

Decreased Activity in Neuropathic Pain Form and Gene Expression of Cyclin-Dependent Kinase5 and Glycogen Synthase Kinase-3 Beta in Soleus Muscle of Wistar Male Rats

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Background: The relationship between decreased activity/neuropathic pain and gene expression alterations in soleus muscle has remained elusive.

Objectives: In this experimental study, we investigated the effects of decreased activity in neuropathic pain form on Cyclin-Dependent Kinase 5 (*CDK5*) and Glycogen Synthase Kinase-3 β (*GSK-3\beta*) gene expression in soleus muscle of rats.

Materials and Methods: Twelve male Wistar rats were randomly divided into three groups: (1) tight ligation of the L₅ spinal nerve (SNL: n = 4); (2) sham surgery (Sham: n = 4), and (3) control (C: n = 4). The threshold to produce a withdrawal response to a mechanical and thermal stimulus was measured using von Frey filaments and radiation heat apparatus, respectively. Following 4 weeks after surgery, the left soleus muscle was removed and mRNA levels were determined by real-time Polymerase Chain Reaction (PCR).

Results: Compared to control animals, L₅ ligated animals developed mechanical and heat hypersensitivity during total period of study. Soleus muscle weight as well as *CDK5* mRNA levels (less than ~ 0.4 fold) was decreased and *GSK-3\beta* mRNA levels (up to ~ 7 folds) increased in L₅ ligated animals.

Conclusions: These results showed enhanced muscle atrophy processes following peripheral nerve damage and might provide a useful approach to study underlying muscle mechanisms associated with clinical neuropathic pain syndromes.

Keywords: Neuropathic Pain; Soleus Muscle; *CDK5* Protein Kinase

1. Background

Neuropathic pain is a chronic pain defined as a pain caused by damages to or dysfunction of somatosensory system and can express itself in the following forms: allodynia, hyperalgesia, and spontaneous pain (1). Along with causing changes to nervous system, neuropathic pain can decrease physical activity levels (2, 3). Furthermore, neuropathic pain can affect structure and function of muscles through muscular atrophy (4, 5). Many studies have proven that models of neuropathic pains will be followed with muscle atrophy (5-8).

However, exact cellular mechanisms, which cause changes after nervous damages are still unknown (8). Cyclin-Dependent Kinases (Cdks) are serine/threonine protein kinases that play key roles in the regulation of cell cycle, initiation of transcription, and control of certain metabolic cascades in mammalian cells (9, 10). Cdks activity has a vital role in different functions of nerve cells such as neuronal growth and migration, cell secretion, dopamine signaling, and cytoskeletal dynamics (10). Glycogen Synthase Kinase-3 (*GSK3*) is also a serine/threonine kinase that exists in two isoforms of alpha and beta (11) and participate in the modulation of various functions

such as cell signaling, growth metabolism, and many other transcriptional regulating survival and death factors in organisms (12). In general, *Cdk5* and *GSK-3\beta* are two of the most important protein kinases involved in neuropathic pain signaling (13, 14). Furthermore, the role of these two proteins in the regulation of muscle plasticity has been clearly proved. For example, it has been shown that *GSK-3\beta* is necessary in myofibrillar protein loss system and its reduction causes preserving of contractile protein against proteolysis (15). *Cdk5* also has a key role in skeletal muscle's structure and its function so that it is effective in changing muscle phenotype and essentially these changes come from its effects on cytoskeleton proteins' structure such as actin and microtubules and regulation of myogenesis (16, 17). On the other hand, changes to expression of *Cdk5* and *GSK-3\beta* in muscular and nervous systems after numerous interferences infer such an insight that related systems change through following conditions: chronic state of motor unit activity, afferent activity level toward motor neurons, number of innervated muscle fibers by motor neurons, and metabolic level of target tissue.

2. Objectives

We set the goal of this study on considering the effects of decreased activity in the form of neuropathic pain caused by ligation of the L₅ spinal nerve on *CDK5* and *GSK-3β* gene expression in soleus muscle of Wistar male rats.

3. Materials and Methods

3.1. Animals

In the present study, 12 mature 10-week-old Wistar male rats, weighted 200 - 240 g were provided by Animal Maintenance Unit of Razi Research Center (Razi Institute Animal Center, Karaj, Iran) and transferred to Animal Laboratory of Tarbiat Modares University. All rats were kept under a controlled environment condition with mean temperature of 22 ± 3°C, dark-light cycle of 12:12 hours, relative humidity of 40% and free access to food and water ad libitum. The experimental protocols to perform this study were approved by the Ethics Committee on the use of animals of Tarbiat Modares University, Tehran, Iran. All efforts were made to minimize discomfort of the animals and reduce the number of experimental animals. All procedures conformed to the ethical guidelines regarding the care and use of laboratory animals, published by the International Association for the Study of Pain and the National Institutes of Health. After two weeks of acclimatization of the animals with new environment, experimental protocols were initiated and the rats were randomly (simple randomization) allocated in 3 groups (4 rats in each group): (1) tight ligation of the L₅ spinal nerve (SNL: n = 4); (2) sham surgery (Sham: n = 4), and (3) control (C: n = 4). Calculated sample size by the following Equation resulted in 3 animals in each group:

$$1) \quad \left[(Z_{\alpha} + Z_{\beta}) 2Sd^2 \right] d^2$$

Where $Z_{\alpha} = 1.96$, $Z_{\beta} = 0.84$, $Sd = 0.18$ and $d = 0.4$. The expected power was considered at 80%.

3.2. Induction of Neuropathic Pain

Animals were anesthetized by pentobarbital sodium (60 mg/kg, intraperitoneal). Then, L₅ spinal nerve was tightly ligated according to the method of Kim and Chung (1992) (18). Briefly, the left paraspinal muscles were separated at L₅-S₂ level and the left transverse process of L₆ vertebra was removed. The left L₅ spinal nerve was identified and gently separated from adjacent L₄ spinal nerve. Then, L₅ spinal nerve was tightly ligated using silk threads (6/0) and was transected just distal to ligature to ensure that all fibers were interrupted. Next, the wound was closed with silk threads (3/0). Great care was taken to avoid any damages to L₄ nerve. In the control sham group, the surgical procedure was identical to that described above,

except for the left L₅ spinal nerve that was not ligated and transected. Only animals showing no signs of motor deficiencies were considered to be used for further experimentations, those animals without neuropathic pains in their behavioral tests. Then the rats were divided in 3 groups with 5 members in each: non-operated, sham, and neuropathic pain (SNL) groups. After 4 weeks, rats were anesthetized by injecting intraperitoneal ketamine (90 mL/kg) and xylazine (10 mL/kg) and the muscle tissue samples were separated from the left soleus muscle and situated in -80°C nitrogen for future analysis.

3.3. Behavioral Tests for Measuring Neuropathic Pain

Radiation heat apparatus was used to measure hyperalgesia in which the middle part of animal's paw from Plexiglas level was exposed to thermal constant radiations and paw withdraw threshold time (PWTs) was calculated. Heat excitations repeated 3 times, with 5 to 10 minutes intervals. To measure the mechanical allodynia, animal was located on the wired network and inside the Plexiglas capsule with 20 × 20 × 30 cm dimensions. After acclimatization with new environment, von Frey fibers with weight range of 2 to 60 g (2, 4, 6, 8, 15, 26, and 60 g) (manufactured by Stolting Inc.) were used. The experiment started with the lightest fiber and in the non-response cases, heavier fibers were gradually employed. For excitation initiation, each fiber was inserted into animal's paw in 3 consecutive times for 1 s with 5 s intervals. If we had positive response (animal raising its foot) in 2 consecutive performances, that weight of fiber would have been selected as the response threshold. If animal had no response to all of the fibers, number 60 would have been considered as the response threshold (19).

3.4. Soleus Removing and Weighting

Soleus muscles from the left leg were quantitatively dissected from the bone, immediately weighed, and frozen at 20°C. Later, these muscles were lyophilized and weighed for their dry weight. After excision of muscles, the left tibia was removed and freed from connective tissue, and the maximal length was measured. Yin et al. (20) showed that expressing muscle mass per unit of tibial length is a valid way to normalize mass when body weight differs between experimental groups.

3.5. RNA Extraction and cDNA Synthesis

RNA extraction was done by QIAzol® Lysis Reagent (Germany, Qiagen) and chloroform (Germany, Qiagen) according to manufacturer's instructions. So, about 50 mg of the muscle tissue was separately homogenized in 1 to 10 portions in QIAzol® Lysis Reagent for total RNA extraction and removing protein components. The final product was centrifuged at 12000 ×g for 10 minutes at 4°C. Then, it mixed with chloroform in 1 to 5 portions and was

shaken severely for 15 s. Then the supernatant was centrifuged at 12000 ×g for 10 minutes at 4°C and its water and mineral part were removed. Finally, its RNA contained portion was removed and mixed with isopropanol in 1 to 5 portions. It was left for 10 minutes at room temperature and then centrifuged at 12000 ×g at 4°C for 10 minutes. RNA contained Pellet was washed and resolved in 20 µL RNsa-free water. RNA concentration was measured by UV spectrophotometry method (Eppendorf, Germany), and 260 to 280 portions in 1.8 - 2 were determined as the desired purification. cDNA synthesis was done by using Quanti Tect Reverse Transcription Kit (Qiagen, Germany) in accordance to the manufacturer's manual.

3.6. Real-Time Polymerase Chain Reaction

Real-Time PCR quantity method was used by Premix SYBR Green II (Qiagen, Germany) for measuring *CDK5* and *GSK-3β* mRNA expression levels (Applied Biosystems StepOne, America). Reaction mixture was done in final volume in 20 µL (includes 1 µL of cDNA, 1 µL of forward primer, 1 µL of reverse primer, 7 µL of DEPC water and 10 µL of Syber Green) and each reaction in duplicate. Designing of primers was done according to *CDK5*, *GSK-3β* and *GAPDH* genes in gene bank of NCBI and by German company, Qiagen. Usable primer sequences have been reported in Table 1. Furthermore, *GAPDH* was used as the reference gene. Thermal program used in Real Time-PCR included: 95°C for 10 minutes, 95°C for 15 s, 60°C for 1 min (40 cycle repetitions). Melt curve and standard curve were drawn and considered for evaluating data authenticity and optimization experiment conditions, respectively and *CDK5* and *GSK-3β* expression data were normalized using *GAPDH* (reference gene). Fold change of genes was measured by Equation 2 (21).

$$2) \quad R = 2^{-ct\Delta\Delta}$$

Table 1. Primer Sequences Used for Real-Time Polymerase Chain Reaction^a

| Genes, Primer Sequence | Gen Bank Code |
|--------------------------------------|---------------|
| GSK-3β | NM_032080.1 |
| For: 5'-CAAAGCAGCTGGTCCGAGG-3' | |
| Rev: 5'-TCCACCAACTGATCCACACCAC-3' | |
| CDK5 | NM_001100673 |
| For: 5'-GGC TTCATGATGTCCTGCATA-3' | |
| Rev: 5'-GAC AGA ATC CCA GGC CTTTC-3' | |
| GAPDH | NM_017008 |
| For: 5'-GACATGCCGCTGGAGAAAC-3' | |
| Rev: 5'-AGCCCAGGATGCCCTTAGT-3' | |

^a Abbreviations: For, forward; *CDK5*, Cyclin-dependent kinase 5; *GAPDH*, Glyceraldehyde-3-phosphate dehydrogenase; *GSK-3β*, Glycogen synthase kinase-3β; Rev, Reverse.

3.7. Statistical Analysis

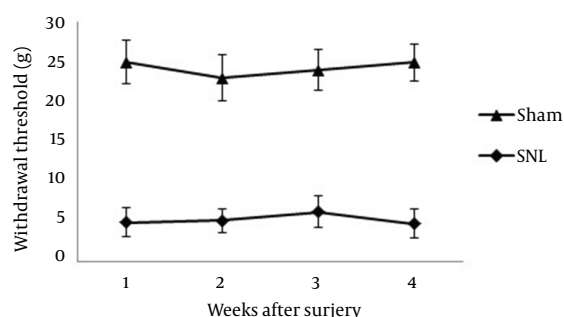
All statistical analyses were done by using SPSS software (version 19, SPSS Inc., Chicago, IL, USA). Normal assumption was examined using 1-sample Kolmogorov-Smirnov test. One-way repeated measures ANOVA and Independent t tests were used to compare groups regarding under study variables and followed by Tukey HSD post hoc. Significant level was determined at $P \geq 0.05$.

4. Results

4.1. Neuropathic Pain Behavior

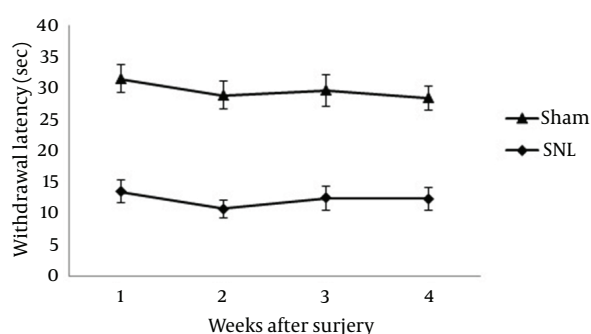
All rats that received L₅ SNL developed mechanical and heat hypersensitivity on the ipsilateral hind paw. As shown in Figure 1 and 2, PWT as well as PWL in the SNL group were lower postoperatively on the first, second, third and fourth weeks ($P < 0.05$ versus sham group), indicating that mechanical allodynia and thermal hyperalgesia has been induced by SNL operation.

Figure 1. Mechanical Withdrawal Threshold of the Left Hind Paws



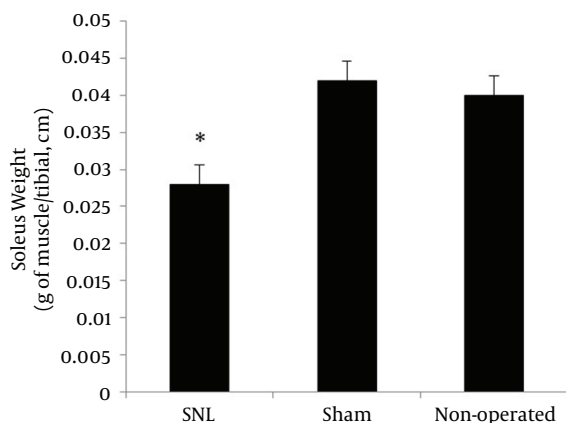
PWT decreased in the SNL rats postoperatively on the first, second, third and fourth weeks ($P < 0.05$ versus sham group). Data are presented as the mean ± SD.

Figure 2. Thermal Withdrawal Latency of the Left Hind Paws



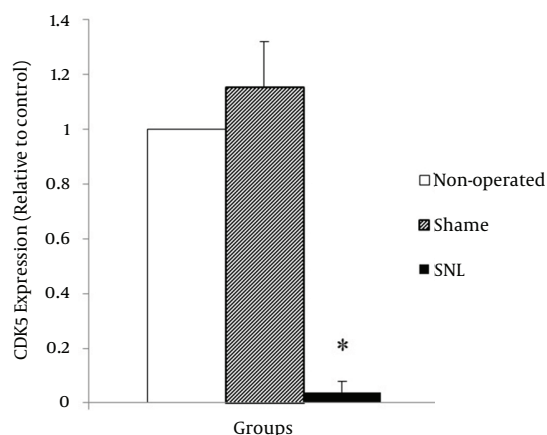
PWT decreased in the SNL rats postoperatively on the first, second, third, and fourth weeks ($P < 0.05$ versus sham group). Data are presented as the mean ± SD.

Figure 3. Weight of Soleus Muscle



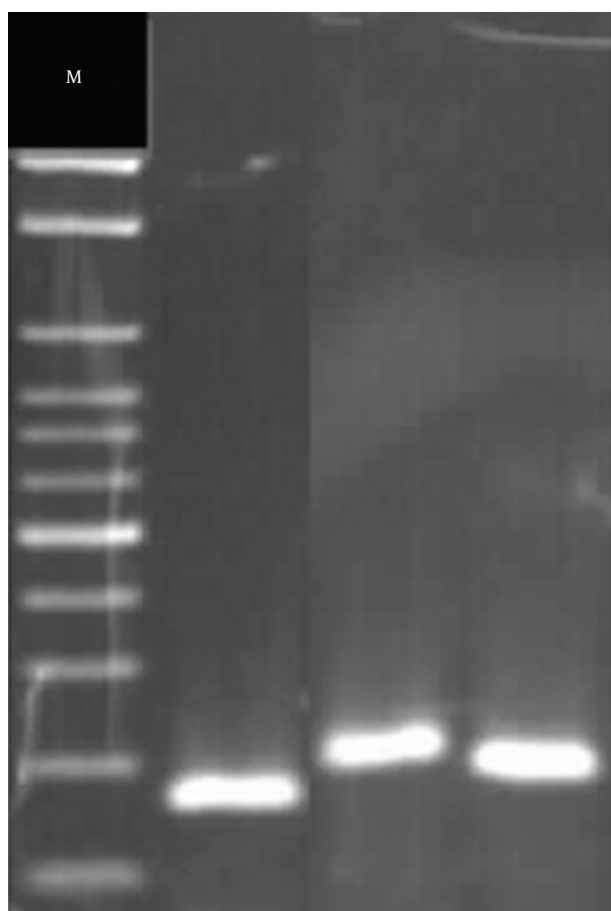
Muscle mass was reduced in SNL rats. Data are presented as mean \pm SD, which indicate significant differences with other groups ($P < 0.05$).

Figure 5. Real-Time Amplification of *CDK5* mRNA in Soleus Muscle



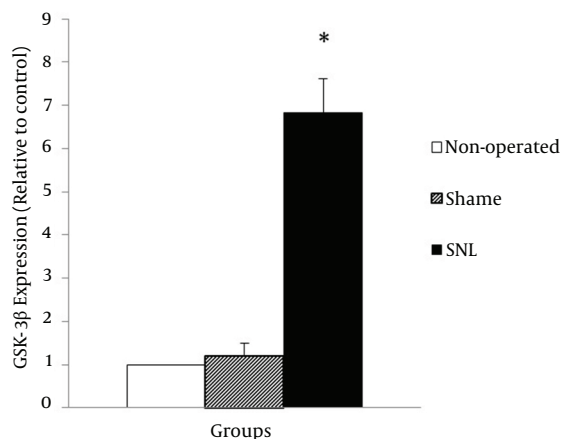
Data are presented as mean \pm SD, which indicate significant differences with other groups ($P < 0.05$).

Figure 4. Agarose Gel Electrophoresis of Polymerase Chain Reaction Products Amplified With a Multiplex Polymerase Chain Reaction Method



For the *GAPDH* (186 bp), *CDK5* (210 bp), and *GSK-3 β* (206 bp) genes, M = 100-bp DNA Ladder.

Figure 6. Real-Time Amplification of *GSK-3 β* mRNA in Soleus Muscle



Data are presented as mean \pm SD, which indicate significant differences with other groups ($P < 0.05$).

4.2. Soleus Atrophy and Down Regulation of *CDK5* and up Regulation of *GSK-3 β* in the Spinal Cord at the Mrna Level

Weight of the soleus muscle decreased in SNL rats and this result indicates muscular atrophy as a consequence of tight ligation of the L_5 spinal nerve (Figure 3). To elucidate the possible regulation of *CDK5* and *GSK-3 β* at mRNA level in rat models with neuropathic pain, we examined *CDK5* and *GSK-3 β* at mRNA level in soleus muscle after 4 weeks of tight ligation of L_5 spinal nerves. Agarose gel electrophoresis presented a single band with the expected size (Figure 4) and as indicated in Figure 5 and 6 *CDK5* mRNA levels were down regulated and *GSK-3 β* mRNA levels were up regulated.

5. Discussion

Candidate gene selection followed by decreased activity in neuropathic pain form studies is greatly influenced by the current knowledge of the pathogenic mechanisms involved in this disease. In the present study, it was observed that decreased activity in form of tight ligation of the L₅ spinal nerve causes decreased *CDK5* mRNA gene expression in soleus muscle, while amount of *GSK-3β* mRNA gene expression increases. Moreover, we show that tight ligation of the L₅ spinal nerve causes soleus muscle atrophy that could be attributed to this changes. Intracellular signaling cascades, which result in immediate early gene induction and maintenance, can control widespread changes in gene and protein expression. These changes lead to changes in muscle plasticity. Careful study in the relationship of different gene regulations is important for further understanding of pathogenesis of neuropathic pain. Previous studies showed that neuropathic pain increased *Cdk5* protein (22) and decreased *GSK-3β* protein (23). However, they did not show whether this increase in *Cdk5* protein levels and decrease in *GSK-3β* protein levels are due to the changes in their mRNA or degradation. The present data led us to conclude that down regulation of *CDK5* and up regulation of *GSK-3β* may cause decrease and increase in their protein level, respectively, a conclusion that previously reported by others.

Neuropathic pain is related to decrease and disorders in physical activities. Such decreases in the activity cause inability in physical fitness, increase in blood pressure, and decrease in musculoskeletal functions. Improper physical activity leads to structural and functional reduction of skeletal muscle (24). Besides, damages to motor neurons lead to many morphological and biochemical changes in muscles (25, 26). Spinal cord injury is related to atrophy and decrease in the size of muscle mass, which pathologists related it to nerve damages and decrease in activity and movement because of pain existence (27). Decreased activity because of hyperalgesia and allodynia increases proteolysis and decreases protein synthesis in muscles (28). Furthermore, existing evidence shows that tight ligation of the L₅ spinal nerve causes limitation but not complete inhibition of muscle electrical activity. Greensmith et al. (1997) (29) showed that by exciting motor neurons of L₄ segment, only 30% of maximum generated force of soleus muscle was recruited; however, by exciting motor neuron of L₅ segment, 70% of maximum generated force of the muscle was recruited. Denervation causes effective intracellular environment changes, decrease in muscle mass, number and volume of mitochondria, and increase in apoptosis proteins (30). *Cdk5* activity and its receptor (p35) in afferent neurons are very important in understanding the reason for its acute pain (14). In muscles, *CDK5* expression is very necessary in its natural amount and unnatural *CDK5* expression in non-nervous cells. Both expressions are done mainly through its direct or indirect effects on organizing cytoskeleton structures

such as actin and microtubules, which seems to cause such structure changes and these changes on cytoskeleton structure leads to cellular disorders and disease (16). Fu et al. (2002) (26) reported *CDK5* expression in skeletal muscles of rats after nervous damages showed an unnatural increase, which followed by increase in acetylcholine receptors in neuromuscular junction. Such increase may have been due to inhibition of motor nerves electrical activity. Differences in the findings of the present study to that of Fu and his colleagues can be due to different denervation methods. However, both unnatural decrease and increase in *CDK5* expression can be related to some complications and diseases (17, 31).

On the other hand, *GSK-3β* is an enzyme with the ability to do several tasks and involved in numerous cellular processes. Along with its metabolic role, *GSK-3β* is involved in maintenance and plasticity of skeletal muscle (15). Even *GSK-3β* has a role in regulation of apoptosis signaling (32). It has been shown that in pain receptors, *GSK-3β* has a role in morphine bearing and modulation of its pain relieving effects (13). In muscular tissue, *GSK-3β* has a key role in regulation of muscle atrophy (33, 34). Also it has been shown that inhibition of *GSK-3β* causes decrease in muscular atrophy (33) and in absence of *GSK-3β*, myofibrils are protected against atrophy (15). *GSK-3β* controls expression of atrogen-1 and MuRf1 (Muscle Ring Finger-1) due to effects of atrophy activators. Expression of atrogen-1 and MuRf1 in response to IGF-1 (Insulin-Like Growth Factor-1) or insulin, which are protein synthesis activators, decreases (35). Evenson et al. (2005) (33) reported that atrogen-1 and MuRf1 levels decrease by inhibiting *GSK-3β* drugs while in absence of *GSK-3β*, dexamethasone injection (which is an industrial glucocorticoid that its injection causes induction of proteolysis) modulates amount of myofibrillar atrophy. Activation of signaling cascades of phosphatidylinositol 3-kinase (PI-3K)/protein kinase B (Akt) is due to insulin or IGF-I effects (the main regulators of *GSK-3β* activity), which causes Akt activation/protein kinase B (PKB) directly through *GSK-3β* phosphorylation (36, 37). Akt blockage causes increase in *GSK-3β* activity and obvious increase in atrogen-1 and MuRf1 levels (35, 38). Phosphorylated-Akt activates (Mammalian Target of Rapamycin) mTOR 1 complex and inactivates *GSK-3β* simultaneously that leads to increase in mRNA transcription and protein synthesis capacity (39, 40). Akt decreases *GSK-3β* activity through phosphorylation and leads to increased expression of mRNA derived from eIF2B (Eukaryotic Translation Initiation Factor 2B) increased activity because of decreased inhibition of eIF2Bε through *GSK-3β* (41). This finding shows that *GSK-3β* activity is very essential in muscular atrophy initiation.

GSK-3β has an important role in regulation of protein synthesis and muscular hypertrophy (42). Suppression of *GSK-3β* expression leads to trivial increases in contractile protein levels. However, acute decrease in *GSK-3β* of contractile proteins maintains proteolysis of contractile proteins more than excitation of protein synthesis. Initiation

of protein transcription is apparently the time-limiting stage in muscular protein synthesis and controlled mainly by Eukaryote Initiation Factors (eIFs). Activated GSK-3 β is responsible for eIF2B ϵ inhibition, which eventually causes decrease in the initiation of protein synthesis. On the other hand, inactivation of GSK-3 β by upstream kinase of it, Akt, causes decrease of eIF2B ϵ phosphorylation and facilitation of transcription initiation of Mrna (41). It was shown that Akt is the critical point in signaling cascade of hypertrophy and atrophy (38). Akt was activated after activation of a series of intracellular signaling cascades like IGF-1 and PI3K (Phosphatidylinositol-4, 5-bisphosphate 3-kinase) (43). Downstream target of Akt is GSK-3 β , which is phosphorylated and inactivated (44) and as a consequence, eIF2B ϵ would be exited from its inhibitive state (45). Akt may take part in muscular atrophy involvement processes; it is also probable that muscular atrophy will be followed by phosphorylated Akt reduction (46).

In the present study, it was shown that decreased activity and tight ligation of the L₅ spinal nerve in neuropathic pain cause decrease and increase in CDK5 and GSK-3 β mRNA levels, respectively and these changes are followed by pain-related disorders and soleus muscle atrophy. Generally increase in GSK-3 β may be related to muscular atrophy and decrease in protein synthesis, while decreases in Cdk5 expression is related to structural disorders and function of skeletal muscle. Thus, it is likely that increased activity in the form of strength and endurance training can contradict unnatural expression of these two proteins in this kind of neuropathic pain. Likewise, Lee et al. (2013) (47) showed that abovementioned types of training can lead to optimization of Cdk5 expression in brain. However, Leger et al. (2006) (46) reported increased activity in form of 8 weeks strength training causes muscular hypertrophy by increasing phosphorylated state of Akt, GSK-3 β , mTOR proteins. However it needs more study to state it as a certain fact.

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Authors' Contributions

Study concept and design: Masoud Rahmati; Analysis and interpretation of data: Masoud Rahmati and Seyed Jalal Taherabadi; Drafting of the manuscript: Masoud Rahmati, S. Jalal Taherabadi, and Mahmoud Mehrabi; Critical revision of the manuscript for important intellectual content: Masoud Rahmati, S. Jalal Taherabadi, and Mahmoud Mehrabi; and Statistical analysis: Masoud Rahmati

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