

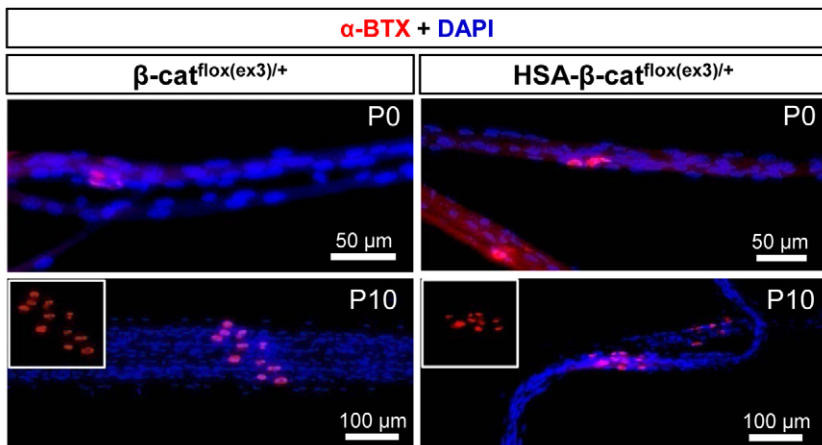
β -Catenin gain of function in muscles impairs neuromuscular junction formation

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In Fig. 5F, two of the panels (bottom left and top right) were misplaced and therefore attributed to the wrong genotype. The correct Fig. 5F appears below.

The authors apologise to readers for this mistake.

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β -Catenin gain of function in muscles impairs neuromuscular junction formation

Haitao Wu^{1,3,*}, Yisheng Lu^{1,*}, Arnab Barik¹, Anish Joseph¹, Makoto Mark Taketo⁴, Wen-Cheng Xiong^{1,2} and Lin Mei^{1,2,†}

SUMMARY

Neuromuscular junction (NMJ) formation requires proper interaction between motoneurons and muscle cells. β -Catenin is required in muscle cells for NMJ formation. To understand underlying mechanisms, we investigated the effect of β -catenin gain of function (GOF) on NMJ development. In HSA- β -cat^{flox(ex3)/+} mice, which express stable β -catenin specifically in muscles, motor nerve terminals became extensively defasciculated and arborized. Ectopic muscles were observed in the diaphragm and were innervated by ectopic phrenic nerve branches. Moreover, extensive outgrowth and branching of spinal axons were evident in the GOF mice. These results indicate that increased β -catenin in muscles alters presynaptic differentiation. Postsynaptically, AChR clusters in HSA- β -cat^{flox(ex3)/+} diaphragms were distributed in a wider region, suggesting that muscle β -catenin GOF disrupted the signal that restricts AChR clustering to the middle region of muscle fibers. Expression of stable β -catenin in motoneurons, however, had no effect on NMJ formation. These observations provide additional genetic evidence that pre- and postsynaptic development of the NMJ requires an intricate balance of β -catenin activity in muscles.

KEY WORDS: Neuromuscular junction, β -Catenin, AChR, Axon outgrowth, Retrograde signals, Mouse

INTRODUCTION

The neuromuscular junction (NMJ), a synapse between motoneurons and muscle fibers, is crucial for control of muscle contraction. This classic chemical synapse has served as an informative model of synapse function and synaptogenesis (Wu et al., 2010). NMJ formation is regulated by interactions between motoneurons and muscles (Sanes and Lichtman, 2001; Wu et al., 2010). Motoneurons release agrin, which binds directly to LRP4 and thus activates MuSK, both of which are required for NMJ formation (McMahan, 1990; DeChiara et al., 1996; Gautam et al., 1996; Glass et al., 1996; Weatherbee et al., 2006; Kim, N. et al., 2008; Zhang et al., 2008; Zong et al., 2012). By contrast, activation of muscles by acetylcholine (ACh) from motoneurons suppresses the expression of acetylcholine receptors (AChRs) and disassembles AChR clusters (Schaeffer et al., 2001; Misgeld et al., 2002; Brandon et al., 2003; Chen et al., 2007).

Less is known about retrograde signals from muscle fibers for presynaptic differentiation. Muscles are known to produce necessary factors for motoneurons (Hamburger, 1934). Several factors have been implicated in motoneuron survival or synapse elimination including glial cell line-derived neurotrophic factor (GDNF) and brain-derived neurotrophic factor (BDNF) (Oppenheim et al., 1995; Nguyen et al., 1998; Keller-Peck et al., 2001; Lu and Je, 2003). Transforming growth factor β (TGF β), fibroblast growth factor (FGF), laminin and collagen appear to play

a role in orchestrating presynaptic development at the NMJ (McCabe et al., 2003; Rawson et al., 2003; Nishimune et al., 2004; Fox et al., 2007; Feng and Ko, 2008; Nishimune et al., 2008). Previously, we demonstrated that phrenic nerve branches are mislocated and synaptic vesicle release is compromised in mice lacking β -catenin in muscles (Li et al., 2008), suggesting a necessary role of muscle β -catenin for presynaptic differentiation or function.

In this study, we investigated the GOF effects of β -catenin on NMJ formation. When a conditional GOF mutation of β -catenin was introduced by HSA Cre in muscle cells, both presynaptic and postsynaptic deficits were observed. Intriguingly, some phenotypes were similar whereas others were converse, compared with those in muscle-specific β -catenin loss-of-function (LOF) mutants. However, when the GOF mutation of β -catenin was introduced by HB9 Cre in motoneurons, it had no apparent effect on NMJ development. We studied the effects of muscle β -catenin by multiple approaches. Results provide additional genetic evidence that an intricate balance of muscle β -catenin activity is crucial for both pre- and postsynaptic differentiation during NMJ development.

MATERIALS AND METHODS

Generation and genotyping of mouse lines

β -Catenin^{flox(ex3)} loxP P85 (PGK reverse) [β -cat^{flox(ex3)}] mice harbor two loxP sites flanking exon 3 of the β -catenin gene (*Ctnnb1* – Mouse Genome Informatics) (Harada et al., 1999). This exon encodes a domain that contains crucial Ser/Thr residues phosphorylation of which promotes the degradation of β -catenin (Fig. 1A). β -Cat^{flox(ex3)} mice were crossed with human skeletal α -actin (HSA)-Cre and HB9-Cre mice, which express Cre specifically in skeletal muscles and motoneurons, respectively (Miniou et al., 1999; Yang et al., 2001; Li et al., 2008), to generate HSA- β -cat^{flox(ex3)/+} and HB9- β -cat^{flox(ex3)/+}. The Wnt signaling TOP-EGFP reporter mice were purchased from RIKEN BioResource Center (RBRC02229) (Moriyama et al., 2007) and crossed with β -cat^{flox(ex3)} and HSA-Cre mice to produce TOP-EGFP; β -cat^{flox(ex3)/+} and TOP-EGFP;HSA- β -cat^{flox(ex3)/+} mice.

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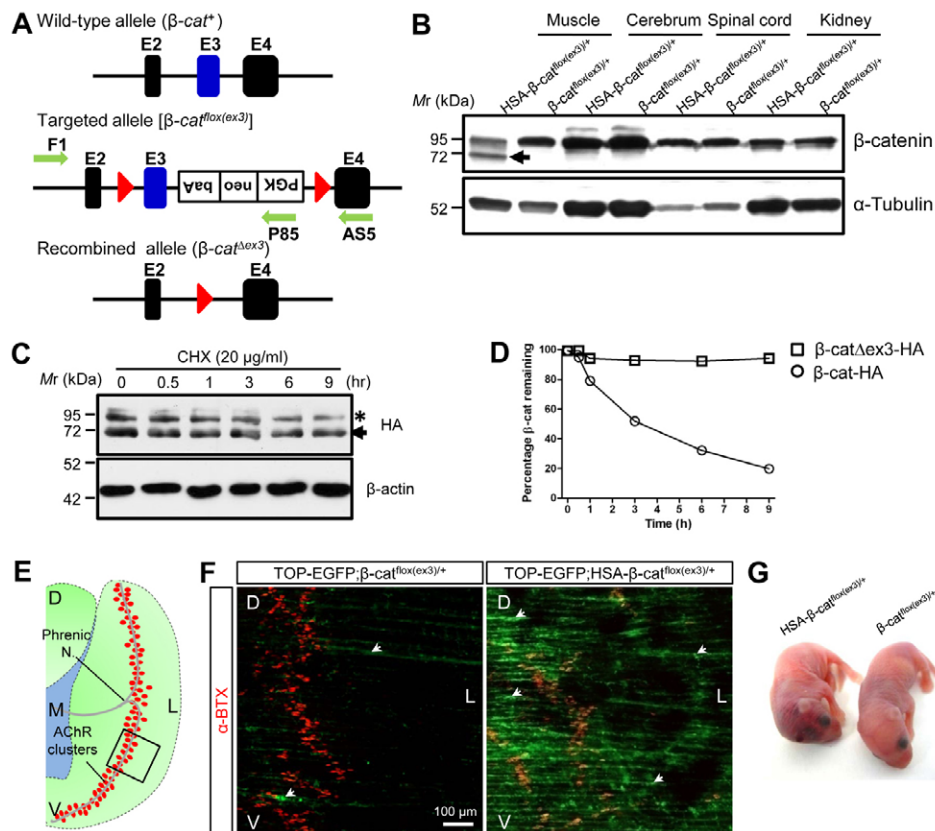


Fig. 1. Increased stability and activity of β-catenin in skeletal muscles in HSA-β-cat^{flox(ex3)}/+ mice. (A) Schematic of the wild-type and targeted β-catenin allele. E, exon; red triangle, loXp sequence; green arrows, primers for genotyping. (B) Specific expression of exon 3-deleted β-catenin (arrow) in skeletal muscles of HSA-β-cat^{flox(ex3)}/+ mice, but not in other tissues. (C) Increased stability of exon 3-deleted β-catenin compared with wild type. HEK293 cells were transfected with HA-tagged wild-type (asterisk) and mutant (arrow) β-catenin, and treated with 20 μg/ml CHX for various times. β-Actin was used as a loading control. (D) Quantification of data in C (mean ± SEM, n=3). (E) Schematic of the left hemi-diaphragm. Red dots, AChR clusters; rectangle, area shown in F; D, dorsal; V, ventral; L, lateral; M, medial. (F) Expression of TOP-EGFP in diaphragm muscles by β-catenin GOF mutant. Muscles of control (TOP-EGFP;β-cat^{flox(ex3)}/+) and TOP-EGFP;HSA-β-cat^{flox(ex3)}/+ mice were stained with Alexa Fluor 594-conjugated α-BTX (red). Image was taken by confocal fluorescence microscope. Area shown is indicated by rectangle in E. (G) Neonatal mice of indicated genotypes.

Genotyping was performed as described in the legend to supplementary material Fig. S1. Animal experimental procedures were approved by the Institutional Animal Care and Use Committee (IACUC) at the Georgia Health Sciences University.

Reagents, constructs and antibodies

Chemicals were purchased from Sigma-Aldrich unless otherwise indicated. Alexa Fluor 594-conjugated α-bungarotoxin (α-BTX) was purchased from Invitrogen (B-13423; 1:3000 for staining). HA-tagged mouse *Ctnnb1* (pKH3-β-catenin) was constructed as described previously (Kim, C. H. et al., 2008). *Ctnnb1* exon3 deletion mutant, β-catΔex3, was generated by using the Quick Change Site-Directed Mutagenesis Kit (Stratagene). The authenticity of all constructs was verified by DNA sequencing. Antibodies used were as follows [antigen (company, catalog number; dilution)]: neurofilament (NF) (Millipore, AB1991; 1:1000 for staining); synaptophysin (Dako, A0010; 1:2000 for staining); SV2 (Developmental Studies Hybridoma Bank, SV2; 1:500 for staining); β-catenin (BD Biosciences, 610154; 1:2000 for western); α-Tubulin (Santa Cruz, sc-23948; 1:3000 for western); β-actin (Novus, NB600-501; 1:3000 for western). Rabbit anti-rapsyn (2741) and anti-MuSK antibodies were described previously (1:1000 for western) (Luo et al., 2002). Antibodies raised against AChRα-subunit (mAb35) and β-subunit (mAb124) were gifts from Dr Richard Rotundo (Miller School of Medicine, Miami, FL, USA) and Dr Jon Lindstrom (Perelman School of Medicine, Philadelphia, PA, USA), respectively (1:1000 for western). Rabbit anti-HB9 (C-terminal 307-403) antibody was a gift from Dr Samuel Pfaff (The Salk Institute, La Jolla, CA, USA) (1:4000 for staining) (Thaler et al., 1999). Alexa Fluor 488-conjugated goat anti-rabbit IgG was purchased from Invitrogen (A-11034, 1:1000 for staining); horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (32260) and goat anti-mouse IgG (32230) antibodies were purchased from Pierce (Thermo Scientific) (1:3000 for western).

Western blotting

Western blotting was performed as described previously (Luo et al., 2002; Zhu et al., 2006).

Analysis of NMJ morphology and function

Whole-mount staining of muscles for AChR, nerve terminals and acetylcholinesterase (AChE); muscle section staining; analysis of NMJs in individual muscle fibers; and electrophysiological recordings were performed as described previously (Li et al., 2008).

Analysis of motoneuron survival

To study the effect of muscle β-catenin on motoneuron survival, spinal cords between C3 and C5 were dissected from postnatal day (P)0 mice, fixed in 4% paraformaldehyde in PBS (pH 7.3) overnight and immersed in 0.1 M phosphate buffer (pH 7.3) containing 30% sucrose for 24 hours. Tissues were embedded in OCT compound (TissueTek), and sectioned on a cryostat. Motoneurons were identified by staining spinal cord sections (14 μm) with anti-HB9 antibody as described previously (Arber et al., 1999). For quantification, HB9-positive motoneurons from hemiventral columns in every fourth section were counted by individuals blind to the genotypes.

Analysis of axons in brachial plexus

To examine axon projection from spinal cords, whole-mount staining was performed on embryos as previously described (Qiu et al., 1997). Briefly, embryos were fixed in methanol:DMSO (4:1) overnight at 4°C, bleached in methanol:DMSO:30% H₂O₂ (4:1:1) for 4-5 hours at 25°C, and rehydrated by incubation in 50% methanol, 15% methanol and PBS (30 minutes each). After being incubated in PBSMT (2% instant skim milk powder, 0.1% Triton X-100 in PBS) for 2 hours at 25°C, they were incubated with anti-NF antibody (1:1000) in PBSMT at 4°C overnight. Embryos were washed twice (1 hour each) in PBSMT at 4°C and three times at 25°C then incubated at 4°C with HRP-conjugated goat anti-mouse IgG (Thermo Fisher, 32230; 1:1000) in PBSMT. After washing, embryos were incubated at 25°C in PBT (0.2% BSA, 0.1% Triton X-100 in PBS) for 20 minutes and in PBT containing 0.3 mg/ml diaminobenzidine tetrahydrochloride (DAB) for 30-60 minutes. Nerve axons were visualized by addition of 0.0003% H₂O₂. Embryos were then rinsed in PBT to stop the reaction, dehydrated through a methanol series (30%, 50%, 80% and

100% for 30-60 minutes each) and cleared in benzyl alcohol:benzyl benzoate (1:2). They were photographed using a Zeiss stemi 200 dissecting microscope (Carl Zeiss).

Electron microscopy

Muscles were fixed in 2% glutaraldehyde and 2% paraformaldehyde in 0.1 M phosphate buffer for 1 hour at 25°C and further fixed in sodium cacodylate-buffered (pH 7.3) 1% osmium tetroxide for 1 hour at 25°C. After washing three times with phosphate buffer (10 minutes each) tissues were dehydrated through a series of ethanol: 30%, 50%, 70%, 80%, 90%, 100%; the 100% ethanol step was followed with three changes of 100% propylene oxide and embedded in plastic resin (EM-bed 812, EM Sciences). Serial thick sections (1-2 μ m) of tissue blocks were stained with 1% Toluidine Blue for light microscopy, from which sections with phrenic nerves were identified and cut into ultra-thin sections. They were mounted on 200-mesh unsupported copper grids and stained with uranyl acetate (3% in 50% methanol) and lead citrate (2.6% lead nitrate and 3.5% sodium citrate, pH 12.0). Electron micrographs were taken using a JEOL 100CXII operated at 80 KeV.

Electrophysiology

Neonatal mice diaphragm with ribs and intact phrenic nerves were dissected in oxygenated (95% O₂/5% CO₂) Ringer's solution (136.8 mM NaCl, 5 mM KCl, 12 mM NaHCO₃, 1 mM NaH₂PO₄, 1 mM MgCl₂, 2 mM CaCl₂ and 11 mM D-glucose, pH 7.3), and pinned on Sylgard gel in a dish perfused with oxygenated Ringer's solution (Li et al., 2008). Microelectrodes, 20-50 M Ω when filled with 3 M KCl, were pierced into the center of muscle fibers. To elicit endplate potential (EPP), phrenic nerves were sucked into a suction electrode and stimulated (0.1 msecond, 2-5 V) with the resting potential between -45 and -55 mV (Liu et al., 2008). Data were collected with Axopatch 200B amplifier (Molecular Devices, Sunnyvale, CA, USA), digitized with Digidata 1322A (Molecular Devices) and analyzed using pClamp 9.2 (Molecular Devices).

Quantitative RT-PCR analysis

Quantitative RT-PCR analysis was performed as described previously (Liu et al., 2011). Primers used for analysis are listed in supplementary material Table S1.

Statistical analysis

Data are presented as mean \pm s.e.m. and analyzed using Student's *t*-test or one-way ANOVA analysis, as appropriate. In the hypertonic sucrose-evoked mEPP experiment, the variability of frequency was analyzed using repeated measures ANOVA analysis. Graphs were generated by GraphPad Prism 5.0 software. *P*<0.05 was considered to be significant.

RESULTS

Expression of stable β -catenin in skeletal muscles

To study the effect of β -catenin GOF in skeletal muscles on NMJ formation, we introduced a conditional mutation of stable β -catenin in muscle cells by crossing HSA-Cre mice with β -cat^{flox(ex3)} mice, in which exon 3 was floxed (Harada et al., 1999) (Fig. 1A). Resulting HSA- β -cat^{flox(ex3)/+} mice express a β -catenin mutant that lacks the critical phosphorylation residues encoded by exon 3 in skeletal muscles, but not in other tissues, including spinal cord (Fig. 1B). The mutant β -catenin is believed to be resistant to degradation mediated by GSK3 β -dependent phosphorylation and is able to activate constitutively Wnt canonical signaling (Rubinfeld et al., 1997; Iwao et al., 1998). In agreement, we found that exon3-deleted β -catenin was more stable compared with wild-type β -catenin in cells treated with cycloheximide (CHX) (Fig. 1C,D). To demonstrate that β -catenin-dependent transcription activity was increased in HSA- β -cat^{flox(ex3)/+} muscles, we crossed them with TOP-EGFP mice, a reporter line of Wnt canonical signaling (Moriyama et al., 2007). As shown in Fig. 1F, EGFP was increased in muscles of TOP-EGFP;HSA- β -cat^{flox(ex3)/+} mice, compared with control mice, indicating the GOF effect of β -catenin in HSA- β -

cat^{flox(ex3)/+} mice. Unless otherwise indicated, β -cat^{flox(ex3)/+} littermates were used as control in the present study. They were viable and fertile and showed no difference in life span, general behavior, or NMJ morphology and function, compared with wild-type controls (data not shown). HSA- β -cat^{flox(ex3)/+} mice were also viable at birth, but they were smaller in body size compared with control littermates (Fig. 1G), and appeared to have subcutaneous hemorrhage at the posterior fontanel site of the skull (Fig. 1G).

β -Catenin is localized on the muscle membrane without specific enrichment at the NMJ in both control and HSA- β -cat^{flox(ex3)/+} mice, except at higher levels in the latter (supplementary material Fig. S1). β -Catenin GOF seemed to have no effect on diaphragm thickness and the cross-sectional area of muscle fibers (supplementary material Fig. S2A-C). Moreover, the nuclei of muscle fibers in HSA- β -cat^{flox(ex3)/+} mice were located beneath the membrane, not in the central region, suggesting that muscle fibers were not regenerated (supplementary material Fig. S2A). Finally, electron microscopic studies indicated that muscle contractile units in HSA- β -cat^{flox(ex3)/+} mice were similar, compared with controls (supplementary material Fig. S2D). These results suggest that muscle β -catenin GOF did not change the overall structure of muscle fibers. Nevertheless, to avoid potential long-term effect on muscle health, we focused on muscles of P0 neonatal mice.

Defasciculation and arborization of axons in HSA- β -cat^{flox(ex3)/+} diaphragms

Whole-mount diaphragms were stained with a mixture of antibodies against neurofilament (NF) and synaptophysin to label both nerve branches and terminals (Li et al., 2008). In control diaphragms, primary phrenic branches traveled in the middle region of muscle fibers and sent out numerous, but short, secondary or intramuscular branches that form NMJs close the primary branches (Fig. 2A). In HSA- β -cat^{flox(ex3)/+} diaphragms, however, the nerve branches were extensively defasciculated, which generated a net of axons covering a broader central region (Fig. 2A). The length of secondary branches was increased but their number was reduced in HSA- β -cat^{flox(ex3)/+} mice (Fig. 2B,C). By contrast, tertiary and quaternary branches, which were few and short in control diaphragms, were apparent and long in HSA- β -cat^{flox(ex3)/+} mice. The 5th branches, which did not exist in control diaphragms, were observable in HSA- β -cat^{flox(ex3)/+} diaphragms (Fig. 2C). Similar deficits were observed in diaphragms of HSA- β -cat^{flox(ex3)/flox(ex3)} homozygous mice (supplementary material Fig. S3).

To determine when the arborization of phrenic nerve terminals occurred, we examined diaphragms at different embryonic ages. At embryonic day (E)13, the primary branches were located in the middle region of muscle fibers in HSA- β -cat^{flox(ex3)/+} embryos, suggesting no problem in axon projection. Compared with controls, in which secondary branches were hardly detectable, branches were readily visible in HSA- β -cat^{flox(ex3)/+} diaphragms (Fig. 2D, arrowheads). The length of secondary branches and the endplate band width continued to increase in both control and HSA- β -cat^{flox(ex3)/+} mice during development, but the rate was significantly faster for the latter (Fig. 2E,G). The number of secondary branches increased more rapidly in control than in HSA- β -cat^{flox(ex3)/+} embryos (Fig. 2F). These results demonstrate that increased β -catenin activity in muscle fibers leads to arborization or defasciculation of phrenic nerve terminals. Similar phenotypes, including wider central endplate band and axon arborization, were observed in intercostal muscles and tibialis anterior muscles in HSA- β -cat^{flox(ex3)/+} mice (supplementary material Fig. S4).

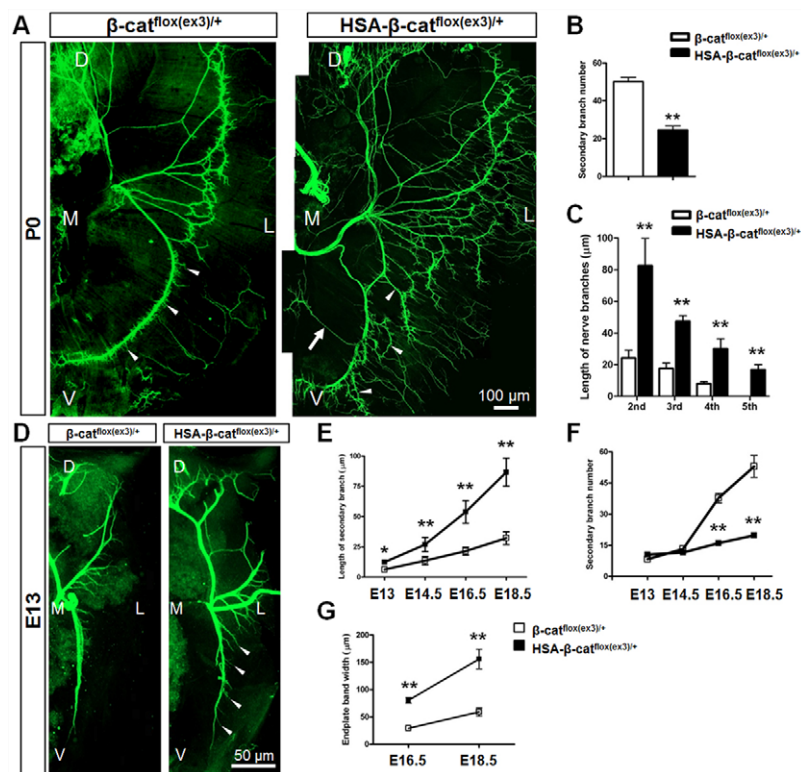


Fig. 2. Aberrant innervation of motor axons in HSA-β-cat^{fllox(ex3)/+} mice. (A) P0 left hemi-diaphragms of indicated genotypes. Phrenic nerves and terminals were stained with anti-NF/synaptophysin antibodies, which were visualized with Alexa Fluor 488-conjugated goat anti-rabbit antibodies. (B) Decreased number of secondary/intramuscular nerve branches in HSA-β-cat^{fllox(ex3)/+} muscles (***P*<0.01, *n*=7, *t*-test). (C) Increased length of secondary, tertiary, quaternary and 5th branches in HSA-β-cat^{fllox(ex3)/+} muscles (***P*<0.01, *n*=6, one-way ANOVA). (D) E13 left hemi-diaphragms of indicated genotypes. (E) Increased secondary branch length in developing HSA-β-cat^{fllox(ex3)/+} embryos (**P*<0.05, ***P*<0.01, *n*=5, one-way ANOVA). (F) Developmental changes of secondary branches in HSA-β-cat^{fllox(ex3)/+} embryos (***P*<0.01, *n*=5, one-way ANOVA). (G) Increased endplate band width in E16.5 and E18.5 HSA-β-cat^{fllox(ex3)/+} embryos (***P*<0.01, *n*=10, one-way ANOVA). In A and D, arrowheads indicate secondary nerve branches; arrow indicates ectopic axon. D, dorsal; V, ventral; L, lateral; M, medial. Error bars indicate s.e.m.

Formation and innervation of ectopic muscles in HSA-β-cat^{fllox(ex3)/+} mice

Although the Wnt/β-catenin-dependent pathway has been implicated in muscle development (Borello et al., 1999; Parker et al., 2003; Perez-Ruiz et al., 2008; Gros et al., 2009), the *in vivo* effect of muscle β-catenin GOF remained unclear. Intriguingly, new muscles were found in the central tendoneous region of HSA-β-cat^{fllox(ex3)/+} diaphragms, a region that usually is clean and transparent in wild-type and control mice. Quantitatively, HSA-β-

cat^{fllox(ex3)/+} mice formed at least one, and as many as three to four, ectopic muscles (Fig. 3D). These results indicate that β-catenin expression under the control of the HSA promoter promotes the formation of ectopic muscles in tendoneous regions.

The ectopic muscles were innervated by ectopic branches of phrenic nerves (Fig. 2A, Fig. 3A). In developing diaphragms, phrenic nerves send branches towards the ribs and rarely (if any) towards the central cavity. However, axon branches were evident towards the central cavity in HSA-β-cat^{fllox(ex3)/+} diaphragms as

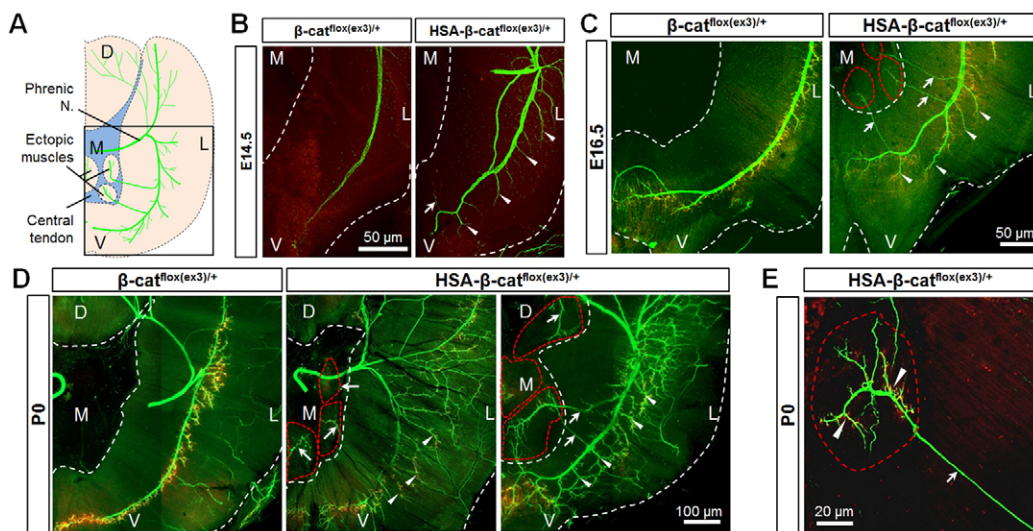


Fig. 3. Formation and innervation of ectopic muscles in HSA-β-cat^{fllox(ex3)/+} mice. (A) Schematic of left hemi-diaphragm with ectopic muscles. Square frame indicates parts of diaphragms analyzed in B-D. D, dorsal; V, ventral; M, medial; L, left. (B-D) Staining of diaphragms of different genotypes at indicated ages (NF/synaptophysin, green; AChR, red). White dashed lines indicate the edge of diaphragms. Red dashed lines encircle ectopic muscles. Arrowheads indicate extensive and longer secondary branches. Arrows indicate secondary nerve branches. (E) Enlarged image of an ectopic muscle. Arrow, ectopic axons; arrowheads, AChR clusters.

early as E14.5, when ectopic muscle mass was barely visible in the central tendoneous region (Fig. 3B). By E16.5, the overshooting phrenic axons arrived at ectopic muscles in the central tendoneous region (Fig. 3C). At birth, the terminals of ectopic branches were fully arborized and formed NMJs (AChR clusters) on ectopic muscle fibers (Fig. 3D,E). These results demonstrate that ectopic muscles in the central tendoneous region were able to attract phrenic nerves to form ectopic branches for innervation, suggesting the existence of an axon attraction signal by muscle β -catenin.

Axon attraction activity from muscles expressing stable β -catenin

To determine whether muscle β -catenin alters axon extension, we examined the primary ventral phrenic branch of developing left hemi-diaphragms, a branch with better spatial resolution. As shown in Fig. 2D and Fig. 3B, it extended more towards the ventral region in E13 and E14.5 HSA- β -cat^{flox(ex3)/+} mice compared with controls (Fig. 2D, Fig. 3B), suggesting that increased β -catenin activity in muscles produces a signal that promotes motor axon extension or outgrowth. To examine this hypothesis further, we looked at nerve projections from early embryonic spinal cords. In brachial plexus of control embryos with 45 somites, radial and axillary nerves (RN and AN, respectively), but not ulnar nerves (UN), were visible (Qiu et al., 1997; Arber et al., 1999; Vickerman et al., 2011) (Fig. 4A,B).

By comparison, RN and AN were more prominent and UN were visible in HSA- β -cat^{flox(ex3)/+} embryos. RN terminals appeared to defasciculate prematurely and extend further in the mutant (Fig. 4B). Immature branching and extension of RN axons were more obvious in embryos with 54 somites (Fig. 4C). Quantitatively, the number of tertiary branch points and lengths of RN branches were increased in embryos with 54 somites (Fig. 4D-F). These results indicate that β -catenin GOF in muscle fibers promotes the outgrowth of spinal nerve axons in HSA- β -cat^{flox(ex3)/+} mice.

Motoneurons are known to be regulated by signals from muscle cells (Oppenheim, 1991). To determine whether muscle β -catenin GOF alters motoneurons, we looked at spinal cords at cervical levels (C3-C5), where phrenic nerves arise from. Spinal cord sections were stained for HB9, a motoneuron-specific homeodomain transcription factor (Saha et al., 1997; Arber et al., 1999; Thaler et al., 1999). As shown in supplementary material Fig. S5A, HB9-positive motoneurons were located in the hemiventral segment of the spinal cord (Arber et al., 1999) (supplementary material Fig. S5A). The numbers of motoneurons in this segment were similar between control and HSA- β -cat^{flox(ex3)/+} mice (supplementary material Fig. S5B). HB9-positive motoneurons of phrenic nerves were primarily located in the lateral and medial portions of the anterior horn [termed the lateral motor column (LMC) and medial motor column (MMC), respectively]

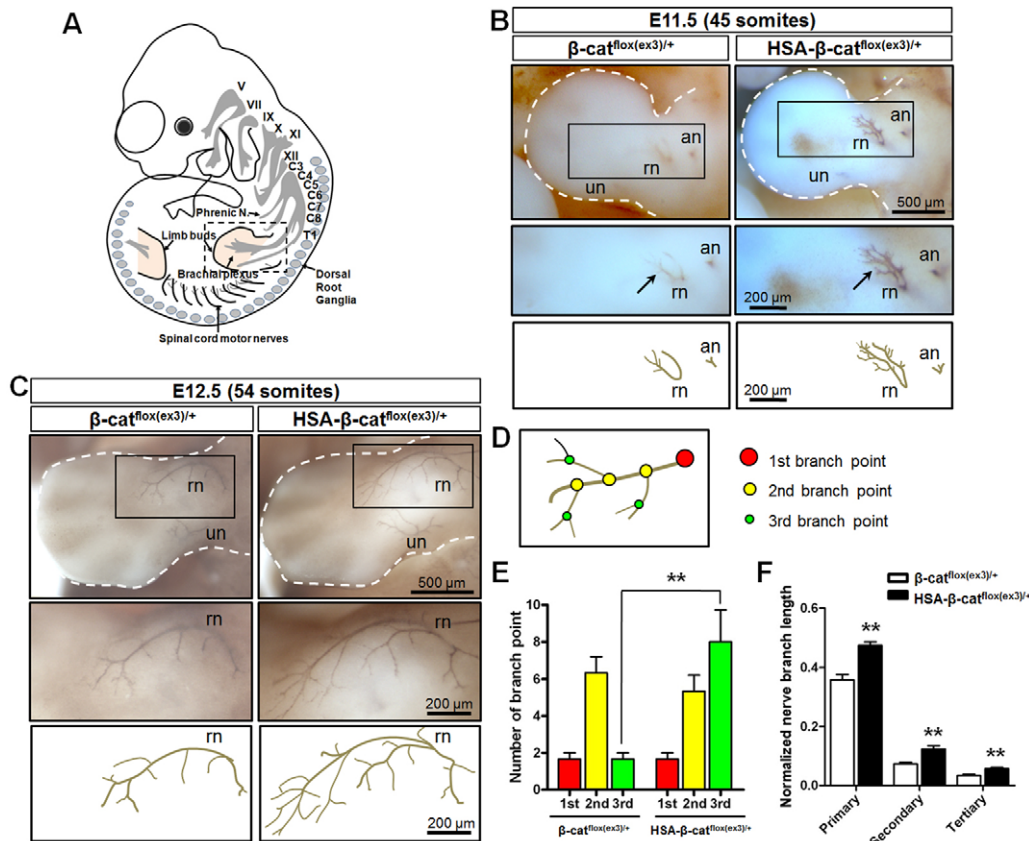


Fig. 4. Augmented extension and branching of brachial plexuses in HSA- β -cat^{flox(ex3)/+} embryos. (A) Schematic lateral view of mouse embryo. Roman numerals indicate cranial nerves. C, cervical; T, thoracic. (B) Embryos at 45-somite stage stained whole-mount with anti-NF antibody, which was visualized with diaminobenzidine (DAB). Arrows indicate brachial plexus. Areas in rectangles in top panels are enlarged in middle panels. Lower panels show camera lucida drawing of axons. an, axillary nerve; rn, radial nerve; un, ulnar nerve. (C) Embryos at 54-somite stage stained as described in B. Areas in rectangles in top panels are enlarged in middle panels. Lower panels show camera lucida drawing of axons. (D) Schematic of axon branches. Color matches data shown in E. (E) Increased numbers of rn tertiary branch points in HSA- β -cat^{flox(ex3)/+} embryos shown in C. ** $P < 0.01$, $n = 3$, one-way ANOVA. (F) Increased length of rn branches in HSA- β -cat^{flox(ex3)/+} embryos shown in C. Shown are ratios of rn branch length over limb bud length. ** $P < 0.01$, $n = 3$, one-way ANOVA. Error bars indicate s.e.m.

(supplementary material Fig. S5A) (Polleux et al., 2007). Muscle β-catenin GOF did not change the numbers of motoneurons in these two subpopulations (supplementary material Fig. S5C,D). These observations suggest that enhanced β-catenin activity in muscles alters presynaptic differentiation without apparent effect on motoneuron number and distribution.

Postjunctional deficits in HSA-β-cat^{flox(ex3)/+} NMJs

To determine whether aberrant expression of stable β-catenin alters the postsynaptic differentiation, diaphragms were whole-mount stained with α-BTX. In advance of innervation, muscle fibers are ‘pre-patterned’ with aneural AChR clusters that are distributed in the central region and form a band perpendicular to muscle fibers of diaphragms (Lin et al., 2001; Yang et al., 2001). Muscle fiber pre-patterning in E13.5 HSA-β-cat^{flox(ex3)/+} diaphragms was similar to that of littermate controls, suggesting that expression of stable β-catenin does not alter the formation and localization of aneural AChR clusters (data not shown). However, the endplate band width in HSA-β-cat^{flox(ex3)/+} embryos became significantly larger than that in controls as early as E16.5 (Fig. 2G, Fig. 3C). In P0 control diaphragms, almost all AChR clusters were restricted to the central region, forming a narrow endplate band width. However, in HSA-β-cat^{flox(ex3)/+} diaphragms, the band width is 2.6-fold larger (Fig. 5A,B). No change was observed in the areas or length of individual clusters (Fig. 5C,D). Moreover, the total number of AChR clusters was similar between HSA-β-cat^{flox(ex3)/+} and control diaphragms (Fig. 5E), indicating expression of stable β-catenin had no effect on the number of NMJs, but altered their location. In addition, we

also determined the number of AChR clusters per muscle fiber and found no difference between the two genotypes (Fig. 5F,G). AChE is accumulated in the synaptic cleft via a distinct mechanism (Arikawa-Hirasawa et al., 2002). Its distribution, assayed by in situ enzymatic activity, was also in a broader central region in HSA-β-cat^{flox(ex3)/+} mice than in control littermates (Fig. 5H). Together, these observations indicate that expression of stable β-catenin in developing muscles alters the location where NMJs form, but does not change the size or the number of AChR clusters per muscle fiber.

Impairment of synaptic transmission at HSA-β-cat^{flox(ex3)/+} NMJs

To determine whether neurotransmission was altered at HSA-β-cat^{flox(ex3)/+} NMJs, we measured miniature endplate potentials (mEPPs), events generated by spontaneous vesicle release. The amplitudes of mEPPs were similar in control and HSA-β-cat^{flox(ex3)/+} mice (2.74±0.42 mV and 2.50±0.13 mV, respectively; *P*>0.05, *n*=6 mice) (Fig. 6A,C), suggesting that AChR density was not altered. This observation is in agreement with results of light microscopic characterization that AChR clusters were apparently normal (Fig. 5C-G) and biochemical studies that expression of postsynaptic proteins was normal in HSA-β-cat^{flox(ex3)/+} muscles, including AChRα- and β-subunits, MuSK and rapsyn (Fig. 6F-J). By contrast, mEPP frequencies were markedly reduced in HSA-β-cat^{flox(ex3)/+} NMJs (Fig. 6B), suggesting potential defects in spontaneous ACh release from presynaptic terminals. Presynaptic vesicle release deficiency might result from a decrease in ready

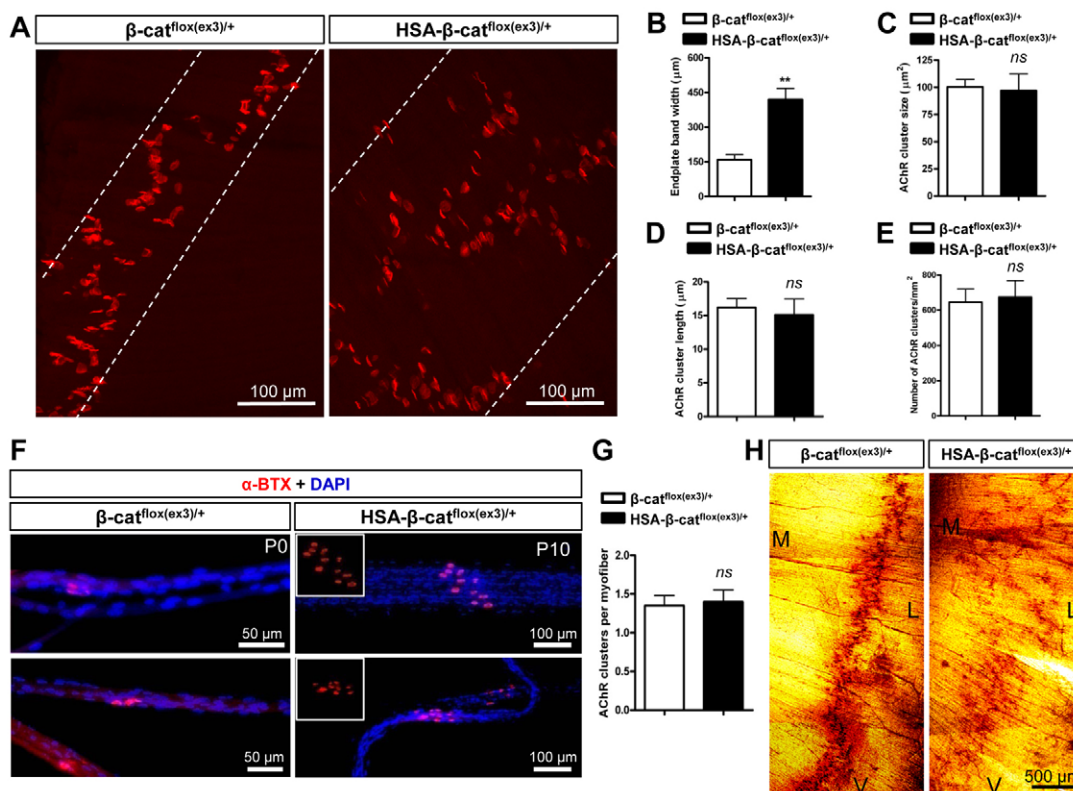


Fig. 5. Postsynaptic deficits of HSA-β-cat^{flox(ex3)/+} NMJs. (A) P0, left hemi-diaphragms were stained for AChR. White dashed lines were drawn to include most AChR clusters. (B) Increased endplate band width in HSA-β-cat^{flox(ex3)/+} diaphragms (***P*<0.01, *n*=6, *t*-test). (C-F) No change in AChR cluster size (C), length (D) or density (E) in HSA-β-cat^{flox(ex3)/+} diaphragms (*P*>0.05, *n*≥5, *t*-test). (F) Individual muscle fibers with AChR clusters (red). Nuclei were stained with DAPI (blue). (G) Quantitative analysis of data shown in F (*P*>0.05, *n*=22, *t*-test). (H) Scattered AChE clusters in HSA-β-cat^{flox(ex3)/+} diaphragms (P0). V, ventral; L, lateral; M, medial. Error bars indicate s.e.m.

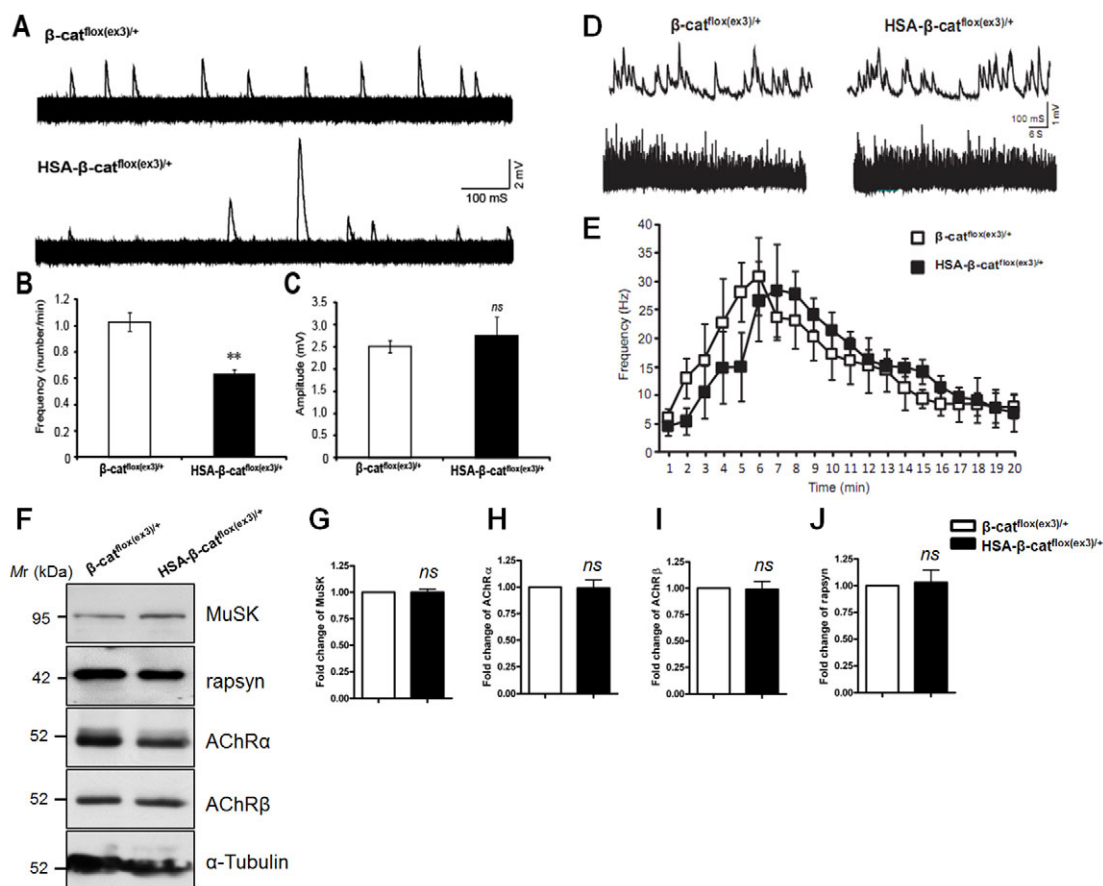


Fig. 6. Impaired synaptic transmission in HSA- β -cat^{flox(ex3)/+} NMJs. (A) Representative superimposed mEPP sample traces. Traces were recorded from neonatal mice at 2 mM Ca²⁺ (1-second \times 600 traces). (B) Reduced mEPP frequency in HSA- β -cat^{flox(ex3)/+} NMJs (** P <0.01, n =6, t -test). (C) mEPP amplitude was not changed at mutant NMJs (P >0.05, n =6, t -test). (D) Representative mEPP traces in response to 0.5 M sucrose. Shown are traces 7 minutes after the addition of sucrose. Time scale: 1 second upper panel; 1 minute lower panel. (E) Quantitative analysis of hypertonic evoked mEPP frequency shown in D (P >0.05, n =5, repeat measure). (F–J) No change in synaptic proteins in HSA- β -cat^{flox(ex3)/+} muscle lysates. Data are mean \pm s.e.m. (P >0.05, n =3 each group, t -test).

release pool (RRP) or a change in vesicle fusion probability. To address these questions, we determined whether RRP was altered by recording mEPPs in response to hypertonic stimulation with 0.5 M sucrose, which causes rapid release of vesicles in the RRP (Stevens and Tsujimoto, 1995). As shown in Fig. 6D,E, there was no difference in the frequencies of sucrose-evoked mEPPs between the two genotypes (Fig. 6D,E), indicating that the size of RRP was not altered in HSA- β -cat^{flox(ex3)/+} mice. Endplate potentials (EPPs) were similar between the genotypes (data not shown), suggesting normal vesicle release in response to action potentials.

Altered NMJ structures in muscle β -catenin LOF, but not GOF, mice

The reduction in mEPP frequency and apparently normal RRP suggest that the reduced ACh release might be due to functional, but not structural, deficits. To test this hypothesis, we determined expression of SV2, a presynaptic vesicle protein, in HSA- β -cat^{flox(ex3)/+} NMJs. SV2 staining was in good registration with AChR clusters (supplementary material Fig. S6A). The areas of AChR clusters covered by SV2 were similar in control and HSA- β -cat^{flox(ex3)/+} NMJs (supplementary material Fig. S6B). These results suggest that presynaptic specialization is normal in HSA- β -cat^{flox(ex3)/+} NMJs. Next, we examined NMJ structures by electron

microscopic analysis. As for NMJs in control littermates, axon terminals in HSA- β -cat^{flox(ex3)/+} skeletal muscles were capped by processes of perisynaptic Schwann cells; they were filled with abundant synaptic vesicles (Fig. 7A). Quantitative studies revealed no difference between control and HSA- β -cat^{flox(ex3)/+} NMJs in numbers of nerve terminals, numbers of active zones, vesicle densities, and synaptic vesicle diameters (Fig. 7B–E). These results demonstrate that the presynaptic structures of HSA- β -cat^{flox(ex3)/+} NMJs were normal. Moreover, synaptic clefts had similar width between the two genotypes (Fig. 7F) and were both filled with synaptic basal lamina (Fig. 7A). On the postsynaptic side, muscle membrane folds were observable, and there was no difference in the average length of junctional folds between control and HSA- β -cat^{flox(ex3)/+} (data not shown). Together, these results indicate that NMJ structures were largely normal in HSA- β -cat^{flox(ex3)/+} mice, in support of the functional deficit hypothesis.

Interestingly, mEPP frequency was reduced in muscle-specific β -catenin mutant (HSA- β -cat^{-/-}) mice (Li et al., 2008). Does β -catenin LOF regulate neurotransmission by a similar mechanism as GOF? Unlike control (β -cat^{flox/flox}) terminals, which were filled with abundant SVs, HSA- β -cat^{-/-} terminals contained sparsely populated SVs (Fig. 7G). Quantitatively, SV densities and active zones were decreased in HSA- β -cat^{-/-} mutant compared with

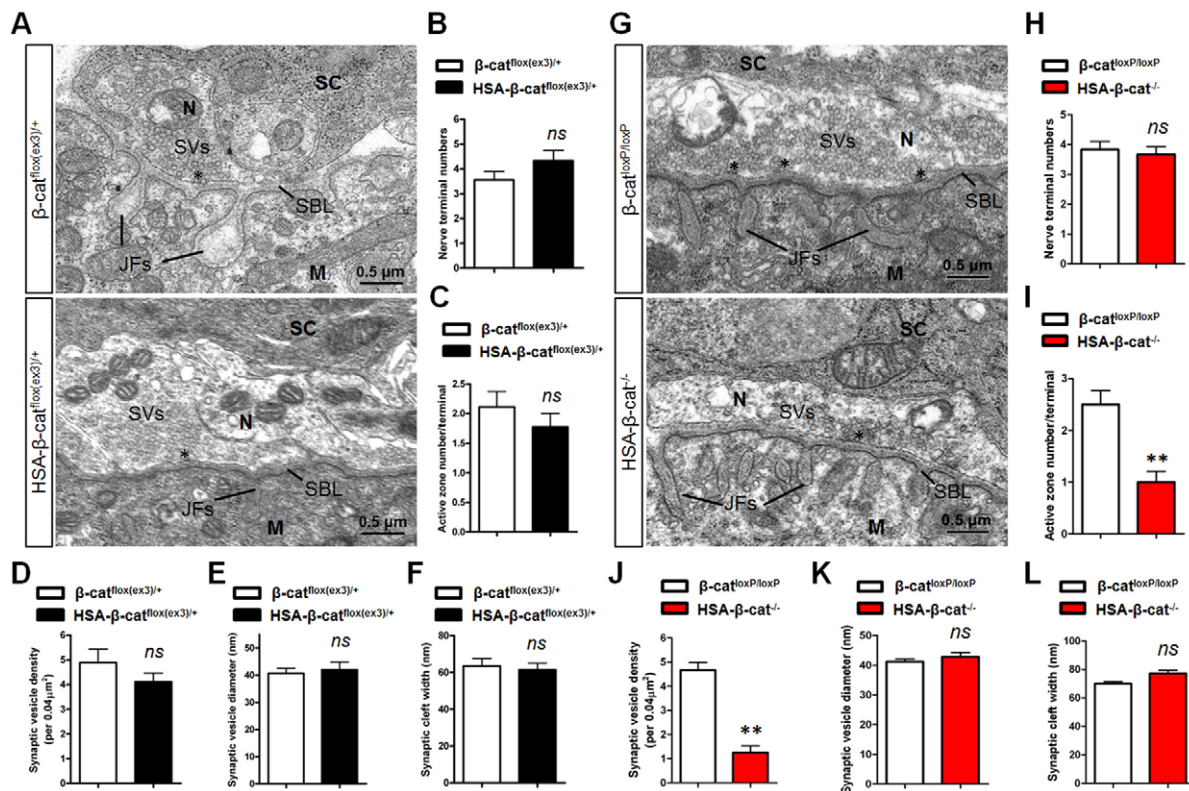


Fig. 7. Reduced number of synaptic vesicles and active zones in muscle β-catenin LOF, but not GOF, NMJs. (A) Representative electron micrographic images of NMJs in β-cat^{fllox(ex3)/+} control (upper panel) and HSA-β-cat^{fllox(ex3)/+} (lower panel) mice. N, nerve terminal; M, muscle fiber; SC, Schwann cell; SVs, synaptic vesicles; SBL, synaptic basal lamina; JFs, junctional folds; Asterisks mark active zones. (B-F) Quantitative analysis showed no difference in nerve terminal numbers (B), active zone numbers per nerve terminal (C), synaptic vesicle density (D), synaptic vesicle diameter (E) or synaptic cleft width (F) ($P > 0.05$, $n = 10$, t -test). (G) Representative electron micrographic images of NMJs in β-cat^{flloxPloxP} control (upper panel) and HSA-β-cat^{-/-} (lower panel) mice. (H-L) Quantitative data are shown for nerve terminal numbers (H), synaptic vesicle diameter (K), synaptic cleft width (L), active zone numbers per nerve terminal (I) and synaptic vesicle density (J) (** $P < 0.01$, $n = 10$, t -test). Error bars indicate s.e.m.

controls, although the numbers of nerve terminals were similar between the two genotypes (Fig. 7H-J). However, no difference was observed in SV diameters, synaptic cleft width (Fig. 7K,L) or junctional folds (data not shown). These results indicate that β-catenin LOF impairs NMJ development, in particular presynaptic differentiation, and provides a mechanism of compromised neurotransmission. Together with results from studies of GOF NMJs, they indicate distinct mechanisms by which β-catenin LOF and GOF regulate NMJ development.

Changes in gene transcription in HSA-β-cat^{fllox(ex3)/+} muscles

Mechanisms by which muscle β-catenin regulates NMJ development could be complex (see Discussion for details). It is possible that some presynaptic deficits occur because of overexpression of a factor(s) by increased β-catenin activity in muscles (Fig. 1F). In an initial step towards its identification, quantitative RT-PCR was performed. Fig. 8 shows increased expression of genes that are known to be β-catenin targets in GOF muscles, including *Axin2*, cyclin D1 (*Ccnd1* – Mouse Genome Informatics), *Myc*, *Lef1* and *Dkk1* (He et al., 1998; Tetsu and McCormick, 1999; Hovanes et al., 2001; Yan et al., 2001). This result confirmed the increased β-catenin activity in muscles, as indicated by TOP-EGFP (Fig. 1F). Next, we compared the expression of morphogens, neurotrophic factors and axon guidance

molecules between β-catenin GOF muscles and controls. Interestingly, several were found to be increased, including sonic hedgehog (*Shh*), *Wnt7a*, *Fgf20*, *Gdnf*, *Hgf* and *reelin*, whereas others were reduced, including *Bmp9* (*Gdf2* – Mouse Genome Informatics), *Wnt1*, *Wnt7b* and *Nt3* (*Ntf3* – Mouse Genome Informatics) (Fig. 8). These results are in agreement with the hypothesis that β-catenin GOF alters expression of various secreted factors to regulate motoneuron differentiation.

Normal NMJ development in motoneuron β-catenin GOF mice

To investigate whether enhanced levels of stable β-catenin in motoneurons alter NMJ development, HB9-β-cat^{fllox(ex3)/+} mice were generated by crossing β-cat^{fllox(ex3)/+} mice with HB9-Cre mice that specifically express Cre in developing spinal cords from E9.5 (Arber et al., 1999; Li et al., 2008). HB9-β-cat^{fllox(ex3)/+} mice expressed the truncated β-catenin specifically in the spinal cord, but not in other tissues, including muscles and the cerebral cortex (supplementary material Fig. S7A). However, HB9-β-cat^{fllox(ex3)/+} mice were viable at birth and normal in body size during development to adult. NMJs in HB9-β-cat^{fllox(ex3)/+} mice did not show any observable abnormality in phrenic nerves, including primary branch location or secondary branch number and length (supplementary material Fig. S7B-E). No difference was observed in the number or size of AChR clusters, or endplate band width

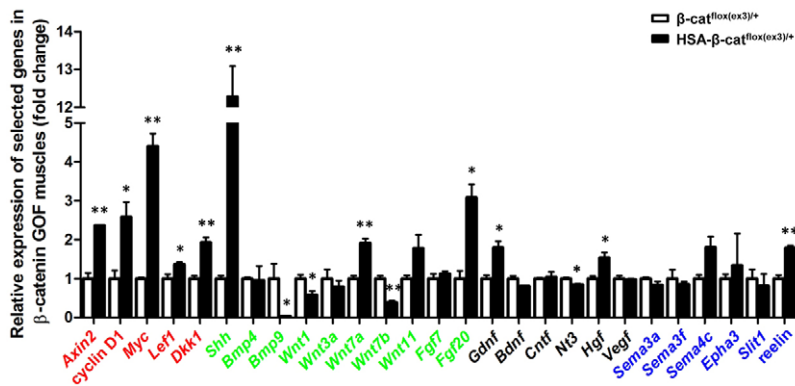


Fig. 8. Different gene expression in HSA- β -cat^{lox(ex3)/+} muscles. Total RNA was isolated and subjected to quantitative real-time PCR for indicated genes. mRNA levels were calibrated to *Gapdh* mRNA levels and normalized to mRNAs from control mice (** $P < 0.01$, * $P < 0.05$, $n = 3$, t -test). Known Wnt/ β -catenin target genes are shown in red; morphogens in green; neurotrophic factors in black; and axon guidance molecules in blue. Error bars indicate s.e.m.

between control and HB9- β -cat^{lox(ex3)/+} mice (supplementary material Fig. S7B,F-H). Finally, mEPP amplitudes and frequencies at HB9- β -cat^{lox(ex3)/+} NMJs were similar to those at control, suggesting normal neurotransmission (data not shown). These results indicate that motoneuron β -catenin GOF had no effect on NMJ formation or function.

DISCUSSION

This study provides evidence that NMJs do not form properly when muscle β -catenin activity is increased. Intriguingly, β -catenin LOF and GOF in muscles generate converse as well as similar NMJ deficits (Table 1). These observations suggest that NMJ formation requires a critical level of β -catenin activity in the muscle.

AChR cluster localization

A similar phenotype between muscle β -catenin LOF and GOF is the increased central area in which AChR clusters are distributed (Fig. 5A,B) (Table 1). How would both muscle β -catenin ablation and expression cause mislocation of AChR clusters? In the simplest

scenario, β -catenin regulates cluster localization and this function can be disrupted by either elevated or decreased levels. β -Catenin interacts with cadherins to regulate cytoskeleton (Kemler, 1993; Huber et al., 1996; Orsulic et al., 1999; Nelson and Nusse, 2004), a mechanism that has been implicated in synaptogenesis in the CNS. For example, the N-cadherin- β -catenin complex accumulates at synapses onto dendritic spines and regulates dendritic spine morphogenesis, synaptic vesicle recycling, and recruitment of vesicles to active zones (Murase et al., 2002; Togashi et al., 2002; Bamji et al., 2003; Vitteira et al., 2012). Two major types of cadherins, N- and M-type, are expressed in muscles (Moore and Walsh, 1993; Cifuentes-Diaz et al., 1994; Kaufmann et al., 1999); however, their function in NMJ formation remains unclear.

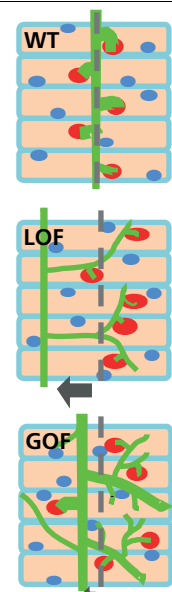
β -Catenin could interact with rapsyn, which is necessary for agrin-induced AChR clustering (Zhang et al., 2007). Rapsyn is an adapter protein that interacts directly with AChR and is required for AChR clusters (Gautam et al., 1995). The fact that clusters form in both LOF and GOF mice suggests a modulatory role. Moreover, mEPP amplitudes of these mice were similar (Fig. 6C) (Table 1), suggesting that β -catenin might not regulate AChR density in the

Table 1. Comparison of NMJ phenotypes in muscle β -catenin LOF and GOF mice

Phenotypes	LOF	GOF
AChR clusters		
Size	↑	–
Band width	↑	↑
Motor axons		
Diameter of axons	↓	–
Primary branch mislocation	↑↑↑	↑
Complexity of arborization	↓	↑
Nerve terminals		
Nerve terminal number	–	–
Active zone number	↓	–
Synaptic vesicle number	↓	–
Synaptic cleft width	–	–
Synaptic transmission		
mEPP frequency	↓	↓
mEPP amplitude	–	–
EPP	↓	–
Ectopic muscles		
	none	↑

–, no change; ↑, increased; ↓, decreased.

Diagrams summarize NMJ morphological phenotypes of wild type (WT), β -catenin loss-of-function (LOF) and gain-of-function (GOF) mice. Green, nerve; red, AChR clusters; blue ovals, nuclei. Dashed lines mark the middle region of muscle fibers. Arrows indicate the shift direction of primary branches.



clusters. Finally, β -catenin-dependent transcription is elevated in entire muscle fibers, as indicated by TOP-EGFP (Fig. 1F); however, most AChR clusters remained in the central, albeit widened, region. These results suggest a scaffold function of β -catenin in controlling where AChR clusters form.

Prior to innervation, muscle fibers are pre-patterned with aneural AChR (Lin et al., 2001; Yang et al., 2001). However, the GOF seemed to have no effect on the pre-patterning (data not shown). Innervation, probably via ACh, disperses AChR clusters in non-synaptic areas (Misgeld et al., 2002; Brandon et al., 2003). This function might be impaired in the LOF and GOF mice because of reduced mEPP frequency. However, the number of AChR clusters per muscle fiber was not altered by the GOF (Fig. 5F,G). Therefore, the phenotype of scattered clusters might not be due to a problem with cluster dispersal. Recently, dihydropyridine receptor (an L-type calcium channel) was shown to control AChR cluster location (Chen et al., 2011). However, this mechanism might not be involved because dihydropyridine receptor mutation impaired mobility, in contrast to β -catenin GOF.

Motor axon outgrowth, navigation and arborization

In the LOF mice, phrenic nerve bundles are smaller with reduced axon size, but not number (Li et al., 2008) (Table 1). This phenotype suggests a trophic function of muscle β -catenin for motor axons. This notion is supported by increased extension and outgrowth of phrenic nerves (Fig. 2A,D, Fig. 3B-D) and spinal axons (Fig. 4C) in GOF mice. Intriguingly, aberrant activation of β -catenin stimulates the formation of ectopic muscles (Fig. 3), which were apparently able to induce and attract ectopic branches of phrenic nerves (Fig. 2A, Fig. 3C-E). This phenotype suggests that muscle β -catenin GOF might direct a guidance activity for innervation. However, in muscle β -catenin LOF mice, phrenic axons are able to navigate to the middle region, suggesting that attractive signals from muscles are not altered. However, the mislocation of primary branches is more severe in LOF mice compared with that in GOF mice (Fig. 2A, Fig. 3D) (Table 1) (Li et al., 2008).

Converse phenotypes of phrenic axon defasciculation and arborization were observed in LOF and GOF muscles. In the LOF mice, secondary branches are scarce, but elongated (Li et al., 2008), whereas in GOF diaphragms, short secondary branches were abundant in addition to long ones (Fig. 2A, Fig. 3D). GOF also increased the length and/or number of tertiary and quaternary branches (Fig. 2). Arborization was also observed in axons of brachial plexus (Fig. 4). Nevertheless, phrenic terminals were confined within the area where AChR clusters are located (Fig. 3), suggesting that the axon stop signal from muscle fibers is locally expressed or delivered although β -catenin activity was increased in entire muscle fibers (Fig. 1F). Several factors have been implicated in promoting axon branching of various neurons including BDNF, CNTF, netrin 1, Wnt3a and Sema3A (Gurney et al., 1992; Cohen-Cory and Fraser, 1995; Krylova et al., 2002; Bagri et al., 2003; Dent et al., 2004; Tang and Kalil, 2005; Singh et al., 2008). However, their expression was not altered by muscle β -catenin GOF (Fig. 8).

Terminal differentiation and function

The frequency of mEPPs is reduced in both GOF and LOF mice. This superficially similar phenotype is apparently caused by distinct mechanisms. Electron microscopic analysis demonstrates that the numbers of synaptic vesicles and active zones were

significantly decreased in β -catenin LOF NMJs (Fig. 7G-J), suggesting that the production and docking of synaptic vesicles were compromised in the absence of muscle β -catenin. By contrast, the ultrastructure of NMJs in GOF mice appears normal (Fig. 7A-F). These observations suggest that neurotransmission deficits in β -catenin LOF NMJs were due to impaired development, whereas those in GOF NMJs were not structural, but functional. In support of this notion, action potentials are able to elicit normal EPPs in GOF, but not LOF NMJs (Table 1) (Li et al., 2008).

Wnt signaling or cadherin signaling in NMJ formation

Recent studies suggest a role of Wnt signaling in synapse formation. In *C. elegans*, Wnt signaling determines the position of NMJs by inhibiting synaptogenesis (Klassen and Shen, 2007), whereas in *Drosophila*, Wnt promotes NMJ formation (Packard et al., 2002; Wu et al., 2010; Budnik and Salinas, 2011). The extracellular domain of MuSK contains a cysteine-rich domain (CRD) that is homologous to that in Fz (Valenzuela et al., 1995; Glass et al., 1996). MuSK also interacts with Dvl, which regulates agrin-induced AChR clustering (Luo et al., 2002). APC, an adaptor downstream of Dvl in the canonical pathway, was shown to regulate AChR clustering (Wang et al., 2003). In zebrafish, Wnt11r binds to *unplugged*, the zebrafish MuSK homolog, to guide motor axons (Jing et al., 2009). In mammalian muscle cells, five Wnts (Wnt9a, Wnt9b, Wnt10b, Wnt11 and Wnt16), presumably by direct binding to MuSK, are able to stimulate AChR clustering, in a manner dependent on LRP4 (Zhang et al., 2012). Recently, WNT4 has been implicated in muscle pre-patterning and to stimulate AChR clustering via activation of MuSK (Strochlic et al., 2012). In addition, agrin-induced AChR clustering was enhanced by WNT3, but reduced by WNT3a (Henriquez et al., 2008; Wang et al., 2008).

β -Catenin-dependent transcription seemed not required for agrin- or Wnt-induced AChR clustering (Zhang et al., 2007) (B. Zhang and L.M., unpublished observations). How muscle β -catenin regulates presynaptic differentiation and/or function remains unclear. It is possible that muscle β -catenin acts by regulating the transcription of genes encoding retrograde factors involved in presynaptic differentiation. In favor of this hypothesis is the observation that various neural morphogens, neurotrophic factors and axon guidance cues are increased in expression (Fig. 8). Overexpression of GDNF has been shown to promote axon growth and delay synapse elimination (Nguyen et al., 1998). However, the number of NMJs per muscle fibers was apparently normal in β -catenin GOF mice (Fig. 5F,G). Sema3A has been implicated in the controlled growth of motor axons into the developing limb territory (Huber et al., 2005); however, its expression was not altered by β -catenin GOF and no apparent deficits in motor axon navigation were observed in β -catenin GOF mice. HGF was shown to promote neurite outgrowth of motoneurons (Ebens et al., 1996; Yamamoto et al., 1997) and, intriguingly, its level was elevated in β -catenin GOF muscles. Further study is needed to determine if HGF is crucial for NMJ formation in vivo.

Whether muscle β -catenin regulates presynaptic differentiation via cadherin signaling remains an outstanding question. N-cadherin could stimulate neurite outgrowth (Matsunaga et al., 1988; Bixby and Zhang, 1990; Payne et al., 1992). The N-cadherin- β -catenin complex was shown to promote dendritic arborization in hippocampal neurons, and the effect does not require β -catenin-dependent transcription (Yu and Malenka, 2003). A recent study reports a trans-synaptic activity of N-cadherin/ β -catenin signaling in

regulating synapse function in hippocampal neurons (Vitureira et al., 2012). Intriguingly, only disruption of postsynaptic, but not presynaptic, N-cadherin activity modulates vesicle release probability (Vitureira et al., 2012), a phenomenon similar to our findings that only muscle, not motoneuron, β -catenin may be crucial. Notice that exon 3 of the β -catenin gene does not encode the motif critical for interaction with α -catenin. Therefore, the β -catenin GOF mutant might function in a dominant-negative manner to disrupt interaction with α -catenin. However, such a cadherin-dependent mechanism is believed to be infeasible at mature NMJs because of large synaptic clefts (Wu et al., 2010). It is possible that this mechanism is involved in early development of the NMJ. In fact, N-cadherin was recently shown to regulate primary motor axon growth and branching during zebrafish embryonic development (Bruses, 2011). It is worth pointing out that NMJ formation or function is apparently not altered by both β -catenin LOF and GOF in motoneurons (Li et al., 2008) (supplementary material Fig. S7), which suggests that β -catenin in motoneurons is dispensable. Recent studies indicate a redundant function of γ -catenin in motoneurons. Positioning or migration of motoneurons in the spinal cord does not require Wnt canonical signaling, but N-cadherin signaling that can be mediated by either β -catenin or γ -catenin (Demireva et al., 2011; Bello et al., 2012). Phenotypes were observed only when both β -catenin and γ -catenin were lost. However, the loss seemed to have no effect on the ability of axons to select appropriate dorsoventral trajectories upon entering the limb.

In summary, this study provides further genetic evidence that muscle β -catenin is crucial for NMJ development and function. Considering multiple functions of β -catenin, it is likely that different NMJ deficits in LOF as well as GOF mice are caused by complex cell autonomous and non-autonomous mechanisms. Muscle lacking β -catenin and expressing stable β -catenin could provide a niche to identify muscle-derived factors by gene array analysis. However, extensive work (including in vivo rescue) is needed for validation and functional characterization of the candidates.

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Competing interests statement

The authors declare no competing financial interests.

Supplementary material

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