Post-transcriptional Inhibition of Hsc70-4/HSPA8 Expression Leads to Synaptic Vesicle Cycling Defects in Multiple Models of ALS

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Graphical Abstract

Highlights

- The Hsc70-4/HSPA8 chaperone is post-transcriptionally reduced in ALS models
- ALS-associated mutant TDP-43 inhibits hsc70-4 mRNA translation via sequestration
- TDP-43 impairs the synaptic CSP/Hsc70-4 chaperone complex impacting dynamin

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In Brief

Amyotrophic lateral sclerosis (ALS) is a fatal disease characterized by synaptic failure. Coyne et al. show that in multiple models of ALS, ranging from Drosophila to mice to patient-derived motor neurons, deficits in synaptic vesicle cycling can be explained by dysregulation of the Hsc70-4/HSPA8 chaperone.
Post-transcriptional Inhibition of Hsc70-4/HSPA8 Expression Leads to Synaptic Vesicle Cycling Defects in Multiple Models of ALS

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SUMMARY
Amyotrophic lateral sclerosis (ALS) is a synaptopathy accompanied by the presence of cytoplasmic aggregates containing TDP-43, an RNA-binding protein linked to ~97% of ALS cases. Using a Drosophila model of ALS, we show that TDP-43 overexpression (OE) in motor neurons results in decreased expression of the Hsc70-4 chaperone at the neuromuscular junction (NMJ). Mechanistically, mutant TDP-43 sequesters hsc70-4 mRNA and impairs its translation. Expression of the Hsc70-4 ortholog, HSPA8, is also reduced in primary motor neurons and NMJs of mice expressing mutant TDP-43. Electrophysiology, imaging, and genetic interaction experiments reveal TDP-43-dependent defects in synaptic vesicle endocytosis. These deficits can be partially restored by OE of Hsc70-4, cysteine-string protein (Csp), or dynamin. This suggests that TDP-43 toxicity results in part from impaired activity of the synaptic CSP/Hsc70 chaperone complex impacting dynamin function. Finally, Hsc70-4/HSPA8 expression is also post-transcriptionally reduced in fly and human induced pluripotent stem cell (iPSC) C9orf72 models, suggesting a common disease pathomechanism.

INTRODUCTION
Amyotrophic lateral sclerosis (ALS) is a progressive neurodegenerative disease linked to several genes, including SOD1 (Rosen et al., 1993), TARDBP (TDP-43) (Kabashi et al., 2008; Sreedharan et al., 2008), FUS (Kwiatkowski et al., 2009; Vance et al., 2009), and C9orf72 (DeJesus-Hernandez et al., 2011; Renton et al., 2011), suggesting the presence of multiple disease mechanisms (Robberecht and Philips, 2013). Synaptic degeneration is a culminating point in ALS, however, the molecular mechanisms that trigger this remain poorly understood (Gillingwater and Wishart, 2013).

With 2%–3% of ALS patients harboring mutations in TDP-43 (Kabashi et al., 2008; Neumann, 2009; Sreedharan et al., 2008; Van Deerlin et al., 2008) and ~97% of all ALS cases exhibiting TDP-43 pathology (Ling et al., 2013), TDP-43 is a common denominator in ALS. Thus, elucidating the mechanisms by which both wild-type and mutant TDP-43 cause neurodegeneration is essential to our understanding of ALS pathogenesis.

TDP-43 is a nucleo-cytoplasmic shuttling, DNA/RNA-binding protein containing a prion-like C-terminal domain where the majority of ALS-causing mutations cluster, causing an increase in TDP-43's intrinsic propensity for aggregation (Johnson et al., 2009). TDP-43 was shown to bind UG-rich sequences and regulate the splicing of several transcripts encoding synaptic proteins (Polymenidou et al., 2011; Sephton et al., 2011; Tollervey et al., 2011). In addition, TDP-43 regulates the localization and translation of specific mRNAs (Alami et al., 2014; Coyne et al., 2014; Falolini et al., 2012). TDP-43 also associates with RNA stress granules (SGs), which sequester specific mRNAs, resulting in translation inhibition (Dewey et al., 2011; Kim et al., 2014; Liu-Yesucevitz et al., 2010; McDonald et al., 2011).

TDP-43 has previously been shown to associate with human Hsc70 (HSPA8) in mammalian cells (Freibaum et al., 2010). Hsc70 proteins comprise a family of constitutive, ubiquitous molecular chaperones with roles in protein folding and degradation, stress response, endosomal microautophagy, and chaperone-mediated autophagy (Liu et al., 2012). Consistent with its role in proteostasis, HSPA8 has been implicated in neurodegeneration as it colocalizes with ubiquitin-positive inclusions in sporadic ALS (Watanabe et al., 2001). Among the multiple Hsc70 proteins in Drosophila (Palter et al., 1986; Perkins et al., 1990), the clathrin-uncoating ATPase Hsc70-4 has roles in synaptic vesicle
(SV) exo- and endocytosis, and it is highly expressed in neurons and at the larval neuromuscular junction (NMJ) (Bronk et al., 2001; Chang et al., 2002; Zinsmaier and Bronk, 2001).

The critical role of Hsc70-4 in SV cycling (SVC) led us to hypothesize that abnormal interactions between TDP-43 and Hsc70-4 may lead to synaptic impairments prior to degeneration. Here we show that motor neuron overexpression (OE) of TDP-43 causes a reduction in synaptic Hsc70-4 levels and impairments in SVC at the Drosophila NMJ. Using electrophysiology and FM1-43 dye uptake experiments, we show that defects in SVC are due to deficits in SV endocytosis. Similar to our findings in Drosophila, we show that HSPA8 protein is reduced post-transcriptionally at the NMJ in TDP-43^A315T mice and in TDP-43^G298S, patient-derived, induced pluripotent stem cell (iPSC) motor neurons. Furthermore, flies harboring 36 G^C2 repeats (Mizielska et al., 2014) and C9orf72 patient-derived iPSC motor neurons (Zhang et al., 2015) exhibit a post-transcriptional reduction in Hsc70-4/HSPA8 protein. Together, our findings uncover defects in SV endocytosis as a source of synaptic dysfunction in TDP-43- and C9orf72-mediated ALS.

RESULTS

Hsc70-4 Protein and mRNA Form a Complex with TDP-43 in Motor Neurons In Vivo

To test whether TDP-43 associates with Hsc70-4, we performed immunoprecipitations (IPs) from Drosophila adults expressing TDP-43 in motor neurons (Figure 1A) followed by mass spectrometry (data not shown). We found that while Hsc70-4 associates with both wild-type TDP-43 (TDP-43^WT) and ALS-associated mutant TDP-43 (TDP-43^G298S; 22.9% and 30.1% sequence coverage, respectively), its interaction with TDP-43^G298S is significantly stronger (Figure 1A). These results confirm that endogenous Hsc70-4 associates with human wild-type and mutant TDP-43 in Drosophila motor neurons.

To determine whether hsc70-4 mRNA is present in TDP-43 complexes, we performed RNA IPs (RIP) from Drosophila adults expressing TDP-43^WT or TDP-43^G298S in motor neurons followed by real-time qPCR. hsc70-4 mRNA was strongly enriched in TDP-43^G298S while less was present in TDP-43^WT complexes (Figure 1B).

It has been proposed that, under prolonged stress, TDP-43 may sequester its mRNA targets into insoluble complexes, thereby preventing their translation and normal function (Ramaswami et al., 2013). To test whether TDP-43 sequesters hsc70-4 mRNA, we performed subcellular fractionations from Drosophila larvae expressing TDP-43 in motor neurons. Using real-time qPCR, we found an enrichment of hsc70-4 mRNA in the insoluble fraction of animals expressing TDP-43^G298S, but not TDP-43^WT (Figure 1C), consistent with hsc70-4 mRNA sequestration specifically by TDP-43^G298S. These data show that both Hsc70-4 protein and its cognate mRNA associate with TDP-43, albeit preferentially with TDP-43^G298S.

OE of TDP-43^G298S Inhibits the Translation of hsc70-4 mRNA

The sequestration of hsc70-4 mRNA by TDP-43^G298S may impair its translation. To test this possibility, we performed polysome fractionations from adult Drosophila overexpressing TDP-43^WT or TDP-43^G298S in motor neurons. Using real-time qPCR, we found that TDP-43^G298S OE resulted in a shift for hsc70-4 mRNA from actively translating polysomes to non-translated (ribonucleoprotein particle [RNP]) fractions, while TDP-43^WT had no effect (Figures 1E and 1F). Given that no changes in overall hsc70-4 mRNA levels were detected in whole animals (Figure 1D), these data suggest that TDP-43^G298S impairs the translation of hsc70-4 mRNA by sequestering it into insoluble RNPs. In contrast, TDP-43^WT did not affect hsc70-4 mRNA translation, at least within the limits of sensitivity provided by this assay.

TDP-43 Expression Reduces Hsc70-4 Levels at the Larval NMJ in Drosophila

The translation inhibition of hsc70-4 mRNA by TDP-43^G298S predicts that Hsc70-4 levels are reduced, which in turn may impair Hsc70-4’s function (Stricher et al., 2013; Uytterhoeven et al., 2015; Zinsmaier and Bronk, 2001). No changes in protein expression were found in extracts of whole larvae or dissected ventral nerve cords (VNCs) from animals overexpressing TDP-43^WT or TDP-43^G298S in motor neurons (Figures 1G, 1H, 1J,

Figure 1. hsc70-4 mRNA Is a Translation Target of TDP-43^G298S, and Hsc70-4 Protein Expression Is Reduced at Synaptic Terminals of TDP-43-Expressing Animals

(A) Immunoprecipitation (IP) of TDP-43 expressed in motor neurons. Genotypes and antibodies used for IP are indicated on the top. Antibodies used for western blot (WB) are indicated on the right.

(B) qPCR for hsc70-4 mRNA in TDP-43 complexes.

(C) qPCR for hsc70-4 mRNA in soluble and insoluble TDP-43 complexes.

(D–F) qPCR for hsc70-4 mRNA in input (D), RNP (E), and polysomes (F) from flies expressing TDP-43^WT and TDP-43^G298S in motor neurons compared to controls.

(G) WB for Hsc70-4 levels in whole larva expressing TDP-43 in motor neurons. Genotypes are indicated on the bottom. Actin was used as a loading control.

(H) Quantification of Hsc70-4 protein levels from WBs represented as a ratio to w^1118 controls.

(I) qPCR for hsc70-4 mRNA in whole larva of animals expressing TDP-43^WT or TDP-43^G298S in motor neurons versus controls.

(J) WB for Hsc70-4 levels in VNCs of TDP-43-expressing larvae. Genotypes are indicated on the bottom. Actin was used as a loading control.

(K) Quantification of Hsc70-4 protein levels from WBs represented as a ratio to w^1118 controls.

(L) qPCR for hsc70-4 mRNA in VNCs of animals expressing TDP-43^WT or TDP-43^G298S in motor neurons versus controls.

(M–O) Single confocal sections of synaptic boutons in NMJ preparations immunostained for Hsc70-4 and the neuronal membrane marker Hp from larvae expressing TDP-43^WT (N) and TDP-43^G298S (O) compared to w^1118 controls (M). Antibodies are indicated on the left.

(P) Quantification of Hsc70-4 intensity in synaptic boutons normalized to bouton area. Note p < 0.001 for TDP-43^WT versus TDP-43^G298S.

(Q) qPCR for hsc70-4 mRNA in NMJ preparations of animals expressing TDP-43^WT or TDP-43^G298S in motor neurons versus controls.

(R) Quantification of Hsc70-4 intensity in muscle normalized to muscle area.

All data shown are mean ± SEM. Scale bars, 5 and 1 μm (M–O).
Figure 2. Hsc70/HSPA8 Expression Is Reduced in Mutant TDP-43-Expressing Mouse Primary Motor Neurons, at Synaptic Terminals of Mouse NMJs, and in TDP-43G298S Human iPSC Neurons

(A–D) Representative fluorescence images of cell bodies from primary motor neurons transfected with expression constructs for GFP (A–A’’) or GFP-tagged TDP-43WT (B–B’’), TDP-43G331K (C–C’’), or TDP-43M337V (D–D’’), as indicated on the left. Antibodies and stains are indicated on the top.

(E–H) Representative fluorescence images of growth cones from primary motor neurons expressing GFP (E and E’) or GFP-tagged TDP-43WT (F and F’), TDP-43G331K (G and G’), or TDP-43M337V (H and H’), as indicated on the left. Antibodies are indicated on the top.

(I and J) Quantification of fluorescent intensity (a.u.) of Hsc70 in the cell body (I) and growth cones (J). Genotypes are indicated on the left and antibodies are indicated on the top.

(K and L) Epifluorescent images of mouse NMJs immunostained for Hsc70/HSPA8 (K’ and L’) and AChR (K’’ and L’’). Genotypes are indicated on the left and antibodies are indicated on the top.

(M) Quantification of Hsc70/HSPA8 intensity from NMJs.

(N) Confocal images of control and TDP-43G298S human iPSC motor neurons labeled with DAPI, the dendritic marker Map2, and Hsc70/HSPA8. Genotypes are indicated on the left and antibodies are indicated on the top.

(legend continued on next page)
OE of Disease-Associated Mutant TDP-43 Reduces HSPA8 Levels in Mouse Primary Motor Neurons

Having identified reduced synaptic expression of Hsc70-4 in the fly, we next asked if HSPA8 levels were also altered in a mammalian model of ALS. To evaluate HSPA8 levels, we transfected mouse primary motor neurons with GFP-tagged TDP-43WT, TDP-43G298S, or GFP alone, and we performed immunostainings for HSPA8. OE of TDP-43 WT had no effect on HSPA8 levels in growth cones or cell bodies (p > 0.05; Figures 2B, 2F, 2I, and 2J). In contrast, TDP-43G298S OE significantly reduced HSPA8 levels in growth cones (by 39% and 41%, respectively, p < 0.01; Figures 2G, 2H, and 2J) and cell bodies (12% decrease for TDP-43G298S, p < 0.05; Figures 2C and 2I; 33% decrease for TDP-43M337V, p < 0.001; Figures 2D and 2I). These data indicate that, similar to what was observed in the fly model, HSPA8 levels are reduced in mammalian primary motor neurons overexpressing disease-associated mutant TDP-43.

HSPA8 Expression Is Reduced at the NMJ in TDP-43A315T Mutant Mice

We next asked whether levels of HSPA8 are also reduced at NMJs of TDP-43 mutant mice. To address this, dissected mouse NMJs were immunostained for HSPA8 and postsynaptic acetylcholine receptors (ACHRs). In comparison to controls, synaptic HSPA8 immunoreactivity at NMJs expressing TDP-43A315T (Wegorzewska et al., 2009) was decreased by 21% (p < 0.05; Figures 2K–2M). Taken together, these data indicate that disease-associated mutant TDP-43 reduces levels of HSPA8 at NMJs of both flies and mice.

HSPA8 Expression Is Reduced in TDP-43G298S Human iPSC Motor Neurons

Next, we sought to determine if expression of HSPA8 was also altered in human iPSC-derived motor neurons harboring the TDP-43G298S mutation. Immunofluorescence experiments showed a significant decrease in HSPA8 expression in the soma and dendrites of TDP-43G298S iPSC motor neurons compared to controls (Figures 2N and 2O). Notably, hspa8 mRNA expression was unchanged (Figure 2P), consistent with post-transcriptional regulation of HSPA8 expression, as in the fly.

| Table 1 |

(O) Quantification of Hsc70/HSPA8 intensity in the soma and dendrites of control and TDP-43G298S human iPSC motor neurons normalized to control. Symbols represent mean of 25–30 neurons from each line (control) or differentiation (TDP-43) and indicate differentiation pairs. (P) qPCR for Hsc70/HSPA8 mRNA in control and TDP-43G298S human iPSC motor neurons.

All data shown are mean ± SEM. Scale bars, 10 μm (A and N) and 20 μm (K).
we hypothesized that the reduced synaptic levels of Hsc70-4 induced by wild-type and mutant TDP-43 may impair synaptic function.

We have previously shown that OE of wild-type or mutant TDP-43 decreases the number of synaptic boutons at larval NMJs (Coyne et al., 2014; Estes et al., 2013) and increases bouton size (Figures S4Q and S4R), similar to mutations that affect SV endocytosis (Verstreken et al., 2002). We did not observe a reduction in the number of active zones for neurotransmitter release per bouton area when TDP-43 was expressed in motor neurons (Figures S4A–S4H). Furthermore, TDP-43 OE in motor neurons did not alter the intensity or area occupied by Csp within synaptic boutons (Figures S4I–S4P), suggesting that the total number of SVs was within normal range.

To test whether wild-type or mutant TDP-43 OE in motor neurons affects synaptic transmission, we recorded spontaneous

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Figure 3. TDP-43 Expression Results in Defects in SV Endocytosis that Are Suppressed by Hsc70-4 in a Variant-Dependent Manner

(A) Motor neuron expression of TDP-43WT or TDP-43G298S results in increased larval turning time, which is mitigated by OE of Hsc70-4.

(B and C) Motor neuron expression of TDP-43 variants WT (B) or G298S (C) leads to reduced lifespan. OE of Hsc70-4 increases lifespan for both TDP-43WT (B) and TDP-43G298S (C).

(D) Representative electrophysiology traces of EJPs. Genotypes are indicated on the top.

(E–G) EJP amplitude (E), miniature EJP (mEJP) amplitude (F), and quantal content (G) measurements from electrophysiology recordings. Genotypes are indicated on the bottom.

(H–P) Confocal images of FM1-43 dye uptake in synaptic boutons of Drosophila larvae after 5 min of stimulation in HL-3 saline containing 90 mM KCl and 2 mM Ca2+. Genotypes are indicated on the top and left.

(Q) Quantification of FM1-43 dye uptake normalized to total FM1-43 uptake area. Note an 18% ± 1.8% reduction in FM1-43 dye uptake for TDP-43G298S, Hsc70-4 OE compared to w1118 controls (p < 0.001).

All data shown are mean ± SEM. Scale bar, 10 μm (D).
and evoked excitatory postsynaptic potentials (EPSPs) from larval NMJs of muscle 6. In comparison to controls, neither TDP-43WT nor TDP-43G298S OE had an effect on spontaneous or stimulus-evoked EPSP amplitudes (Figures 3D–3F). Accordingly, quantal content of evoked neurotransmitter release was normal (Figure 3G).

Next, we examined Svc by performing FM1-43 dye uptake assays at the larval NMJ (see Experimental Procedures; Kuromi and Kidokoro, 2005). In comparison to controls, TDP-43WT and TDP-43G298S OE significantly reduced K+-induced FM1-43 dye uptake by 33% ± 1.8% and 39% ± 2.1%, respectively (p < 0.001; Figures 3H–3J and 3Q). Subsequent K+ stimulation to unload FM1-43 dye from boutons showed no difference between TDP-43 mutant NMJs and controls (Figures S5A–S5C). Taken together, these results indicate that TDP-43 OE does not affect exocytosis but causes defects in SV endocytosis.

The Chaperone Activity of Hsc70-4 Is Required to Mitigate SVC Defects Caused by TDP-43G298S

We next asked whether Hsc70-4 OE can mitigate TDP-43-induced defects in FM1-43 dye uptake. When OE was in a wild-type background or together with TDP-43WT, Hsc70-4 had no effect on FM1-43 dye uptake at the NMJ (p > 0.05; Figures 3K, 3L, and 3Q). In contrast, co-OE of Hsc70-4 with TDP-43G298S strongly suppressed the defects in FM1-43 dye uptake induced by TDP-43G298S OE (p < 0.001; Figures 3M and 3Q).

To determine whether the chaperone activity of Hsc70-4 is required for its protective effect, we co-overexpressed chaperone-dead Hsc70-4 (Hsc70-4D10N; Uytterhoeven et al., 2015), and we found that it does not alter FM1-43 dye uptake deficits induced by TDP-43 (p > 0.05; Figures 3O–3Q). It also had no effect on its own when OE was in a wild-type background (p > 0.05; Figures 3N and 3Q). These data suggest that TDP-43WT OE induces SV endocytosis defects that are, at least in part, independent of Hsc70-4. In contrast, mutant TDP-43G298S induces defects in SV endocytosis that are mechanistically linked to the chaperone activity of Hsc70-4.

Auxilin OE or RNAi-Mediated Knockdown Does Not Alter TDP-43 Locomotor Phenotypes

Hsc70’s molecular chaperone activity critically requires DNAJ proteins, which transition Hsc70-ATP to the ADP-bound state to stabilize client interactions (Jiang et al., 2007; Kampina and Craig, 2010). At synaptic terminals, Hsc70’s chaperone function in SV exo- and endocytosis is facilitated by the DNAJ proteins auxilin and Csp on SVs (Kampina and Craig, 2010; Liu et al., 2012; Zinmaier and Bronk, 2001). To determine the mechanisms underlying Hsc70-4-dependent phenotypes of TDP-43 OE, we used larval turning assays to examine the effects of mutations affecting critical steps of SVC (Figure 4A).

We first tested whether TDP-43 OE impairs auxilin/Hsc70-mediated clathrin uncoating of SV. Auxilin OE or RNAi-mediated knockdown (55% decrease in expression by qPCR, data not shown) had no effect on larval turning on its own or in combination with wild-type and mutant TDP-43 (Figures 4B and 4C). This suggests that the synaptic defects induced by wild-type or mutant TDP-43 OE are unlikely due to a compromised auxilin/Hsc70 clathrin-uncoating activity.

Csp Levels and Chaperone Activity Modulate TDP-43-Mediated Locomotor Dysfunction

Next, we tested for genetic interactions with Csp, which recruits Hsc70-4 to a chaperone complex on SVs, ensuring proper assembly of SNARE and dynamin complexes during SV exo- and endocytosis, respectively (Chandra et al., 2005; Tobaben et al., 2001; Zhang et al., 2012; Zinmaier, 2001). OE of Csp in a wild-type background from a genomic transgene (increasing Csp gene dosage by one copy) (Zinmaier et al., 1994) had no effect on larval turning time (Figure 4D), while co-OE with TDP-43WT or TDP-43G298S restored larval turning time to control levels (Figure 4D). OE of Csp with a UAS transgene had a significant effect on larval turning time on its own (data not shown), likely due to high levels of Csp causing neurodegeneration (Nie et al., 1999); therefore, we did not pursue this line for further genetic interactions with TDP-43.

Csp’s J domain promotes the ATPase activity of both Hsc70 and Hsp70, which is abolished by mutations in the HPD motif and J domain (Braun et al., 1996; Bronk et al., 2005; Chamberlain and Burgoyne, 1997; Zhang et al., 1999). OE of J domain mutant CspH45Galone (Bronk et al., 2005) had no effect on larval turning (Figure 4E); however, co-OE with TDP-43WT or TDP-43G298S strongly enhanced locomotor dysfunction (Figure 4E). Taken together, these results suggest that OE of wild-type or mutant TDP-43 impairs the activity of the Csp/Hsc70-4 chaperone complex.

OE of Csp/Hsc70-4’s Client Dynamin Mitigates TDP-43-Induced Locomotor Dysfunction

Csp/Hsc70-4 chaperone activity maintains SV endocytosis by ensuring proper oligomerization of the GTPase dynamin (Rozas et al., 2012; Zhang et al., 2012). Once a critical mass is reached, oligomerization activates dynamin’s GTPase activity, which in turn pinches or pops off newly formed SVs from the plasma membrane (Haucke et al., 2011; Jahn and Faushauer, 2012; Ramachandran, 2011). To determine whether TDP-43 OE affects Csp/Hsc70-4’s chaperone activity on dynamin, we tested whether OE of Drosophila dynamin (shibire) may restore TDP-43-induced locomotor defects, and we found that, while OE of shibire alone had no effect on larval turning (Figure 4F), co-OE with TDP-43WT or TDP-43G298S in motor neurons restored locomotor function (Figure 4F). Notably, none of the mRNAs encoding the examined SVC proteins was enriched in wild-type or mutant TDP-43 complexes or were translationally dysregulated, as suggested by polysome fractionations (Figure S6).

OE of Hsc70-4’s Co-chaperone Csp or Its Client Dynamin Suppresses SVC Defects Induced by TDP-43 in Motor Neurons

To determine whether OE of Hsc70-4’s co-chaperone Csp or their client dynamin can mitigate TDP-43-dependent defects in SVC, we tested if the genomic Csp construct or dynamin OE can restore FM1-43 dye uptake defects induced by TDP-43. We found that OE of either Csp from a genomic transgene or dynamin alone had no effect on K+-induced FM1-43 dye uptake in comparison to control (p > 0.05; Figures 4J, 4M, and 4P). Co-OE of Csp fully restored the defect in FM1-43 dye uptake induced by TDP-43WT or TDP-43G298S OE (Figures 4G–4I) to control levels (Figures 4K, 4L, and 4P). Similarly, co-OE of dynamin also fully suppressed...
the defect in FM1-43 dye uptake to control levels (Figures 4N–4P).

Taken together, these data suggest that OE of wild-type or mutant TDP-43 impairs the activity of the Csp/Hsc70 complex in maintaining normal dynamin function for SV endocytosis.

OE of Hsc70-4 Attenuates the Aggregation of TDP-43WT but Not TDP-43G298S

Given the general role of Hsc70-4 in clearance of misfolded or aggregated proteins (Liu et al., 2012; Stricher et al., 2013; Zetterström et al., 2011), we hypothesized that Hsc70-4 OE may influence TDP-43 aggregation. To examine this, we performed subcellular fractionations from Drosophila larvae co-overexpressing wild-type or mutant TDP-43 with Hsc70-4 in motor neurons, and we quantified the amount of TDP-43 in the soluble (low-salt [LS]), sarkosyl (Sark), and urea (Urea) fractions. Interestingly, Hsc70-4 co-OE significantly reduced the amount of TDP-43WT in the Urea fraction and increased the amount of TDP-43WT in the Sark fraction ($p < 0.01$ and $p < 0.05$ respectively; Figure 4. Factors Involved in Hsc70-4-Independent and -Dependent Steps of SVC Modulate TDP-43 Toxicity

(A) Diagram of the SV cycle.

(B–F) Motor neuron expression of TDP-43 WT or G298S results in increased larval turning time, which is enhanced upon OE of the Csp J domain mutant CspH45Q (E) and suppressed upon genomic expression of Csp (D) or OE of dynamin (shibire, F). OE (B) or reduction in auxilin (Auxilin RNAi, C) does not alter TDP-43-mediated locomotor defects.

(G–O) Confocal images of FM1-43 dye uptake after 5 min of stimulation in HL-3 saline containing 90 mM KCl and 2 mM Ca2+. Genotypes are indicated on the top and left.

(P) Quantification of FM1-43 dye uptake normalized to total FM1-43 uptake area. All data shown are mean ± SEM. Scale bar, 10 μm (G).
Figures 5 A and 5E). Notably, Hsc70-4 OE had no significant effect on the solubility of ALS-associated mutant TDP-43 G298S (Figures 5B and 5E), despite its neuroprotective effects on TDP-43G298S-induced functional phenotypes (Figures 3 A–3C and 3H–3M).

Conversely, TDP-43 WT or TDP-43 G298S OE in motor neurons did not affect the solubility of endogenous Hsc70-4 at the larval stage (Figures 5C and 5F). However, adult flies (7 days) exhibited a significant increase in insoluble Hsc70-4 protein for both wild-type and mutant TDP-43 (p < 0.001; Figures 5D and 5G). Adults expressing TDP-43WT also showed a corresponding decrease of Hsc70-4 protein in the Sark fraction in comparison to controls (p < 0.001; Figures 5D and 5G). These data suggest an age-dependent sequestration of Hsc70-4 by insoluble cytoplasmic RNA/protein complexes, as previously observed in sporadic ALS patient samples (Watanabe et al., 2001).

Co-OE of Hsc70-4 with TDP-43 WT significantly decreased the amount of insoluble Hsc70-4 in the Sark fraction compared to TDP-43 WT expression alone.
G4C2 repeats back to control levels (Figures 6L and 6M). These
findings indicate that C9 repeat expansions may also disrupt SVC.

To test this possibility, we performed FM1-43 dye uptake experiments
at the fly NMJ. OE of 36 G4C2 repeats in motor neurons signifi-
cantly decreased FM1-43 dye uptake compared to controls (21%, p < 0.001; Figures 6E-6G). There was no change in hsc70-4 mRNA transcript levels in VNCs or NMJ prepa-
ration (Figures 6C and 6D), and Hsc70-4 protein levels in muscles
were normal (Figure 6H). Together, these data indicate that OE of G4C2 repeats impairs Hsc70-4 protein expression post-
transcriptionally and cell autonomously in motor neurons.

To determine if HSPA8 expression was altered in a human C9
model of ALS, we assayed iPSC-derived motor neurons. In compar-
ison to controls, HSPA8 levels were reduced in the soma and
dendrites of C9 iPSC motor neurons by 57% and 32%, respec-
tively (p < 0.001; Figures 6I and 6J). Using a fly model of C9
ALS (Mzielinska et al., 2014) containing 36 G4C2 repeats, we de-
ected no alterations in Hsc70-4 expression in dissected VNCs (Figures 6A and 6B). However, at synaptic boutons of larval
NMJs, Hsc70-4 levels were significantly decreased compared
to controls (21%, p < 0.001; Figures 6E-6G). There was no change
in hsc70-4 mRNA transcript levels in VNCs or NMJ prepa-
ration (Figures 6C and 6D), and Hsc70-4 protein levels in muscles
were normal (Figure 6H). Together, these data indicate that OE of G4C2 repeats impairs Hsc70-4 protein expression post-
transcriptionally and cell autonomously in motor neurons.

Since the reduction in Hsc70-4 levels in C9 models of ALS is
reminiscent of that seen in the TDP-43 fly model, we hypotho-
ized that C9 repeat expansions may also disrupt SVC. To test
this possibility, we performed FM1-43 dye uptake experiments
at the fly NMJ. OE of 36 G4C2 repeats in motor neurons signifi-
cantly decreased FM1-43 dye uptake compared to G4C2 3X con-
trols (53% ± 4.9%, p < 0.001; Figures 6L and 6M). To determine
the contribution of reduced synaptic expression of Hsc70-4 to
SVC deficits, we co-overexpressed Hsc70-4, and we found
that it restores FM1-43 dye uptake in flies expressing expanded
G4C2 repeats back to control levels (Figures 6L and 6M). These
data indicate that C9-mediated FM1-43 dye uptake deficits are,
at least in part, the result of decreased Hsc70-4 expression at
the NMJ. Furthermore, these findings suggest that both
TDP-43 and C9orf72 models of ALS share common features of
post-transcriptionally reduced Hsc70 expression that are
accompanied by defects in SVC.

**DISCUSSION**

ALS is a synaptopathy like other neurodegenerative disorders,
including Parkinson’s and Alzheimer’s diseases (Gillingwater
and Wishart, 2013). However, the mechanisms underlying syn-
aptic dysfunction and neurodegeneration remain poorly under-
stood. TDP-43 has been linked to a majority of ALS cases and
is known to associate with RNA SGs, which sequester specific
mRNAs and reduce their translation during stress (Dewey
et al., 2011; Kim et al., 2014; Liu-Yesucevitz et al., 2010; McDonald et al., 2011). Collectively, these studies have suggested that
TDP-43 association with RNA SGs plays a critical role in main-
taining both proteostasis and ribostasis by controlling the
expression of target mRNAs and sequestration of protein part-
ers. Because of its central role in ALS pathogenesis, under-
standing the molecular mechanisms underlying TDP-43-mediated
neurodegeneration and defining its mRNA targets and their
mode of regulation are expected to provide a deeper under-
standing of disease pathophysiology and identify novel therapeu-
tic targets.

**Distinct Mechanisms Govern TDP-43-Dependent
Toxicity and Regulation of Hsc70-4 Expression**

Using a Drosophila model of ALS based on TDP-43, we show
that OE of ALS-associated mutant TDP-43 in motor neurons
sequestrers hsc70-4 mRNA and reduces synaptic Hsc70-4 protein
levels at larval NMJs. HSPA8 expression is also reduced in cell
bodies and growth cones of cultured primary motor neurons,
at NMJs of mice expressing mutant TDP-43, and in human
iPSC motor neurons, suggesting that the relationship between
TDP-43 and Hsc70 is evolutionarily conserved.

TDP-43WT OE in fly motor neurons also leads to a post-trans-
criptional reduction of synaptic Hsc70-4 levels at larval NMJs.
However, IPs and cellular fractionations suggest that this is un-
likely due to hsc70-4 mRNA sequestration and reduced transla-
tion. It is possible that the sensitivity of this assay is limited or that
subtle defects in other aspects of RNA or protein processing
result in reduced Hsc70-4 protein expression at the NMJ.
Supporting this, bioinformatics analyses of published crosslinking
IP (CLIP) data suggest that endogenous TDP-43 can bind
HSPA8 introns and exons in mouse brains and human K562 cells
(Polymenidou et al., 2011; Van Nostrand et al., 2016) (Figure S7).

Hsc70 belongs to a neuroprotective synaptic network that
maintains SV exo- and endocytosis and prevents neurodegener-
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In the context of mutant TDP-43, co-OE of Hsc70-4 provides
significant rescue of the FM1-43 dye uptake defect, although we
note that this is not a full restoration to the levels observed in controls.
Co-OE of chaperone-dead Hsc70-4D10N fails to mitigate
TDP-43G298S-induced FM1-43 dye uptake deficits, suggesting
that motor neuron expression of mutant TDP-43 reduces
Hsc70-4 levels below a threshold that is necessary to maintain
a molecular chaperone activity facilitating SV endocytosis/cycling.

In contrast, for TDP-43WT, co-OE of Hsc70-4, while not suffi-
cient to rescue SVC, appears to mitigate toxicity by generally
Figure 6. C9orf72 Repeat Expansions Cause Reduced Hsc70-4/HSPA8 Expression and Defects in SVC

(A) WB for Hsc70-4 levels in VNCs of G_{c2}-expressing larvae. Genotypes are indicated on the bottom. Actin was used as a loading control.

(B) Quantification of Hsc70-4 protein levels from WBs represented as a ratio to G_{c2} 3X controls.

(C) qPCR for hsc70-4 mRNA in VNCs from animals expressing G_{c2} 36X in motor neurons versus G_{c2} 3X controls.

(D) qPCR for hsc70-4 mRNA in NMJ preparations from animals expressing G_{c2} 36X in motor neurons versus G_{c2} 3X controls.

(E and F) Single confocal sections of synaptic boutons in NMJ preparations immunostained for Hsc70-4 and the neuronal membrane marker Hrp from larvae expressing G_{c2} 36X (F) compared to G_{c2} 3X controls (E). Antibodies are indicated on the left.

(G) Quantification of Hsc70-4 intensity in synaptic boutons, normalized to bouton area.

(H) Quantification of Hsc70-4 intensity in muscle, normalized to muscle area.

(I) Confocal images of control and C9 ALS human iPSC motor neurons immunostained for DAPI, the dendritic marker Map2, and Hsc70/HSPA8. Genotypes are indicated on the left and antibodies are indicated on the top.

(J) Quantification of Hsc70/HSPA8 intensity in the soma and dendrites of control and C9 ALS human iPS motor neurons.

(K) qPCR for Hsc70/HSPA8 mRNA in control and C9 ALS human iPS motor neurons.

(L) Confocal images of FM1-43 dye uptake in Drosophila NMJs after 5 min of stimulation in HL-3 saline containing 90 mM KCl and 2 mM Ca^{2+}. Genotypes are indicated on the top and left.

(M) Quantification of FM1-43 dye uptake, normalized to total FM1-43 uptake area.

All data shown are mean ± SEM. Scale bars, 5 and 1 μm (E and F) and 10 μm (I and L).
improving proteostasis and specifically reducing the aggregation of TDP-43<sup>WT</sup> itself, which correlates with improved locomotor function and increased lifespan. This suggests that the mechanism by which TDP-43<sup>WT</sup> and TDP-43<sup>G258S</sup> cause defects in FM1-43 dye uptake is, at least in part, distinct; but, ultimately, reduced Hsc70-4 levels at the NMJ impact the activity of its co-chaperone complexes and clients, many of which control critical steps in SVC.

**Perturbations in the SV Cycle Highlight Key TDP-43-Dependent Functional Interactions between Hsc70-4 Co-chaperone and Client Partners**

Co-OE of Csp, an Hsc70-4 co-chaperone critical for stabilizing client binding (Jiang et al., 2007; Kampinga and Craig, 2010), restores FM1-43 dye uptake to control levels for both TDP-43<sup>WT</sup> and TDP-43<sup>G258S</sup>. This is surprising given that Hsc70-4 co-OE mitigated the effects of TDP-43<sup>G258S</sup> but failed to rescue TDP-43<sup>WT</sup>-mediated defects in endocytosis. A reasonable explanation for this result is that Csp has synaptic functions that are independent of Hsc70-4, as previously suggested (Bronk et al., 2005). These additional functions may contribute to improved SVC when Csp and TDP-43<sup>WT</sup> are co-overexpressed in motor neurons. Co-OE of the GTPase dynamin, a client of the Csp/Hsc70-4 chaperone complex (Zhang et al., 2012), also significantly restores FM1-43 dye uptake to control levels for wild-type and mutant TDP-43. Together, these findings are consistent with a scenario whereby TDP-43 impairs the activity of the Csp/Hsc70 chaperone complex to maintain normal dynamin function.

**Toward a Common, Synaptic Pathomechanism of Disease**

Similar to what we found for TDP-43, we show that HSPA8 expression is reduced at synaptic terminals of C9 repeat-expressing flies and in cell bodies and dendrites of human C9 iPSC motor neurons. Furthermore, we show that C9 repeat expansions in flies lead to defects in FM1-43 dye uptake. However, the precise mechanism by which C9 repeat expansions regulate Hsc70 expression and SVC remains to be determined. As previous studies have reported a decrease in excitatory junction potential (EJP) amplitude and quantal content in a fly model of C9 ALS (Zhang et al., 2015), it is possible that C9-mediated reduced synaptic expression of Hsc70 may elicit FM1-43 dye uptake defects that arise due to deficits in both SV exo- and endocytosis. Recent reports suggest that splicing and transcriptome alterations, including changes in synaptic gene expression, contribute to C9 ALS pathology (Cooper-Knock et al., 2015; Prudencio et al., 2015). Thus, it is likely that multiple mechanisms may be at play to alter SVC in C9-mediated ALS.

Although OE of either TDP-43<sup>WT</sup> or disease-associated TDP-43<sup>G258S</sup> leads to similar phenotypes, our findings of <i>hsc70-4</i> mRNA as a mutant-specific sequestration and translation target suggest that the molecular mechanisms utilized by different TDP-43 variants are, at least in part, distinct (see Figure 7). This knowledge could impact the development of therapeutic strategies that ought to take into consideration the specific mechanisms of toxicity associated with TDP-43<sup>WT</sup>, which is linked to 97% of ALS cases, as opposed to ALS-associated mutant TDP-43 that only represents 2%–3% of ALS cases. Taken together, our data suggest that post-transcriptional dysregulation of Hsc70-4 expression connects defects in ribostasis and proteostasis at synapses in ALS across multiple models. Improving SVC through OE of Hsc70, its co-chaperone Csp, or their client dynamin highlights the SV cycle as a unifying therapeutic target in ALS and related neurodegenerative disorders, such as frontotemporal dementia (FTD) and Alzheimer’s disease, in which TDP-43 pathology has been reported.

**EXPERIMENTAL PROCEDURES**

**Drosophila Genetics**

All Drosophila stocks and crosses were kept on standard yeast/cornmeal/molasses food at 22° C unless otherwise noted.

**Larval Turning Assays**

Wandering third-instar larvae were placed on a grape juice plate. After a 30-s acclimation, larvae were gently turned ventral side up and observed until they turned over and began making a forward motion.

**Lifespan Analysis**

Newly eclosed males and females (100 per genotype) were separated and placed in a new vial every 7 days. Survival plots were generated using R and RStudio.

**Electrophysiology**

Current-clamp recordings were made from muscle 6 of third-instar larvae using a modified Ca<sup>2+</sup> free HL-3 saline, as described (Dawson-Scully et al., 2000).

**FM1-43 Dye Uptake Assays**

FM1-43 dye uptake assays using 4 μM FM1-43FX, 90 mM KCl, and 2 mM Ca<sup>2+</sup> for stimulation were adapted from Kuromi and Kodokoro (2005) and Verstreken et al. (2008).

**Cellular Fractionations and Western Blotting**

Cellular fractionations were previously described (Estes et al., 2013; Liachko et al., 2010).

**Western blots (WBs)** were performed as described (Coyne et al., 2014, 2015). Primary antibodies were 1:3,000 rabbit Anti-Hsc70-4 (from K. Zinsmaier), 1:6,000 rabbit Anti-GFP (Invitrogen), and 1:5,000 rabbit Anti-β-actin (Cell Signaling Technology).

**IPs**

IPs were performed using published protocols (Coyne et al., 2014).

**Polysome Fractionations**

Polysome fractionations were previously described (Coyne et al., 2014, 2015).

**Real-Time qPCR**

Real-time qPCR is described in the Supplemental Experimental Procedures. An RNaseasy kit (QIAGEN) was used for RNA isolation. First-strand cDNA synthesis was performed with a Superscript III cDNA synthesis kit (Invitrogen). qPCR reactions were carried out using SYBR Select Master Mix (Applied Biosystems) and an ABI 7300 Real Time PCR System (Applied Biosystems).

**Immunohistochemistry and Confocal Imaging**

Larval NMJs were prepared as described (Estes et al., 2011, 2013). Primary antibodies were 1:50 rabbit Anti-Hsc70-4 (from K. Zinsmaier), 1:300 mouse Anti-DCSP2 (DSHB), and 1:50 mouse Anti-Bruchpilot, NCB2 (DSHB).
Primary Neuron Cultures: Transfection, Staining, Image Acquisition, and Analysis

All procedures for animal experiments were approved by the Emory University Institutional Animal Care and Use Committee (IACUC) and the Emory University Institutional Review Board. Primary motor neurons from spinal cords of mouse embryos were isolated and plated at day 13.5 (Fallini et al., 2010).

Human iPSC Motor Neuron Differentiation and Immunocytochemistry for Hsc70/HSPA8

Control iPSCs and C9 iPSCs (>30 repeats) were differentiated to motor neurons as described by previous studies (Donnelly et al., 2013; Zhang et al., 2015).

Mouse NMJs

Animal lines and procedures are described in the Supplemental Experimental Procedures. All procedures were approved by the University of Kansas Medical Center IACUC. Immunohistochemistry, image analysis, and statistics were described in Chen et al. (2011, 2012) and Nishimune et al. (2004).

Bioinformatics Analyses

See the Supplemental Experimental Procedures.

Statistical Analyses

Larval turning assays used 30 larvae/genotype. Lifespan analysis used 100 flies/genotype. At least 6 animals/genotype were used for electrophysiology. FM1-43 dye uptake experiments were performed on 12 animals, and a total of 24 NMJs (2/animal) were used for analysis. Cellular fractionation, IPs, WBs, polysome fractionations, and qPCR were performed in triplicate. For analysis of mouse primary motor neurons, cell bodies and 10-μm segments of the axonal growth cones were analyzed (cell body: GFP n = 57, TDP-43 WT n = 49, TDP-43 Q331K n = 42, and TDP-43 M337V n = 49; growth cone: GFP n = 30, TDP-43 WT n = 24, TDP-43 Q331K n = 35, and TDP-43 M337V n = 36). For immunocytochemistry in TDP-43 iPSC motor neurons, two control lines and two differentiations of one TDP-43 line were used. For immunocytochemistry in C9orf72 iPSC motor neurons, one control line and one C9 line were used. Statistical analyses were performed using the appropriate t test or ANOVA model with Tukey’s post hoc test. For cellular fractionations, Fisher’s test was used. Statistics were performed using GraphPad Prism software version 7.0. Lifespan statistical analysis was done using the log-rank test in R. All data shown are mean ± SEM (*p < 0.05, **p < 0.01, and ***p < 0.001; n.s., not significant).

Additional details on fly lines, iPSC demographic information, and experimental procedures can be found in the Supplemental Experimental Procedures.
SUPPLEMENTAL INFORMATION

Supplemental Information includes Supplemental Experimental Procedures and seven figures and can be found with this article online at https://doi.org/10.1016/j.celrep.2017.09.028.

AUTHOR CONTRIBUTIONS


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