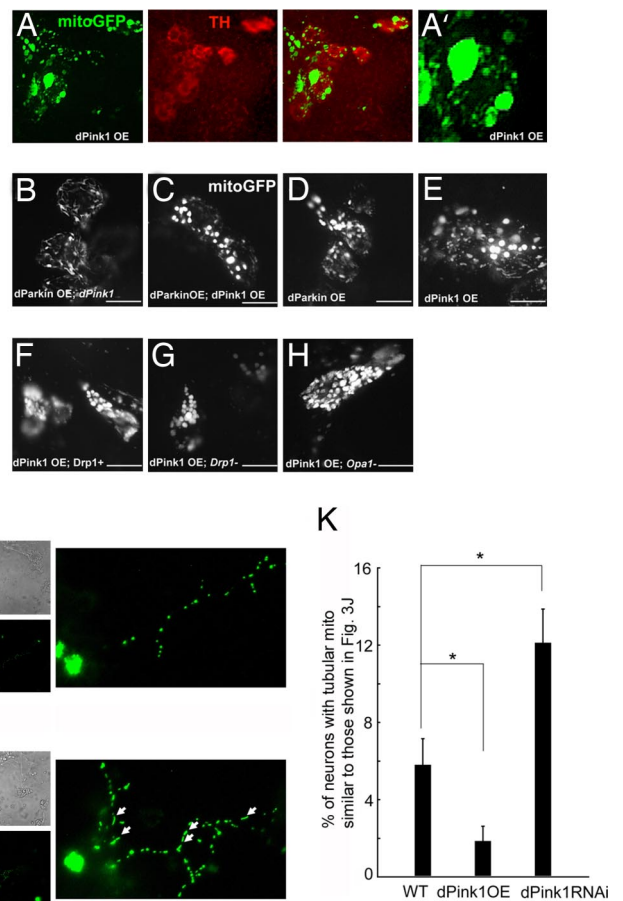


Fig. 3. *dPink1* overexpression causes mitochondrial morphological changes in DA neurons. (A) Whole-mount immunostaining of PPL1 cluster DA neurons in 5- to ≈7-day-old adult flies of the genotype *TH-Gal4>UAS-mitoGFP, UAS-dPink1*. (Left) Immunostaining of mitoGFP. (Center) Immunostaining of TH. (Right) Merged image. (B–H) Live imaging of mitochondria within the PPL1 cluster neurons in 5- to ≈7-day-old adult flies of the following genotypes: *TH-Gal4>UAS-mitoGFP; dPink1^{B9}; UAS-dParkin* B, *TH-Gal4>UAS-mitoGFP; UAS-dPink1; UAS-dParkin* C, *TH-Gal4>UAS-mitoGFP; UAS-dParkin* D, *TH-Gal4>UAS-mitoGFP; UAS-dPink1* (E), *TH-Gal4>UAS-mitoGFP; UAS-dPink1; Drp1 extra copy* (F), *TH-Gal4>UAS-mitoGFP; UAS-dPink1; Drp1/+* (G), *TH-Gal4>UAS-mitoGFP; UAS-dPink1, Opa1-like/+* (H). A' shows higher magnification view of the mitochondrial network in A. (I and J) representative images of 1-week-old cultured wild-type fly DA neurons exhibiting punctate-like (I) and thread-like (J) mitochondria in neuronal processes. Left images show bright-field microscopy views of the general neuronal culture (Upper) and TH⁺ neurons with mitochondria labeled with mito-GFP (Lower); the right images display magnified views of neuritic mitochondria of TH⁺ neurons. Arrows point to typical thread-like tubular mitochondria in the neurites. (K) Statistical analysis of the percentage of TH⁺ neurons with thread-like tubular mitochondria in their processes in the indicated genotypes. On average, ≈200 TH⁺ neurons for each genotype were counted. *, $P < 0.01$, one-way ANOVA test.

dPink1 mutant (Fig. 3B), but not in *dPink1* overexpression background (Fig. 3C). Interestingly, Parkin overexpression alone in wild-type background also led to a mitochondrial phenotype similar to that seen in *dPink1* overexpression (Fig. 3D). To rule out the possibility that this phenotype was nonspecifically caused by overexpression, we overexpressed DJ-1a in DA neurons. No obvious effect on mitochondrial morphology was seen in this situation (Fig. S3B). The fact that mitochondrial morphology was restored by the combined actions of *dParkin* overexpression and *dPink1* loss of function, despite the abnormal mitochondrial morphology in either condition alone, strongly suggests that the

two manipulations exert opposite effects on mitochondrial dynamics. Further supporting a functional difference between the two mitochondrial states, overexpression of Pink1 or Parkin has no detrimental effect on DA neuron function (8), whereas inhibition of Pink1 does. Overexpression of Pink1 in IFM or the eye also has no effect (data not shown). A recent study showed that Pink1 overexpression induced a mild rough eye phenotype (16), the physiological relevance of which remains to be tested, because dPink1-null mutant flies do not show any eye phenotype.

The mitochondrial aggregation phenotype seen in dPink1 overexpression background could be caused by excessive mitochondrial fission, as observed in mammalian cells overexpressing the pro-fission protein Fis1 (17, 18). We began to test this possibility by attenuating fission via removal of one copy of Drp1 in dPink1 overexpression background, but did not observe an obvious effect (Fig. 3G). Stronger attenuation of mitochondrial fission by other means might be needed to mitigate the dPink1 overexpression effect.

To further test the activity of dPink1 in regulating mitochondrial dynamics, we analyzed mitochondrial morphology in cultured DA neurons, which allow better resolution of mitochondrial networks and units, especially in neuronal processes. In control neuronal cultures that specifically express mito-GFP in DA neurons, most neurons possess punctate, short mitochondria (Fig. 3J), and only a small number of them contain tubular mitochondria (Fig. 3J). In dPink1 mutant neuronal culture, however, there were more neurons possessing tubular mitochondria (Fig. 3K). In contrast, the number of neurons having long tubular mitochondria was significantly reduced after dPink1 overexpression (Fig. 3K). These results support the notion that dPink1 promotes mitochondrial fission.

Characterization of the Relationships Between Pink1 and Drp1 in Cultured S2R Cells. To gain more information on the activity of dPink1 and its relationship with the fission machinery, we used *Drosophila* S2R cells, which allow efficient gene knockdown by RNAi. The larger size of S2R cells also allows better imaging of the mitochondrial network in the soma. In *gfp* dsRNA-transfected control cells, mitochondria were uniformly distributed as punctate units and tubular segments (Fig. 4A). Transfection of Drp1 dsRNA knocked down Drp1 mRNA level (Fig. S4), and caused the mitochondrial network to collapse into one large perinuclear aggregate (Fig. 4C). Transfection of dPink1 dsRNA led to a similar phenotype (Fig. 4B and Fig. S4), albeit at a lower frequency ($35.6 \pm 5.2\%$ for dPink1 RNAi vs. $97.5 \pm 3.6\%$ for Drp1 RNAi). To further test the functional relationship between Drp1 and Pink1, we performed Drp1 and dPink1 double RNAi. In these cells, perinuclear aggregation of mitochondria persisted. Remarkably, a distinct phenotype not seen in single RNAi condition was observed, which was manifested as long continuous threads of mitochondria indicative of extreme fusion ($27.6 \pm 5.4\%$) (Fig. 4D). Double RNAi of Drp1 and another key mitochondrial gene *MnSODII* did not have the same effect (Fig. S5), suggesting that mere disruption of mitochondrial function does not lead to extreme mitochondrial fusion, and that the interaction between Pink1 and Drp1 in regulating mitochondrial morphology is specific.

Pink1 Functionally Interacts with Fis1 to Regulate Mitochondrial Morphogenesis in *Drosophila* and Mammalian Cells. To further examine the role of Pink1 in mitochondrial fission, we explored the functional relationship between Pink1 and Fis1, another key player in the fission pathway. Transfection of an EGFP-dFis1 fusion protein but not EGFP efficiently suppressed the mitochondrial aggregation phenotype caused by dPink1 RNAi in *Drosophila* S2R cells (Fig. 4E and F) ($3.6 \pm 1.2\%$ for EGFP-dFis1 vs. $41.8 \pm 6.6\%$ for EGFP). In contrast, EGFP-dFis1 was

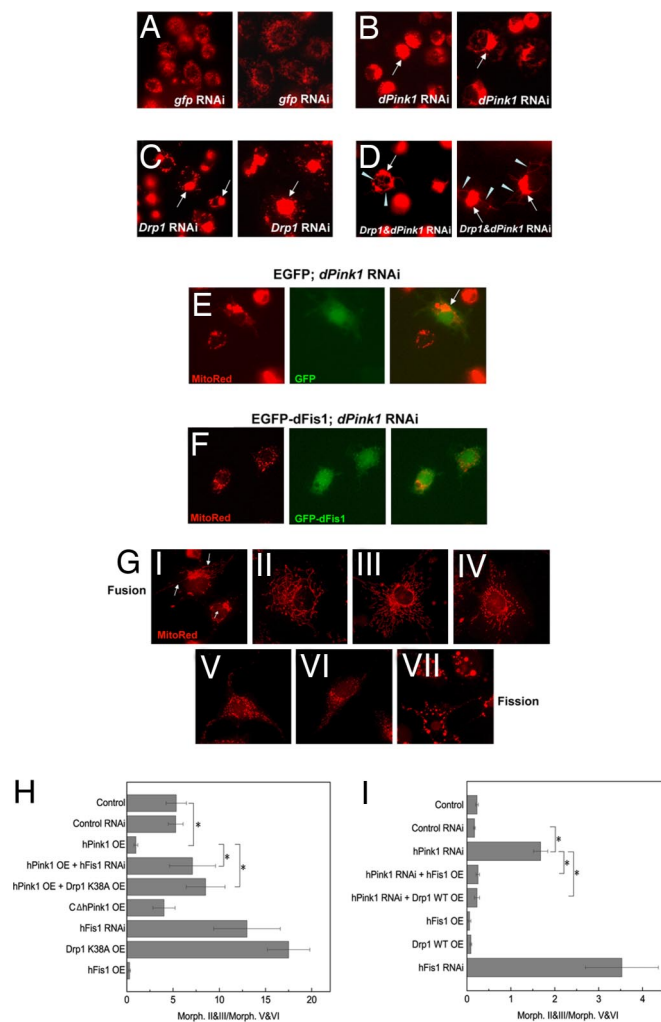


Fig. 4. Genetic interactions between Pink1 and fission genes in modulating mitochondrial dynamics in *Drosophila* and mammalian cells. (A–D) Live imaging of mitochondria in S2R cells subjected to the following genetic manipulations: *gfp* RNAi (A), *dPink1* RNAi (B), *Drp1* RNAi (C), *dPink1* and *Drp1* double RNAi (D). Note the similar phenotype of mitochondrial aggregation (arrows) caused by *dPink1* RNAi and *Drp1* RNAi despite the lower percentage in the case of *dPink1* RNAi. The arrowheads point to thin continuous mitochondrial tubules extending beyond mitochondrial clusters, which appeared exclusively in D. S2R cells are not homogenous in size. Two types of cells, smaller-sized (Left) and bigger-sized (Right), are shown for each genetic manipulation. The effects are similar in both sized cells. (E and F) Live imaging of mitochondria in S2R cells subject to the following genetic manipulations: *dPink1* RNAi plus EGFP transfection (E); *dPink1* RNAi plus EGFP-dFis1 transfection (F). EGFP-dFis1 (E) but not EGFP (F) suppressed mitochondrial clustering induced by *dPink1* RNAi. The arrow marks aggregated mitochondria. (G) Seven categories of mitochondrial morphology in mammalian COS-7 cells ranging from extreme fusion (type I) to extreme fission (type VII), as revealed by MitoTracker Red staining. The categories are arbitrarily defined based on the following mitochondrial characteristics: I, perinuclear aggregates together with long continuous tubules; II, extensive, long tubular network; III, intermediate-length tubular network; IV, a mixture of short tubular mitochondria and punctate units; V, small round tubules with holes in the center; VI, small punctate dots; VII, spherical aggregates similar to that seen in DA neurons overexpressing Pink1 or Parkin. To highlight the difference between categories I and VII, the long tubular mitochondria threads present in I but not VII are marked by arrows. (H and I) Statistical analysis showing that hPink1 overexpression promotes mitochondrial fission, which is prevented by the overexpression of Drp1^{K38A} (H), whereas hPink1 RNAi promotes mitochondrial fusion in COS-7 cells, which is rescued by hFis1 or hDrp1 overexpression (I). The ratio of cells exhibiting type II and III morphology typical of fusion versus type V and VI morphology typical of fission were quantified. *, $P < 0.01$, one-way ANOVA test.

unable to suppress similar phenotypes caused by Drp1 RNAi (Fig. S6).

To test whether the functional relationship between Pink1 and Drp1/Fis1 is conserved in mammals, we used COS-7 cells, which possess dynamic mitochondrial networks (Fig. 4G and Fig. S7), and can be efficiently transfected. Overexpression of hPink1 but not hPink1 Δ C resulted in more cells possessing punctate mitochondria, whereas hFis1 RNAi or overexpression of dominant-negative Drp1 (Drp1K38A) resulted in more cells having long tubular mitochondria (Fig. 4H). Overexpression of Drp1K38A in hPink1 overexpression background also resulted in more cells having long tubular mitochondria (Fig. 4H), suggesting that Drp1 may act downstream of Pink1. We then examined the effect of Pink1 knockdown in COS-7 cells. Expression of Pink1 shRNA but not a control shRNA knocked down hPink1 mRNA level (Fig. S8) and resulted in more cells having long tubular mitochondria (Fig. 4I). This effect was significantly suppressed by the overexpression of hFis1 or Drp1 (Fig. 4I). These results are consistent with Pink1, Fis1, and Drp1 being conserved positive regulators of mitochondrial fission and that Drp1 and Fis1 are epistatic to Pink1.

Discussion

Mitochondrial fission and fusion are membrane-remodeling processes that control the *in vivo* dynamic changes of the distribution and structure of the mitochondria network. These processes respond to energy status of the cell and are necessary for proper mitochondrial function. Dysfunction of mitochondrial fission/fusion has been linked to the pathogenesis of neurodegenerative diseases. For example, mutations in Opa-1 are associated with autosomal dominant optic atrophy (19), whereas mutations in Mfn2 cause the autosomal dominant disease Charcot-Marie-Tooth type 2A (20).

Our results demonstrate that Pink1 regulates mitochondrial morphogenesis and function through the fission/fusion pathway in IFM and DA neurons. These results link defects in mitochondrial fission/fusion to the pathogenesis of PD, a major neurodegenerative disease. A recent study reported similar genetic interactions between Pink1 and genes in the fission/fusion pathway in *Drosophila*, although their relationship in DA neurons, the PD-relevant cell type, was not shown in that study (16). We further show that Pink1 appears to act through Fis1 and Drp1 to regulate mitochondrial fission. Because Drp1 is recruited from the cytosol to fission sites on the outer surface of mitochondria (21, 22), whereas Pink1 is localized on the inner membrane cristae facing the intermembrane space (23), the action of Drp1 and Pink1 is likely coordinated by some intermediate protein(s) such as Fis1. Consistent with this model, direct interaction between Fis1 and Drp1 has been reported before (18). The biochemical relationship between Pink1 and Fis1 awaits further investigation. Studies in yeast have identified other proteins that interact with Drp1 or Fis1 and participate in mitochondrial fission/fusion, such as Mdv1p and Caf4p (12). Further studies will test the relationships between Pink1 and the *Drosophila* or mammalian counterparts of these proteins in regulating mitochondrial dynamics and DA neuron maintenance and function. Our results show that mammalian Pink1 interacts with components of the mitochondrial fission machinery and regulates mitochondrial fission/fusion, in a manner similar to that in *Drosophila*. This finding supports a conserved role for Pink1 in mitochondrial morphogenesis. A recent study in mammalian HeLa-7 cells showed that inhibition of Pink1 resulted in abnormal mitochondrial morphology and that this effect was rescued by Parkin overexpression (24). Intriguingly, the mitochondria in Pink1 knockdown cells appear fragmented, although some of the fragments appeared to have larger diameter. It is unclear in that case whether the phenotype was due to excessive fission or fusion. It is possible that the differential appearance of

the mitochondrial network in HeLa cells and Cos-7 cells after Pink1 knockdown is due to a cell type-specific effect, or variations in the degree of Pink1 inhibition by RNAi in the two studies.

Mitochondrial fission and fusion has been implicated in cell death regulation, with Drp1-mediated fission being proapoptotic (25–27) and Mitofusin-mediated fusion being antiapoptotic (28, 29). In dPink1 mutant background, Drp1-mediated mitochondrial fission is protective. Drp1-dependent mitochondrial fission also protects against Ca²⁺-mediated apoptosis (30). The role of mitochondrial fission/fusion in regulating cell survival is therefore context and likely cell type dependent. This is consistent with dPink1 mutant showing phenotypes in select tissues. Neurons may be particularly vulnerable to the imbalance of mitochondrial fission/fusion because of their unique morphology and the requirement for synaptic function, in which Drp1-regulated mitochondrial fission is involved (13, 31). For example, in cultured mammalian hippocampal neurons, it has been shown that the extension or movement of mitochondria into dendritic protrusions correlates with the development and morphological plasticity of spines. Inhibition of mitochondrial fission through the expression of dominant-negative Drp1 resulted in loss of synapses and dendritic spines (31). This raises the possibility that dysfunction of Pink1 may also result in loss of dopaminergic synapses, which ultimately leads to DA neuron degeneration. In this regard, it is interesting to note that knockout of mouse Pink1 leads to impairment of dopamine release and synaptic plasticity in the striatum (32). Further studies of the regulation and function of Pink1 in synaptogenesis promise to shed light on the newly recognized role of mitochondrial dynamics in PD pathogenesis.

Materials and Methods

Drosophila Genetics and Fly Stocks. Fly culture and crosses were performed according to standard procedures and raised at indicated temperatures. All general fly stocks and *Gal4* lines were obtained from *Drosophila* stock centers. *dPink1*-null mutant line *dPink1*^{B9} was a gift from Jongkeong Chung (Korea Advanced Institute of Science and Technology, Taejeon, Korea) (9). The *TH-GAL4* driver was a gift from Serge Birman (Centre National de la Recherche Scientifique–Université de la Méditerranée, Marseille, France). *UAS-mitoGFP* was a gift from William Saxton (Indiana University, Bloomington, IN). *Drp1* (or *fratboy*)-null mutant line *drp1*² and extra copy *Drp1* line FLAG-FLASH-HA-*drp1*⁺ were gifts from Patrick Verstreken and Hugo Bellen (Baylor College of Medicine, Baylor, TX). The other fly stocks were described previously: *UAS-dPink1* and *UAS-dPink1 RNAi* (8), *UAS-dParkin* (33), *UAS-DJ-1a* and *UAS-DJ-1a RNAi* (15), and *Opa1-like* mutant line P{EPgy2}CG8479 (14).

Molecular Biology. For cell culture transfection experiment, the *hPink1* and rhesus monkey *Drp1* cDNAs were cloned into pCDNA3 expression vectors containing the indicated epitope tags (Invitrogen). Human full-length *Pink1* (*hPink1*) and C-terminal truncated form (*hPink1* Δ C) were tagged with C-terminal FLAG epitope (8). *hFis1* RNAi construct was a gift from Michael T. Ryan (La Trobe University, Melbourne, Australia) (17). *hPink1* shRNA clones (ID TRCN0000007099 and TRCN0000007101) were purchased from Open Biosystems. *hFis1* construct was a gift from Yisang Yoon (Mayo Clinic and Foundation, Rochester, MN) (18). Rhesus monkey wild-type Drp1 and dominant-negative Drp1^{K38A} were gifts from E. Smirnova and A. M. van der Bliek (University of California, Los Angeles, CA) (34). dFis1 cDNA was acquired from DGRC. N-terminal EGFP was fused in-frame into dFis1 cDNA and subcloned into pAc-5.1 vector (Invitrogen). EGFP was similarly subcloned into pAc-5.1 vector, taking advantage of the built-in *actin* promoter. pCMV-Venus was a gift from Yuzuru Imai (Stanford University, Stanford, CA). Each construct was fully sequenced before performing transfection. dsRNAs of *Drp1*, *dPink1*, *MnSODII* (mitochondrial manganese superoxide dismutase), and *gfp* were generated according to standard protocols using *in vitro* transcription systems (Ambion and Epicenter). Primer sequences for individual genes are available on request. For RT-PCR analysis, total RNA was prepared by using an RNeasy kit (Qiagen). Details of the quantitative RT-PCR procedure were essentially as described in ref. 35.

Live Imaging of Mitochondrial Morphology in COS-7 Cells and S2R Cells: Visualizing Mitochondria in Fly DA Neurons. For details of these procedures, see *SI Materials and Methods*. Note that to facilitate the monitoring of effects of genetic manipulations on mitochondrial fission in COS-7 cells, observation was started 2–3 h after the cells were removed from the 37°C, 5% CO₂ chamber and placed at ambient temperature in the presence of MTRed dye. This treatment resulted in more control cells with tubular (or fused) mitochondria (Fig. S7), making it easier to score fission events induced by the genetic manipulations. This change in mitochondrial morphology is adaptive and transient, because when the control cells were transferred back to the 37°C, 5% CO₂ chamber, their mitochondrial morphology also showed corresponding changes.

Muscle Histology and Transmission Electron Microscopy Analysis. Muscle histology and transmission electron microscopy analysis were performed as described in ref. 33, except that Epon resin was used for embedding.

Dopamine Measurements. HPLC analysis of catecholamine levels was performed essentially as described in ref. 15. For sample preparation, adult male fly heads were dissected out and homogenized in 0.1 M perchloric acid (generally 50 μ l per four or five heads) by using a motorized hand-held tissue grinder. The homogenate was frozen immediately on dry ice and stored at –80°C before HPLC analysis.

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