ABSTRACT

Satellite cells are maintained in an undifferentiated quiescent state, but during muscle regeneration they acquire an activated stage, and initiate to proliferate and differentiate as myoblasts. The transmembrane protein teneurin-4 (Ten-4) is specifically expressed in the quiescent satellite cells; however, its cellular and molecular functions remain unknown. We therefore aimed to elucidate the function of Ten-4 in muscle satellite cells. In the tibialis anterior (TA) muscle of Ten-4-deficient mice, the number and the size of myofibers, as well as the population of satellite cells, were reduced with/without induction of muscle regeneration. Furthermore, we found an accelerated activation of satellite cells in the regenerated Ten-4-deficient TA muscle. The cell culture analysis using primary satellite cells showed that Ten-4 suppressed the progression of myogenic differentiation. Together, our findings revealed that Ten-4 functions as a crucial player in maintaining the quiescence of muscle satellite cells.

SIGNIFICANCE STATEMENT

The transmembrane protein teneurin-4 (Ten-4) is specifically expressed in the quiescent satellite cells; however, its cellular and molecular function remains unknown. Our results provided evidences that Ten-4 possesses a suppressive function in the satellite cell quiescence and myogenic differentiation. This is the first report to our knowledge that demonstrates the biological function of Ten-4 in muscle satellite cells. Our findings will facilitate a better understanding of the molecular mechanism of muscle satellite cell biology.

INTRODUCTION

Skeletal muscle is essential for the physical support of the animal body and for the proper body movement. In human, the mass of the skeletal muscle occupies ~38% and ~30% of the total body mass of men and women, respectively [1]. Defects in the formation and/or maintenance of skeletal muscle tissues, in some cases, cause severe disorders, such as muscular dystrophy and muscle atrophy [2]. During embryonic development, somites differentiate to the sclerotome and dermomyotome, and the dermomyotome tissue gives rise to myogenic progenitor cells, which are specified with the expression of paired box transcription factors Pax3 and Pax7 [3–5]. Subsequently, myogenic progenitor cells differentiate into myoblasts, positive for the transcription factor myogenic differentiation 1, Myod1/MyoD. The fusion of numerous myoblasts results in the formation of cylindrical and multinucleated myofibers abundant with filaments of actin and myosin heavy chains (MHCs). The myofibers are eventually surrounded by basement membranes and matured as functional units [6].

Muscle satellite cells, myogenic stem cells, are found between a myofiber sarcolemma and the basement membrane, accompanying with the expression of its marker Pax7 [6, 7]. Satellite cells exert their function at two main occasions, the postnatal development and the regeneration process. During postnatal stages, satellite cells contribute to the growth and maturation of myofibers, by supplying additional myonuclei [8]. Satellite cell function is critical at the stages. In this sense, in Pax7 knockout mice, satellite cell number is dramatically reduced a few weeks after birth, and as a result, myofibers have small diameter and the muscle weakens [9–11]. In the adult skeletal muscle, satellite cells remain in the quiescent phase. However, once the muscle tissue is damaged, satellite cells become activated and begin to proliferate, followed by differentiation into myoblasts. Then, multinucleated myofibers are formed to complete the tissue repair [12]. Some of satellite cells do not
commit to the differentiation process, but self-renew to maintain their own population [12]. Ablation of CSL (CBF1/suppressor of hairless/LAG-1) (RBP-J) or Hesr1 and Hesr3, which are key molecules in Notch signaling for the quiescence of satellite cells, forces to exit from the quiescent phase, and promotes proliferation and differentiation. This causes impairment in the tissue repair [13–16]. In addition, microRNA-489 also maintains the quiescent phase of satellite cells [17]. However, the detailed molecular mechanism for the satellite cell quiescence is poorly understood.

Teneurin-4 (Ten-4) is a member of the teneurin (Ten-m/Odz) family that encodes type II transmembrane proteins [18]. Ten-4 is highly expressed in the central nervous system, but its expression is observed in various tissues, including mesenchymal tissues [19]. During cartilage development, Ten-4 is expressed in the mesenchymal condensation area, where chondrogenic stem cells reside, and suppresses chondrogenic differentiation [20]. In skeletal muscle, the specific expression of Ten-4 is found in quiescent satellite cells, and it disappears after the activation of satellite cells during the tissue regeneration process [21, 22]. However, the cellular and molecular functions of Ten-4 in satellite cells remain unknown.

In this study, we elucidated the biological function of Ten-4 in muscle satellite cells. We analyzed the musculature in the skeletal muscle tissue of Ten-4-deficient (∼/−) mice with/without cardiotoxin (CTX) injection and the character of cultured satellite cells isolated from the mutant mice. We found defects in the myofiber formation in Ten-4/− mice and increased activation of Ten-4-deficient satellite cells. Our findings present Ten-4 as a novel player in the satellite cell biology.

## Materials and Methods

### Mice

The Ten-4-deficient furue mouse line was kindly provided by Dr. Yoshihiko Yamada from NIDCR, NIH [23]. Littermates or age-matched mice between different genotypes were used for experiments 8–12 weeks after birth. All procedures for experimental animals were approved by the Institutional Animal Care and Use Committees of Tokyo Medical and Dental University and Keio University.

### Cryosections

Tibialis anterior (TA) muscles were dissected out and frozen in liquid nitrogen-cooled isopentane (Wako, Osaka, Japan, www.wako-chem.co.jp). Using a cryostat (Leica, Wetzlar, Germany, www.leica-microsystems.com), the frozen TA muscles were sectioned transversely at a 10 μm thickness, and sections from the widest part in the TA muscles were attached on MAS-coated slide glasses (MATSUNAMI, Kishiwada, Japan, www.matsunami-glass.co.jp). The cryosections were kept at −80°C until they were used for immunostaining.

### Immunostaining

Cryosections described above were used for immunohistochemistry. For immunocytochemistry, primary satellite cells were cultured on eight-well chamber slides (MATSUNAMI) coated with Matrigel (BD Biosciences, San Jose, California, www.bd.com). Tissue sections or cells were fixed in 4% formaldehyde in PBS for 10 minutes at room temperature, and then permeabilized with 0.2% Triton X-100 (Sigma-Aldrich, St. Louis, Missouri, www.sigmaaldrich.com) in phosphate buffered saline (PBS) for 15 minutes at room temperature. After blocking with Power Block Universal Blocking Reagent (BioGenex, Fremont, California, http://biogenex.com Laboratories) or M.O.M. kit (Vector Laboratories, Burlingame, California, www.vectorlabs.com), the fixed cells were incubated with primary antibodies overnight at 4°C. After washing, bound primary antibodies were labeled with fluorescence-conjugated secondary antibodies for 1 hour at room temperature. The immunostained samples were mounted with Mounting medium for fluorescence with DAPI (Vector Laboratories). Primary and secondary antibodies were as follows: anti-laminin α2 (Sigma-Aldrich), anti-Pax7 (Developmental Studies Hybridoma Bank, Iowa City, Iowa, http://dshb.biology.uiowa.edu), anti-Ki67 (Leica or BD Biosciences), anti-MHC (Leica), and mouse/rabbit/rat IgG–Alexa488, -Alexa594, or Alexa647 (Life Technologies, St. Aubin, France, www.lifetech.com).

### Muscle Injury

To induce regeneration of skeletal muscle, mice were anesthetized with isoflurane, and hairs in their hind limbs were shaved. One hundred microliters of CTX (10 μM in 0.9% NaCl; Sigma-Aldrich) was injected into the TA muscle using a 29-gauge needle. Seven or fourteen days after injections, mice were euthanized and the frozen tissue sections were prepared for the analysis as described above.

### Quantification of Myofibers and Satellite Cells on Immunostained Tissue Sections

Immunofluorescent images of laminin α2 chain were taken, and myofibers surrounded by the laminin α2 signal in TA cross-sections were analyzed using the MetaMorph 7.5 software (Molecular Devices, Wokingham-Berkshire, United Kingdom, www.moleculardevices.com). The signal was thresholded, and the number of myofibers in whole areas of the cross-sections was counted using the integrated Morphometry Analysis program of the software. The thresholded images were also used for measurement of areas of individual myofibers by the program. Five hundreds to one thousand fibers per mouse were analyzed for the measurement of individual fiber areas. For counting satellite cells, cells positive for Pax7 and DAPI staining and located between a myofiber and the laminin α2 signal were counted as satellite cells. The number of satellite cells per 100 myofibers was measured in each genotype.

### Flow Cytometric Analysis of Muscle Satellite Cells

Skeletal muscles from both fore-limbs and hind limbs were dissected out and digested with 0.2% collagenase type II (Worthington Biochemical CorporaAon, Lakewood, Washington, www.worthington-biochem.com) for 1 hour at 37°C. Then, the digested tissue was filtered through 100 μm- and 40 μm-cell strainers (BD Biosciences). The filtered mononuclear cells were stained with phycoerythrin (PE)-conjugated anti-CD31 (BD Biosciences), PE-conjugated anti-CD45 (BD Biosciences), FITC-conjugated anti-Sca-1 (BD Biosciences), and biotinylated SM/C-2.6 antibodies [24] on ice for 30 minutes.
After washing, streptavidin-allophycocyanin (BD Biosciences) was added to the cells labeled with biotinylated SM/C-2.6 antibody and incubated on ice for 30 minutes. All the cells were resuspended in HBSS (–) and propidium iodide. Cell sorting was performed using MoFlo flow cytometer (Beckman, Brea, California, www.beckmancoulter.com), and CD31+CD45−cells containing 20% fetal bovine serum (Sigma-Aldrich), 1% Chick embryo extract (U.S. Biological, Swampscott, Massachusetts, www.usbio.net), 100 units/ml penicillin and 100 µg/ml streptomycin (Life Technologies), and 5 ng/ml basic-FGF (ReproCell, Yokohama, Japan, www.reprocell.com) under 5% CO2 at 37°C. For differentiation assay, satellite cells were cultured in differentiation medium consisting of DMEM with GlutaMAX (Life Technologies) containing 20% fetal bovine serum (Sigma-Aldrich), 1% Chick Embryo Extract (U.S. Biological, Swampscott, Massachusetts, www.usbio.net), 100 units/ml penicillin and 100 µg/ml streptomycin (Life Technologies), and 5 ng/ml basic-FGF (ReproCell, Yokohama, Japan, www.reprocell.com) under 5% CO2 at 37°C. For differentiation assay, satellite cells were cultured in differentiation medium consisting of DMEM with GlutaMAX supplemented with 5% horse serum (Life Technologies), 100 units/ml penicillin, and 100 µg/ml streptomycin.

Primary Culture
Isolated satellite cells were plated on plastic dishes or glass chamber slides coated with Matrigel. For proliferative condition, satellite cells were cultured in Dulbecco’s modified Eagle’s medium (DMEM) with GlutaMAX (Life Technologies) containing 20% fetal bovine serum (Sigma-Aldrich), 1% Chick Embryo Extract (U.S. Biological, Swampscott, Massachusetts, www.usbio.net), 100 units/ml penicillin and 100 µg/ml streptomycin (Life Technologies), and 5 ng/ml basic-FGF (ReproCell, Yokohama, Japan, www.reprocell.com) under 5% CO2 at 37°C. For differentiation assay, satellite cells were cultured in differentiation medium consisting of DMEM with GlutaMAX supplemented with 5% horse serum (Life Technologies), 100 units/ml penicillin, and 100 µg/ml streptomycin.

RESULTS
Defects in the Formation or Maintenance of Myofibers and Satellite Cells in Ten-4-Deficient Mice
For analyses of the Ten-4 function in satellite cells, we used the furue (fur) mutant mouse line whose Ten-4 expression is absent because of a transgene insertion in the Ten-4 gene (Odz4/Tenm4) [23]. The mutant fur/fur mice are hereafter referred to as Ten-4−/− mice. Since one of main functions of satellite cells is promoting muscle growth after birth, we first analyzed the body size and weight of Ten-4−/− mice during postnatal stages. At postnatal day (P)0–1, no difference in body weight between wild-type (WT), Ten-4 heterozygous (+/−), and Ten-4−/− mice was observed. However, a delay of the body growth was found at P2 and became obvious in Ten-4−/− mice during postnatal stages (Fig. 1A). The body size of Ten-4−/− mice was smaller than that of WT and Ten-4+/− mice at P5 (Fig. 1B). These results suggest that the postnatal body growth of Ten-4−/− mice was interfered.

We next examined weight and musculature of the TA muscle of adult Ten-4−/− mice (8–12-weeks old). The TA weight of Ten-4−/− mice was lighter than that of WT mice (Fig. 2A). To analyze the myofiber formation in the TA muscle tissue, immunostaining of laminin a2 chain, a basement membrane protein, on transverse sections of TA was performed. The whole area of the TA sections of Ten-4−/− mice was smaller, compared with that of WT (Fig. 2B: left images, Fig. 2C: left graph). Less number of the myofiber number was observed in the whole area of the Ten-4−/− TA section, relative to that in the WT section (Fig. 2B: left images, Fig. 2C: center graph). In addition, the area of individual myofibers was reduced in Ten-4−/− mice (Fig. 2B: right images, Fig. 2C: right graph). These observations indicate that Ten-4 was required for production of the normal number and growth of myofibers.

We further analyzed the population of satellite cells by immunostaining of Pax7, a marker for satellite cells, on TA transverse sections. In Ten-4+/− and Ten-4−/− tissues, the number of Pax7-positive satellite cells per 100 myofibers was attenuated (Fig. 3A). Moreover, the population of TA satellite cells was assessed by flow cytometry using the antibody SM/C-2.6, which recognizes satellite cells specifically
Figure 2. Defects in the muscle tissue of Ten-4-deficient mice. (A): Size and weight of the TA muscle from WT, Ten-4+/−, and Ten-4−/− mice. The TA muscle was dissected out and weighed. The weight of the WT TA muscle was set as 1.0. Error bars, SEM; **, p < .01. Scale bar = 10 mm. (B): Immunostaining of laminin α2 (green) in the TA muscle tissue from WT, Ten-4+/−, and Ten-4−/− mice. DAPI staining (blue) was used to visualize nuclei. Scale bar = 500 μm in lower magnification images (left); 100 μm in higher magnification images (right). (C): The whole area (left), the total myofiber number (center), and the area of individual myofibers (right) in transverse TA sections. The individual myofiber area of WT mice was set as 1.0. Error bars, SEM; **, p < .01. Abbreviations: TA, tibialis anterior; WT, wild type.
After mononuclear cells negative for CD31 and CD45, markers for endothelial cells and lymphocytes/leukocytes, respectively, were gated in the total cells from TA tissues, the population of satellite cells, positive and negative for SM/C-2.6 antibody and Sca-1, respectively, was analyzed. We found that the percentage of Ten-4−/− satellite cell population per the total mononuclear cells, except for endothelial cells and lymphocytes/leukocytes, was significantly decreased, compared to that of WT (Fig. 3B). These data indicate that Ten-4 was required for myofiber formation and production or maintenance of satellite cells during normal development.

Defective Tissue Regeneration in the Ten-4-Deficient Muscle

The regeneration of injured tissues in skeletal muscle is another important function of satellite cells. Therefore, we carried out a muscle regeneration experiment by injection of CTX. After CTX was injected into TA muscle, myofiber formation and satellite cell population were analyzed. Seven days after the CTX injection, the weight of the TA muscle after the regeneration was diminished in Ten-4+/− and Ten-4−/− mice, in comparison with WT (Fig. 4A). The whole area of transverse TA sections and the number of regenerated myofibers, whose nuclei were centralized in the fibers, were reduced in Ten-4−/− TA muscles (Fig. 4B: left images, Fig. 4C: left and center graphs). The area of individual regenerated myofibers was smaller in Ten-4+/− and Ten-4−/− TA muscles (Fig. 4B: right images, Fig. 4C: right graph). From these results, we concluded that the deficiency of Ten-4 perturbed the tissue repair.

To address the self-renewal capacity of satellite cells, we analyzed the population of satellite cells in regenerated TA tissues after the CTX injection. Immunostaining of Pax7 showed the decreased number of satellite cells in the Ten-4−/− regenerated tissue, relative to the WT tissue (Fig. 5A). Similarly, by flow cytometric analysis using SM/C-2.6 antibody, a decrease of the satellite cell population was observed in the regenerated TA tissue of Ten-4−/− mice (Fig. 5B). In the both cases of with and without CTX injection, however, the percentages of Ten-4−/− satellite cells to WT satellite cells set as 100% were similar, and no statistical difference was observed between the TA tissues with and without CTX injection (immunostaining: with CTX: 60.7% ± 4.1%, without CTX: 57.4% ± 9.1%; flow cytometry: with CTX: 89.5% ± 3.1%, without CTX: 63.9% ± 11.5%, in Ten-4−/− mice, relative to 100% for in WT mice) (Figs. 3, 5). This evidence indicates that the satellite cell population was maintained in the Ten-4−/− muscle after the regeneration. From these observations, we found that in the Ten-4−/− TA muscle, the tissue repair was defective, albeit the self-renewal of satellite cells occurred normally.

Promoted Activation of Ten-4-Deficient Satellite Cells

To address the mechanism of Ten-4 function in satellite cells, we analyzed activation of satellite cells in Ten-4-deficient mice by immunostaining of Pax7 and Ki67, a marker for proliferative cells. We counted Pax7- and Ki67-dual positive cells as activated satellite cells (Fig. 6, arrows) in TA tissues, 7 days after CTX injection. The ratio of Pax7- and Ki67-dual positive cells to total Pax7-positive cells in Ten-4+/− and Ten-4−/−
Figure 4. Abnormalities in the regenerated muscle tissue of Ten-4-deficient mice. (A): Weight of TA muscles from WT, Ten-4+/−, and Ten-4−/− mice. TA muscles were dissected out 7 days after the injection of CTX and were analyzed. The weight of the WT TA muscle was set as 1.0. Error bars, SEM; **, p < .01. Scale bar = 10 mm. (B): Immunostaining of laminin α2 (green) in the TA muscle tissue from WT, Ten-4+/−, and Ten-4−/− mice, 7 days after the injection of CTX. DAPI staining (blue) was used to visualize nuclei. Scale bar = 500 μm in lower magnification images (left); 100 μm in higher magnification images (right). (C): The whole area (left), the total myofiber number (center), and the area of individual myofibers (right) in transverse TA sections consisting of regenerated muscle fibers, whose nuclei were in the center. The individual myofiber area of WT tissue was set as 1.0. Error bars, SEM; *, p < .05; **, p < .01. Abbreviations: CTX, cardiotoxin; TA, tibialis anterior; WT, wild type.
Promotion of Proliferation and Differentiation in the Ten-4-Deficient Cell Culture

We further analyzed Ten-4 function in proliferation and differentiation of primary satellite cell culture. Satellite cells were purified from fore-limb and hind limb skeletal muscles by flow cytometry using the SM/C-2.6 antibody (Fig. 7A). The population of Ten-4+/− and Ten-4−/− satellite cells in the skeletal muscles from fore-limb and hind limbs was less than that of WT cells (Fig. 7A), suggesting that satellite cells were reduced in not only the TA muscle but also other skeletal muscles of mutant mice. We then cultured the purified satellite cells and analyzed cell proliferation and differentiation into myotubes. When cells were cultured under the proliferative condition for 3 days, ~70% of WT cells were positive for Ki67, indicating that these Ki67-positive cells were in the activated phase, as reported previously [21]. In the Ten-4−/− culture, the number of proliferating cells, labeled with anti-Ki67 antibody, was increased, compared with in the WT culture (Fig. 7B). This data show an accelerated activation of Ten-4−/− satellite cells, and agrees with the in vivo immunohistochemistry result (Fig. 6). At this stage, most proliferating cells were positive for MyoD and represented spread and typical myoblast-like morphology (data not shown), meaning that activated satellite cells were differentiating into myoblasts. We further performed the differentiation assay into myotubes. Two days after induction of differentiation, the MHC-positive myotubes with multiple nuclei from Ten-4+/− and Ten-4−/− cells were increased compared with those from WT cells (Fig. 7C). In contrast, a decrease of MHC-positive cells with single nucleus was observed in the Ten-4+/− and −/− cultures (Fig. 7C). This abnormally early differentiation of Ten-4-deficient cells may be due to the accelerated activation in satellite cells, since Ten-4 is specifically expressed in satellite cells [22]. However, it is also possible that Ten-4 deficiency might have promoted differentiation of myoblasts into myotubes. Taken all together, Ten-4 is a novel suppressor of satellite cell activation and possibly of myogenic differentiation as well, and is required for the formation of the normal and regenerated muscle tissues.

DISCUSSION

In this study, we showed a critical role of Ten-4 in skeletal muscle satellite cells. In Ten-4−/− mice with/without the
Figure 6. Increased activated satellite cells in regenerated TA muscle from WT, Ten-4+/−, and Ten-4−/− mice. Immunostaining of laminin α2 (white), Pax7 (green), and Ki67 (red) in the TA muscle tissue of WT, Ten-4+/−, and Ten-4−/− mice, 7 days after CTX injection. Arrows indicate Pax7- and Ki67-double positive cells, and arrowheads represent Pax7-single positive cells. DAPI staining (blue) was used to visualize nuclei. The ratio of the numbers of Pax7- and Ki67-dual positive cells/Pax7-single positive cells was quantified. Error bars, SEM; *, p < .05. Scale bar = 50 μm. Abbreviation: CTX, cardiotoxin; TA, tibialis anterior; WT, wild type.
injection of CTX, the number and size of muscle fibers were smaller, and the satellite cell population was decreased. In addition, activation of satellite cells was promoted in Ten-4 

A genome-wide gene expression analysis revealed that a specific expression of Ten-4 is detected in quiescent satellite cells, and that the expression is diminished in activated satellite cells, similar to that of calcitonin receptor, a marker for quiescent satellite cells [21]. Analysis of Ten-4 expression pattern by immunostaining showed that all the quiescent satellite cells expressed Ten-4 at the neonatal stage, while the calcitonin receptor was expressed only in part of the cells [22]. These evidences indicate that Ten-4 is a new marker of quiescent satellite cells, and its expression pattern is broader than that of calcitonin receptor. Here, we found that the population of satellite cells was lower in Ten-4−/− mice than in WT mice (Figs. 3, 5). Therefore, Ten-4 was required for production and/or maintenance of the normal satellite cell population. The loss of satellite cells perturbs the postnatal muscle growth, because of defects in the myonuclear fusion, in Pax7 null mice [9–11]. Furthermore, a reduced number and size of myofibers are observed in Pax7 knockout mice after the regeneration process [9, 25]. In this study, the caliber size of myofibers and the weight of the TA muscle were decreased in both normal and regenerated cases, due to the deficiency of Ten-4 (Figs. 2, 4). Also, we found an attenuation of the postnatal body growth of Ten-4−/− mice (Fig. 1). These defects were presumably caused by the declined number of Ten-4−/− satellite cells, similar to the phenotype in Pax7 knockout mice.

There are several studies regarding the mechanism of maintaining quiescent satellite cells. In the conditional knockout mice of CSL, a key molecule of Notch signaling, and the double knockout mice of Hes1 and Hes3, target genes of Notch, a decrease of satellite cells is observed, which results in the impairment of both normal and regenerated muscle formation [13–15]. Moreover, in the satellite cells of the both cases, a spontaneous exit from the quiescent phase occurs, indicating that these pathways are necessary for maintaining the quiescence. These mutant satellite cells undergo abnormally early differentiation and fail to self-renew properly [13–15]. Our data in this study showed the increased number of Pax7- and Ki67-dual positive cells in Ten-4−/− tissue.
Ten-4+/− mice exhibited defects in several analyses shown in this study, while significant abnormalities in Ten-4+/− mice were found in all the analyses (Figs. 1–7). In regenerating Ten-4+/− TA, for instance, the weight was lighter than WT, whereas the size and the entire cross-sectioned area of TA were normal (Fig. 4). This may be due to a delay of muscle growth and maturation in the regenerating Ten-4+/− tissue. There is presumably a difference in an increase/decrease of downstream signaling and/or compensational feedback discussed below between hetero- and homozygosity of Ten-4.

The Notch signaling participates in the satellite cell quiescence [26]. Inhibition of the Notch signaling in satellite cells, as well as the deficiency of Ten-4 as shown in this study, forces to exit the quiescent state [13–15]. Brohl et al. reported that the Ten-4 expression level was significantly reduced in the double knockout mice of RBP-J and MyoD, where Notch signaling was deactivated. Furthermore, Ten-4 was upregulated in satellite cells by stimulating the Notch signaling with its ligand Dll1, suggesting that Ten-4 is a downstream molecule of the Notch signaling [27]. Our preliminary data showed that the expression levels of Notch pathway molecules, including Hesr1 and Hesr3, were elevated in Ten-4−/− satellite cells. This increased expression may have been due to a feedback. In addition, genes encoding cell adhesion molecules and basement membrane components, such as integrin alpha-7, dystroglycan, alpha-1 chain of type XVIII collagen, and alpha-2 chain of type IV collagen, were also upregulated and downregulated in satellite cells by the stimulus with Dll1 and by the deficiency of RBP-J and MyoD, respectively [27]. Adhesion to the basement membrane and the myofiber plasma membrane is crucial for the quiescence of satellite cells in their niche [28]. We have previously reported that Ten-4 regulates the activation of focal adhesion kinase, a critical downstream molecule of integrins and dystroglycan, in neural and glial cells [23, 28, 29]. Therefore, Ten-4 may play a key role in satellite cells at the quiescent state together with these proteins.

There are four members in the vertebrate teneurin family (Ten-1 to Ten-4). Some cell types express specific teneurin members, but the others express all the members [20, 29–31]. In muscle development, no detailed expression analysis of teneurin family members, except for Ten-4, has been reported. However, in a genome-wide expression analysis by microarray, Ten-3 is expressed in quiescent satellite cells, and an attenuation of its expression level is observed in activated satellite cells [21]. This expression pattern of Ten-3 is similar to that of Ten-4. In addition, both, Ten-4 and Ten-3, are expressed in somites during somitogenesis, where and when Pax3- and Pax7-positive myogenic progenitor cells appear [6, 30]. Since teneurins share a high homology in their sequence and bind homophilically or heterophilically between their family members [32], it is possible that Ten-4 and Ten-3 redundantly regulate critical signaling for the quiescence of satellite cells. The absence of Ten-4 significantly diminished the function of satellite cells, but did not eliminate it completely (Figs. 2–6). Ten-3 and/or the other teneurins may have partially compensated for the lack of Ten-4.

**SUMMARY**

In summary, our results provided evidences that Ten-4 possesses a suppressive function in the satellite cell quiescence and myogenic differentiation. This is the first report to our knowledge that demonstrates the biological function of Ten-4 in muscle satellite cells. Our findings will facilitate a better understanding of the molecular mechanism of muscle satellite cell biology.

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**AUTHOR CONTRIBUTIONS**

K.I.: conception and design, collection and/or assembly of data, data analysis and interpretation, and manuscript writing; N.S.: conception and design, financial support, collection and/or assembly of data, data analysis and interpretation, and manuscript writing; Y.M.: collection and/or assembly of data, financial support, and data analysis and interpretation; N.I. and S.T.: conception and design and data analysis and interpretation; N.K.: collection and/or assembly of data; S.F.: provision of study material or patients and data analysis and interpretation; H.O.: provision of study material or patients; C.A.: conception and design, financial
support, data analysis and interpretation, manuscript writing, and final approval of manuscript. K.I. and N.S. contributed equally to this work.

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