ERRATUM

Restoration of synapse formation in Musk mutant mice expressing a Musk/Trk chimeric receptor
Herbst, R., Avetisova, E. and Burden, S. J. Development 129, 5449-5460

On page 5454 of this article, the first paragraph in the section ‘Motor axons extend…’ should read ‘Based on the expression of transgenes containing the MCK enhancer and promoter, the endogenous MCK gene is activated in skeletal muscle at ~E13.5 (S. Hauschka, personal communication), 1 day after motor axons first enter the muscle.’

We apologise to the authors and readers for this mistake.

CORRIGENDUM

Kremen proteins interact with Dickkopf1 to regulate anteroposterior CNS patterning

In the reference list and text, one of the references was mis-spelled.


The authors apologise to readers for this mistake.

CORRIGENDUM

Quantitative developmental anatomy of definitive haematopoietic stem cells/long-term repopulating units (HSC/RUs): role of the aorta-gonad-mesonephros (AGM) region and the yolk sac in colonisation of the mouse embryonic liver

There is an error in Fig. 1A of this article. The correct version of the figure is printed below.

The authors apologise to readers for this mistake.
INTRODUCTION

Neuromuscular synapses form following a series of complex interactions between motor neurons, muscle fibers and Schwann cells (Burden, 1998; Sanes and Lichtman, 1999; Schaeffer et al., 2001; Son and Thompson, 1995). Agrin, a 200 kDa protein synthesized by motor neurons, is a critical synaptic signaling molecule that organizes postsynaptic differentiation by stimulating Musk, a receptor tyrosine kinase (RTK) expressed selectively in skeletal muscle (Glass and Yancopoulos, 1997; McMahan, 1990). Embryos lacking agrin or Musk fail to form neuromuscular synapses and consequently die at birth because of their failure to move or breathe. We produced mice that express a chimeric receptor, containing the juxtamembrane region of Musk and the kinase domain of TrkA, selectively in muscle, and we crossed this transgene into Musk mutant mice. Expression of this chimeric receptor restores presynaptic and postsynaptic differentiation, including the formation of nerve terminal arbors, synapse-specific transcription, and clustering of postsynaptic proteins, allowing Musk mutant mice to move, breathe and survive as adults. These results show that the juxtamembrane region of Musk, including a single phosphotyrosine docking site, even in the context of a different kinase domain, is sufficient to activate the multiple pathways leading to presynaptic and postsynaptic differentiation in vivo. In addition, we find that Musk protein can be clustered at synaptic sites, even if Musk mRNA is expressed uniformly in muscle. Moreover, acetylcholine receptor clustering and motor terminal branching are restored in parallel, indicating that the extent of presynaptic differentiation is matched to the extent of postsynaptic differentiation.

SUMMARY

Mice lacking Musk, a muscle-specific receptor tyrosine kinase that is activated by agrin, fail to form neuromuscular synapses and consequently die at birth because of their failure to move or breathe. We produced mice that express a chimeric receptor, containing the juxtamembrane region of Musk and the kinase domain of TrkA, selectively in muscle, and we crossed this transgene into Musk mutant mice. Expression of this chimeric receptor restores presynaptic and postsynaptic differentiation, including the formation of nerve terminal arbors, synapse-specific transcription, and clustering of postsynaptic proteins, allowing Musk mutant mice to move, breathe and survive as adults. These results show that the juxtamembrane region of Musk, including a single phosphotyrosine docking site, even in the context of a different kinase domain, is sufficient to activate the multiple pathways leading to presynaptic and postsynaptic differentiation in vivo. In addition, we find that Musk protein can be clustered at synaptic sites, even if Musk mRNA is expressed uniformly in muscle. Moreover, acetylcholine receptor clustering and motor terminal branching are restored in parallel, indicating that the extent of presynaptic differentiation is matched to the extent of postsynaptic differentiation.

Supplemental data available on-line

Key words: Agrin, Acetylcholine receptor, Synapse-specific transcription, Receptor tyrosine kinase, Neuromuscular synapse, Mouse
AChR clustering or tyrosine phosphorylation in muscle cell lines. The same Musk/TrkA chimera, but including a substitution of thirteen amino acids from the juxtamembrane region of Musk, including Y553, with the comparable region in TrkA is similarly tyrosine phosphorylated by agrin stimulation, but tyrosine residue phosphorylation of this chimera leads to clustering and tyrosine phosphorylation of AChRs in cultured myotubes (Herbst and Burden, 2000). These experiments indicate that the juxtamembrane region of Musk, even in the context of a different kinase domain, is sufficient to activate a signaling pathway leading to the clustering and tyrosine phosphorylation of AChRs in muscle cell lines. Nonetheless, as cultured muscle cells are not amenable to study all aspects of synaptic differentiation, these experiments could not address whether this juxtamembrane region of Musk is sufficient to confer additional aspects of Musk signaling, including clustering of additional postsynaptic proteins, synapse-specific transcription, presynaptic differentiation and synapse formation. Moreover, requirements for synaptic proteins may differ in cell culture and in vivo; for example, rapsyn is required to cluster Musk in cultured cells but not at synapses in vivo (Gillespie et al., 1996; Moscoso et al., 1995), and the ectodomain of Musk is required to cluster AChRs in cultured cells but not in vivo (Apel et al., 1997; Sander et al., 2001). Thus, in vivo studies are required to delineate the signaling mechanisms that lead to the complex biological response initiated by Musk at synapses.

In addition to Y553 in the juxtamembrane region of Musk, agrin stimulates the phosphorylation of five tyrosine residues in the kinase domain of Musk (Watty et al., 2000). In other RTKs, recruitment of different adaptor proteins to distinct phosphorytrosine docking sites leads to activation of disparate signaling pathways, which are often coupled to different biological responses (Madhani, 2001; Pawson and Scott, 1997). For example, in TrkA, the receptor for nerve growth factor, phosphorylation of a juxtamembrane tyrosine leads to Ras and PI3-kinase activation, which are important for cell survival and neurite outgrowth (Greene and Kaplan, 1995; Huang and Reichardt, 2001), whereas phosphorylation of a tyrosine in the C-terminal region of TrkA leads to PLCγ activation, which is crucial for NGF-dependent Na+ channel and VRL channel regulation (Choi et al., 2001; Chuang et al., 2001). Moreover, in TrkB, phosphorylation of a single juxtamembrane tyrosine is required for nearly all of NT4-dependent signaling whereas phosphorylation of other tyrosine residues are required to mediate BDNF-dependent signaling in vivo (Minichiello et al., 1998). Similarly, in Met, the receptor for hepatocyte growth factor, a binding site for Grb2 is essential for late steps in myogenesis (Maina et al., 1996), a binding-site for PI3-kinase is essential for placental development, hepatocyte survival and myoblast migration, and a Src binding-site is essential for motor axon outgrowth (Maina et al., 2001). To determine whether clustering of postsynaptic proteins, synapse-specific transcription and presynaptic differentiation depend upon multiple, distinct docking sites in Musk, we produced mice that expressed the Musk/TrkA chimera, containing thirteen amino acids from the juxtamembrane region of Musk and the kinase domain of TrkA, and we crossed this transgene into Musk mutant mice. We found that expression of this chimeric receptor could restore all aspects of postsynaptic and presynaptic differentiation that were defective in Musk mutant mice. Moreover, we found that accumulation of Musk protein at synaptic sites was not dependent upon synaptic localization of Musk mRNA. These results indicate that the juxtamembrane region of Musk, including a single phosphorytrosine docking site, even in the context of a different kinase domain, is sufficient to activate the multiple pathways that lead to presynaptic and postsynaptic differentiation in vivo.

**MATERIALS AND METHODS**

**Generation of MCK-Musk and MCK-MMT transgenic mice**

GFp was ligated in frame to the C terminus of the rat Musk cDNA using the pEGFP N1 vector (Clontech). Lithmus 29 (New England Biolabs) was modified by introducing an AgeI/KpnI/SacI/AgeI cassette into the AgeI site and removing the original KpnI and SacI sites from the polylinker; the MCK promoter/enhancer/intron cassette was excised from pBS-MCK (Bruning et al., 1998) and subcloned into the modified, KpnI/SacI-digested Litusm 29 plasmid. The Musk-GFP and the MMT (Herbst and Burden, 2000) constructs were subcloned into the Litusm 29/MCK. Both the Musk and MMT constructs contain a FLAG epitope tag in the extracellular domain (Herbst and Burden, 2000). The MCK-Musk and MCK-MMT constructs were excised from Lithmus 29 using AgeI and gel purified. DNA was injected into the male pronucleus of B6/D2 fertilized eggs, which were subsequently transferred into pseudopregnant foster mice; progeny were genotyped by Southern blotting and PCR, and transgene-containing founder mice were crossed to Musk mice.

**Mouse genotyping**

Tail DNA of F1 mice was digested with EcoRI and analyzed by Southern blotting using probes from the rat Musk (nucleotides 2080-2708) and rat TrkA (nucleotides 2191-2572) kinase domains (Meakin et al., 1992). Subsequent generations of mice were genotyped by PCR using primers that are specific for sequences in the extracellular region of the mouse Musk cDNA (5′-GAAAGCACCCTTTTCTCTGAG-3′ and 5′-ATTGAGCAGGAGCTTGC3′) using the following conditions: one cycle of 94°C for 2 minutes, 40 cycles of 94°C for 30 seconds, 58°C for 30 seconds and 72°C for 40 seconds followed by 1 cycle of 72°C for 2 minutes. The wild-type and mutant Musk alleles (DeChiara et al., 1996) were detected by PCR, using primers that are specific for sequences in the Musk kinase domain and the neo gene (5′-ATGCCGCCCCAGTCTAGGCTAC-3′, 5′-TTCCTGCCC- AAAACATCAACTG-3′ and 5′-CATAACGCTGAGAAGCTAGA- TCAGCAGC-3′), using the following conditions: one cycle of 94°C for 2 minutes, 40 cycles of 94°C for 30 seconds, 56°C for 30 seconds and 72°C for 40 seconds followed by one cycle of 72°C for 2 minutes. We examined three MCK-Musk lines and two MCK-MMT lines; analyzed mice were heterozygous for MCK-Musk or MCK-MMT. We noted attrition in large litters up to the time of weaning, indicating that Musk+/-; MCK-MMT and, in particular, Musk+/-; MCK-Musk pups competed poorly with their wild-type littermates for nourishment. Musk+/-; MCK-MMT and Musk+/-; MCK-Musk mice are smaller than their littermates during this period, but attain the same weight as their littermates by one month after birth. We assessed presynaptic and postsynaptic differentiation in embryos and in 3-week postnatal mice, when neuromuscular synapses are fully mature. Musk+/-; MCK-Musk and Musk+/-; MCK-MMT adult mice appeared to have normal motility as they performed identically to wild-type mice on a Rotarod at a constant speed (32 rpm). Rescued adult mice show no signs of a shortened longevity, as they have been maintained for an excess of 1 year.
**Immunohistochemistry**

Diaphragm muscles were dissected from embryos or three week old mice, fixed for 90 minutes in 1% formaldehyde, rinsed in PBS and incubated with 0.1 M glycine in PBS for 15 minutes. After dissection of the overlying connective tissue, the muscles were permeabilized in 0.5% Triton X-100 in PBS (PBST) for 5 minutes, incubated overnight at 4°C with rabbit polyclonal antibodies against neurofilament (1:500, Chemicon) and synaptophysin (1:5, Zymed) in 4% goat serum/2% BSA/PBST, washed three times for 20 minutes in PBST, incubated overnight at 4°C with fluorescein-conjugated goat anti-rabbit IgG (1:200, Jackson Immunoresearch) and Texas Red-conjugated a-BGT (Molecular Probes). The muscles were washed twice for 20 minutes in PBST, twice for 20 minutes in PBS, postfixed in 1% formaldehyde for 10 minutes, rinsed in PBS, flat mounted in Vectashield (Vector Labs) and viewed with optics selective for either fluorescein or Texas Red.

Frozen sections (10 μm) from fixed, adult leg muscles were stained with antibodies as described previously (DeChiara et al., 1996; Zhu et al., 1995). The following primary antibodies were used: affinity-purified rabbit anti-rapsyn (1:500) (Herbst and Burden, 2000), mouse et al., 1995). The following primary antibodies were used: affinity-purified rabbit anti-rapsyn (1:500) (Herbst and Burden, 2000), mouse anti-α-BGT (Molecular Probes). The muscles were washed twice for 20 minutes in PBST, twice for 20 minutes in PBS, postfixed in 1% formaldehyde for 10 minutes, rinsed in PBS, flat mounted in Vectashield (Vector Labs) and viewed with optics selective for either fluorescein or Texas Red.

**Whole-mount in situ hybridization**

Intercostal muscles were fixed in 4% formaldehyde, dehydrated in methanol, digested with Proteinase K, probed with a digoxigenin-labeled riboprobe transcribed from an AChR α (DeChiara et al., 1996), AChR δ subunit (Simon et al., 1992) or a rat Musk cDNA (nucleotides 1-1663) and processed as described elsewhere (Wilkinson, 1992). A low level of uniform staining was observed with control, sense probes for the AChR α and δ subunits.

**Immunoprecipitation and western blotting**

Tissues were homogenized and lysed as described previously (Bruning et al., 1998). Tissues were homogenized in lysis buffer

![Image](Image.png)

**Fig. 1.** MCK-Musk and MCK-MMT transgenes are expressed and active in skeletal muscle. (A) Expression of Musk and MMT are controlled by the 5’ regulatory region of the muscle creatine kinase (MCK) gene. MCK-Musk and MCK-MMT transgenes each contain a FLAG epitope-tag in the extracellular domain of Musk (Herbst and Burden, 2000). The MCK-Musk transgene encodes a fusion protein between Musk and GFP. MMT, a chimera between the extracellular and transmembrane regions of Musk and the intracellular region of TrkA, includes a substitution of 13 amino acids from the juxtamembrane (JM) of Musk with the comparable region in TrkA (Herbst and Burden, 2000). (B) Transgene expression was measured by immunoprecipitation followed by western blotting. MCK-Musk and MCK-MMT are expressed strongly in skeletal muscle; heart expression is detected in some, but not all lines (arrows). TrkA is expressed strongly in brain. (C) Transgene expression, determined by western blotting of skeletal muscle lysates, differs among the different transgenic (Tg) lines: MCK-MMT, line 5; expresses more MMT in skeletal muscle than MCK-MMT (line 29), and MCK-Musk (line 23) expresses more Musk in skeletal muscle than MCK-Musk, line 47. (D) Musk as well as MMT are tyrosine phosphorylated. Musk and MMT were immunoprecipitated from skeletal muscle with antibodies to Musk and antibodies to TrkA, respectively, and western blots were probed with antibodies to phosphotyrosine (pY) and FLAG.
Metamorph. We measured the total number of pixels in two thresholded areas of the left hemi-diaphragm muscle, each adjacent to the main intramuscular nerve. The value in the first area, which is immediately adjacent to the main intramuscular nerve, provides an estimate of the number of axons that branched from the main intramuscular nerve, while the value in the second area, which is adjacent to the main nerve but extends further toward the edge of the muscle, provides an estimate of the extent of axon growth away from the main intramuscular nerve. The value in each area of wild-type muscle was assigned as 100%, and the values in mutant mice have reached the edge of the muscle; owing to the irregular shape of the muscle, however, we measured axon outgrowth in an area that did not extend to the edge of the muscle. Consequently, our procedure underestimates the increase in axon outgrowth in mutant mice. Images from at least three mice were analyzed per age and per genotype. The total number of pixels did not differ significantly (Fig. 1A). Moreover, synapses are maintained and differentiate and that AChRs cluster in the rescued newborn mice (Fig. 2B). These experiments demonstrate that the juxtamembrane region of Musk, even in the context of a kinase domain from a different tyrosine kinase, is sufficient to activate a signaling cascade that leads to clustering of AChRs in vivo. Furthermore, as MMT expression, selectively in muscle, restores presynaptic differentiation, these results indicate that the juxtamembrane region of Musk has a central role in producing and/or organizing retrograde signals required for presynaptic differentiation and synapse formation.

Antibodies
Antibodies (anti-GFP) to GFP were produced by immunizing rabbits with a conjugate between KLH and an extracellular sequence (SCGALQVKMKPKTRPPINV) in Musk. To produce transgenic mice that express this chimera (MMT) under the control of the muscle-specific creatine kinase [MCK (Ckmm – Mouse Genome Informatics)] 5′ regulatory region (Fig. 1), and we subsequently crossed this transgene into Musk mutant mice. We produced and analyzed two MCK-MMT transgenic lines, as well as three MCK-MusK control transgenic lines. These mice express the transgenes in a muscle-specific manner, as assessed by western blotting (Fig. 1), and the protein encoded by each transgene is tyrosine phosphorylated (Fig. 1). MCK-MMT and MCK-MusK transgenic mice were crossed to MusK+/– mice, and progeny were crossed to MusK+/– mice to generate mice that lack endogenous Musk but express either MCK-MMT or MCK-MusK. Unlike MusK mutant mice, which lack neuromuscular synapses and die at birth (DeChiara et al., 1996), MusK mutant mice expressing either MCK-MMT or MCK-MusK move, breathe and survive after birth (Table 1).

We stained wholemounts of diaphragm muscle from E18.5 mice with a mixture of antibodies to neurofilament and synaptophysin, to label axons and terminals, respectively, and with Texas Red α-Bungarotoxin (TR α-BGT) to label acetyl choline receptors (AChRs). We found that nerve terminals differentiate and that AChRs cluster in the rescued newborn mice (Fig. 2A). Moreover, synapses are maintained and continue to mature postnatally (Fig. 2B). These experiments demonstrate that the juxtamembrane region of Musk, even in the context of a kinase domain from a different tyrosine kinase, is sufficient to activate a signaling cascade that leads to clustering of AChRs in vivo. Furthermore, as MMT expression, selectively in muscle, restores presynaptic differentiation, these results indicate that the juxtamembrane region of Musk has a central role in producing and/or organizing retrograde signals required for presynaptic differentiation and synapse formation.

RESULTS
A Musk/TrkA chimera can substitute for endogenous Musk in vivo

To determine whether the Musk/TrkA chimera, including the juxtamembrane region of Musk, contains sequences that are sufficient for presynaptic and postsynaptic differentiation, we produced transgenic mice that express this chimera (MMT) under the control of the muscle-specific creatine kinase [MCK (Ckmm – Mouse Genome Informatics)] 5′ regulatory region (Fig. 1), and we subsequently crossed this transgene into MusK mutant mice. We produced and analyzed two MCK-MMT transgenic lines, as well as three MCK-MusK control transgenic lines. These mice express the transgenes in a muscle-specific manner, as assessed by western blotting (Fig. 1), and the protein encoded by each transgene is tyrosine phosphorylated (Fig. 1). MCK-MMT and MCK-MusK transgenic mice were crossed to MusK+/– mice, and progeny were crossed to MusK+/– mice to generate mice that lack endogenous Musk but express either MCK-MMT or MCK-MusK. Unlike MusK mutant mice, which lack neuromuscular synapses and die at birth (DeChiara et al., 1996), MusK mutant mice expressing either MCK-MMT or MCK-MusK move, breathe and survive after birth (Table 1).

Table 1. MCK-MusK and MCK-MMT rescue MusK mutant mice

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<thead>
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<th>MCK-MMT (#29)</th>
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<th>MCK-MusK (#23)</th>
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<td></td>
<td>Observed</td>
<td>Expected</td>
<td>χ-square</td>
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<td>MusK+/–; MCK-Tg</td>
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<td>11.7</td>
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<tr>
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<tr>
<td>MusK+/–</td>
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<tr>
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<td>3</td>
<td>5.8</td>
<td>1.35</td>
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MusK+/– mice were crossed with MusK+/–; MCK-MusK or MusK+/–; MCK-MMT mice, and we determined the genotype of the surviving progeny at P0 or P1. χ-square analysis of progeny from a cross of MusK+/– × MusK+/–; MCK-MusK (line #23) mice or MusK+/– × MusK+/–; MCK-MMT (line #5, line #29) mice shows that the occurrence of genotypes is unlikely (P>0.05) to occur by chance, indicating that the neonatal lethality of MusK mutant mice can be fully rescued by expression of MCK-MusK or MCK-MMT.
The density of AChRs at synaptic sites in rescued mice

We measured the density of AChRs at synaptic sites (Fig. 2B) and found that the density of synaptic AChRs is modestly lower in \( \text{Musk}^{-/-}; \ MCK-\text{Musk} \) mice than in wild-type mice (line #23, \( 79\pm3.2\% \), \( n=3 \); line #54, 77\%, \( n=1 \)). These results indicate that the MCK regulatory region confers sufficient Musk expression to restore AChR clustering at synapses but that the level and/or pattern of \( MCK-\text{Musk} \) expression differs from wild-type Musk expression, resulting in a 20\% reduction in AChR density at synapses. In \( \text{Musk}^{-/-}; \ MCK-\text{MMT} \) mice, AChRs are likewise clustered at synapses, but at a reduced density (line #5, \( 66\pm3.4\% \), \( n=3 \); line #29, \( 48\pm0.8\% \), \( n=3 \)). The lower density of synaptic AChRs in rescued mice expressing MMT may be due to a modestly (~twofold) lower expression of the MMT transgene (see supplementary data at http://dev.biologists.org/supplemental/) or to sequences in Musk that are absent from MMT. Nonetheless, the MMT chimera also restores AChR clustering at synapses.

MMT chimera contains sequences that are sufficient for clustering postsynaptic proteins

Agrin/Musk signaling is necessary and sufficient to cluster several muscle-derived proteins in addition to AChRs (DeChiara et al., 1996; Gautam et al., 1996; Meier et al., 1997; Rimer et al., 1998). We therefore examined whether the MMT chimera contains sequences sufficient for clustering these proteins at synaptic sites. We stained frozen sections of muscle from three week old mice with antibodies to: (1) rapsyn, an intracellular, peripheral membrane protein associated with AChRs; (2) utrophin, a component of the dystrophin-associated glycoprotein complex; (3) ErbB4, a receptor for neuregulin; and (4) Na\(^+\) channels (Sanes and Lichtman, 1999). Fig. 3 shows that rapsyn, ErbB4, utrophin and Na\(^+\) channels are each concentrated at synapses in \( \text{Musk}^{-/-}; \ MCK-\text{MMT} \) mice. These results indicate that the MMT chimera contains sequences that are sufficient to restore the clustering of most, if not all postsynaptic proteins.

MMT chimera induces extrasynaptic AChR clusters

In innervated muscle fibers from wild-type and \( \text{Musk}^{-/-}; \ MCK-\text{Musk} \) mice, AChR clusters are restricted to synaptic sites. In \( \text{Musk}^{-/-}; \ MCK-\text{MMT} \) mice, however, AChR clusters are readily evident at ectopic sites in addition to neuromuscular synapses (Fig. 4). Because these ectopic AChR clusters are found on most muscle fibers and are often located immediately adjacent to synaptic sites, marked by terminal arbors and synaptic AChR clusters, these ectopic AChR clusters are not associated with muscle denervation. Moreover, other synaptic proteins, including rapsyn, ErbB4, utrophin and Na\(^+\) channels, are each co-clustered with ectopic AChR clusters in \( \text{Musk}^{-/-}; \ MCK-\text{MMT} \) mice (Fig. 4).

Synapse-specific transcription is activated by MMT

Agrin/Musk signaling is necessary to activate synapse-specific transcription, leading to an enrichment of AChR mRNA in the central, synapse-rich region of muscle (DeChiara et al., 1996; Gautam et al., 1996). We therefore used in-situ hybridization to determine whether AChR genes are expressed selectively in synaptic nuclei of \( \text{Musk}^{-/-}; \ MCK-\text{MMT} \) mice. Fig. 5 shows that
AChR α and δ subunit mRNAs are each enriched in the central region of intercostal muscle from Musk−/−; MCK-MMT newborn mice, demonstrating that the Musk juxtamembrane region has a key role in activating a pathway leading to synapse-specific gene expression.

Musk and MMT proteins are concentrated at synaptic sites despite uniform Musk and MMT RNA expression

In wild-type mice, Musk RNA, like AChR mRNA, is expressed preferentially at synaptic sites (Valenzuela et al., 1995). Because the MCK gene is expressed in nuclei throughout the myofiber (Tang et al., 1994), the MCK regulatory region, as expected, confers Musk and MMT mRNA expression uniformly throughout the muscle (Fig. 6). We examined whether Musk protein, encoded by the transgene, is expressed uniformly in the muscle or concentrated at synaptic sites in Musk−/−; MCK-Musk mice. We stained sections of muscle from Musk−/−; MCK-Musk mice with antibodies to Musk and found that Musk is clustered at synaptic sites (Fig. 6). Likewise, we found that MMT is clustered at synaptic sites in Musk−/−; MCK-MMT mice (Fig. 6). Thus, Musk protein can be clustered at synaptic sites even if Musk mRNA is expressed uniformly in muscle.

Motor axons extend until Musk is expressed, yet postsynaptic differentiation is limited to a discrete site on the muscle fiber

Motor axons enter developing skeletal muscle at E12.5 and form a main intramuscular nerve in the middle of the muscle (Sanes and Lichtman, 1999). In wild-type mice, motor axons branch and terminate adjacent to the main intramuscular nerve, resulting in a narrow, distinct endplate zone in the middle of the muscle, marked by presynaptic nerve terminals and a high concentration of AChRs (Fig. 7). In Musk mutant mice, motor axons branch from the main intramuscular nerve, but these axons fail to terminate and instead wander across the muscle surface without forming specialized nerve terminals (DeChiara et al., 1996).

Based on the expression of transgenes containing the MCK enhancer and promoter, the endogenous MCK gene is activated in skeletal muscle at ~E15.5 (S. Hauschka, personal communication), 1 day after motor axons first enter the muscle. Expression of the MCK gene increases modestly (~20-fold) between E13.5 and birth, and dramatically after birth (>300-fold between birth and P21) (S. Hauschka, personal communication). We therefore reasoned that axon growth would be exuberant in Musk−/−; MCK-Musk mice, as in Musk mutant mice, prior to the onset of Musk expression. Fig. 7 shows that motor axon growth is similarly exuberant in Musk−/−; MCK-Musk and Musk−/−; MCK-MMT mice at E14.5 and E16.5. In addition, postsynaptic clustering of AChRs is absent at E14.5 and only weakly detectable at E16.5 in Musk−/−; MCK-Musk and Musk−/−; MCK-MMT mice. Between E16.5 and E18.5 in Musk−/−; MCK-Musk mice, the rate of axon growth reverts to normal and postsynaptic differentiation commences (Fig. 7). In Musk−/−;
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MCK-MMT mice, the rate of axon outgrowth also reverts between E16.5 and E18.5, but incompletely, like AChR clustering. Owing to the late expression of the MCK gene, motor axons extend well beyond their normal termination zone prior to forming neuromuscular synapses, resulting in a substantially wider end-plate zone, which persists in rescued adult mice (Fig. 8).

Because motor axons in Musk−/−; MCK-Musk mice grow over a substantial region of muscle prior to E16.5, we reasoned that postsynaptic differentiation, initiated by neurally deposited agrin, might occur over an unusually large region of each muscle fiber once Musk expression began. We found, however, that the size of AChR clusters was normal (102±4%, n=3) in muscle from Musk−/−; MCK-Musk newborn mice (Fig. 2). These results raise the possibility that agrin is preferentially available, or active, at or near the growth cone of the motor axon, and that the resulting bias in Musk activation consolidates Musk and AChR clustering to a discrete patch on the muscle fiber surface (see Discussion).
Terminal arbors are immature in \textit{Musk}^{−/−}; \textit{MCK-MM}T mice

Neuromuscular synapses undergo several structural and functional transitions during the first few weeks after birth, as the number of presynaptic inputs at individual synapses is reduced to one, and the single, remaining nerve terminal arbor becomes more complex (Sanes and Lichtman, 1999). To determine whether synaptic sites become singly innervated and nerve terminal arbors become more complex in \textit{Musk} mutant mice expressing MMT, we stained wholemounts of diaphragm muscle from 3-week-old mice with antibodies to neurofilament/synaptophysin and with TR-\(\alpha\)-BGT. In \textit{Musk}^{+/−}; \textit{MCK-Musk} and \textit{Musk}^{+/−}; \textit{MCK-MM}T mice, as in wild-type mice, synaptic sites are singly innervated (Fig. 2B, Fig. 9). Thus, synapse elimination appears normal. In \textit{Musk}^{+/-}; \textit{MCK-Musk} mice, motor axon terminals arborize and invariably form a complex, branched endplate (Fig. 2B, Fig. 9). In muscle from \textit{Musk}^{+/−}; \textit{MCK-MM}T mice, however, terminal branching is less extensive, resulting in less complex terminal arbors (Fig. 2B, Fig. 9). Some (~10%) terminal arbors are remarkably simplified, as terminal branches are fragmented into individual boutons (Fig. 9). These results indicate that MMT can restore presynaptic differentiation, although the extent of terminal arborization is incomplete. Because terminal arbor differentiation and AChR clustering are restored to a similar degree in \textit{Musk}^{+/−}; \textit{MCK-MM}T mice, \textit{Musk} activity in the postsynaptic cell appears to be limiting for presynaptic differentiation.

DISCUSSION

Agrin stimulates the phosphorylation of five tyrosine residues in the Musk kinase domain and one tyrosine residue, Y553, in the juxtamembrane region of Musk (Watty et al., 2000). Previously, we generated chimeric proteins, composed of the extracellular and transmembrane domains of Musk, and the intracellular domain of TrkA, and we found that inclusion of thirteen amino acids from the juxtamembrane region of Musk, including Y553, is sufficient, even in the context of the TrkA kinase domain, to activate a signaling pathway leading to the clustering and tyrosine phosphorylation of AChRs in cultured myotubes (Herbst and Burden, 2000). We show that muscle-specific expression of this chimeric protein rescues \textit{Musk} mutant mice, as mice that lack endogenous Musk but express the chimeric protein move, breathe and survive after birth. Expression of the chimeric protein restores neuromuscular synapse formation: nerve terminal arbors develop, postsynaptic

\begin{figure}[h]
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\includegraphics[width=\textwidth]{fig6.png}
\caption{Musk and MMT protein are concentrated at synaptic sites, despite uniform mRNA expression in \textit{Musk}^{+/−}; \textit{MCK-Musk} and \textit{Musk}^{+/−}; \textit{MCK-MM}T mice. (A) \textit{Musk} mRNA is expressed preferentially in the central, synaptic region of muscle in wild-type mice and at uniform, low levels in muscle from \textit{Musk} mutant embryos. In \textit{Musk}^{+/−}; \textit{MCK-Musk} and \textit{Musk}^{+/−}; \textit{MCK-MM}T embryos, \textit{Musk} and MMT mRNA, respectively, are expressed uniformly in muscle. The pattern of \textit{Musk} or MMT transcription was assessed by in situ hybridization of intercostal muscle from E18.5 embryos. (B) Sections from muscle of 3-week-old mice were stained with antibodies to Musk (EC, antibodies, 83033, to an extracellular epitope; IC, antibodies, 24908, to an intracellular epitope) and with Texas Red \(\alpha\)-Bungarotoxin to label AChRs. Musk and MMT proteins are concentrated at postsynaptic sites despite uniform expression of \textit{Musk} and MMT RNAs throughout the muscle. IC does not recognize MMT, as MMT lacks the epitope from the C terminus of Musk.}
\end{figure}
proteins, including AChRs, cluster at synaptic sites, and AChR genes are transcribed selectively in synaptic nuclei. Kinase activity per se is insufficient to activate synaptic differentiation, as chimeric RTKs lacking the Musk juxtamembrane region fail to activate synaptic differentiation in cultured muscle cells and at ectopic sites in muscle in vivo (Jones et al., 1999; Herbst and Burden, 2000). Thus, the Musk juxtamembrane region has a key role in activating the multiple pathways leading to presynaptic and postsynaptic differentiation in vivo.

The precise role for Musk in synapse-specific expression is poorly understood. Analysis of rapsyn mutant mice indicates that Musk may activate a signaling pathway that directly stimulates synapse-specific transcription (Gautam et al., 1995). Expression of ectopic agrin or activated Musk in adult myofibers, however, stimulates AChR transcription in an ErbB-dependent manner, suggesting that synapse-specific expression may require the Musk-dependent recruitment of a Nrg-1/ErbB signaling complex to synaptic sites (Jones et al., 1999; Meier et al., 1997; Moore et al., 2001; Rimer et al., 1998). Our experiments indicate that the Musk juxtamembrane region contains sequences that are crucial for synapse-specific transcription, but they do not shed light on whether these sequences act in a manner that is independent or dependent on Nrg-1/ErbB signaling. Thus, the Musk juxtamembrane region could recruit a signaling complex that directly regulates transcription in synaptic nuclei, or the Musk juxtamembrane region could regulate synapse-specific transcription indirectly by regulating the distribution of a Nrg-1/ErbB signaling complex, analogous to the action of Musk on AChR protein. In either case, our experiments indicate that the Musk juxtamembrane region, rather than other potential phosphotyrosine docking sites in the Musk kinase domain, has a crucial role in synapse-specific transcription.

In wild-type mice, Musk mRNA and protein are concentrated at neuromuscular synapses (Valenzuela et al., 1995). In Musk mutant mice carrying MCK-Musk, Musk mRNA is expressed throughout the muscle, yet Musk protein is concentrated at synaptic sites. These experiments demonstrate that accumulation of Musk protein at synaptic sites is not dependent upon synaptic localization of Musk mRNA. Thus, it may be more important that muscle fibers express an adequate level of mRNAs encoding synaptic proteins, such as Musk and AChR, rather than restricting these mRNAs to synaptic sites. Moreover, in Musk mutant mice carrying MCK-MMT, MMT mRNA is likewise expressed throughout the muscle, yet the chimeric protein is concentrated at synaptic sites. These results suggest that activation of the chimeric protein by agrin is sufficient to recruit additional chimeric protein from non-synaptic regions, via a positive feedback mechanism (Jones et al., 1999). Our findings, both...
from in vitro and in vivo experiments, therefore, underscore the crucial role of the Musk juxtamembrane domain in clustering Musk as well as AChRs. The mechanisms by which activated Musk recruits Musk are not understood, but this positive feedback loop may be important for achieving an adequate level of Musk expression at the synapse, sufficient to cluster more than ten million AChR molecules per synapse. Nonetheless, it seems likely that this positive feedback mechanism is restrained, as Musk activation and clustering would otherwise proceed beyond the synaptic site.

We showed previously that the MMT chimera is less responsive than Musk to agrin, as agrin stimulates half the number of AChR clusters in MMT-expressing myotubes than in Musk-expressing myotubes (Herbst and Burden, 2000). We found that the density of synaptic AChRs is lower in Musk mutant mice rescued with MMT than in Musk mutant mice rescued with Musk. The lower density of synaptic AChRs could be due to the reduced responsiveness of MMT to agrin, poorer (approx. half) expression of the MMT transgene, or both. Because MMT is less responsive to agrin in vitro, we favor the idea that sequences in the intracellular domain of Musk, not present in the MMT chimera, contribute to clustering of AChRs at synapses as well. Consistent with this idea, Y576, in the N-terminal lobe of the kinase domain, is phosphorylated in activated Musk and contributes to AChR clustering in cultured muscle cells (Herbst and Burden, 2000). Alternatively, maximal AChR clustering may depend upon sequences in Musk, not represented in the MMT chimera, that bind proteins independently of tyrosine phosphorylation (Strochlic et al., 2001).
We showed previously that MMT-expressing myotubes have more agrin-independent AChR clusters than Musk-expressing myotubes (Herbst and Burden, 2000). Consistent with these results, we found ectopic AChR clusters in innervated myofibers from Musk mutant mice rescued with MMT but not in Musk mutant mice rescued with Musk. These data support the idea that sequences in Musk restrain Musk activation in vivo, and that the absence of this auto-inhibitory sequence in the MMT chimera results in adventitious Musk activity (Till et al., 2002).

During the first few weeks after birth, the structure and function of the neuromuscular synapse is modified (Sanes and Lichtman, 1999). At birth, multiple motor axons terminate at a single, elliptical synaptic site on each myofiber. During the next few weeks, all but one of these motor axons is withdrawn, leading to innervation of the single synaptic site by a single motor axon. In addition, the shape of the synaptic site becomes more complex, as the presynaptic terminal grows and branches, and the shape of synaptic AChR clusters, ovoid at birth, becomes correspondingly complex. Although the elimination of polynuclear innervation appears to occur normally in Musk mutant mice expressing MCK-MMT, presynaptic terminal arbors are often less complex and lack extensive terminal branching. Indeed, some terminal arbors are so simplified that the synaptic site is composed of only a few boutons. These aberrations in terminal branching could arise from a failure to branch adequately when synapses first form or from remodeling and simplification, rather than growth and elaboration of branches later in development (Balice-Gordon and Lichtman, 1990). In either case, presynaptic differentiation, like postsynaptic differentiation, is incomplete in Musk mutant mice expressing MCK-MMT, and these results reinforce the idea that sequences other than the critical juxtamembrane region of Musk, though not essential for synapse formation, contribute to postsynaptic differentiation and presynaptic terminal arborization. Moreover, although prior studies of rapsyn and Musk mutant mice demonstrated that presynaptic differentiation is linked to postsynaptic differentiation (DeChiara et al., 1996; Gautam et al., 1995), the experiments described demonstrate that the extent of presynaptic and postsynaptic differentiation are matched, indicating that Muscle activity in the postsynaptic cell is limiting for presynaptic differentiation. Despite the simplified structure of synapses in Musk−/−; MCK-MMT mice, these mice are viable, fertile and behave normally in simple behavioral paradigms (see Materials and Methods).

Musk mutant mice carrying a MCK-Musk transgene first express Musk when motor axons have already extended well beyond their normal termination zone. Thus, it seemed possible that agrin, which is deposited by motor axons growing exuberantly along the muscle, might initiate postsynaptic differentiation over an unusually extensive area of the muscle once Musk expression begins. The shape and size of AChR clusters, however, are similar in wild-type mice and in Musk mutant mice that express MCK-Musk. The mechanisms that regulate release and retention of agrin from motor axons are poorly understood (Cohen et al., 1994; Ma et al., 2000), but our results are consistent with the idea that agrin accumulates, or is more active, at or near the growth cone and that an ensuing bias in Musk activation leads to focal clustering of Musk and AChRs.

The timing of MCK-Musk expression correlates well with the cessation of motor axon growth and the onset of nerve terminal differentiation in Musk−/−; MCK-Musk mice. Prior to transgene expression, motor axon growth proceeds well beyond the normal synaptic zone in the muscle. Motor axon growth halts and the differentiation of nerve terminals begins in Musk−/−; MCK-Musk and Musk+/−; MCK-MMT mice, only after Musk, or MMT, is expressed from the MCK regulatory region. These results confirm prior studies of Musk and agrin mutant mice and provide direct evidence that Musk regulates the organization, or synthesis of a stop signal(s) for axon growth and nerve terminal differentiation.

The C terminus of Musk contains a binding-site for PDZ domain-containing proteins. Clustering of AChRs, at least in cultured muscle cells, however, is not dependent upon this sequence, as agrin stimulates AChR clustering in myotubes expressing a C-terminal Musk mutant (Zhou et al., 1999). As the MCK-Musk transgene, which is studied here, is a fusion between Musk and GFP, this gene fusion encodes a protein that lacks a C-terminal binding site for PDZ domains. Because this transgene fully rescues the presynaptic and postsynaptic deficits of Musk mutant mice, our experiments indicate that recruitment of PDZ domain-containing proteins to the C-terminus of Musk is not required for Musk to stimulate presynaptic and postsynaptic differentiation in vivo.

In chick and Xenopus, Musk expression is not restricted to skeletal muscle. In chick embryos, Musk is expressed transiently in the brain and liver (Ip et al., 2000), and in Xenopus embryos, Musk is expressed widely in the CNS (Fu et al., 1999). These findings raised the possibility that low and/or transient Musk expression in motoneurons of mice may have escaped attention and that a loss of motoneuron-derived Musk could be responsible for the presynaptic deficits in Musk mutant mice. We find that muscle-specific expression of Musk is sufficient to restore presynaptic as well as postsynaptic differentiation in Musk mutant mice. Thus, these data strongly support the idea that activation of Musk in skeletal muscle is required to initiate a signaling pathway that leads to production and/or clustering of a retrograde signal for presynaptic differentiation.

The steps that follow Musk activation and that lead to neuromuscular synapse formation are poorly understood (Mittaud et al., 2001; Mohamed et al., 2001; Smith et al., 2001; Weston et al., 2000). Our results indicate that phosphorylation of the Musk juxtamembrane region initiates a signaling pathway that regulates nerve terminal differentiation, synapse-specific transcription and clustering of postsynaptic proteins. Thus, the Musk juxtamembrane region, including a single phosphotyrosine docking site, controls multiple pathways leading to presynaptic and postsynaptic differentiation in vivo.

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