The Changes of Muscle FGF-6 Expression at Different Time Points after Exerciseinduced Muscle Damage

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ABSTRACT

Objective: To investigate the changes of fibroblast growth factor (FGF)-6 expression in the regeneration and repair process after exercise-induced muscle damage (EIMD) and the relationship with skeletal muscle regeneration and repair.

Methods: The expression of FGF-6 at different time points was examined by reverse transcriptionpolymerase chain reaction (RT-PCR) and immunohistochemistry staining after a downhill treadmill exercise. Skeletal muscle injury and regeneration at different times after EIMD was assessed by haematoxylin and eosin (H&E) staining.

Results: The FGF-6 protein expression was initially elevated, followed by a gradual reduction, while the changes of FGF-6 mRNA were almost all raised after the treadmill exercise.

Conclusion: The results point out that FGF-6 is closely related to skeletal muscle regeneration and repair, probably implying a dual function in muscle regeneration.

Keywords: Downhill treadmill, exercise, exercise-induced muscle damage, FGF-6, gastrocnemius, regeneration

Cambios de Expresión de FGF-6 Muscular en Diferentes Puntos del Tiempo después del Daño Muscular Inducido por Ejercicio

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RESUMEN

Objetivo: Investigar los cambios de expresión del factor de crecimiento fibroblástico (FGF)-6 en el proceso de regeneración y reparación después de daño muscular inducido por el ejercicio (DMIE) y la relación con la reparación y regeneración del músculo esquelético.

Métodos: La expresión de FGF-6 en diferentes tiempos fue examinada mediante reacción en cadena de la polimerasa con transcriptasa inversa (RT-PCR) y tinción inmunohistoquímica, después de un ejercicio de carrera descendente en cinta rodante. La lesión del músculo esquelético y la regeneración en diferentes momentos después del DMIE, fueron evaluadas mediante hematoxilina y eosina (H & E). *Resultados:* La expresión de la proteína FGF-6 fue elevada al principio, seguida por una reducción gradual, mientras que los cambios de FGF-6 mRNA fueron casi todos incrementados tras los ejercicios en la cinta rodante.

Conclusión: Nuestros resultados señalan que FGF-6 se relaciona estrechamente con la regeneración del músculo esquelético, lo que probablemente implica una función dual en la regeneración muscular.

Palabras claves: Carrera descendente en cinta rodante, ejercicio, daño muscular inducido por el ejercicio, FGF-6, gastrocnemio, regeneración

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INTRODUCTION

Exercise-induced muscle damage (EIMD) is a common phenomenon in unaccustomed eccentric exercise, which is embodied in impairment of neuromuscular function, reduction of motion range, increase of muscle soreness, localized limb swelling/oedema and the leakage of intramuscular proteins (1–3). These effects can last for several days after the initial bout and may negatively impact the athletes' ability to perform subsequent exercise training schedule or competitions (4, 5). Although EIMD can have detrimental short-term effects, it has been hypothesized that a moderate amount of damage would be most appropriate for maximizing the long-term hypertrophic adaptations (4).

The regeneration and repair of muscle is performed by satellite cells, which are situated between the basal lamina and sarcolemma of muscle fibres (6). Satellite cells are the quiescent stem cells in mature muscle and could be dramatically activated in response to exercise, muscle damage or degenerative muscle diseases for the sake of muscle growth, regeneration and repair (7). Once activated, they proliferate and evolve along with the myogenic lineage pathway to turn into myogenic precursor cells (Mpc), also called myoblasts (8). They eventually differentiate into fusion-competent myocytes that form multinucleated syncytia (myotubes), the immediate precursors of myofibres. Proliferation of activated satellite cells is a fundamental step of the regeneration process, serving the function of increasing the population of myoblasts and reconstituting the pool of reserve satellite cells (6). Many different growth factors may be involved in the regulation of the process such as members of the fibroblast growth factor (FGF) family, human growth factor (HGF), insulin-like growth factor (IGF1), leukaemia inhibitory factor (LIH) and others. Interestingly, FGF's activity is preferentially upregulated by eccentric exercise (4). Different members of the FGF family, comprising 10 subtypes, are present in muscle tissue and could be released after injury (9). The FGFs play a critical role in the recruitment of satellite cells to break quiescence and enter the proliferative phase (10). Among the FGF family, FGF-6 is the only member of this family showing relatively high expression which is restricted to the myogenic lineage in adult and developing skeletal muscle (11). Importantly, muscle regeneration was poor in FGF-6 gene knockout mice (12), whereas injection of FGF-6 accelerated regeneration of the soleus muscle (13).

There is some lack of clarity about the changes of FGF-6 expression in regeneration and repair process after EIMD. Therefore, in the present study, we caused injury to the gastrocnemius with eccentric exercise to investigate the FGF-6 gene expression at different times after EIMD.

SUBJECTS AND METHODS Animals and treatment

Male Sprague-Dawley rats, two months old and weighing 264 ± 11.94 g, supplied by Shanghai SIPPR-BK Co, were acclimatized for one week in an air-conditioned room (temperature 22 °C ± 2 °C, relative humidity $55\% \pm 5\%$ and 12-hour light/dark cycle). Drinking water and standard chow diet were supplied *ad libitum*. All experimental procedures were conducted in conformity with the National Institutes of Health's Guide for the Care and Use of Laboratory Animals (NIH Publications No. 80-23), revised in 1996, and approved by the Animal Ethics Committee of Shanghai University of Sport.

The rats were randomly divided into a non-exercise control group and an exercise group and then sub-divided by the different times of sacrifice (n = 8 in each group). Experimental procedures of the exercise group are described in the Table. Rats in the exercise group performed a downhill treadmill (-16° incline) exercise at the speed of 16 m/minute for 90 minutes, according to the method described in a previous paper (14). Subsequently, the rats were placed back in their cages with free access to food and water until the time of death. At different time points after exercise, rats were anaesthetized with 10% chloral hydrate (400 mg/kg body weight) and bilateral gastrocnemius were rapidly dissected. The left was fixed in 4% paraformaldehyde for paraffin section, while the right was frozen in liquid nitrogen and stored at -70 °C for later mRNA analysis. The gastrocnemius of the control animals was treated to the same process.

Preparation of paraffin section, histological analysis and immunohistochemical staining

After 24 hours of fixation, the specimen was then washed thoroughly with tap water and trimmed to suitable size, in the correct orientation, using a band saw. Thereafter, the specimen was dehydrated through a graded alcohol series and infiltrated with paraffin. Serial sections were taken at 5 μ m. For histological analysis, haematoxylin and eosin (H&E) stainings were processed by standard procedures (15). For

Table: Experimental procedures of the exercise group

Exercise group	Exercise time (day) 8:00 am~9:30 am	Recovery time (day)	Sampling time (day) after 9:30 am
Control			10
6 hours	12		12 (3:30 pm~5:00 pm)
12 hours	7 (8:00 pm~9:00 pm)		8
24 hours	13		14
48 hours	5	6	7
72 hours	3	4~5	6
1 week	2	3~8	9
2 weeks	1	2~14	15

immunohistochemistry, sections were stained with a rabbit polyclonal antibody against FGF-6 (purchased from Santa Cruz Biotechnology Inc, Santa Cruz, CA, USA). Five sections were randomly selected from each group and five areas were randomly captured by Olympus DX70. Subsequently, integrated optical density (IOD) was analysed by Image-Pro Plus 4.1 software.

Reverse transcription-polymerase chain reaction

Total RNA was extracted using the TRIzol reagent (Invitrogen) according to the manufacturer's instructions. The RNA concentration was determined spectrophotometrically at 260 nm and the quality of RNA was checked spectrophotometrically by using the ratio of absorbances at 260/280 nm. After extraction, the total RNA $(2 \mu g)$ was used to synthesize first-strand cDNA and the cDNA product (200 ng) was amplified by polymerase chain reaction (PCR) Express Thermal Cycler (Bio-Rad) using the following primers: GAPDH (forward) 5' - ACC ACA GTC CAT GCC ATC AC -3', GAPDH (reverse), 5'- TCC ACC ACC CTG TTG CTG TA -3', FGF-6 (forward), 5'- CAT TAA GCG ACA GCG GAG ACT -3', FGF-6 (reverse), 5'- CGT CCA TAT TTG CTC AGT GCA -3'. The amplified PCR products were separated by 1.2% agarose gel electrophoresis, and the product was finally visualized with an image analyser (Bio-Rad Gel-Doc, 2000). The GAPDH mRNA was used as an internal control and the expression of FGF-6 was normalized to respective GAPDH.

All results were expressed as means and standard errors. Differences among means were assessed by one-way analysis of variance (ANOVA) followed by StudentNewman-Keuls post hoc test. Significance level was set at p < 0.05. All calculations were performed using the SPSS 17.0 software.

RESULTS

Haematoxylin and eosin staining on the serial cross-sections of paraffin blocks of muscles from the non-exercise group displayed cell polygon, normal size, sarcolemma integrated and no inflammatory reaction in muscle interstitium (Fig. 1A). Six hours after exercise, some cells swelled and appeared round (Fig. 1B). Seventy-two hours after exercise, swelling of cells was gradually increased and aggregations of mononuclear cells were observed in the muscle interstitium and some muscle fibres (Fig. 1F). Inflammatory cell infiltration almost disappeared (Fig. 1G) and the structure of muscle fibres generally recovered normally two weeks post exercise (Fig. 1H).

As shown in Fig. 2A, the expression of FGF-6 was visualized by immunohistochemistry staining. The statistical comparison revealed that the IODs of FGF-6 were significantly elevated at the 6th and 12th hour after downhill treadmill exercise, almost close to twice the control 12-hour group (p < 0.05, Fig. 2B). Twenty-four hours after exercise, the IODs displayed a downtrend. In the 48-hour, 72-hour, one-week and two-week groups, the IODs were obviously decreased (p < 0.05; Fig. 2B).

Different from the changes of FGF-6 protein, the levels of FGF-6 mRNA were obviously increased at various time points after exercise, except for one week after exercise (Fig. 3). At the 6th and 48th hour after exercise, the levels of FGF-6 mRNA were markedly elevated, up to three times the

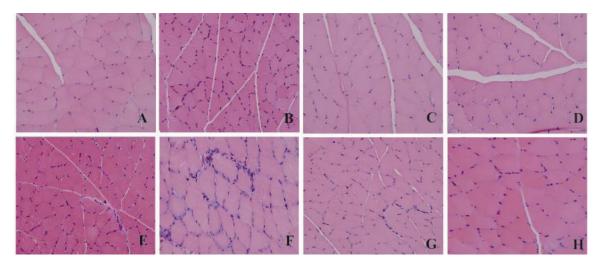


Fig. 1: Skeletal muscle injury and regeneration at different times after exercise-induced muscle damage (EIMD). Paraffin sections were stained with haematoxylin and eosin (H&E) at six hours (B), 12 hours (C), 24 hours (D), 48 hours (E), 72 hours (F), one week (G) and two weeks (H) after exercise. Muscle fibre showed sarcolemma integrated, interstice clear, no swelling and pycnosis in the non-exercise control group (A). In the exercise group, swelling was visible six hours after exercise (B) and gradually increased at 72 hours (F), accompanying inflammatory cell infiltration and muscle fibrinolysis. Regeneration was obvious after one week (G) and the sign of previous damage was scarcely detected after two weeks (H). ×400 magnifications.

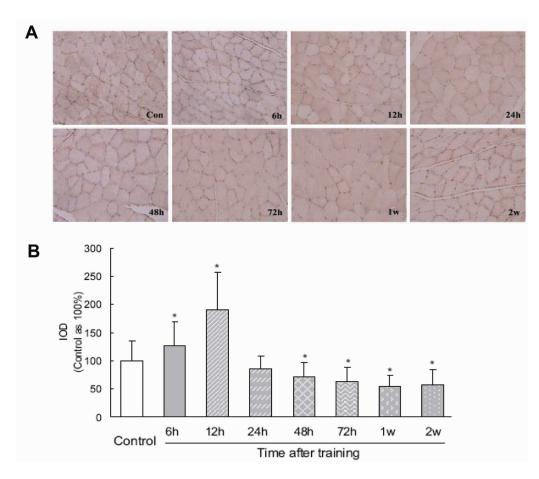


Fig. 2: The expression of FGF-6 protein in gastrocnemius at different times after exercise-induced muscle damage (EIMD). After six hours, 12 hours, 24 hours, 48 hours, 72 hours, one week and two weeks of exercise, gastrocnemius was taken and then immunohistochemistry staining for FGF-6 was performed. (A) Representative images of FGF-6 staining at ×400 magnification. (B) Statistical analysis of IOD in various groups. Data were expressed as means \pm SEM, n = 5. *indicates p < 0.05, when compared with the control group.

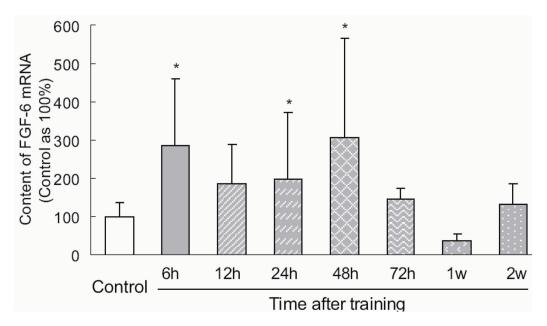


Fig. 3: The expression of FGF-6 mRNA in gastrocnemius at different times after exercise-induced muscle damage (EIMD). Data were expressed as means \pm SEM, n = 5. *indicates p < 0.05, when compared with the control group.

control group (p < 0.05; Fig. 3). As can be seen in Fig. 3, there was also a significant difference between the 24-hour group and the control group. However, the levels of FGF-6 mRNA did not significantly rise in the 12-hour, 72-hour and two-week groups.

DISCUSSION

We were interested in the gastrocnemius because this muscle could provide the driving force for many daily and sports activities, eg walking, running and jumping. Therefore, in the present study, we damaged the gastrocnemius with downhill treadmill exercise, an unaccustomed eccentric exercise followed by muscle damage (16). The morphological alterations of the muscle fibres after exercise reflected by histological observation displayed the approximate process of muscle injury and recovery (Fig. 1). At different time periods after exercise, we investigated the gene and protein changes of FGF-6. Interestingly, the tendency of change of FGF-6 mRNA disaccorded with that of FGF-6 protein. The changes of FGF-6 protein looked like a parabola, *ie* initial elevation following reduction, while the changes of FGF-6 mRNA were always raised except for the group of seven days. There was some disparity with a previous report. Armand et al (13) reported the endogenous FGF-6 protein in soleus regenerating muscle peaked at 72 to 84 hours after injection of cardiotoxin, a single injection which caused an almost complete degeneration of the myofibres within 24 hours. The disparity may be due to the different kinds and extent of muscle injury.

Activated satellite cells express several myogenic markers, such as MyoD. Our previous study suggested that the number of MyoD positive nuclei was significantly increased 24 hours after downhill running, and reached peak level after 48 hours (17). A number of studies have reported the relations between specific FGFs and satellite cell activation during muscle regeneration and hypertrophy (18). Three days after freeze-crush injury, FGF-6 as well as myogenic transcription factor MyoD mRNAs were concomitantly upregulated with the regeneration of skeletal muscle (12). Fibroblast growth factor-6 mRNA levels were significantly increased 72 hours after synergist ablation in the plantaris muscles, while MyoD mRNA levels started to increase 48 hours after synergist ablation (18). While FGF-6 mRNA levels did not change after bupivacaine administration in the plantaris muscles, MyoD mRNA started to increase 24 hours after bupivacaine administration (18). The increase of FGF-6 mRNA was very small and the expression pattern of FGF-6 mRNA did not overlap with MyoD and other myogenic or cell cycle markers. Tanaka et al (18) presumed that FGF-6 was unlikely to relate to satellite cell activation during muscle regeneration and hypertrophy.

The analyses of Fgf6 (-/-) mutant mice also gave contradictory results. In Fgf6 (-/-) mutant mice, the number of MyoD-expressing activated satellite cells was significantly reduced after injury, presumably due to a lack of

activation or proliferation of quiescent satellite cells (12). In contrast, Fiore *et al* (19) reported that skeletal muscle regeneration is not impaired in Fgf6 (-/-) mutant mice after notexin injection or crush injury. Different hypotheses had been suggested to explain the differences observed between the two mutant mice lacking FGF-6, including compensation of FGF gene expression in different genetic backgrounds (20), the different design of the targeted mutation (21) and stochastic variations (22), even in a uniform genetic background.

Recent reports suggested that FGF-6 had a biphasic effect on muscle regeneration, stimulating myoblast proliferation/migration and muscle differentiation/hypertrophy in a dose-dependent manner (13, 23). The dual role of FGF-6 in myogenesis would be the result of a balance of its action on the two receptors FGFR1 and FGFR4 (23, 24). In the present study, the expression of FGF-6 protein after EIMD was gradually increased up to a peak and subsequently reduced. Further study should be performed to clarify whether the expression pattern of FGF-6 protein after EIMD is related to the dual role of FGF-6 in myogenesis.

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