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Expression of zinc-deficient human SOD1 in *Drosophila* neurons produces a locomotor defect linked to mitochondrial dysfunction

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Abstract

More than 130 different mutations in the Cu/Zn superoxide dismutase (SOD1) gene have been associated with Amyotrophic lateral sclerosis (ALS) but the mechanism of this toxicity remains controversial. To gain insight into the importance of the zinc site in the pathogenesis of SOD1 *in vivo*, we generated a *Drosophila* model with transgenic expression of a zinc-deficient human SOD1. Expression of zinc-deficient SOD1 in *Drosophila* resulted in a progressive movement defect with associated mitochondrial cristae vacuolization and reductions in ATP levels. Furthermore, these flies are sensitized to mitochondrial toxins, paraquat and zinc. Importantly, we show that the zinc-deficient SOD1-induced motor defect can be ameliorated by supplementing the endogenous fly respiratory chain machinery with the single-subunit NADH-ubiquinone oxidoreductase (*Ndi1*) from yeast. These results demonstrate that zinc-deficient SOD1 is neurotoxic *in vivo* and suggest that mitochondrial dysfunction plays a critical role in this toxicity. The robust behavioral, pathological and biochemical phenotypes conferred by zinc-deficient SOD1 in *Drosophila* have general implications for the role of the zinc ion in both familial and sporadic ALS.

Keywords

Amyotrophic Lateral Sclerosis (ALS); fly; *Ndi1*; paraquat; mitochondria; respiration; superoxide dismutase (SOD1)

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Introduction

Amyotrophic lateral sclerosis (ALS) patients suffer from progressive paralysis and muscle weakness that will eventually lead to respiratory failure and death. About 90% of ALS patients are sporadic cases whereas the other 10% are familial cases, 20% of which inherit mutations in the copper, zinc-superoxide dismutase (*Cu, Zn SOD* or *SOD1*) gene (Rosen et al., 1993; Vucic and Kiernan, 2009). The pathology and symptoms of SOD1-patients closely resemble those of sporadic ALS cases. Therefore, transgenic animal models expressing SOD1 mutations may provide not only insight into the pathogenic mechanisms of both familial and sporadic ALS, but also valuable tools to develop and test therapeutic approaches. Although it is known that mutations result in a toxic gain of function of the SOD1 enzyme, which normally functions as a free radical scavenger, the mechanisms underlying disease progression have not been clearly elucidated (Beckman et al., 2001; Bruijn et al., 2004).

In the last decade, perturbations in mitochondrial energy metabolism have emerged as a focal point in studies to elucidate the molecular mechanisms of aging and age-related diseases (Lin and Beal, 2006; Wallace, 2005). Mitochondria dysfunction has also been implicated in the pathophysiology of ALS (Hervias et al., 2006). Biochemical and morphological mitochondrial abnormalities have been demonstrated in postmortem spinal cords of ALS patients (Borthwick et al., 1999; Wiedemann et al., 2002; Wiedemann et al., 1998) and in rodent models of SOD1-linked ALS (Kong and Xu, 1998; Wong et al., 1995). Although it is not fully understood how mutant SOD1 could damage mitochondria, a number of recent studies have shown that ALS-linked SOD1 mutations are selectively recruited to mitochondria and induce respiratory defects (Ferri et al., 2006; Liu et al., 2004; Pasinelli et al., 2004; Vande Velde et al., 2008).

The SOD1 mutations associated with ALS occur in all of the functional elements of the protein structure, and many SOD ALS mutants have superoxide-scavenging activity comparable to that of wild-type SOD (Vucic and Kiernan, 2009; Bruijn et al., 2004). However, ALS mutations make the SOD1 mutants more susceptible to the loss of their metal cofactors (Crow et al., 1997; Lyons et al., 1996; Trumbull and Beckman, 2009). SOD has approximately a 7000-fold lower affinity for zinc than it does for copper (Crow et al., 1997) and, therefore, zinc is more likely to disassociate than copper. Zinc deficiency dramatically increases the reduction of the active-site copper by low molecular weight reductants such as ascorbate (Estévez et al., 1999). Delivery of zinc-deficient SOD into cultured motor neurons is sufficient to induce apoptosis, associated with increased redox activity of the remaining copper (Estévez et al., 1999). Furthermore, zinc-deficient wild-type SOD was shown to be just as toxic to cultured motor neurons as zinc-deficient mutant SOD. This observation leads to the idea that loss of zinc from wild-type SOD could be involved in the 98% of ALS patients without SOD mutations (Crow et al., 1997; Trumbull and Beckman, 2009). In the zinc-deficient SOD hypothesis, the mutations to SOD do not directly cause the toxic gain-of-function but rather increase the propensity of SOD to become zinc deficient (Crow et al., 1997).

A significant limitation of the ‘zinc deficient’ hypothesis has been a lack of *in vivo* data. Moreover, although mitochondrial dysfunction has been widely reported as an early event in ALS pathogenesis (Hervias et al., 2006), it is not yet known whether strategies to enhance respiratory chain function can improve the behavioral defects associated with ALS. In this study, we set out to examine the physiological and phenotypic consequences of expression of zinc-deficient SOD1 in the fruit fly *Drosophila*. A previous attempt to construct a *Drosophila* model of ALS demonstrated that expression of G41S mutant allele of human SOD in *Drosophila* motorneurons had no adverse effects on life-span but, somewhat surprisingly, enhanced survival both under normal and under oxidative stress conditions (Elia et al., 1999). Later studies also confirmed that expression of mutant SOD alleles in *Drosophila* does not result in a dominant gain-of-function as observed in humans and mice (Mockett et al., 2003). Recently, a *Drosophila* model of ALS was constructed by transgenic expression of mutant SOD1 (A4V and G85R) in motorneurons using *D42-GAL4* driver; the resulting strains exhibit a progressive loss of locomotor function along with an impaired neural circuit electrophysiology (Watson et al., 2008). While the latter models mimic certain phenotypes of ALS, these SOD1 alleles are not ideal for the purpose of our study in that the relatively short lifespan of the fruit flies may either be insufficient for these mutant alleles of SOD1 to lose their zinc-cofactor or it may simply be too late in the life cycle to induce any visible mitochondrial phenotypes. Consequently, we have constructed transgenic flies expressing zinc-deficient SOD throughout their entire lifespan. We report that zinc-deficient SOD1 expression confers an age-related movement disorder, a hallmark of ALS in humans. Moreover, zinc-deficient SOD1 expression produces alterations in mitochondrial structure as well as ATP production and sensitizes flies to mitochondrial toxins such as paraquat and zinc. Finally, we demonstrate that transgenic enhancement of respiratory chain function ameliorates the motor defect associated with zinc-deficient SOD1, suggesting that mitochondrial dysfunction may be an important intermediate in development of ALS associated phenotypes.

Results

Transgenic expression of D83S SOD (zinc-deficient SOD1 or CuSOD) in *Drosophila*

D83S is a zinc-deficient mutant SOD in which the zinc-binding residue Asp83 was mutated to Ser using PCR mutagenesis (Roberts et al., 2007). In this mutant protein, the zinc-binding site has no electron density for zinc; the copper-binding site, however, maintains its electron density for copper and the coordination of copper remains similar to that of wild-type SOD. Here, we expressed *D83S SOD* in flies using the GAL4/UAS system. We transformed flies with UAS-constructs containing the *D83S SOD* gene and performed ten rounds of backcrossing into a *w¹¹¹⁸* background. In all subsequent experiments, *w¹¹¹⁸* was used as a control strain. Except for the insertion of the *UAS-D83S SOD* transgene, these control flies were genetically very similar to the experimental flies. Activation of *UAS-D83S SOD* transgene under the control of the ubiquitous *da-GAL4* driver strongly (more than 5-fold) induces expression of SOD protein in transgenic flies, as assayed by Western blot analysis (Figure 1A). Next, we examined SOD activity in *UAS-D83S SOD* transgenic flies to determine whether or not *D83S* mutant SOD maintains its catalytic activity. As Figure 1B illustrates, expression of *D83S SOD* under the control of *da-GAL4* confers additional SOD activities in adult flies, as

evaluated by an in-gel NBT SOD assay. These activities are attributed to production of Cu, Zn SOD-CuSOD heterodimers and CuSOD-CuSOD dimers upon expression of *D83S* SOD1 transgene.

Effects of zinc-deficient SOD1 on fly lifespan, neurodegeneration and motor activity

Next, we tested the effects of zinc-deficient SOD1 on longevity, neurodegeneration and motor performance of adult flies. As Figure 1C and Table S1 illustrate, ubiquitous expression of wild-type human SOD1 under the control of the *da-GAL4* driver slightly shortens lifespan of adult flies (5%-7%). Importantly, expression of zinc-deficient SOD1 with the same GAL4 drives had no impact, negative or positive, on adult longevity (Figure 1C).

Next, we investigated whether expression of zinc-deficient SOD1 resulted in gross neurodegeneration in the fly brain or retina. As figures 1D and 1E illustrate, no gross anatomical degeneration of the brain or retina was observed in response to zinc-deficient SOD1 expression even at 60 days of age. Finally, we assessed motor activity in zinc-deficient SOD1 transgenic flies by measuring and comparing their locomotor activity to control lines; physical activity was quantified using a *Drosophila* Activity Monitor (DAM) system, in which two rings of infrared beams record fly movement in a small chamber. As Figure 1F and 1G illustrates, expression of zinc-deficient SOD1 produces an age-related movement disorder. At 10 days of age, zinc-deficient SOD1 expressing flies did not display reduced movement compared to isogenic controls. At 30 days of age, however, ubiquitous expression of zinc-deficient SOD1 reduces physical activity by 43% compared to isogenic controls; ubiquitous expression of wild-type human SOD1 had no adverse effect on physical activity. Superficially at least, this phenotype of our fly model is similar to that of ALS patients who suffer from a progressive paralysis.

Finally, to determine whether expression of zinc-deficient SOD resulted in a respiratory deficit, we measured ATP production in these flies. In doing so, we observed that ATP levels were decreased by >60% in the heads of flies expressing zinc-deficient SOD1 (Figure 1H).

Expression of zinc-deficient SOD leads to alterations in mitochondrial structure

A growing body of evidence suggests that mitochondrial dysfunction may play an important role in the pathophysiology of ALS (Hervias et al., 2006). Here, we examined the impact of zinc-deficient SOD1 expression on mitochondrial structure and function in *Drosophila*. Using electron microscopy, we identified a striking mitochondrial pathology within the flight muscle in response to zinc-deficient SOD1 expression. Insect flight muscle is well-suited for studies of mitochondrial ultrastructure. The mitochondria typically lie in columns, aligned longitudinally between the myofibrils (Figure 2). The cristae within these mitochondria are packed very densely so that any changes in their configuration are easily seen. Zinc-deficient SOD1 expression did not affect the structure of the myofibrils. Expression of wild-type human SOD1 (hSOD1) did not affect mitochondrial ultrastructure (Figure 2B, 2E, 2H, 2K). However, we observed a localized rearrangement of the cristae within individual mitochondria of zinc-deficient SOD1 flies (Figure 2C, 2F, 2I, 2L). These

observations were consistent for both 10-day-old and 30-day-old flies, with cristae derangement in zinc-deficient SOD1 flies being more severe on day 30 compared to day 10 (Figure 2).

Expression of zinc-deficient SOD1 in *Drosophila* neurons and glial cells produces locomotor defects

To test the tissue-specific effects of zinc-deficient SOD1 in *Drosophila*, we analyzed locomotor function and lifespan of flies expressing zinc-deficient SOD1 in neurons, glial cells, and muscle cells respectively (Figure 3A-H). As with ubiquitous expression, expression of zinc-deficient SOD1 mediated by the pan-neuronal GAL4 driver *Elav-GAL4* had no major impact on longevity (Figure 3A; Table S1). Using the *D42-GAL4* driver, we observed that expression of wild-type human SOD in motor neurons produces a moderate decrease in lifespan but we observed no further deficit in zinc-deficient SOD1-expressing flies (Figure 3B; Table S1). Expression of zinc-deficient SOD1 mediated by the glial driver *Repo-GAL4* (Figure 3C; Table S1) or the muscle driver *MHC-GAL4* (Figure 3D; Table S1) did not produce major effects on longevity.

Next, using the *Drosophila* Activity Monitor, we examined the tissue-specific requirements for the zinc-deficient SOD1-induced movement defect. We assayed physical activity in 30-day-old flies and observed that expression of zinc-deficient SOD1 in neurons and glial cells, but not muscles, reduces physical activity by about 40% to 50% compared to wild-type hSOD1 (Figure 3E-3H). Expression of zinc-deficient SOD1 mediated by the pan-neuronal *Elav-GAL4* (Figure 3E) and the glial driver *Repo-GAL4* (Figure 3G) produced a motor defect compared to isogenic controls. As expression of zinc-deficient SOD1 mediated by the pan-neuronal *Elav-GAL4* produced the most robust movement defect, we focused on characterizing this model. In comparison to aged (30-day-old) flies where physical activity was reduced by almost 48% (Figures 3E and S1B), pan-neuronal expression of zinc-deficient SOD1 in young (10-day-old) flies reduced physical activity by only 13% (Figure S1A), indicating that neuronal expression of zinc-deficient SOD1 causes an age-related decline in motor activity.

Neuronal expression of zinc-deficient SOD1 reduces ATP levels and sensitizes flies to paraquat and zinc toxicity

To investigate the mechanism by which neuronal zinc-deficient SOD1 expression results in a locomotor defect, we focused on the role of mitochondrial energy metabolism. Firstly, to determine whether expression of zinc-deficient SOD1 affects overall energy production in neuronal tissue, we examined ATP levels in heads of zinc-deficient SOD1-expressing flies and controls. In doing so, we observed a significant decrease in ATP levels in the heads of flies expressing zinc-deficient SOD1 (Figure 4A). Next, we examined whether neuronal expression of zinc-deficient SOD1 affected tolerance to dietary paraquat (PQ), which is widely used as a redox cyler to stimulate superoxide production in organisms, cells, and mitochondria. We observed that neuronal expression of zinc-deficient SOD1 greatly sensitized flies to PQ toxicity (Figure 4B). In contrast, neuronal expression of wild-type hSOD1 increased flies' resistance to PQ (Figure 4C).

Next, we assayed whether zinc-deficient SOD1 affects tolerance to the metals copper, iron and zinc. Neuronal expression of zinc-deficient SOD1 results in decreased tolerance to zinc but not to copper or iron (Figure 4D). In contrast, neuronal expression of wild-type hSOD1 did not alter sensitivity to zinc (Figure 4E).

Neuronal expression of *Ndi1* restores ATP levels and motor function of zinc-deficient SOD1 flies

To further investigate the relationship between mitochondrial dysfunction and the locomotor phenotype associated with zinc-deficient SOD1, we set out to enhance respiratory chain activity in zinc-deficient SOD1 flies. Recently, we reported the generation of a novel transgenic approach to supplement the endogenous respiratory chain machinery of the fly with the alternative single-subunit NADH–ubiquinone oxidoreductase (*Ndi1*) of the baker's yeast *Saccharomyces cerevisiae* (Bahadorani et al., 2010). Expression of *Ndi1* in wild-type flies leads to an increase in NADH–ubiquinone oxidoreductase activity, oxygen consumption and ATP levels (Bahadorani et al., 2010). Here, we have extended this study to zinc-deficient SOD1 flies. To eliminate the possible contribution of induction of endogenous genes at the site of insertion of the *Ndi1* transgene, we tested two independent insertions of the *UAS-Ndi1* transgene. We observed that neuronal expression of *Ndi1* can increase ATP levels in heads of zinc-deficient SOD1 flies (Figure 5B). Next, we set out to determine whether *Ndi1* expression could improve motor function in zinc-deficient SOD1 flies. Using the *Drosophila* Activity Monitor, we assayed physical activity in zinc-deficient SOD1 flies with and without *Ndi1* gene expression. In doing so, we observed that *Ndi1* gene expression in neurons is sufficient to ameliorate the movement disorder associated with zinc-deficient SOD1 expression (Figure 5C), suggesting that the zinc-deficient SOD1 locomotor defects are linked to respiratory chain defects.

Discussion

To better understand the mechanism by which mutations in SOD1 leads to the neurodegenerative manifestations of ALS, we have developed a *Drosophila* model that expresses a zinc-deficient form of human SOD1. To date, most studies have failed to demonstrate any significant phenotype arising from the expression of human mutant ALS-associated human ALS SOD1 in *Drosophila*. Gradual loss of the Zn cofactor has been identified as a critical pathogenic event in the neurodegenerative phenotypes associated with ALS. It has been argued that a possible shortcoming of the *Drosophila* model might be that the short adult lifespan is insufficient for the gradual loss of the Zn cofactor and the ensuing pathology to manifest phenotypically. To address this possibility, we have used a mutant form of human SOD1 (D83S SOD) that is structurally incapable of binding Zn but retains the catalytically-important Cu cofactor along with robust enzymatic activity. Expression of zinc-deficient SOD1 (D84S SOD) in *Drosophila* confers a robust, age-dependent movement disorder reminiscent of ALS in humans. Expression of zinc-deficient SOD1 in neurons or glial cells is sufficient to impair motor function in aging flies. The zinc-deficient SOD1-associated movement defect is accompanied by mitochondrial structural damage, presumably due to zinc-deficient SOD1 penetration into mitochondria owing to the absence of the zinc cofactor (Field et al., 2003). Hypothetically, it could be possible that the

consequent penetration and accumulation of zinc-deficient SOD1 in mitochondria may be the cause of cristae disruption and mitochondrial vacuolization in these flies.

We provide genetic evidence that such respiratory defect underlies the movement disorder, and that expression of the alternative single-subunit yeast NADH-ubiquinone oxidoreductase (*Ndi1*) is sufficient to enhance ATP levels and motor activity of zinc-deficient SOD1 flies. These findings support the link between zinc-deficient SOD1 and mitochondrial respiratory chain dysfunction.

Expression of zinc-deficient SOD1 produces an age-related movement defect without shortening lifespan

We did not detect any major effect on longevity or neuronal integrity in response to zinc-deficient SOD1 expression in *Drosophila*. Similar negative findings have been reported with expression of ALS-associated SOD1 mutations in *Drosophila* (Elia et al., 1999; Mockett et al., 2003; Watson et al., 2008; Parkes et al., 1998). Although the reasons for this are unclear, it is possible that the short adult lifespan of *Drosophila* relative to mammals may preclude the manifestation of all but the most catastrophic neurodegenerative phenotypes.

It is worth noting that expression of wild-type human SOD1 (hSOD1) in neurons and glial cells significantly increases physical activity (Figure 3A and 3C). Similar findings were recently reported and attributed to increased production of hydrogen peroxide and the consequent alterations in *Drosophila* activity (Grover et al., 2009). Therefore, it is particularly striking that zinc-deficient SOD1 expression leads to a *decrease* in physical activity.

Mitochondrial respiratory chain dysfunction underlies the zinc-deficient SOD1-induced movement defect: a novel therapeutic strategy to treat ALS

A number of previous studies have reported that ALS-linked mutant forms of SOD1 can produce mitochondrial respiratory defects (Ferri et al., 2006; Liu et al., 2004; Pasinelli et al., 2004; Vande Velde et al., 2008), presumably through mechanisms involving nitric oxide-mediated toxicity (Cassina et al., 2008; Chen et al., 2010). Nitric oxide is synthesized by the enzyme nitric oxide synthase and associates with mitochondrial dysfunction (Park et al., 2011). Previous studies demonstrated that supplementation with nitric oxide synthase inhibitors prevented neuronal loss and ameliorated phenotypes of G93A-mSOD1 ALS mice (Cassina et al., 2008; Chen et al., 2010). These observations suggest that mitochondrial dysfunction may be a central mechanism of mutant SOD toxicity. Considering that zinc-deficient SOD neurotoxicity is also dependent on endogenous production of nitric oxide (Estévez et al., 1999), we hypothesized that mitochondrial dysfunction may be responsible for locomotor defects in our ALS flies. Our results demonstrate that a mutation affecting the zinc-binding site of SOD1 is sufficient to induce mitochondrial vacuolization and to reduce ATP levels. Furthermore, zinc-deficient SOD1-expressing flies are severely sensitized to mitochondrial toxins paraquat and zinc: paraquat lowers mitochondrial complex I activity while zinc ion interacts with nitric oxide to induce mitochondrial dysfunction and neuronal cell death (Bossy-Wetzel et al., 2004; Tawara et al., 1996).

Similar to our findings, one of the major pathological changes in transgenic mouse models of ALS includes mitochondrial vacuolization resulting from expansion of mitochondrial intermembrane space and extension of the outer membrane (Xu et al., 2004). Our interpretation of these findings is that it is the loss of zinc from the ALS-linked mutant SODs that leads to mitochondrial structural damage. While the etiology of this toxicity is unknown, the fact that loss of zinc cofactor enhances SOD1 penetration into mitochondria proposes a potential mechanism for zinc-deficient SOD1-mediated mitochondrial vacuolization (Field et al., 2003). It is also worth noting that when we assayed ATP levels, zinc-deficient SOD1 flies displayed a defect in heads but not in the rest of the body. Thus, it is possible that zinc-deficient SOD1-induced mitochondrial damage is more pronounced in neuronal tissue.

In rodent models of SOD1-ALS, mitochondrial dysfunction precedes behavioral manifestations of the disorder (Hervias et al., 2006). In a similar fashion, while robust behavioral defects were observed in 30-day-old zinc-deficient SOD1 flies, we detected alterations in mitochondrial structure and function in 10-day old flies. An important advance reported in the present study is the finding that the locomotor defect conferred by zinc-deficient SOD1 can be treated by expression of the *Ndi1* enzyme. In the first place, the observation that genetic enhancement of the respiratory chain can suppress zinc-deficient SOD1 toxicity provides a compelling argument that mitochondrial dysfunction underlies the movement disorder. Secondly, this finding suggests that the *Ndi1* gene may be an effective therapeutic target to counteract ALS symptoms. Previously, it has been shown that *Ndi1* expression, mediated by a recombinant adeno-associated virus, can function in mammalian mitochondria *in vivo* and protect against mitochondrial insults (Barber-Singh et al., 2009; Marella et al., 2008; Marella et al., 2009). Our findings in flies lend further weight to the idea that that *Ndi1* gene therapy as well as other approaches to enhancing mitochondrial ATP output in neurons hold promise for ameliorating the devastating effects of ALS.

Experimental Procedures

Fly stocks and maintenance—To ensure that female egg laying behavior did not interfere with survival and behavioral assays, male flies were used throughout the study. Fly food was prepared using the same protocol to that described previously (Bahadorani et al., 2010): 1 liter of stock medium consisted of 19 g of sucrose, 38 g of dextrose, 10 g of agar, 91 g of cornmeal, 30 g of yeast, 11 ml of propionic acid, and 1.5 g of tegosept. Metal-supplemented media were prepared by addition of iron (III) sulfate hydrate (10 mM), zinc chloride (10 mM) or copper (II) sulfate (1 mM) into normal fly food. The selected concentrations are partially lethal to developing *Drosophila* larvae (Bahadorani and Hilliker, 2009).

The *GAL4* drivers *Elav-GAL4*, *Repo-GAL4*, and *MHC-GAL4* are expressed in *Drosophila* neurons, glial cells, and muscles, respectively; *da-GAL4* is expressed ubiquitously. *UAS-SOD1-IR* (or *UAS-SOD1-RNAi*), *UAS-hSOD1*, and *UAS-Ndi1* transgenic flies were described previously (Bahadorani et al., 2010; Parkes et al., 1998; Wicks et al., 2009). *UAS-GFP* was obtained from the Bloomington Stock Center (stock 1521). *UAS-D83S* transgenic flies, also referred to as CuSOD or zinc-deficient SOD1, were constructed by cloning human

D83S mutant SOD cDNA into the *pUAST* transformation vector, after which the *pUAST* transformation vector and the p(2-3) helper plasmid were co-injected into *w¹¹¹⁸* embryos using standard procedures. To ensure that the genetic background was not a confounding factor in our study, *UAS-hSOD1*, *UAS-D83S*, *UAS-GFP* and *UAS-Ndi1* transgenic flies were backcrossed into the *w¹¹¹⁸* background ten times.

Western blot analysis—Twelve adult flies from each genotype were homogenized in 1.5 ml polypropylene centrifuge tubes containing 100 μ l of ice-cold 50 mM Tris-HCl, 1 mM EDTA, pH 7.2. The homogenate was centrifuged at 12,000 g for 5 min and then the protein concentration was determined using the Bio-Rad Protein Dye Reagent (Bio-Rad Canada, Mississauga, Ont) based on the method of Bradford (Bradford, 1976). The protein concentration was adjusted to 2 μ g/ μ l in all samples, then mixed with an equal volume of SDS-PAGE sample loading buffer and boiled for 5 min. The samples were loaded into a 15% separating gel (10 μ g/lane) and ran at 200 V. Following electrophoresis, the proteins were electroblotted onto a PVDF membrane and subsequently, the membrane was dried for 2 h at room temperature. The membrane was probed with a sheep anti-human SOD1 antibody (Stressgen, Victoria, BC) diluted 1:5000 and detected with a donkey anti-sheep HRP conjugate (Millipore, Mississauga, Ont) diluted 1:75,000 and finally visualized with the ECL Plus Detection Reagents (GE Life Sciences). Image was obtained with a digital camera (Minolta DiMAGE X).

SOD activity assay—SOD activity was measured using the nitroblue tetrazolium (NBT) in-gel assay. In brief, a total of 10 flies were homogenized in 100 μ l of homogenizing buffer (1% Triton X-100), centrifuged at 10 000 g for 5 minutes at 4°C, and the supernatant was mixed at 1:1 ratio with 2 \times loading buffer (0.125 M Tris-HCl pH=6.8, 20% Glycerol, and 0.1% bromophenol blue). Afterward, samples were loaded on 10% acrylamide gels and run for 2 hours at 20 mA at 4°C. Each gel was soaked in 25 ml of NBT solution (50 mg of NBT dissolved in 25 ml of ddH₂O) for few minutes and then transferred into 100 ml of stain solution (0.63 g of K₂HPO₄ pH 7.8, 1.3 mg of riboflavin, and 300 μ l of TEMED dissolved in 100 ml of ddH₂O) for 7 minutes while exposed to intense light. Thereafter, the gels were rinsed and soaked in water and were placed on a light box until the desired contrast was obtained. Image was obtained with a digital camera (Minolta DiMAGE X).

Lifespan assay—Lifespan studies were performed at 25°C on normal food on a 12h:12h light:dark cycle. A total of 150-180 newly-eclosed males (initially 30 per vial) were tested for each genotype, with survivors transferred to fresh food every 3-4 days. For each longevity assay, control and experimental lifespans were determined at the same time.

Physical Activity—Physical activity was measured using the same protocol to that described previously (Bahadorani et al., 2010). For measuring motor activity, 30 adult male flies were placed in a *Drosophila* activity meter (TriKinetics Inc., Massachusetts). Movements were recorded continuously under normal culturing conditions for about 24 hours on a 12:12-hour dark:light cycle. Triplicate or more samples were used for each activity measurement.

Electron and light microscopy—Flies were fixed in 1% glutaraldehyde, 1% paraformaldehyde in 0.1M phosphate buffer, 0.9% sodium chloride (PBS) overnight and washed. After postfixation in 1% OsO₄ in PB for 1.5-2 hours, the tissues were dehydrated in a graded series of ethanol, treated with propylene oxide and embedded in Eponate 12 (Ted Pella). Sections of the dorsal IFM were cut on a Reichert-Jung Ultracut E ultramicrotome at approximately 60-70 nm thick and picked up on formvar coated copper grids. The sections were stained with uranyl acetate and Reynolds lead citrate and examined on a JEOL 100CX electron microscope at 80kV. Horizontal sections (1 μ m thick) of fly heads were mounted on glass slides and stained with 1% toluidine blue in 1% sodium borate, after which samples were examined with light microscope. For light microscopy, sections were 1.25 micrometer thick and were analyzed using the Nikon Microphot-FXA microscope.

ATP assay—Fly heads (10 to 20) were homogenized in 100 μ l of 6 M guanidine-HCl in extraction buffer (100 mM Tris and 4 mM EDTA, pH 7.8) and boiled for 5 minutes. The samples were then centrifuged to collect the supernatant, which was then diluted (1/750) with extraction buffer. The luminescence was measured by a luminometer using autoinjection of the luminescent solution (Invitrogen: “ATP determination kit”), and the results were compared to the standards. The ATP level was then calculated by dividing the luminescence by the total protein concentration, which was determined by the Bradford method.

Stress resistance—Paraquat resistance: male flies were aged for 10 days, starved for 6 h and then transferred to vials containing a 2.4cm glass fiber filter circles (Whatman) wetted with 150 μ l of 30 mM paraquat (Sigma) in 5% sucrose solution. Flies were kept in the dark at all times, except for scoring death and the daily addition of 100 μ l 30mM paraquat solution.

Heavy metal resistance: *w¹¹¹⁸* control or *UAS* transgenic female flies were crossed to *Elav-GAL4* males (*GAL4* located on the X-chromosome), aged on normal food for 3-4 days, and then transferred to metal-supplemented medium for 2-3 days. Afterward, adult flies were removed from vials and eclosion rate was monitored by normalizing female eclosion rate to that of males.

Statistical analysis—Unless specified otherwise, all genotypes were compared to *w¹¹¹⁸* control flies. Bars are expressed as mean \pm standard error of the mean (SEM) of triplicate or more samples. The significance of the difference between means was analyzed using one-way analysis of variance (ANOVA). For survival assays, the significance of the difference between longevity curves was analyzed using the log-rank test.

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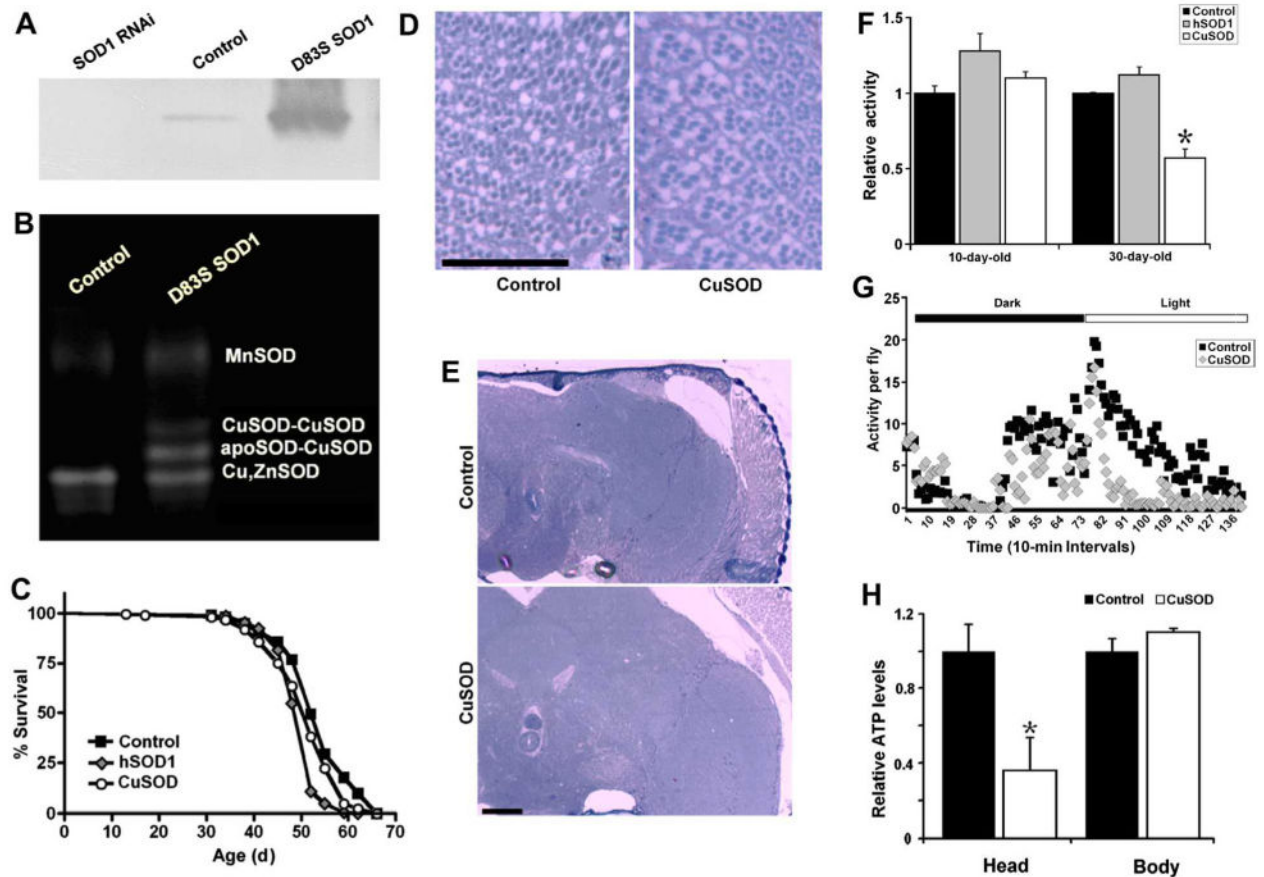


Fig. 1.

Ubiquitous expression of zinc-deficient superoxide dismutase (SOD1, CuSOD, or D83S SOD1) in *Drosophila* produces locomotor defects without neurodegeneration or shortened life span. (A) Western blot analysis of SOD protein level in adult fly extracts confirmed *daGAL4*-mediated expression of the CuSOD transgene (third lane). The first lane indicates the knockdown of SOD protein on *daGAL4*-mediated activation of a *UAS-SOD1-RNAi* transgene. (B) Expression of CuSOD leads to formation of heterodimers between wild type SOD and CuSOD. SOD activity was measured using the in-gel nitroblue tetrazolium assay. (C) Expression of wild type human SOD1 (hSOD1) or CuSOD has a minor negative effect on the adult life span. (D) and (E) Expression of CuSOD does not promote neurodegeneration in *Drosophila* eye or brain (60-day-old flies; scale bar, 50 μ m). (F) Expression of CuSOD significantly compromises locomotor activity ($p=0.002$) in 30-day-old flies. Ubiquitous expression of wild type hSOD1, however, had no adverse effect on physical activity in young ($p=0.067$) or old ($p=0.093$) flies. (G) Representative activity pattern for 30-day-old flies during a 24-hour period. (H) Expression of CuSOD lowers ATP levels in heads ($p=0.029$) but not the rest of the body ($p=0.21$) of 10-day-old flies. Genotypes were as follows: control (*da-Gal4/+*), SOD1 RNAi (*da-Gal4/UAS-SOD1-RNAi*); and D83S SOD1 or CuSOD (*da-Gal4/UAS-D83S*), hSOD1 (*da-Gal4/UAS-hSOD1*). For (F) and (H), the significance of the difference (Note that RNAi is an abbreviation for double stranded interference RNA.) between means was analyzed using 1-way analysis of variance.

Abbreviations: apoSOD, endogenous *Drosophila* SOD1 monomer; MnSOD, manganese superoxide dismutase (SOD2).

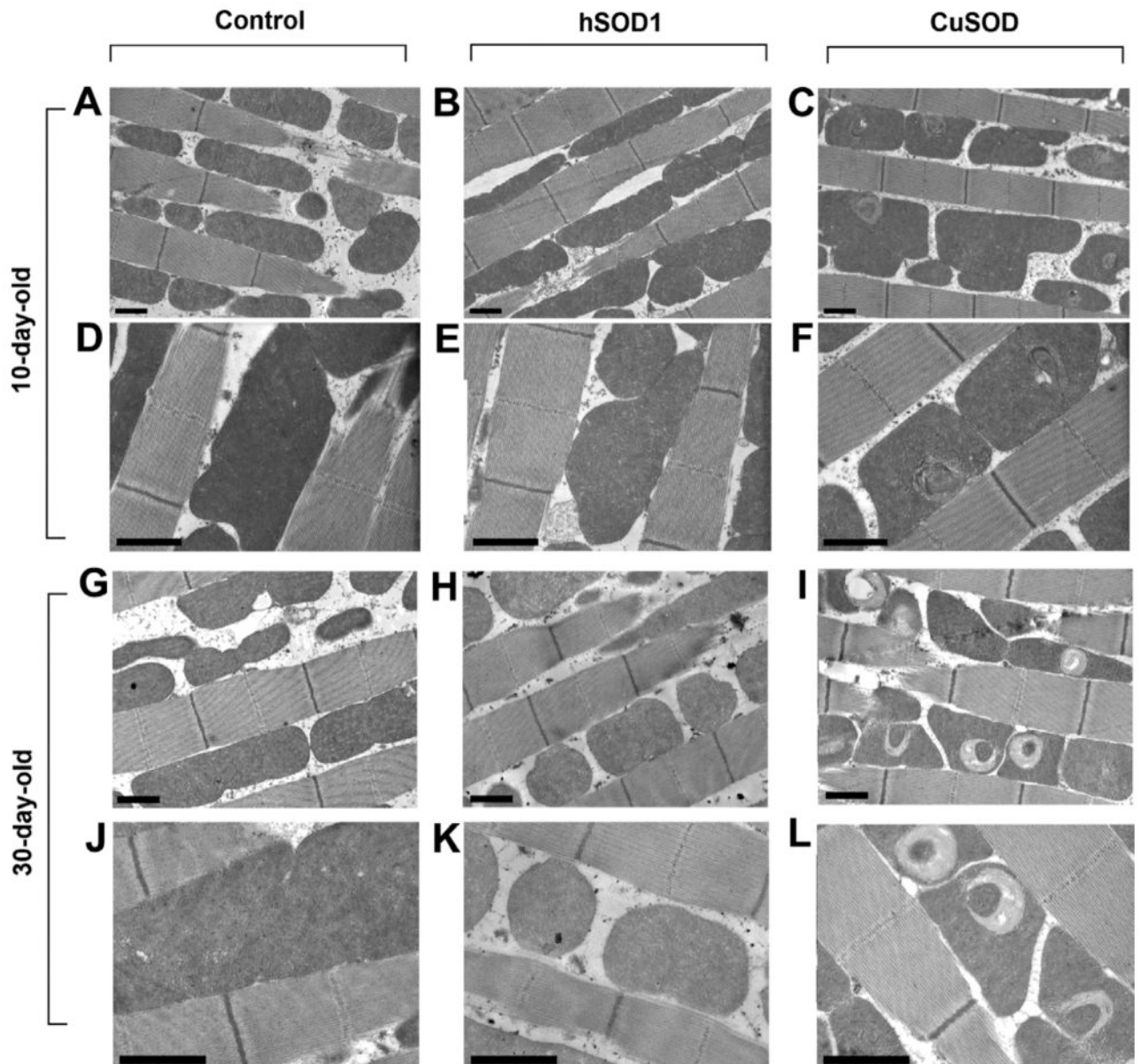


Fig. 2. Ubiquitous expression of zinc-deficient SOD (or CuSOD) in *Drosophila* deteriorates mitochondrial structure. Electron micrographs of flight muscle of 10-day-old flies (A–F) and 30-day-old flies (G–L). (A), (D), (G), and (J) control flies show normal mitochondrial ultrastructure whereas expression of CuSOD (C), (F), (I), (L) but not wild-type human superoxide dismutase (hSOD1) (B), (E), (H), (K) produces a localized rearrangement of the cristae within individual mitochondria (scale bar, 1 μ m). CuSOD phenotypes were more severe on day 30 compared to day 10. Genotypes were as follows: Control (*da-Gal4/+*), hSOD1 (*da-Gal4/UAS-hSOD1*), and CuSOD (*da-Gal4/UAS-D83S*).

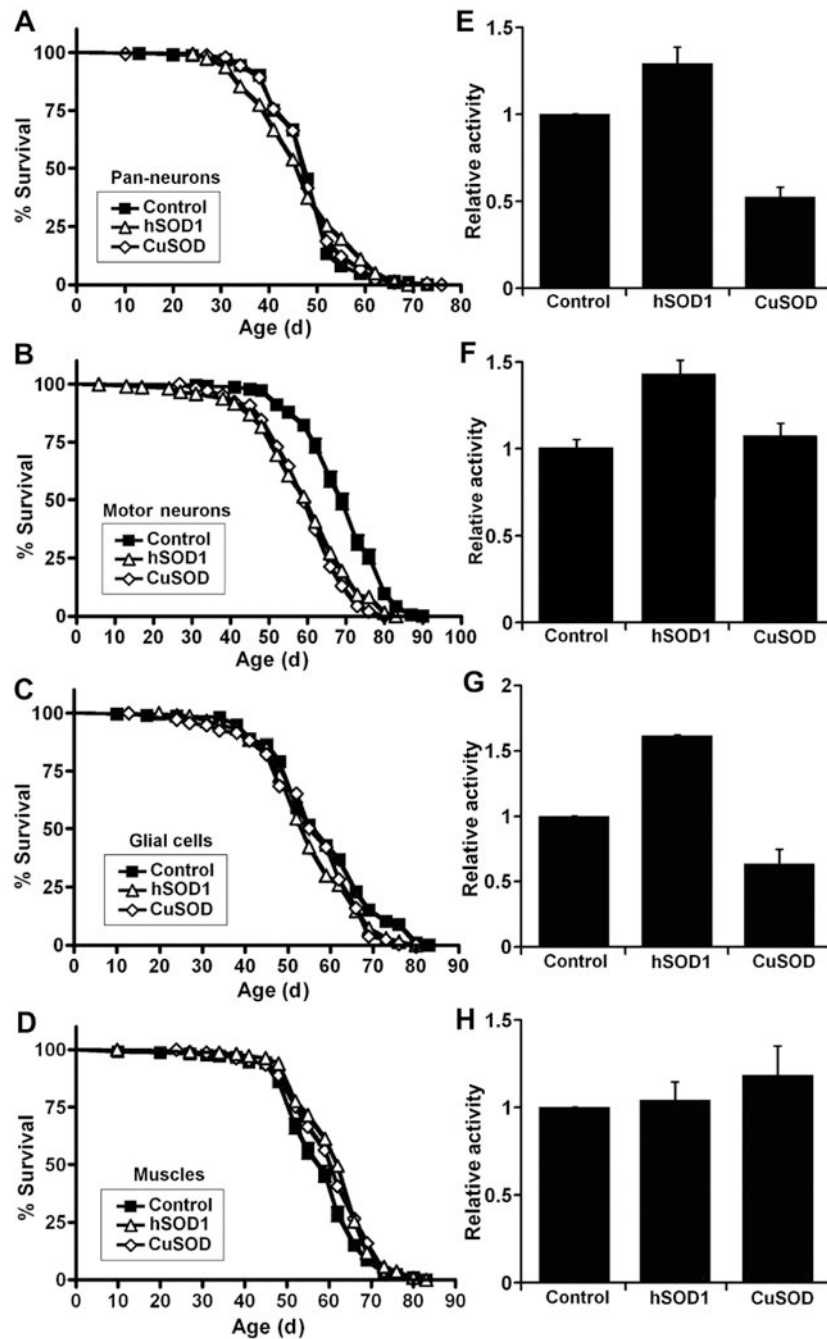


Fig. 3. Expression of zinc-deficient superoxide dismutase (SOD1, CuSOD, or D83S) in *Drosophila* neurons and glial cells produces locomotor defects without major changes in life span. (A–D) Expression of zinc-deficient SOD (CuSOD) or human SOD1 (hSOD1) in *Drosophila* neurons, glial cells, or muscles has a minor effect on adult life span. Expression of CuSOD in (E) neurons ($p=0.0003$), (F) motor neurons ($p=0.014$) and (G) glial cells ($p=0.0001$) but not (H) muscles ($p=0.27$) produces locomotor defects in 30-day-old flies as compared to wild-type hSOD1 flies. Genotypes were as follows: (A) and (E) Control: *Elav-GAL4/+*;

hSOD1: *Elav-GAL4/UAS-hSOD1*; CuSOD: *Elav-GAL4/UAS-D83S*. (B) and (F) Control: *D42-GAL4/+*; hSOD1: *D42-GAL4/UAS-hSOD1*; CuSOD: *D42-GAL4/UAS-D83S*. (C) and (G) Control: *Repo-GAL4/+*; hSOD1: *Repo-GAL4/UAS-hSOD1*; CuSOD: *Repo-GAL4/UAS-D83S*. (D) and (H) Control: *MHC-GAL4/+*; hSOD1: *MHC-GAL4/UAS-hSOD1*; CuSOD: *MHC-GAL4/UAS-D83S*. For (E), (F), (G), and (H), the significance of the difference between means was analyzed using one-way analysis of variance.

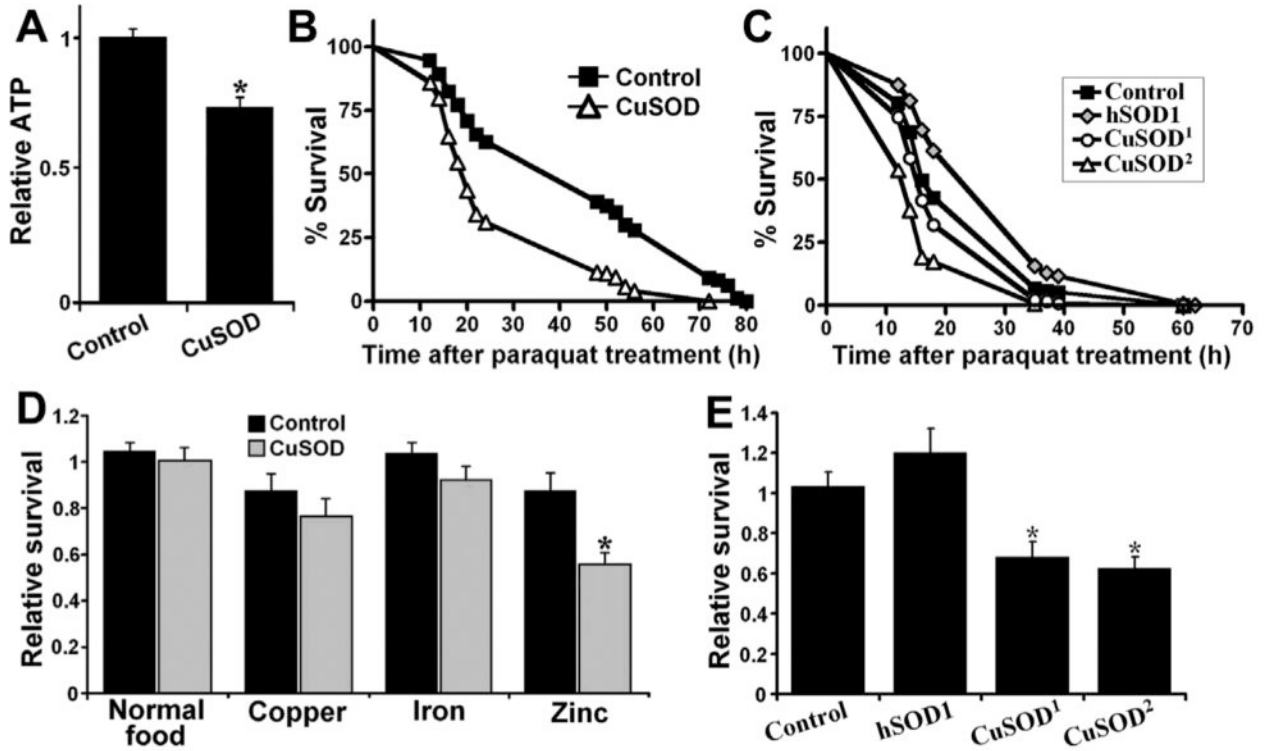


Fig. 4.

Neuronal expression of zinc-deficient SOD (CuSOD) sensitizes flies to zinc and paraquat toxicity. (A) Neuronal expression of zinc-deficient superoxide dismutase (SOD1 or CuSOD) leads to a decrease in ATP levels of fly heads ($p=0.005$). (B) and (C) Neuronal expression of *CuSOD* sensitizes flies to paraquat toxicity, whereas expression of wild-type human SOD1 (hSOD1) enhances resistance to paraquat ($p<0.0001$). (D) Neuronal expression of *CuSOD* sensitizes larvae to zinc ($p=0.003$) but not iron or copper toxicity ($p>0.05$). Relative survival represents eclosion ratio of females (*Elav-GAL4/+* for control, *Elav-GAL4/UAS-D83S* for CuSOD, *Elav-GAL4/UAS-hSOD1* for hSOD1) to male flies (+/+ for control, +/*UAS-D83S* for CuSOD, +/*UAS-hSOD1* for hSOD1) on different media. (E) Expression of hSOD1 does not sensitize flies to zinc toxicity. Genotypes were as follows: control (*Elav-GAL4/+*), hSOD1 (*Elav-GAL4/UAS-hSOD1*), CuSOD¹ (*Elav-GAL4/UAS-D83S₁*), CuSOD or CuSOD² (*Elav-GAL4/UAS-D83S₂*). For panels (A), (D), and (E), the significance of the difference between means was analyzed using 1-way analysis of variance. For (B) and (C), the significance of the difference between survival curves was analyzed using log-rank test.

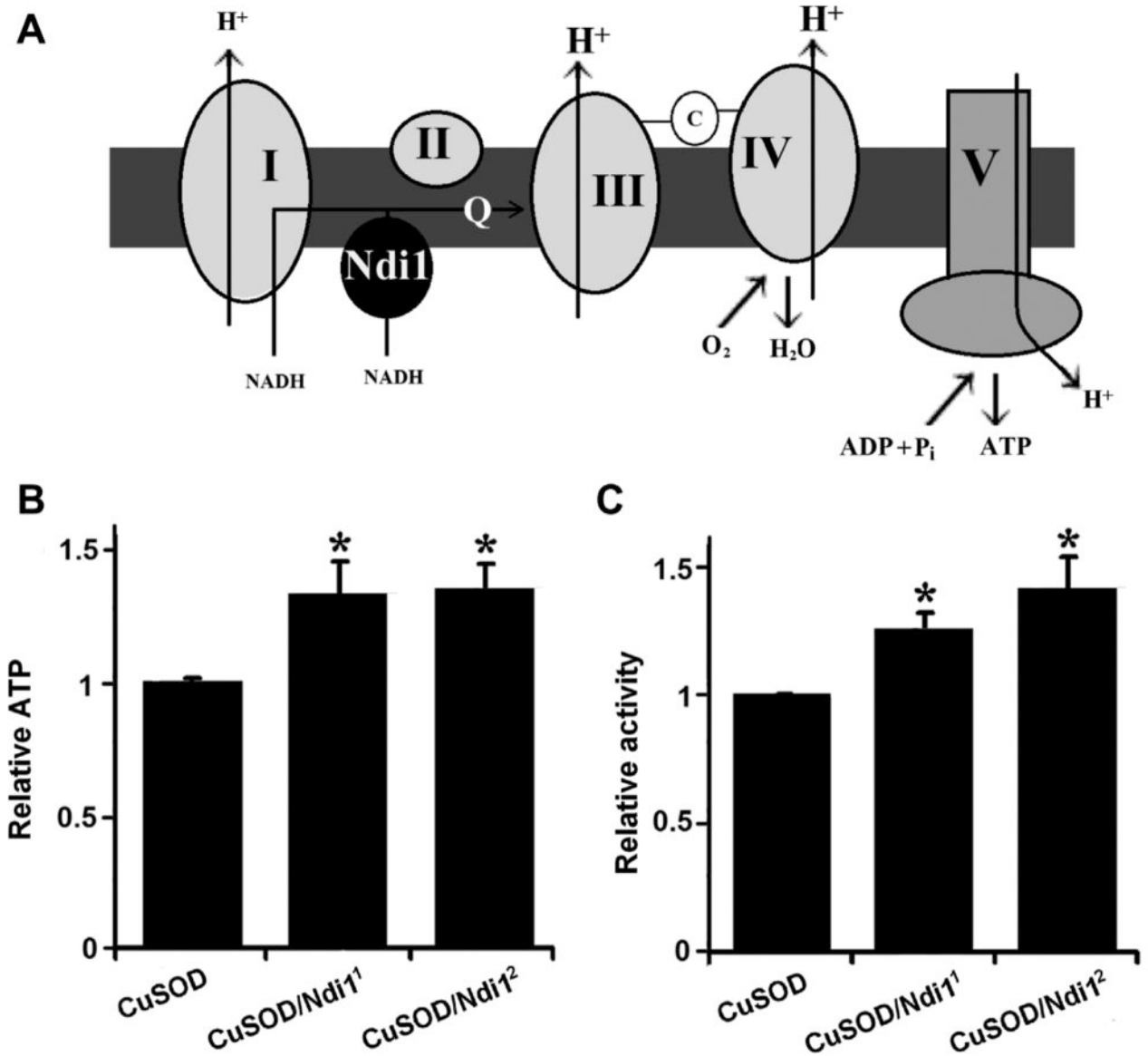


Fig. 5.

Neuronal expression of NADH-ubiquinone oxidoreductase increases ATP production and rescues locomotor defects of zinc-deficient superoxide dismutase (SOD1, CuSOD, or D83S) flies. (A) Schematic drawing of the mechanism of NADH-ubiquinone oxidoreductase (Ndi1) enhanced energy production in the electron transport chain. Coexpression of the single-subunit yeast *Ndi1* with CuSOD (B) increases ATP levels in heads and (C) boosts physical activity in 30-day-old flies. The significance of the difference between means was analyzed using 1-way analysis of variance ($p < 0.05$). Genotypes were as follows: CuSOD (*Elav-GAL4/Y; +/UAS-GFP; +/UAS-D83S*), CuSOD/*Ndi1*¹ (*Elav-GAL4/Y; +/UAS-Ndi1; +/UAS-D83S*), CuSOD/*Ndi1*² (*Elav-GAL4/Y; +; UAS-Ndi1/UAS-D83S*). Abbreviations: ADP, adenosine diphosphate; C, cytochrome c; P_i, inorganic phosphate; Q, coenzyme Q; +, a wild-type chromosome containing no mutations or transgenes.