# Muscle atrophy in Titin M-line deficient mice

J. PENG<sup>1,†</sup>, K. RADDATZ<sup>2,†</sup>, S. LABEIT<sup>3</sup>, H. GRANZIER<sup>1</sup> and M. GOTTHARDT<sup>1,2,\*</sup><br><sup>1</sup> Department of Veteringry and Comparative Anatomy, Pharmacology and Physiology, Washi

<sup>1</sup>Department of Veterinary and Comparative Anatomy, Pharmacology and Physiology, Washington State University, Wegner Hall, Room 205, Pullman, WA, 99164-6520, USA; <sup>2</sup>Max Delbrück Center for Molecular Medicine (MDC) Berlin Buch, Robert Rössel Str. 10, 13125, Berlin, Germany; <sup>3</sup>Institut für Anästhesiologie und Operative Intensivmedizin, Universitätsklinikum, 68135, Mannheim, Germany

#### Abstract

We investigated the response to deletion of the titin M-line region in striated muscle, using a titin knockout model and a range of techniques that include histology, in situ hybridization, electron microscopy, and 2D gel analysis. We found that the loss of titin's kinase domain and binding sites for myomesin and MURF-1 causes structural changes in the sarcomere that proceed from the M-line to the Z-disc and ultimately result in disassembly of the sarcomere. Disassembly goes along with central localization of nuclei (a hallmark for muscular dystrophy), up-regulation of heat-shock proteins, and induction of proteasome activity. While fiber type composition does not change in soleus and extensor digitorum longus muscle, fiber size is reduced. Animals die from complications of muscle atrophy at five weeks of age. In addition to the structural importance of the titin M-line region in any striated muscle, our data show how differences in M-line composition between heart and skeletal muscle affect sarcomere stability and function.

### Introduction

Titin alias connectin is a filamentous protein with a mass of 3.0–3.7 MDa, depending on the isoform type (Labeit and Kolmerer, 1995; Freiburg et al., 2000), that forms a separate myofilament system in both skeletal and cardiac muscle. Titin is, after myosin and actin, the third most abundant muscle protein (Wang, 1984) and forms a continuous elastic scaffold with multiple binding sites for sarcomeric and signaling proteins (Obermann et al., 1996; Gregorio et al., 1998; Young et al., 1998).

Close to its C-terminal M-line region, the titin polypeptide contains a kinase domain with homology to the catalytic serine/threonine kinase domain of smooth muscle myosin light chain kinase (Olson et al., 1990), as well as the kinase domain of the giant invertebrate muscle proteins twitchin (Benian et al., 1989), projectin (Ayme-Southgate et al., 1995), and stretchin-MLCK (Champagne et al., 2000). In vitro studies with a mutant kinase domain have provided evidence that telethonin, alias T-cap, is a titin kinase substrate in embryonic muscle (Mayans et al., 1998).

The ubiquitin ligase MURF-1 binds N-terminal to the titin kinase region (repeats A168/A169) (Centner et al., 2001) and has been shown to be involved in muscle atrophy (Bodine et al., 2001). Through heterodimerization with the microtubule binding protein

MURF-3 and the transcription factor GMEB-1 it is involved in differentiation and signal transduction (Spencer et al., 2000; Centner et al., 2001; McElhinny et al., 2002). A second protein binding site deleted in the titin kinase region knockout is the Ig-domain M4, which binds myomesin, a structural protein that confers stability to the M-line region during contraction (Obermann et al., 1997). Thus, the titin M-line region integrates signaling and structural functions.

To study the titin kinase region, we have generated an inducible knockout of titin's M-line Exons 1 and 2 (MEx1 and 2) using the cre-lox recombination system (Gotthardt et al., 2003). Embryonic lethality as a result of early cardiac depletion of the titin kinase region was circumvented by using the muscle creatine kinase promoter to drive expression of the Cre recombinase (MCKcre). Analysis of the cardiac phenotype revealed progressive sarcomeric disassembly with death at 5 weeks of age.

Here we investigated the skeletal muscle phenotype with progressive atrophy. With deletion of the titin kinase region, skeletal muscle displays sarcomeric disassembly with loss of striation, centralized nuclei, and, in contrast to cardiac muscle, an absence of M-line widening.

### Materials and methods

#### Animal model

Titin kinase region knockout mice have been described previously (Gotthardt et al., 2003). In brief, the

<sup>\*</sup> To whom correspondence should be addressed. Phone: +1-509- 335-7785; Fax: +1-509-335-4650; E-mail: Gotthardt@vetmed. wsu.edu

These authors contributed equally to the study.

MCKcre transgenic animals (Bruning et al., 1998) were used to target deletion of the titin kinase region (M-line exons 1 and 2) to striated muscle cells. Animals heterozygous for the MCKcre allele and homozygous for the floxed titin kinase region (MCKcre<sup>+</sup> Ti MEx1+2<sup>lox/lox</sup>) are referred to as knockout (KO) animals. The MCKcre negative homozygous floxed titin kinase region animals (Ti  $MEx1 + 2^{\log/\log}$ ) contain intronic lox sites and express wildtype titin protein. They are indistinguishable from wildtype animals and were used as controls (WT). Breedings (Ti  $MEx1 + 2^{\log/\log} \times MCK$ cre<sup>+</sup> Ti  $MEx1 + 2$ lox/lox) resulted in KO and control making up 50% each of the offspring.

All animal experiments were carried out under protocols approved by the Washington State University Institutional Review Committee for animal use and followed the NIH Guidelines, ''Using Animals in Intramural Research''.

### SDS-agarose electrophoresis

Vertical SDS agarose gel electrophoresis (VAGE) was used to separate titin isoforms (Warren et al., 2003b).

#### In situ hybridization

In situ hybridization was performed with  $\alpha$ -<sup>33</sup>P labeled probes directed against sequences shown in Figure 2A) as described previously (Trommsdorff et al., 1999).

### Fiber typing

Fibers types were determined using 8-µm serial cryosections of triceps surae and quadriceps muscles that were immuno-stained with a monoclonal antibody that recognizes MHC type I isoform (NOQ7.5.4D, 1:16,000; Sigma, St. Louis, MO), and MHC type IIa (SC-71, 1:1200; generously provided by Dr. S. Schiaffino, Padva, Italy) as described elsewhere (Chin et al., 1998).

#### 2D gel electrophoresis

Quadriceps from 30-day-old mice were dissected and immediately frozen in liquid  $N_2$ . Tissue powder was resuspended in 2D-lysis buffer (8 M urea, 2 M thiourea, 4% w/v CHAPS, 10 mM DTT and 1 mM PMSF). After sonication protein lysates were centrifuged for 30 min at  $4^{\circ}$ C and 50,000 rpm. Isoelectric focusing was performed with IPG strips (Amersham Biosciences, Uppsala, Sweden; 18 cm, pH 3–10L) according to the manufacturer's instructions, with the MultiphorII unit (Amersham Biosciences) employing the following voltage profiles: linear increase from 0 to 300 V for 30 min, 300 V for 30 min, increase from 300 to 500 V for 1 h, 500 V for 1 h, linear increase from 500 to 3500 V for 1.5 h, and a final phase of 3500 V for 4.5 h. After equilibration of the gels strips with DTT and iodoacetamide the second dimension was

performed in polyacrylamide gels of 12.5% T and 2.6% C. The gels were then stained with silver nitrate (Bloom and Goodpasture, 1976) or with Colloidal Coomassie G 250 (Amersham). The gel images were analyzed with the software Delta2D (Decodon) as described previously (Bernhardt et al., 1999). Coomassie stained proteins were identified by MALDI/MS peptide mass fingerprinting at the Institute of Microbiology, EMAU, Greifswald, Germany (Eymann et al., 2004).

#### Results

### Histology

The morphological correlate of muscle weakness in titin KO mice is the disassembly of the sarcomere, which is revealed at the light microscopy level. Fibers that lack titin's kinase region do not maintain their striated appearance (Figure 1A, arrows). The myopathic changes in skeletal muscle at 4 weeks of age range from ripples or vertices in striation to overt myopathy with loss of striation and centralized nuclei (Figure 1A b, c). Fibers are reduced in size when compared to control animals, in all skeletal muscle types investigated (quadriceps, diaphragm, and triceps surae). Overall, the diaphragm is less affected (as judged by a relatively normal striation pattern and peripherally located nuclei; Figure 1A e, f) than triceps surae and quadriceps muscle. The low magnification overview of KO triceps surae muscle shows a reduced and uneven fiber diameter with the majority of the nuclei centralized (Figure 1B).

#### Ultrastructure

Electron microscopy of KO animals revealed areas of intact sarcomeres as well as sarcomeres in various stages of disassembly. Unlike in the sarcomeres of titin deficient hearts, we could not detect progressive M-line widening with increased levels of mutant titin (Gotthardt et al., 2003). Three sarcomeres at different levels of disassembly show the progression of the structural rearrangements (Figure 1C, white arrow). While the left sarcomere (line 1) is largely intact, the right sarcomere (line 3) shows a pale M-line, and the central sarcomere (line 2) displays complete disassembly of the M-line. Major disassembly with filaments at various angles is present at the left. Only some residual Z-discs of various shapes (black areas) remain.

### Correlation of titin kinase expression and myopathy

Expression of titin and proper splicing after the deletion of MEx1 and MEx2 is documented by RT-PCR and sequencing of the splice junction (data not shown) and in situ hybridization with probes that detect the region 5¢, within, and 3¢ of the floxed M-line exons (Figure 2A). In wildtype muscle all three exons are



Fig. 1. Morphology of titin deficient striated muscle. (A) Hematoxylin and eosin staining of longitudinal sections from striated muscle (quadriceps a,b,c; diaphragm d,e,f). Knockout quadriceps at 4 weeks of age displays various stages of disease from atrophy with reduced fiber diameter (b vs. a and d vs. e – arrows) to overt myopathy with centralized nuclei (c, arrows). Disassembly of sarcomeres is revealed by loss of striation in individual muscle fiber segments (small circle, compared to striation in the unaffected area in the large circle). The diaphragm shows signs of atrophy, but overall a less severe phenotype than quadriceps muscle. (B) Lower magnification of triceps surae muscle shows the extend of myopathy reflected in the distribution of nuclei (peripheral vs. centralized). (C) Ultrastructure of knockout quadriceps. Sarcomere disassembly progresses from M-line (M) to Z-disc (Z) and ends in complete disarray with intersecting filaments (black arrows). Three lines of sarcomeres at various stages of disassembly are labeled 1–3.

expressed (Figure 2B). In knockout muscle only nuclei that are not recombined produce titin mRNA containing MEx1 (white dots in Figure 2B f). The Mex-1/2 flanking exons (pre-kinase and post-kinase) are expressed in knockout animals indicating proper splicing of the kinase region deficient mRNA. Compared to the wildtype sections, knockouts show increased levels of titin transcripts (compare Figure 2B a and e / c and g).

# Titin expression

At the protein level, recombination is >95% complete in skeletal muscle of animals of 4 weeks of age (Figure 3). Different splice isoforms are present in skeletal muscle of WT animals (Figure 3A, left). In KO animals all isoforms are truncated (as suggested by their increased mobility) revealing that splicing does not seem to be affected in the kinase deficient animals.

### Fiber typing

We used transversal sections of soleus and extensor digitorum longus (EDL) muscle as examples of muscles that are mainly composed of type 2 fibers





Fig. 2. Deletion of the titin kinase region. (A) Location of in situ probes before, in, and after the kinase domain, which is contained in M-line exon 1 (MEx1). (B) In situ hybridization (dark field) with S35 labeled antisense RNA probes demonstrated that in skeletal muscle the titin kinase region is deleted after cre-lox recombination (f), while the flanking exons (pre- and post-kinase) are expressed at increased levels compared with the wildtype (a vs. e and c vs. g). Only few unrecombined nuclei remain in the knockout (white spots in (f). (C) In situ hybridization – bright field. Fibers that still express the kinase domain (a) show normal localization of nuclei, while fibers that lack the kinase domain contain centralized nuclei, a sign for myopathy and regeneration.

(EDL) and type 1 and type 2 fibers (soleus muscle). With progressive loss of the titin kinase region, there is no obvious change in fiber composition but fiber size is generally reduced in the knockout animals (Figure 4).

#### Biochemical changes

After 30 days, muscle weakness is accompanied by altered expression of heat-shock proteins, components of the proteasome, and cytoskeletal proteins. Three knockout animals and three controls were compared by 2D gel analysis (Figure 5). Immobilin Dry Strips (Amersham biosciences) pH 3–10 were used for iso-

electric focusing to separate  $\sim 600$  proteins, of which 177 were identified by mass spectrometry. For comparison of knockout and wildtype tissue we used silver staining because of its higher sensitivity compared to Coomassie staining. Differentially regulated proteins were pseudocolored using the Decodon Delta2D software (blue – wildtype; orange – knockout; black – no change). We found consistent upregulation of the heat-shock proteins alpha-B crystalline, HSP27 (HSPB1) and cvHSP (HSPB7). The expression of proteasome subunits alpha type 1 and beta type 7 is another late change in development of the knockout phenotype visible at 30 days of age. These changes in protein expression have been associated with denerva-



Fig. 3. SDS agarose gel electrophoresis of control and knockout muscle. Excision of the titin kinase region results in expression of titin with reduced mobility. In quadriceps of 29-day-old animals recombination is almost complete. Both full length titin (T1) and its degradation product (T2) show increased mobility in the knockout. The double band for T1 and T2, respectively represents different isoforms. After 4 weeks, wildtype protein is almost completely replaced by kinase deficient titin in the knockout animals.

tion, starvation, glucocorticoid, and disuse atrophy (Medina et al., 1995; Auclair et al., 1997; Stevenson et al., 2003).

#### **Discussion**

# Titin deficiency

The analysis of titin *in vivo* has mainly relied on differences in titin isoform expression between species, between muscle types, in embryonic development vs. adult tissue, and in health vs. disease (Trombitas et al., 2000; Neagoe et al., 2003; Warren et al., 2003a; Lahmers et al., 2004; Nagueh et al., 2004). Added to this has the availability of naturally occurring mutants such as the MDM mouse (muscular dystrophy with myositis), mutants derived from genetic screens in drosophila or zebrafish, and the recently identified families with cardiomyopathy and muscular dystrophy due to titin deficiency (Machado and Andrew, 2000; van der Ven et al., 2000; Garvey et al., 2002; Gerull et al., 2002; Hackman et al., 2002; Xu et al., 2002). The resulting phenotypes affect skeletal muscle alone (Tibial muscular dystrophy TMD; MDM mice), exclusively the heart (zebrafish mutants Pickwick<sup>m171, m242, m740, m186</sup>, patients with cardiomyopathy due to titin mutations), or both (PickmVO62H mutation in zebrafish). Except for mutations affecting heart specific exons, such as the zebrafish Pickm<sup>171</sup> mutant with a T to G transversion in the N2B exon it has been difficult to predict which type of muscle would be affected.

#### Myopathy in the titin kinase region mutants

Knockout technology has provided a means to target mutations to specific regions in the genome and generate additional model systems to study titin in vivo and in tissue culture cells (Gotthardt et al., 2003; Miller et al., 2003). To circumvent embryonic lethality, we have used the cre-lox recombination system to study titin's M-line region, which combines structural and



Fig. 4. Fiber typing. Immunostaining of striated muscle titin kinase region deficient mice and cre negative control mice at 4 weeks of age. Staining of soleus and extensor digitorum longus (EDL) muscles shows a reduced fiber size in the knockout animals, but essentially no differences in fiber composition between control and knockout mice. Size bar corresponds to 200 um.



Fig. 5. Heat-shock response and signs of atrophy in the titin deficient muscle. Representative 2D gel  $(n=6)$  comparing the wildtype (blue) and knockout (orange) protein expression. Among the identified proteins upregulated in the knockout are heat-shock proteins (HSPB1, HSPB7, alpha-B crystalline [CRYAB]), and proteasome subunits alpha type 2 and beta type 7 (PSMA2, PSMB7).

signaling functions. We have not been able to separate the two in the adult organism, since tissue specific knockout technology is confined to exon/intron boundaries. Thus, we cannot exclusively target the kinase region but produce a combined phenotype with deletion of signaling as well as adaptor domains.

Recombination of the kinase region advances over time so that wildtype titin is replaced by mutant titin. By *in situ* hybridization and sequencing of the splice junction we have verified that the MEx1 and -2 deficient mRNA is spliced properly, maintaining the M-line region, and that recombination is virtually complete at 4 weeks of age. The resulting myopathy with loss of striation in the histological sections and centralized nuclei in affected fibers leads to atrophy and death before the animals reach sexual maturity.

#### Sarcomere disassembly

Our titin M-line deficient animals assemble intact sarcomeres, which disintegrate upon deletion of the kinase region. In contrast to the heart, quadriceps muscle does not show widening of the M-line region with progressive replacement of wildtype with mutant titin (Gotthardt et al., 2003). This can be attributed either to differences in mechanical stress with the continuous beating heart vs. the less frequently used skeletal muscle or to differences in M-line structure: in contrast to skeletal muscle, cardiac muscle expresses the LIM-protein DRAL/FHL-2, which is not only involved in cardiac signal transduction (Purcell *et al.*, 2004), but also coordinates metabolic enzymes at the M-line region (Lange et al., 2002) and is involved in myocyte differentiation (Martin et al., 2002).

Structural differences in skeletal muscle M-lines exist between fiber types, which are classified as type I (oxidative, slow) and type II (glycolytic, fast) and vary in metabolism, fatique, and force generation (Thomason and Booth, 1989; Olson and Williams, 2000). The EH-splice isoform of the M-line protein myomesin is expressed almost exclusively in type I fibers and the embryonic heart (Agarkova et al., 2004) while all M-bands in type II fibers contain M-protein (Grove et al., 1989). Skeletal muscle can switch fiber type i.e. in response to changes in motor nerve activity or calcineurin signaling (Schneider et al., 1999; Olson and Williams, 2000). To determine if the composition of the sarcomeric M-line affects the phenotype, we investigated if different fiber types would show altered phenotypical changes. In the titin kinase region knockout animals, there is atrophy in type I as well as in type II muscle fibers and we do not see major changes in fiber type (Figure 4). The uneven fiber diameter in atrophic muscle may reflect the kinetics of recombination, which results in different levels of deficient titin protein around individual nuclei.

In skeletal muscles disassembly of the sarcomere seems to proceed from M-line to Z-disc, which is maintained even in areas with loss of thick filament and disarray of thin filament structures (as seen in Figure 1C). Thus, disassembly is a local event and depends on the titin M-line, which covers signaling and structural functions. The underlying mechanism might involve the titin binding protein myomesin, a structural M-line protein (Obermann et al., 1997), the titin kinase region, implied in the regulation of sarcomere assembly (Mayans *et al.*, 1998), or the titin binding protein MURF-1, a ubiquitin ligase involved in atrophy (Dehoux et al., 2003).

Myomesin integrates titin into the sarcomeric M-line. The reduced stability of the M-line might be secondary to loss of the myomesin binding site, aggravated by mechanical strain. The relevance of the titin kinase domain for the myopathy phenotype is still unclear. While it has been implied in sarcomere assembly through its possible substrate T-cap/telethonin (Mayans et al., 1998), the titin kinase function could so far not be confirmed in vivo.

# Heat-shock response and protein degradation after excision of the titin M-line

To investigate the mechanism underlying the myopathy phenotype in our M-line deficient animals, we used 2D-PAGE to investigate changes in protein expression.

The most striking changes were upregulation of heatshock proteins such as alpha-B crystalline and HSPB1/ HSPB7 as well as proteasome subunits. These features have been described in various atrophy models ranging from denervation, and disuse atrophy to starvation and glucocorticoid treatment (Medina et al., 1995; Auclair et al., 1997; Stevenson et al., 2003). In denervation atrophy abundant disused proteins are ubiquitinated resulting in both increased degradation (proteasomal activity) and aggregation with induction of heat-shock proteins (Kato et al., 2002). In our knockout animals changes in titin's M-line region would cause destabilization of the sarcomere and failure to properly assembly newly synthesized sarcomeric proteins. The fate of these proteins would be aggregation with induction of heat-shock proteins or degradation with increased proteasome activity.

It is tempting to speculate, that the mislocalization of the ubiquitin ligase MURF-1 in our knockout animals (Gotthardt et al., 2003) could contribute to the development of the phenotype. With deletion of M-line exons 1 and 2, the MURF-1 binding site N-terminal to the kinase region is excised resulting in mislocalization of MURF-1 as demonstrated by electron microscopy (Gotthardt et al., 2003). The increased level of ''free'' MURF-1 might result in increased ubiquitination, activation of the heat-shock response, and increased protein degradation.

In summary, multiple molecular changes in the Mline titin deficient animals affect the development and progression of the myopathy phenotype. These include decreased M-line stability, altered signal transduction, as well as increased protein degradation. Future experiments will have to dissect the respective contribution of structural and signaling functions to understand

how sarcomere assembly, maintenance, and degradation are regulated.

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