

Dystrobrevin Controls Neurotransmitter Release and Muscle Ca²⁺ Transients by Localizing BK Channels in *Caenorhabditis elegans*

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Dystrobrevin is a major component of a dystrophin-associated protein complex. It is widely expressed in mammalian tissues, including the nervous system, in which it is localized to the presynaptic nerve terminal with unknown function. In a genetic screen for suppressors of a lethargic phenotype caused by a gain-of-function isoform of SLO-1 in *Caenorhabditis elegans*, we isolated multiple loss-of-function (*lf*) mutants of the dystrobrevin gene *dyb-1*. *dyb-1(lf)* phenocopied *slo-1(lf)*, causing increased neurotransmitter release at the neuromuscular junction, increased frequency of Ca²⁺ transients in body-wall muscle, and abnormal locomotion behavior. Neuron- and muscle-specific rescue experiments suggest that DYB-1 is required for SLO-1 function in both neurons and muscle cells. DYB-1 colocalized with SLO-1 at presynaptic sites in neurons and dense body regions in muscle cells, and *dyb-1(lf)* caused SLO-1 mislocalization in both types of cells without altering SLO-1 protein level. The neuronal phenotypes of *dyb-1(lf)* were partially rescued by mouse α -dystrobrevin-1. These observations revealed novel functions of the BK channel in regulating muscle Ca²⁺ transients and of dystrobrevin in controlling neurotransmitter release and muscle Ca²⁺ transients by localizing the BK channel.

Introduction

Dystrobrevins are cytosolic proteins that bind to dystrophin and are considered to be a major component of a dystrophin-associated protein complex (DAPC) that links the cytoskeleton to extracellular matrix (Davies and Nowak, 2006). There are two dystrobrevin genes (α DB and β DB) in human and mice (Rees et al., 2007) but only one such gene (*dyb-1*) in *Caenorhabditis elegans* (www.wormbase.org, WS225). In mouse, knock-out of α DB causes skeletal and cardiac myopathies and impaired nitric oxide-mediated signaling (Grady et al., 1999), and knock-out of both α DB and β DB causes synaptic defects and abnormal motor behavior (Grady et al., 2006). In *C. elegans*, mutations of either *dyb-1* or the dystrophin gene *dys-1* cause muscle degeneration in a sensitized genetic background as well as behavioral and pharmacological phenotypes suggestive of enhanced cholinergic transmission (Gieseler et al., 1999; Giugia et al., 1999). Intriguingly, similar phenotypes are observed in mutants of the BK channel gene *slo-1* (Carre-Pierrat et al., 2006). The relationship between SLO-1 and the DAPC is not totally clear. A recent study

shows that a membrane protein known as ISLO-1 is important to SLO-1 subcellular localization in *C. elegans* body-wall muscle by interacting with the DAPC (Kim et al., 2009).

The BK channel is a Ca²⁺ and voltage-gated K⁺ channel expressed in many tissues, including the nervous system (Wang, 2008) and skeletal muscle (Latorre et al., 1982; Blatz and Magleby, 1984; Knaus et al., 1995). In the nervous system, the BK channel colocalizes with voltage-gated Ca²⁺ channels at axon presynaptic sites (Robitaille et al., 1993; Yazejian et al., 2000) and serves as a key negative regulator of neurotransmitter release (Robitaille et al., 1993; Hu et al., 2001; Wang et al., 2001; Raffaelli et al., 2004; Wang, 2008). The localization of the BK channel to the vicinity of voltage-gated Ca²⁺ channels allows it to be activated by Ca²⁺ microdomains, which are hemispheric sites of high [Ca²⁺] at the inner mouth of open Ca²⁺ channels (Roberts et al., 1990; Augustine et al., 2003). In mammalian striated muscle, the BK channel is enriched in the transverse tubule (t-tubule) membrane (Latorre et al., 1982; Knaus et al., 1995) with unknown function. In *C. elegans* body-wall muscle, which is analogous to mammalian striated muscle (Moerman and Fire, 1997), SLO-1 colocalizes with the L-type voltage-gated Ca²⁺ channel EGL-19. SLO-1 in muscle is potentially involved in regulating locomotion and egg-laying behaviors (Kim et al., 2009; Abraham et al., 2010; Chen et al., 2010a,b).

Through a genetic screen for suppressors of a behavioral phenotype caused by a gain-of-function (*gf*) isoform of SLO-1, we identified DYB-1 as a protein essential to SLO-1 function *in vivo*. Analyses of mutant phenotypes revealed that SLO-1 regulates Ca²⁺ transients in body-wall muscle cells and that DYB-1 regulates neurotransmitter release and muscle Ca²⁺ transients by localizing SLO-1 to presynaptic sites and muscle-dense body

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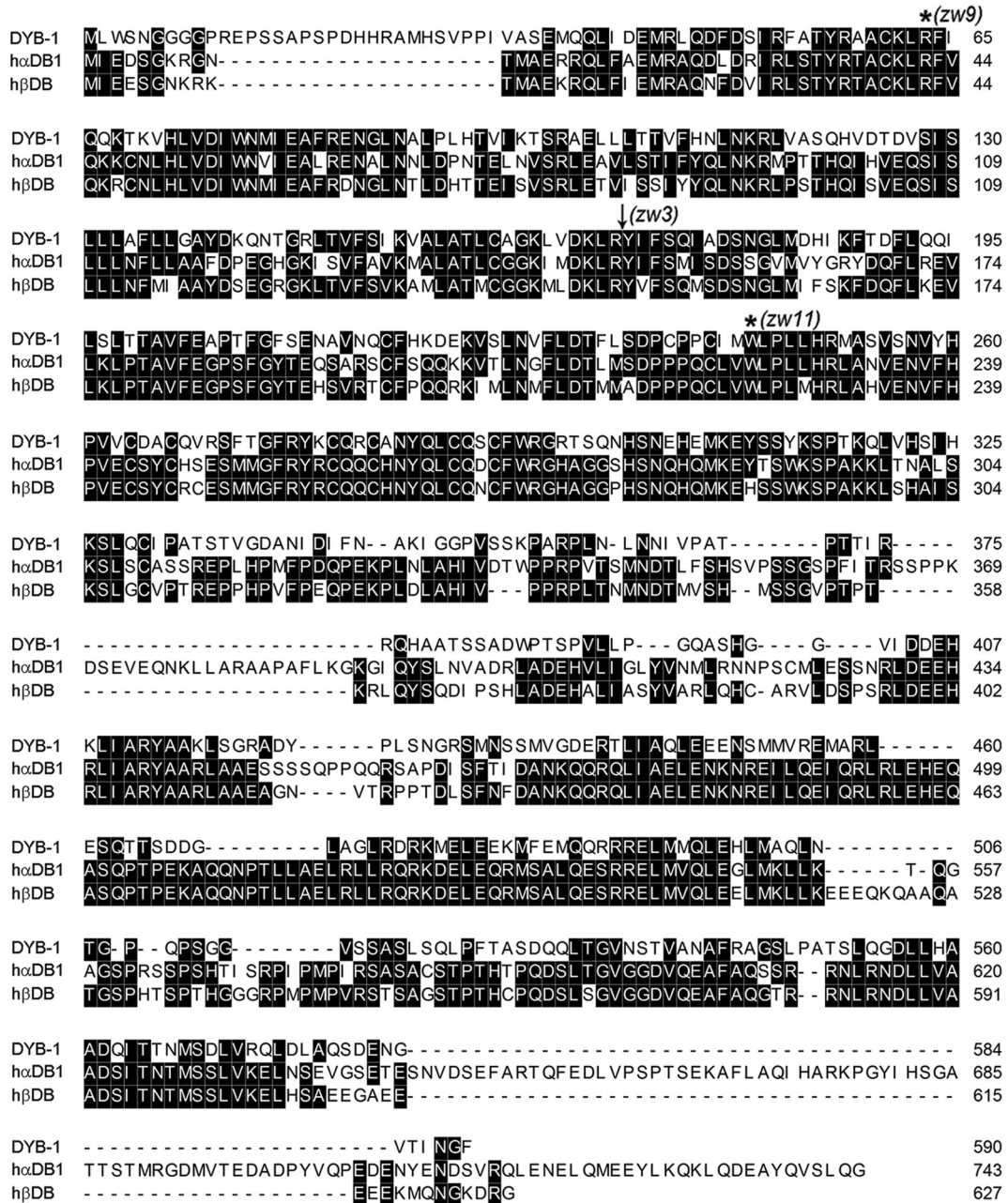


Figure 1. *dyb-1* encodes a homolog of mammalian dystrobrevins. Shown is the alignment of predicted amino acid sequences of DYB-1, human αDB1 (accession number NM_001390), and human βDB (accession number AF022728). Positions of molecular lesions of three *dyb-1* alleles are indicated. Two alleles have mutations leading to premature stop codon (marked by *) and one allele (marked by an arrow) disrupts the splice acceptor site of the fourth intron, leading to the removal of exon 5 and a frame shift after Y171, which was shortly followed by a stop codon.

regions, respectively. These findings potentially help to understand the molecular mechanisms of neurological and muscular defects caused by deficiencies of the DAPC.

Materials and Methods

Growth and culture of *C. elegans*. *C. elegans* hermaphrodites were grown on agar plates with a layer of OP50 *Escherichia coli* at room temperature (21–22°C) or inside an environmental chamber (21°C).

Strains. N2 Bristol was used as the wild-type in all experiments. The other strains used in this study are as follows: ZW083 [*zwIs101* (*Pslo-1::slo-1::GFP*)]; ZW320 [*zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*)]; ZW331 [*zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*); *dyb-1*(*zw11*)]; ZW349 [*dyb-1*(*zw11*)]; ZW352 [*zwIs101* (*Pslo-1::slo-1::GFP*); *dyb-1*(*zw11*)]; ZW385 [*zwEx132* (*Pdyb-1::GFP*; *lin-15*(+)); *lin-15*(*n765*)]; ZW471 [*zwEx150* (*Pdyb-1::αDB1*); *Pmyo-3::GFP*]; *zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*);

dyb-1(*zw11*); ZW495 [*zwIs132* (*Pmyo-3::GCaMP2*; *lin-15*(+))]; ZW527 [*zwIs132* (*Pmyo-3::GCaMP2*; *lin-15*(+)); *slo-1*(*md1745*)]; ZW528 [*zwIs132* (*Pmyo-3::GCaMP2*; *lin-15*(+)); *zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*)]; ZW536 [*zwIs132* (*Pmyo-3::GCaMP2*; *lin-15*(+)); *dyb-1*(*zw11*)]; ZW581 [*slo-1*(*md1745*); *dyb-1*(*zw11*)]; ZW604 [*zwIs101* (*Pslo-1::slo-1::GFP*); *zwIs135* (*Pdyb-1::dyb-1::mStrawberry*; *rol-6*(+))]; ZW605 [*zwEx164* (*Punc-47::slo-1::mStrawberry*; *Punc-25::GFP::unc-2*; *lin-15*(+)); *lin-15*(*n765*)]; ZW608 [*zwEx166* (*Pmyo-3::dyb-1*; *Pmyo-3::GFP*); *zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*); *dyb-1*(*zw11*)]; ZW609 [*zwEx167* (*Prab-3::dyb-1*; *Prab-3::GFP*); *zwIs129* (*Pslo-1::slo-1*(*gf*); *Pmyo-2::YFP*); *dyb-1*(*zw11*)]; ZW610 [*zwEx166* (*Pmyo-3::dyb-1*; *Pmyo-3::GFP*); *dyb-1*(*zw11*)]; ZW611 [*zwEx167* (*Prab-3::dyb-1*; *Prab-3::GFP*); *dyb-1*(*zw11*)]; ZW612 [*zwIs101* (*Pslo-1::slo-1::GFP*); *dys-1*(*cx18*)]; ZW615 [*zwIs136* (*Pdyb-1::dyb-1::GFP*; *lin-15*(+)); *lin-15*(*n765*)]; ZW616 [*zwIs136* (*Pdyb-1::dyb-1::GFP*; *lin-15*(+)); *dys-1*(*cx18*)]; ZW622

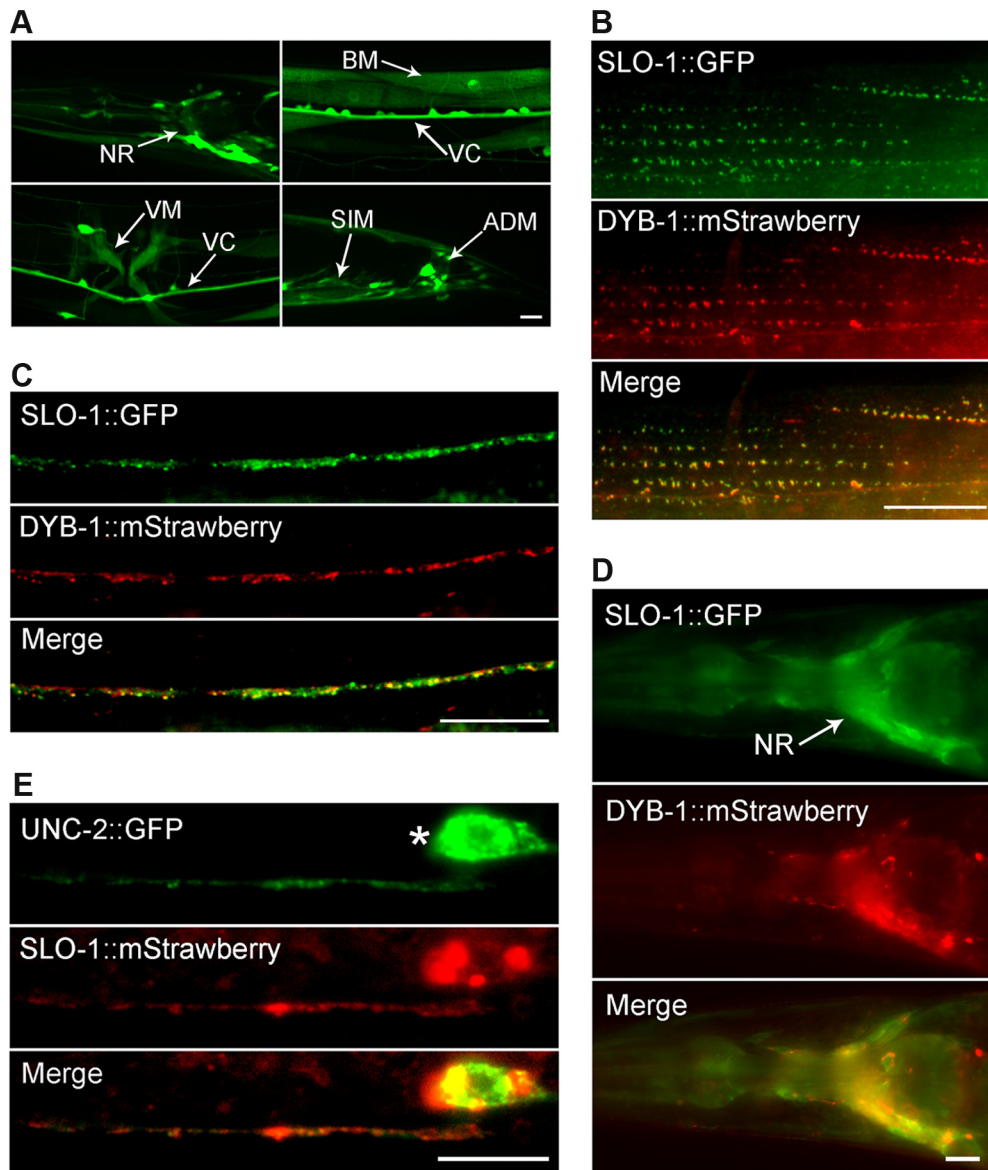


Figure 2. DYB-1 and SLO-1 were coexpressed and colocalized in neurons and muscle cells. **A**, Expression of GFP under the control of *Pdyb-1* resulted in GFP epifluorescence in many neurons in the nerve ring (NR), ventral cord (VC), and tail (not labeled), as well as several types of muscles, including body-wall muscle (BM), vulval muscle (VM), stomatointestinal muscle (SIM), and anal depressor muscle (ADM). **B–D**, SLO-1::GFP and DYB-1::mStrawberry colocalized in body-wall muscle cells (**B**), the dorsal nerve cord (**C**), and the nerve ring (**D**) when they were expressed under the control of their native promoters. **E**, SLO-1::mStrawberry and UNC-2::GFP, which were expressed in GABAergic neurons, colocalized in the ventral nerve cord. * indicates the soma of a GABAergic motoneuron. Scale bars, 10 μm.

[*zwEx168(Pmyo-3::αDB1; Pmyo-3::GFP); zwIs129(Pslo-1::slo-1(gf); Pmyo-2::YFP); dyb-1(zw11)*]; ZW623 [*zwEx169(Prab-3::αDB1; Prab-3::GFP); zwIs129(Pslo-1::slo-1(gf); Pmyo-2::YFP); dyb-1(zw11)*]; ZW624 [*zwEx170(Pdyb-1::αDB1; Pmyo-2::DsRed); zwIs101(Pslo-1::slo-1::GFP); dyb-1(zw11)*]; ZW625 [*zwEx171(Pmyo-3::slo-1; Pmyo-2::DsRed); zwIs132(Pmyo-3::GCaMP2; lin-15(+)); slo-1(md1745)*]; ZW626 [*zwEx172(Prab-3::slo-1; Pmyo-2::DsRed); zwIs132(Pmyo-3::GCaMP2; lin-15(+)); slo-1(md1745)*].

Mutant screen. An integrated transgenic strain expressing *Pslo-1::SLO-1(E^{350Q})* and *Pmyo-2::YFP* in the wild-type genetic background was used for mutant screen. The *Pslo-1::SLO-1(E^{350Q})* transgene caused a lethargic phenotype, whereas the *Pmyo-2::YFP* transgene was used as a genetic marker. Synchronized L4-stage worms were treated with the chemical mutagen ethyl methanesulfonate (50 mM) for 4 h at room temperature. The F2 progeny were screened for animals that moved better than the original *slo-1(gf)* animals.

Behavioral assay. Locomotion velocity and the head-bending angle were quantified using an automated worm tracking and analysis system

(Chen et al., 2010a). Specifically, a single adult hermaphrodite was transferred to an agar plate without food. After allowing ~30 s for recovery from the transfer, snapshots of the worm were taken at 15 frames/s for 30 s using a VGA FireWire camera (XCD-V60; Sony) mounted on a stereomicroscope (SMZ800; Nikon). The worm was constantly kept in the center of the view field with a motorized microscope stage (OptiScan ES111; Prior Scientific). Both the camera and the motorized stage were controlled by a custom program running in MATLAB (MathWorks).

Cloning of *dyb-1*. *dyb-1(zw11)* was used for single nucleotide polymorphism-based genetic mapping (Davis et al., 2005). After mapping the mutation to a small interval, the candidate gene was identified by testing whether mutants of candidate genes would complement with *zw11* in suppressing the lethargic phenotype of *slo-1(gf)*. Full-length *dyb-1* cDNA was obtained by RT-PCR. Molecular lesions of *dyb-1* mutants were identified by sequencing the cDNA or genomic DNA of *dyb-1* gene prepared from three randomly picked mutant alleles.

Analysis of expression pattern and subcellular localization. The expression pattern of *dyb-1* was determined by expressing GFP under the con-

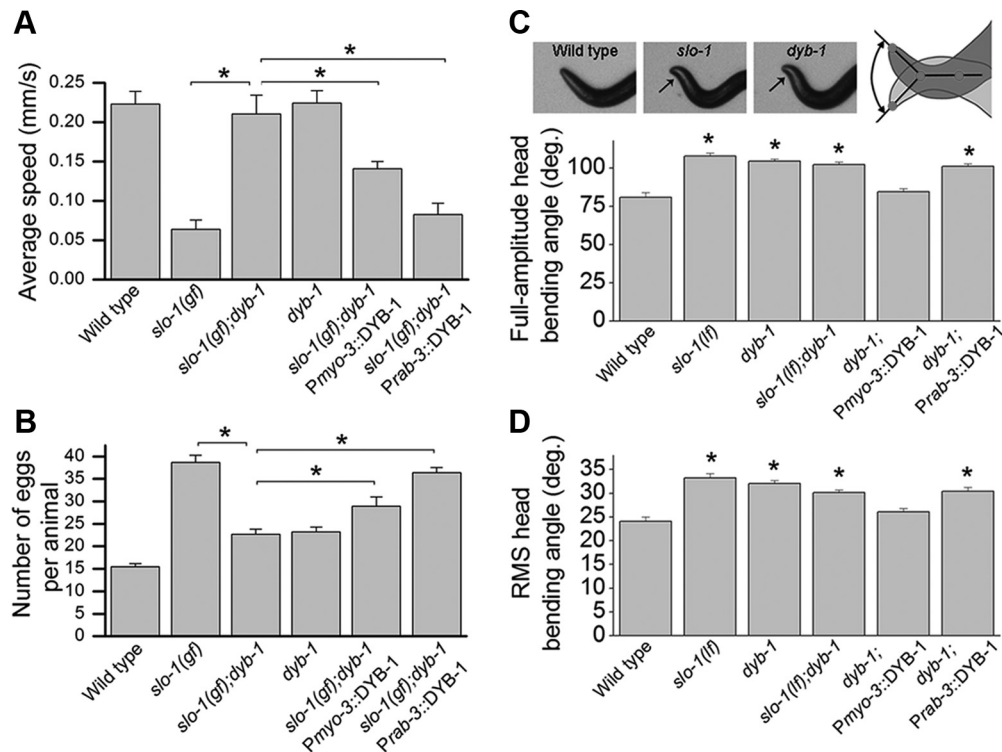


Figure 3. *dyb-1* mutant suppressed locomotion and egg-laying phenotypes of *slo-1(gf)* and phenocopied *slo-1(lf)* in head-bending behavior. **A, B**, *dyb-1* mutant suppressed the locomotion (**A**) and egg-laying (**B**) defects caused by *slo-1(gf)*, which could be reversed by expressing wild-type DYB-1 in either neurons (*Prab-3*) or body-wall muscle cells (*Pmyo-3*). Twenty to 30 worms per group in **A** and 10 worms per group in **B**. **C, D**, Comparisons of full-amplitude head-bending angle and root mean square (RMS) head-bending angle. The head-bending angle was similarly increased in *dyb-1* mutant, *slo-1(lf)*, and the double mutant. This phenotype of *dyb-1* mutant could be rescued by expressing wild-type DYB-1 in muscle but not neurons. Ten worms per group. * $p < 0.01$, significantly different between the indicated groups (**A, B**) or from wild type (**C, D**) (one-way ANOVA with Bonferroni's *post hoc* tests).

tol of a 3.1-kb *dyb-1* promoter (*Pdyb-1::GFP*, wp747). The plasmid was injected into the *lin-15(n765)* strain using a *lin-15* rescue plasmid as a transformation marker. The transgenic animals were photographed using a Carl Zeiss Axiovert 200M fluorescence microscope (40 \times objective) with an apotome device (Carl Zeiss) for optical sectioning.

Subcellular localization of DYB-1 was determined by expressing DYB-1 with GFP tagged to its C terminus under the control of *Pdyb-1::DYB-1::GFP*, wp1132). SLO-1 subcellular localization was examined with an integrated transgenic strain expressing *Pslo-1::SLO-1::GFP* (wp5) (Chen et al., 2010a). To determine whether DYB-1 and SLO-1 are colocalized, mStrawberry-tagged DYB-1 (*Pdyb-1::DYB-1::mStrawberry*, wp1130) was expressed in the transgenic strain expressing *Pslo-1::SLO-1::GFP*. To determine whether SLO-1 colocalized with UNC-2 in neurons, SLO-1::mStrawberry and UNC-2::GFP (Saheki and Bargmann, 2009) fusion proteins were expressed under the control of *Punc-47* and *Punc-25*, respectively. Epifluorescence of the fusion proteins in transgenic animals was visualized and photographed with a Nikon TE2000-U inverted microscope (100 \times objective) and a Peltier cooled digital camera (F-view II; Olympus).

Recording of postsynaptic currents. Evoked postsynaptic currents (ePSCs) and miniature postsynaptic currents (mPSCs) were recorded from the *C. elegans* neuromuscular junction (NMJ) using an established technique (Richmond et al., 1999; Liu et al., 2007). The recording pipette solution contained the following (in mM): 120 KCl, 20 KOH, 5 Tris, 0.25 CaCl₂, 4 MgCl₂, 36 sucrose, 5 EGTA, and 4 Na₂ATP, pH 7.2. Two external solutions with different [Ca²⁺] (5 and 0.5 mM) were used. The external solution with the higher [Ca²⁺] contained the following (in mM): 140 NaCl, 5 KCl, 5 CaCl₂, 5 MgCl₂, 11 dextrose, and 5 HEPES, pH 7.2. This solution was modified by reducing CaCl₂ to 0.5 mM and increasing NaCl to 145 mM to make the external solution with the lower [Ca²⁺].

Ca²⁺ imaging. An integrated *Pmyo-3::GCaMP2* transgene was crossed into different mutant backgrounds. Ca²⁺ transients were monitored by imaging GCaMP2 fluorescence in body-wall muscle cells of dissected worms as described previously (Liu et al., 2011).

Western blot. To examine SLO-1::GFP protein level expressed in worms, worms of mixed stages were homogenized in a lysis buffer containing 2% SDS/100 mM NaCl/10% glycerol/50 mM Tris, pH 6.8. Soluble protein extracts were separated on 4–12% SDS-PAGE gels and probed with GFP (Invitrogen) and α -tubulin (Santa Cruz Biotechnology) antibodies. Anti-mouse IgG HRP (Santa Cruz Biotechnology) was used as the secondary antibody for detection by enhanced chemiluminescence (Pierce).

Data analysis. The averaged locomotion speed and head-bending angle were determined using an automated worm tracking and analysis system (Chen et al., 2010a,b). Briefly, 13 marker points were placed at equal intervals along the spline of each worm, which was imaged at 15 frames/s, and the angle supplementary to the angle formed by the two straight lines connecting the marker points 1 and 2 and the marker points 2 and 3 was determined. Both the full-amplitude dorsal/ventral head-bending angle and the root mean square of head-bending angles were determined. The mean locomotion speed was calculated based on the distance traveled over time.

Amplitude and frequency of mPSCs were analyzed using MiniAnalysis (Synaptosoft). A detection threshold of 10 pA was used in initial automatic analysis, followed by visual inspections to include missed smaller events (5 pA or larger) and to exclude false events resulting from baseline fluctuations. For each recording, we adjusted the position of the stimulating electrode to obtain the largest ePSCs and used an average of the two largest values to derive an amplitude measurement for that recording.

For Ca²⁺ imaging data, all the muscle cells within the camera imaging field were chosen as separate regions of interest for quantification of fluorescence intensity changes over successive frames. Fluorescence intensity was first plotted as absolute signal intensity over time using the NIS-Elements software (Nikon) and then converted to F/F_0 using MATLAB (MathWorks) running a custom module. The frequency of Ca²⁺ transient peaks (defined as $F/F_0 \geq 0.05$) was quantified for each cell. The mean frequency of all cells in the imaging field per preparation except for

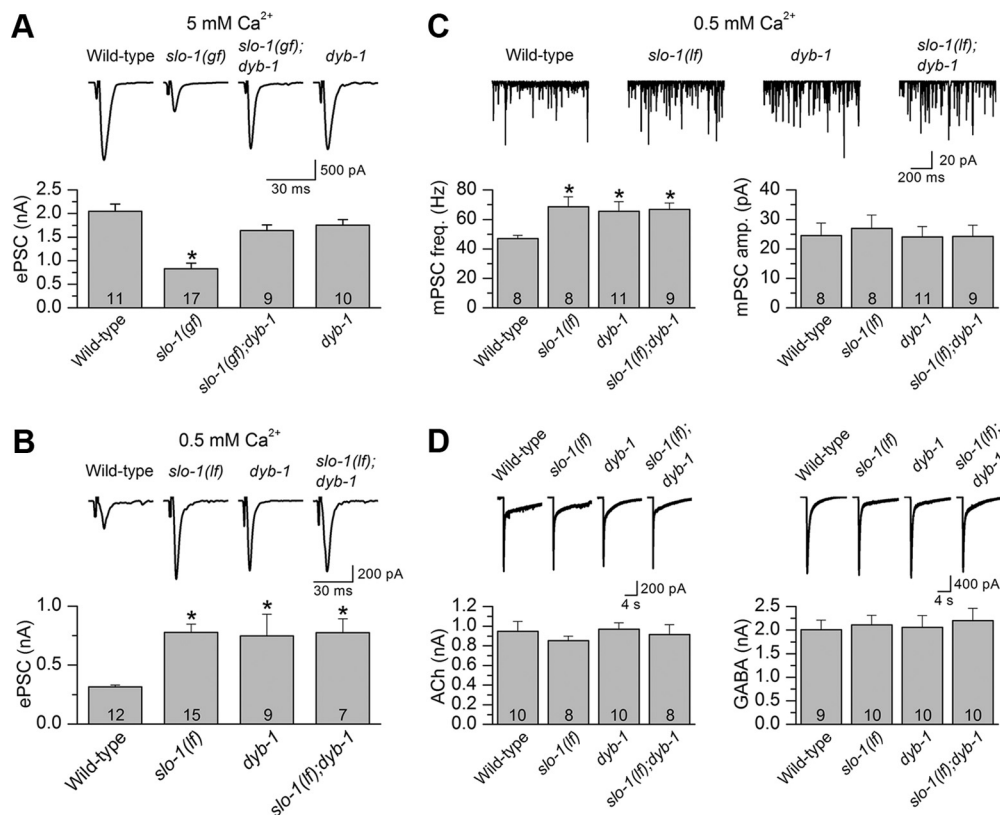


Figure 4. DYB-1 was required by SLO-1 to control neurotransmitter release at NMJs. **A**, *dyb-1* mutant counteracted the inhibitory effect of *slo-1(gf)* on ePSCs at 5 mM $[Ca^{2+}]_o$. **B**, The ePSC amplitude of *dyb-1* mutant was significantly larger than that of the wild type but similar to that of *slo-1(lf)* or the *slo-1(lf); dyb-1* double mutant. **C**, The frequency of mPSCs was increased, whereas the mean amplitude did not change in *slo-1(lf)*, *dyb-1*, and *slo-1(lf); dyb-1* compared with the wild type. **D**, Responses of body-wall muscle cells to exogenous ACh (100 μ M) and GABA (100 μ M) were unaltered in the mutants compared with the wild type. The holding potential for all recordings was -60 mV. * $p < 0.05$, significantly different from the wild type (one-way ANOVA with Bonferroni's *post hoc* test). The number of samples analyzed is indicated inside each column.

the *slo-1(gf)* group was treated as one sample. For the *slo-1(gf)* group, only data from cells with Ca^{2+} transients were included in the analysis.

Data graphing and statistical analyses. Graphing and statistical analyses were performed with Origin (version 8.5; OriginLab). Either unpaired *t* test or one-way ANOVA (followed by Bonferroni's *post hoc* test) was used for statistical comparisons. $p < 0.05$ is considered statistically significant. All values are expressed as mean \pm SEM.

Results

dyb-1 mutants suppressed the lethargic phenotype caused by *slo-1(gf)*

To identify proteins important to BK channel function *in vivo*, we performed a genetic screen for mutants that suppressed the lethargic phenotype of a worm strain expressing SLO-1(*gf*) under the control of *slo-1* promoter (*Pslo-1*) (Chen et al., 2010b). SLO-1(*gf*) was created by mutating glutamate 350 to glutamine, which causes a 40 mV shift toward more hyperpolarized potentials in the SLO-1 conductance–voltage relationship (Chen et al., 2010b). Among 25 mutants isolated from a screen of 24,000 haploid genomes, seven mutants belonged to the dystrobrevin gene *dyb-1* (locus F47G6.1, www.wormbase.org, WS225) based on complementation tests and rescue experiments. Although there are two dystrobrevin genes (α and β) with multiple splice variants in human (Peters et al., 1998; Holzfeind et al., 1999), the *C. elegans* genome has only one dystrobrevin gene with probably a single isoform (www.wormbase.org, WS225). DYB-1 shares 45.5 and 46.9% sequence identity with human α DB1 and β DB, respectively (Fig. 1). Three mutants were randomly chosen for se-

quencing, and all of them were found to be putative nulls (Fig. 1). Subsequent analyses were performed with the *dyb-1(zw11)* allele.

DYB-1 was coexpressed and colocalized with SLO-1 in both neurons and muscle cells

The isolation of *dyb-1* mutants as suppressors of the *slo-1(gf)* lethargic phenotype suggests that DYB-1 might be required for SLO-1 function *in vivo* and have similar expression and subcellular localization patterns as SLO-1. Expression of GFP under the control of the *dyb-1* promoter (*Pdyb-1*) resulted in GFP epifluorescence in many neurons and several muscles, including body-wall muscle (Fig. 2A). This expression pattern of *dyb-1* is indistinguishable from that of *slo-1* (Chen et al., 2010a,b). When mStrawberry-tagged full-length DYB-1 (DYB-1::mStrawberry) was coexpressed with GFP-tagged full-length SLO-1 (SLO-1::GFP) under the controls of their respective promoters, the two fusion proteins appeared as puncta with colocalization at muscle dense body regions and along the dorsal nerve cord (Fig. 2B,C). The two fusion proteins also showed overlapping expression in the nerve ring (Fig. 2D).

To determine whether the SLO-1::GFP puncta in the dorsal nerve cord corresponded to synaptic sites, we coexpressed mStrawberry-tagged SLO-1 (SLO-1::mStrawberry) and GFP-tagged UNC-2 (UNC-2::GFP) in GABAergic neurons under the controls of the *unc-47* (vesicular GABA transporter) and *unc-25* (glutamic acid decarboxylase) promoters, respectively, and analyzed their localization in the ventral nerve cord (SLO-1::mStrawberry signal in the dorsal cord was too weak to be imaged). UNC-2 is an N/P/Q-type

(Ca_v2) voltage-gated Ca²⁺ channel (Bargmann, 1998) and is localized to presynaptic sites in motoneurons (Saheki and Bargmann, 2009). We found that the two fusion proteins colocalized in the ventral nerve cord (Fig. 2E). The separate observations of SLO-1 colocalization with DYB-1 and UNC-2 at the nerve cords collectively suggest that SLO-1 and DYB-1 colocalized at presynaptic sites in motoneurons.

DYB-1 was required for SLO-1 function in both neurons and muscle cells

To determine whether DYB-1 contributes to SLO-1 function in neurons, muscles, or both, we expressed wild-type DYB-1 in the *slo-1(gf);dyb-1* double mutant using either the muscle-specific *myo-3* promoter (*Pmyo-3*) or the pan-neuronal *rab-3* promoter (*Prab-3*) and analyzed its effects on animal behaviors. Expression of wild-type DYB-1 in either neurons or muscle cells primarily reinstated the locomotion and egg-laying defects of *slo-1(gf)* (Fig. 3A,B), suggesting that DYB-1 is required for SLO-1 function in both neurons and muscle cells. In addition, we analyzed a head-bending phenotype associated with *slo-1(lf)* mutant (Kim et al., 2009; Chen et al., 2010a,b). The head-bending angle was increased to a similar degree in the *dyb-1* mutant, *slo-1(lf)*, and the double-mutant, *slo-1(lf);dyb-1* (Fig. 3C,D). The head-bending phenotype of the *dyb-1* mutant could be rescued by expressing wild-type DYB-1 in muscle cells but not neurons (Fig. 3C,D), suggesting that DYB-1 functions together with SLO-1 in muscle to regulate the head-bending angle.

DYB-1 was required by SLO-1 to regulate neurotransmitter release

SLO-1 is a potent negative regulator of neurotransmitter release at the *C. elegans* NMJ (Wang et al., 2001; Liu et al., 2007). To determine whether DYB-1 is required for this function of SLO-1, we analyzed the effect of *dyb-1* on ePSCs at the NMJ at two different extracellular Ca²⁺ concentrations (5 and 0.5 mM). The higher [Ca²⁺]_o is more suitable for determining whether *dyb-1* may reverse an inhibitory effect of *slo-1(gf)* on neurotransmitter release, whereas the lower [Ca²⁺]_o is more suitable for testing whether *dyb-1* could increase ePSC amplitude as *slo-1(lf)* does (Liu et al., 2007; Chen et al., 2010b). At 5 mM [Ca²⁺]_o, the inhibitory effect of *slo-1(gf)* on the ePSC amplitude was reversed by *dyb-1*, whereas *dyb-1* alone did not increase ePSC amplitude (Fig. 4A), which resembles *slo-1(lf)* and is probably attributable to the limited capacity of the readily releasable pool of synaptic vesicles (Wang et al., 2001). At 0.5 mM [Ca²⁺]_o, ePSC amplitude was increased to a similar degree in *dyb-1* and *slo-1(lf)* as well as in the *slo-1(lf);dyb-1* double mutant (Fig. 4B). These observations suggest that DYB-1 is required by SLO-1 to control neurotransmitter release.

To test whether the abnormal ePSCs in *slo-1* and *dyb-1* mutants resulted from altered sensitivity of postsynaptic receptors, we recorded the mPSCs at 0.5 mM [Ca²⁺]_o and compared the frequency and amplitude of mPSCs between the wild-type and the mutants. We found that the frequency of mPSCs was increased, whereas the amplitude of mPSCs did not change in *slo-1(lf)*, *dyb-1*, and *slo-1(lf);dyb-1* (Fig. 4C). We also examined the sensitivities of postsynaptic acetylcholine (ACh) and GABA receptors by puffing exogenous ACh and GABA directly onto the muscle cells. We found that the amplitude of ACh- and GABA-induced currents was similar between wild-type worms and the mutants (Fig. 4D). These observations suggest that SLO-1 and DYB-1 regulated neuromuscular transmission through a presynaptic mechanism.

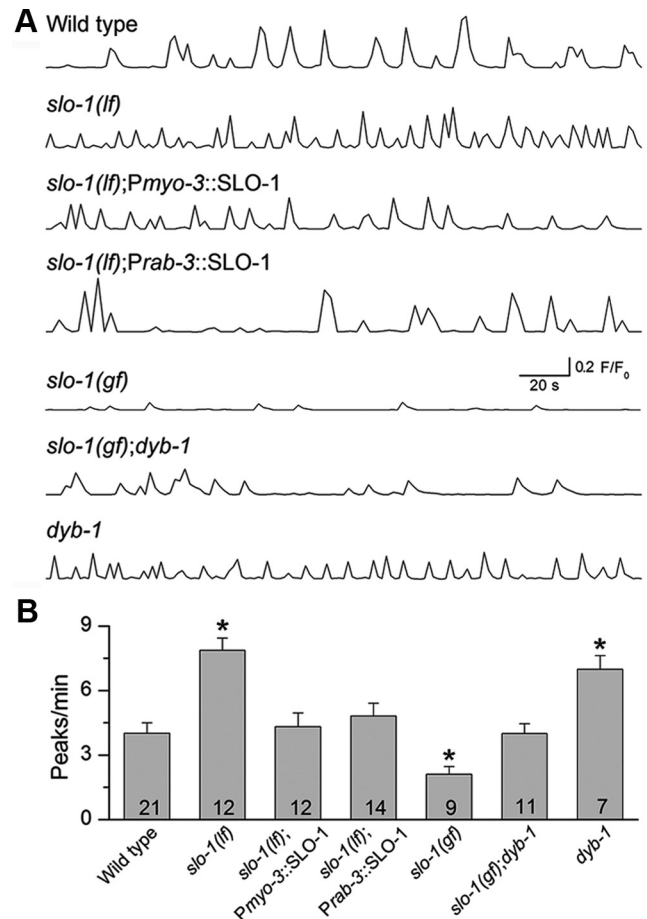


Figure 5. The frequency of Ca²⁺ transients in body-wall muscle cells was altered by mutations of *slo-1* and *dyb-1*. **A**, Representative traces of Ca²⁺ transients from wild-type worms and various mutants. **B**, Quantification of the frequency of Ca²⁺ transients. The frequency of muscle Ca²⁺ transients was increased in *slo-1(lf)*, which was restored by expressing wild-type SLO-1 in either muscles (*Pmyo-3::SLO-1*) or neurons (*Prab-3::SLO-1*). The frequency of Ca²⁺ transients was decreased in *slo-1(gf)*, which was reversed by *dyb-1* mutant. *dyb-1* mutant showed a similar increase in the frequency of Ca²⁺ transients as *slo-1(lf)* did. **p* < 0.05, significantly different from the wild type (one-way ANOVA with Bonferroni's *post hoc* test). The number of samples analyzed is indicated inside each column.

SLO-1 and DYB-1 controlled Ca²⁺ transients in body-wall muscle cells

Because SLO-1 colocalizes with the L-type voltage-gated Ca²⁺ channel EGL-19 (Kim et al., 2009) and EGL-19 is the predominant voltage-gated Ca²⁺ channel in body-wall muscle (Jospin et al., 2002), SLO-1 could potentially be activated by Ca²⁺ entering through EGL-19 and, in turn, regulate intracellular [Ca²⁺]_i. To examine this possibility, we expressed the calcium indicator GCaMP2 in body-wall muscle under the control of *Pmyo-3* and imaged Ca²⁺ transients as described previously (Liu et al., 2011). Ca²⁺ transients were observed in all muscle cells examined in wild-type and *slo-1(lf)* worms, with the frequency being much higher in the *slo-1(lf)* mutant than the wild type (Fig. 5). In *slo-1(gf)* worms, however, Ca²⁺ transients were detected in only approximately one-third of the preps with reduced frequency (Fig. 5). These observations suggest that a physiological function of SLO-1 is to inhibit muscle Ca²⁺ transients. Expression of wild-type SLO-1 in either muscle cells or neurons reversed the effect of *slo-1(lf)* on Ca²⁺ transients (Fig. 5), suggesting that SLO-1 regulates muscle Ca²⁺ transients through both presynaptic and postsynaptic effects.

To determine whether DYB-1 also regulates muscle Ca^{2+} transients by affecting SLO-1, we analyzed Ca^{2+} transients in the *dyb-1* mutant and *slo-1(gf);dyb-1* double mutant. The frequency of muscle Ca^{2+} transients was increased to a similar degree in the *dyb-1* mutant as in the *slo-1(lf)* mutant; furthermore, the *dyb-1* mutant counteracted the effect of *slo-1(gf)* on muscle Ca^{2+} transients (Fig. 5). These observations suggest that DYB-1 is required by SLO-1 to regulate Ca^{2+} transients.

DYB-1 was required for SLO-1 subcellular localization in neurons and body-wall muscle cells

DYB-1 could contribute to SLO-1 function through several potential mechanisms. We found that DYB-1 did not affect channel functional properties when it was coexpressed with SLO-1 in *Xenopus* oocytes (data not shown). In contrast, SLO-1::GFP puncta appeared diffuse and dim in the ventral and dorsal nerve cords (Fig. 6*A,E*) and were essentially absent in body-wall muscle cells (Fig. 6*B*) of the *dyb-1* mutant. The apparent SLO-1::GFP mislocalization did not result from a gross destabilization of presynaptic sites or muscle-dense bodies because the presynaptic marker RIM (Koushika et al., 2001) (Fig. 6*C,E*) and the dense body marker vinculin (Francis and Waterston, 1985) (Fig. 6*D*) were normally localized in the *dyb-1* mutant, nor did it result from a decrease of SLO-1::GFP protein level, which was similar between the wild-type and *dyb-1* mutant (Fig. 6*F*). These observations suggest that DYB-1 is required for proper SLO-1 subcellular localization in neurons and muscle cells and that the observed behavioral, synaptic, and muscle Ca^{2+} transient phenotypes of the *dyb-1* mutant were likely caused by SLO-1 mislocalization in these cells.

DYB-1 subcellular localization did not depend on dystrophin

Dystrophin has been implicated in controlling dystrobrevin expression in mouse (Ohlendieck and Campbell, 1991) and SLO-1 subcellular localization in *C. elegans* (Kim et al., 2009; Chen et al., 2010a). We found that SLO-1::GFP was mislocalized in body-wall muscle but normally localized in the dorsal nerve cord in the dystrophin null mutant *dys-1(cx18)* (Fig. 7*A,B,E*). However, DYB-1 expression and subcellular localization were unaltered in both muscle cells and neurons of the *dys-1* mutant (Fig. 7*C–E*). These observations suggest that in neurons DYB-1 localizes SLO-1 through a mechanism that is independent of DYS-1.

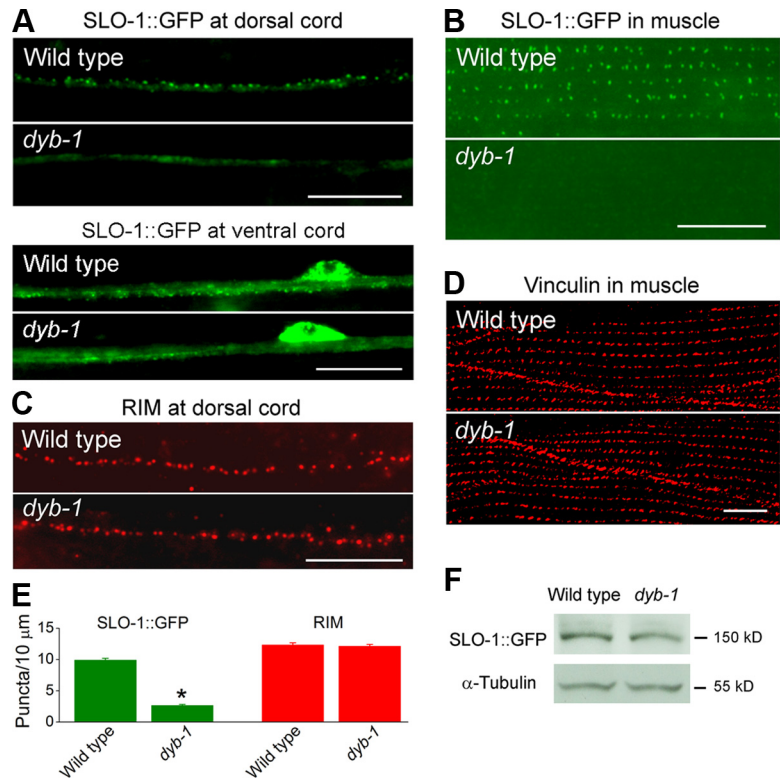


Figure 6. SLO-1 was mislocalized in both body-wall muscle cells and neurons of *dyb-1* mutant. **A**, SLO-1::GFP in the dorsal cord and ventral cord appeared as puncta in the wild type but was diffusely distributed in *dyb-1* mutant. **B**, SLO-1::GFP in muscle was localized to dense body regions in the wild type but undetectable in *dyb-1* mutant. **C**, **D**, The presynaptic marker RIM (**C**) and dense body marker vinculin (**D**) were normally localized in *dyb-1* mutant. **E**, Quantification of the density of dorsal cord puncta shown in **A** and **C**. * $p < 0.0001$, significantly different from the wild type (unpaired *t* test). **F**, Western blot shows that the level of SLO-1::GFP total protein was comparable between the wild-type and *dyb-1* mutant. α -Tubulin was blotted to show equal loading of the protein samples. Scale bars, 10 μm .

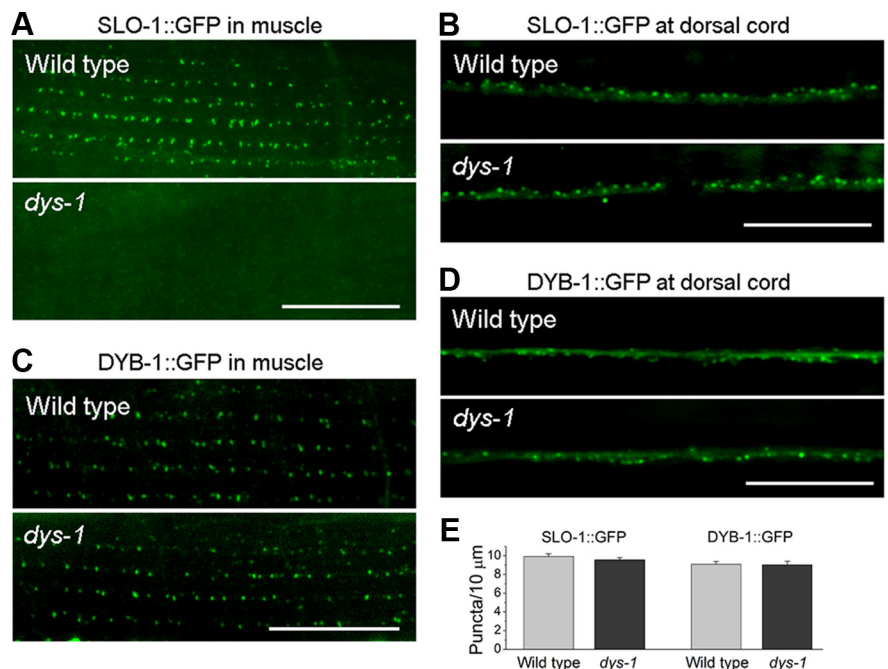


Figure 7. Effects of *dys-1* mutant on SLO-1 and DYB-1 subcellular localization. **A**, **B**, SLO-1::GFP was mislocalized in body-wall muscle but not neurons of the dystrophin mutant *dys-1(cx18)*. **C**, **D**, DYB-1::GFP was normally localized in both muscle cells and neurons of *dys-1* mutant. **E**, Quantification of SLO-1::GFP and DYB-1::GFP puncta density in the dorsal nerve cord. Scale bars, 10 μm .

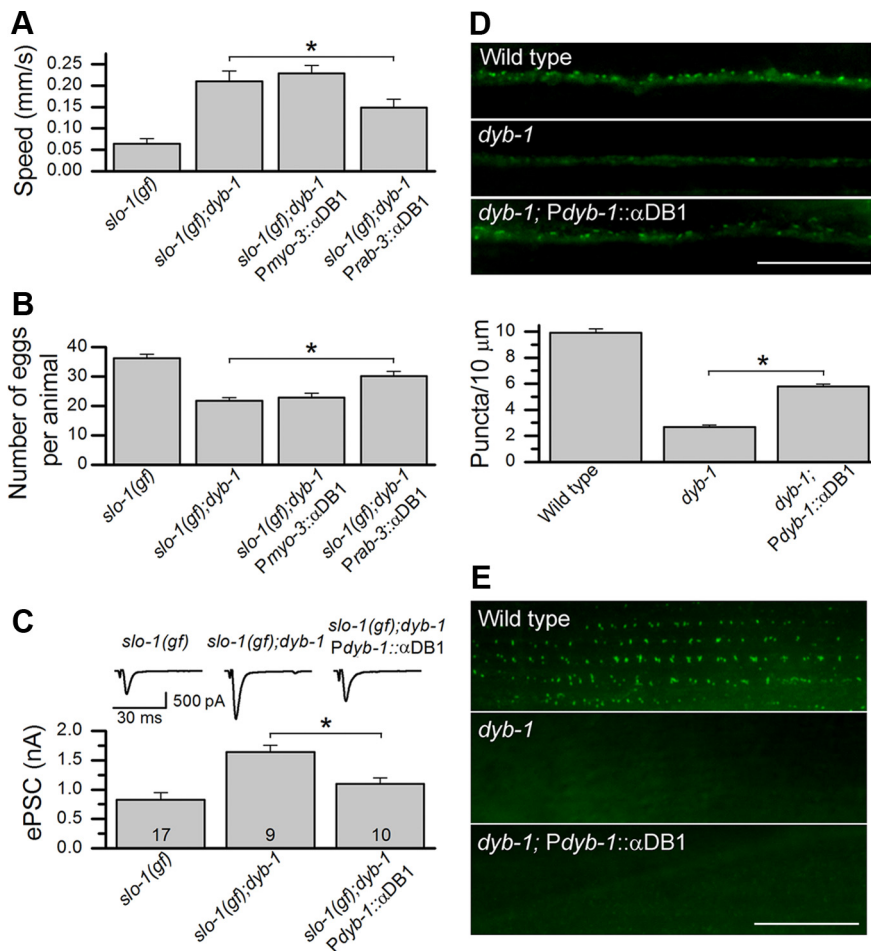


Figure 8. Mouse α DB1 partially rescued the neuronal phenotypes of *dyb-1* mutant. **A, B**, Expression of α DB1 in neurons (*Prab-3::\alpha*DB1) but not muscles (*Pmyo-3::\alpha*DB1) partially reinstated the defective locomotion (**A**) and egg-laying (**B**) phenotypes of *slo-1(gf)* in *slo-1(gf);dyb-1* mutant. The datasets for *slo-1(gf)* and *slo-1(gf);dyb-1* in Figures 3 were replotted here for comparison. **C**, Expression of α DB1 in *slo-1(gf);dyb-1* mutant under the control of *Pdyb-1* essentially restored the small ePSC amplitude of *slo-1(gf)*. The datasets for *slo-1(gf)* and *slo-1(gf);dyb-1* in Figure 4 were replotted here for comparison. The holding potential used for recording ePSCs was -60 mV. **D, E**, Expression of α DB1 under the control of *Pdyb-1* partially rescued SLO-1::GFP localization in the dorsal nerve cord but did not have an effect in body-wall muscle. * $p < 0.05$, significantly different between the indicated groups (one-way ANOVA with Bonferroni's *post hoc* test). Scale bars, $10 \mu\text{m}$.

Mouse α DB1 partially rescued *dyb-1* mutant phenotype

The relatively high level of sequence homology between DYB-1 and mammalian dystrobrevins (Fig. 1) raised the possibility that a mammalian dystrobrevin may be able to substitute DYB-1 with respect to SLO-1 function *in vivo*. Therefore, we analyzed the effect of mouse α DB1 on various phenotypes of the *dyb-1* mutant. Targeted expression of α DB1 in neurons but not muscle cells of the *slo-1(gf);dyb-1* double mutant essentially reinstated behavioral phenotypes of *slo-1(gf)*, including reduced locomotion speed and increased egg retention in the uterus (Fig. 8A,B). Expression of α DB1 in *slo-1(gf);dyb-1* double mutant also essentially restored the small ePSC amplitude of *slo-1(gf)* (Fig. 8C). Furthermore, expression of α DB1 under the control of *Pdyb-1* essentially rescued *dyb-1*-induced SLO-1::GFP mislocalization in the dorsal cord but had no effect in body-wall muscle (Fig. 8D,E). The differential rescuing effects of mouse α DB1 in neurons and muscle cells suggest that DYB-1 likely mediates SLO-1 localization by interacting with different proteins in neurons and muscle cells and that α DB1 may interact with the related protein(s) in neurons but not muscle cells.

Discussion

The present study shows that DYB-1 plays pivotal roles in controlling muscle Ca^{2+} transients and synaptic exocytosis in *C. elegans* by localizing the BK channel to subcellular domains containing a high density of voltage-gated Ca^{2+} channels. These observations are consistent with the previous reports that dystrobrevin is localized to axon terminals (Ueda et al., 2000) and that mutations of either dystrobrevin or the BK channel may contribute to synaptic defects and muscular dystrophy (Grady et al., 1999; Wang et al., 2001; Carre-Pierrat et al., 2006; Grady et al., 2006).

Ca^{2+} mishandling has been suggested as the basis of muscle degeneration in Duchenne muscular dystrophy (DMD) patients (Duncan, 1978). It is thought that sarcolemma damage or enhanced activity of Ca^{2+} leak channels leads to elevation of intracellular $[\text{Ca}^{2+}]$, which activates Ca^{2+} -dependent proteases such as calpains and eventually leads to cell death (Deconinck and Dan, 2007; Allen et al., 2010). However, measurements of Ca^{2+} concentration in skeletal muscle from DMD patients and *mdx* mice have produced controversial results (Constantin et al., 2006). Through Ca^{2+} imaging in live worms, we found that SLO-1 is a potent regulator of muscle Ca^{2+} transients in *C. elegans* body-wall muscle and that its mislocalization in *dyb-1* mutant led to a large increase in Ca^{2+} transient frequency. Given that SLO-1 colocalizes with the L-type voltage-gated Ca^{2+} channel EGL-19 at dense body regions in body-wall muscle (Kim et al., 2009), Ca^{2+} microdomains formed at the inner mouth of EGL-19 in response to action potentials likely activate colocalized SLO-1, which in turn terminates further Ca^{2+} entry by fa-

ilitating membrane repolarization. This function of SLO-1 in regulating muscle Ca^{2+} transients might be an important contributing factor to the muscular degenerative changes related to *slo-1*, *dyb-1*, and *dys-1* mutants.

Contraction of mammalian skeletal muscle requires the functions of both L-type voltage-gated Ca^{2+} channels in the sarcolemma and ryanodine receptors (RyRs) in the sarcoplasmic reticulum (SR) membrane (Rios and Brum, 1987; Takeshima et al., 1994), which is similar to *C. elegans* body-wall muscle (Liu et al., 2011). Although RyR-mediated Ca^{2+} release from the SR serves as the primary source of Ca^{2+} for skeletal muscle contraction, the activation of RyRs is coupled to depolarization-induced conformational changes of voltage-gated Ca^{2+} channels through protein/protein interactions occurring in t-tubules (Ríos et al., 1992; Schneider, 1994). Because the BK channel is also enriched or localized in t-tubules (Latorre et al., 1982; Knaus et al., 1995), whether or not it colocalizes with voltage-gated Ca^{2+} channels to regulate Ca^{2+} concentrations in mammalian skeletal muscle is worthy of investigation.

The similar locomotion and muscle Ca^{2+} transient phenotypes of *dyb-1* and *slo-1* mutants suggest that DYB-1 likely regulates muscle Ca^{2+} transients mainly through SLO-1. Nevertheless, we cannot exclude the potential involvement of other mechanisms. For example, Ca^{2+} mishandling in dystrophic mammalian skeletal muscle has been attributed to weakened membrane integrity and increased opening of other ion channels (e.g., store-operated cation channels and inositol triphosphate receptors) (Constantin et al., 2006). It would be interesting to determine whether these kinds of mechanisms are also implicated in the regulatory effect of DYB-1 on worm muscle Ca^{2+} transients.

The DAPC has also been implicated in synaptic transmission. For example, α DB1 is required for ACh receptor clustering at postsynaptic sites in mouse skeletal muscle (Pawlikowski and Maimone, 2008). In *C. elegans*, mutations of a DAPC component (DYS-1, DYB-1, or STN-1/syntrophin) result in increased cholinergic transmission as suggested by behavioral and pharmacological assays (Gieseler et al., 1999; Giugia et al., 1999). The present study shows that dystrobrevin controls neurotransmitter release by localizing SLO-1 to presynaptic sites, which potentially allows SLO-1 to be functionally coupled with UNC-2, the predominant voltage-gated Ca^{2+} channel at the *C. elegans* NMJ (Richmond et al., 2001; Saheki and Bargmann, 2009). Given that mouse α DB1 is also localized to axon terminals in mammals (Ueda et al., 2000) and may partially substitute DYB-1 in neurons with respect to SLO-1 function in *C. elegans*, it would be interesting to see whether dystrobrevins play a similar role in mammalian brain.

Dystrobrevins are mainly known to function as a component in the DAPC. However, there is evidence that dystrobrevins may also function independently of dystrophin. In mice lacking dystrophin, the assembly of the DAPC is disrupted in skeletal muscle but the dystrobrevin–syntrophin complexes are still formed in the brains (Blake et al., 1999). Although DYB-1 was required for SLO-1 localization in neurons as well as body-wall muscle cells, *DYS-1* was dispensable for normal SLO-1 localization in neurons, suggesting that, at least in neurons, DYB-1 localizes SLO-1 through a mechanism independent of *DYS-1*. This conclusion is consistent with the data showing that mouse α DB1 was able to partially rescue SLO-1 mislocalization in neurons but not muscle cells in *dyb-1* mutant. It remains to be determined what differences exist in the protein partners interacting with DYB-1 to mediate SLO-1 localization in neurons and body-wall muscle cells.

α DB and β DB are widely expressed in mammalian tissues. It has been suggested that mutations of dystrobrevins may underlie some behavioral and cognitive defects (Rees et al., 2007). However, relatively little is known about the functions of dystrobrevins. Patients with α DB deficiency may display severe congenital muscular dystrophy with ophthalmoplegia (Jones et al., 2003). Studies with mice have shown that knock-out of α DB causes mild muscular dystrophy and cardiomyopathy, abnormal distribution and reduced level of ACh receptors at the NMJs, and reduced level of neuronal nitric oxide synthase at the sarcolemma, and knock-out of β DB causes reduction in the number of GABA_A α 1 receptors in cerebellar Purkinje cells (Rees et al., 2007). Our analyses with *C. elegans* revealed new physiological functions of dystrobrevins and raised an interesting question as to whether dystrobrevin-dependent BK channel localization is involved in DAPC-associated diseases.

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