## Original Article miR-124 mediated neuron apoptosis in spinal cord injury by targeting the 3' UTR end of calpain 1

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Abstract: Spinal cord injury (SCI) is a devastating and common neurologic disorder. Different sets of minas are usually deregulated in different cancers. Here, we determined the function of miR-124 in SCI and study showed the expression level of miR-124 in spinal cord tissues of rats treated with miR-124 agomir before the construction of SCI models was significantly higher than rats treated with agomir-NC and saline. Overexpression of miR-124 induced by miR-124 agomir resulted in the activation extent of caspase 3 was inhibited, the decrease of calpain level and the number of apoptotic neurons. MiR-124 targeted the 3' UTR end of calpain 1, which were found to be important for the regulation of cell growth and differentiation. Luciferase report assay showed the normalized luciferase activity of calpin 1-UTR in neurons transfected with miR-124 mimic was markedly decreased compared with the negative control group. Our study also showed the percentage of apoptotic neurons cells decreased after transfection of miR-124 mimic. Moreover, the protein levels of neurons cells isolated from spinal cord tissues after transfection with miR-124 mimic were decreased observably. Therefore, we concluded miR-124 could regulate neuron apoptosis in spinal cord injury by targeting the 3' UTR end of calpain 1.

Keywords: Spinal cord injury, MiR-124, Calpain 1, 3' UTR, Caspase

#### Introduction

Spinal cord injury (SCI) is a devastating and common neurologic disorder that has profound influences on modern society from physical, psychosocial, and socioeconomic perspectives [1]. It results in loss of oligodendrocytes demyelination of surviving axons and severe functional impairment [2]. Many individuals with SCI do not regain their ability to walk, even though it is a primary goal of rehabilitation [3]. SCI is characterized by an immediate, irreversible loss of tissue at the lesion site, as well as a secondary expansion of tissue damage over time [4]. It has been recognized as a complication of cervical transforaminal injections, and at present, there is no universally accepted treatment and the mechanism of injury is uncertain [5].

The recent discovery of microRNAs (miRNAs) suggests a novel regulatory control over gene expression during plant and animal development. MiRNAs play important roles during development and also in adult organisms by

regulating the expression of multiple target genes. They have been identified in various tumor types, showing that different sets of miR-NAs are usually deregulated in different cancers [6]. MicroRNA miR-133b is essential for functional recovery after SCI in adult zebra fish [7]. MicroRNA-223 expression in neutrophils plays an important role in the early phase of SCI after spinal cord injury [8]. The time course of miRNA-223 expression may be related to inflammatory responses after SCI, and the time course of decreased miR-124a expression may reflect cell death [9]. The brain-enriched miR-124 is an important regulator of the temporal progression of adult neurogenesis in mice [10]. It is abundantly expressed in neurons in the mammalian central nervous system, and plays critical roles in the regulation of gene expression during embryonic neurogenesis and postnatal neural differentiation [11]. However, miR-124 expression in neurons is reduced after SCI. and may reflect the severity of SCI [12]. Online software (www.targetscan.org) predicted the 3' UTR end of calpain 1 existed potential target site of miR-124. Therefore, we predicted miR-124 may express its role in regulating neuronal apoptosis by targeting calpain 1.

In this study, we constructed SCI mice models and determined the mRNA levels of miR-124, calpain 1, 68-kD NFP and cleaved caspase 3 to explore the influence of miR-124 on SCI.

## Materials and methods

## Rat models of spinal cord injury

Adult male SD rats (230-270 g; Animal Center of Shandong University) with damage of T10 segment were used for all experiments. The rats were maintained in a climate-controlled vivarium with a 12 h light-dark cycle with free access to food. Before SCI models were constructed by improved Allen for 1 day, 20 nmol miR-124 agomir, 20 nmol agomir-NC and saline were injected into the T10 segment of spinal cord tissues. Then, SCI models were constructed by Allen method: SD rats were injected intraperitoneally with 100 mg/mL chloral hydrate (300 mg/kg) in the left 1 cm of abdomen midline. Rats were prostrate lied, fixed on the operating table and back wool was cut. Then, the back of rats was sterilized with iodophor and cut along spine spines. Vertebral plate and spine on T10 segment were removed. The T10 segment of spinal cord was struck by NYU Blow device (Curtiss-Wright, Grimethorpe) with 60  $gcm (10 g \times 6 cm)$  to result in the injury of spinal cord in T10 segment. Therefore, SD models were divided into 3 groups: miR-124 agomir group, agomir group and sham group.

The original generation of neurons HEK 293T were isolated from spinal cord of newborn rats. Newborn rats were soaked in 75% alcohol for 3~5 min. The head of rats were cut and rats were dissected along median line to expose spinal column. Rats were put into D-hanks and vascular tunic and meninge were stripped by separating spinal cord. Spinal cord tissues were cut into  $0.5 \sim 1 \text{ mm}^3$  and 2 mg/ml papain was added to culture in 37°C incubator for 20~30 min. Isopycnic fetal calf serum of 20% was used to end digestion. After blowing and beating with tubularis for some times, the solution was centrifuged at 1000 rpm for 5 min. The supernatant was cultured in 37°C incubator for 30 min and the cells at exponential phase were used for current experiments. All the samples were stored at -80°C for further use.

RNA extraction and qRT-PCR assay

QRT-PCR was used to determine the relative level of miR-124 in spinal cord in sham, agomir-NC and miR-124 agomir group. The relative levels of miR-124 and calpain1 in spinal cord before SCI and after SCI 1, 3 and 7 days were also determined.

TRIzol solubilization and extraction is a relatively recently developed general method for deproteinizing RNA [10]. In this study, the total RNA was extracted and isolated from BMSCs using the Trizol reagent (Invitrogen, Carlsbad, CA). RNA quality was assessed with the ThermoScientific NanoDrop1000 (Thermo Fisher Scientific, Inc., Waltham, MA, USA). RT-PCR was performed using QuantiTect Primer Assay (Qiagen GmbH, Hilden, Germany) and Quanti-Tect SYBR Green RT-PCR Kit (Qiagen GmbH, Hilden, Germany) on a LightCycler 480 Instrument (Roche Diagnostics, Mannheim, Germany). The detection and quantification contained the following steps: first, reverse transcription was performed at 55°C for 30 min, initial activation for 15 min at 95°C, next 40 cycles of denaturation were conducted at 94°C for 15 s, then annealing for 30 s at 55°C, extension for 30 s at 72°C. Fluorescence data was collected at the extension step. The relative expression of the target gene was determined using the 2- $\Delta$ Ct method.

## Western blot analysis

The protein levels of activated caspase 3, calpain 1 and its specific substrate 68-kD NFP in spinal cord tissues in sham, agomir-NC and miR-124 agomir group were determined by western blot assay. The protein levels of activated caspase 3, calpain 1 and 68-kD NFP in neurons cells HEK 293T in DMSO, negative control and miR-124 mimic group were also determined.

Total proteins were prepared from transfected Ishikawa or HEC-1-B cells using radioimmunoprecipitation assay (RIPA) lysis buffer (Sigma, USA) supplemented with protease inhibitors. An equal amount (50  $\mu$ g) of cellular lysates was separated on 10% SDS-polyacrylamide gel electrophoresis (PAGE) minigels and trans-

ferred to nitrocellulose filter membranes (Hybond, Escondido, CA, USA). The membrane was blocked with Tris-buffered saline Tween-20 (TBST) containing 5% skimmed milk powder for 1 h at room temperature, followed by incubation in TBST containing 5% BSA (Sigma, St. Louis, MO, USA) and primary antibodies overnight at 4°C. Primary antibodies were detected using a peroxidase-coupled goat anti-rabbit secondary antibody (1:8000, ZSBio, Beijing, China) and EZ-ECL chemiluminescence Detection kit for HRP (Biological Industries, Beit-Haemek, Israel). The following primary antibodies were used: rabbit mAb miR-381 (1:1000, Cell Signaling Technology, Danvers, MA, USA) and rabbit pAb B-actin (1:1000, Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA).

## Luciferase reporter assay

A fragment of the 3' UTR segment of calpain 1 containing the predicted miR-124 binding site was amplified by RT-PCR. The primers used were5'-CATACTAGTGTCTTCTGGACAGGCTCTG-3' (sense) and 5'-CTTAAGCTTAGAATCTCTCA CATA-CACAC-3' (antisense). Restriction sites are bolded and underlined. Site-directed mutagenesis of the miR-124 target site was carried out using Stratagene Quik-Change site-directed mutagenesis kit (Stratagene, Heidelberg, Germany). The construct was sequenced and named calpain 1-UTR-Mut. The pMIR-report luciferase vector was used for the construction of calpain 1-UTR or calpain 1-UTR-Mut plasmids (Ambion, USA). HEK293T cells were cultured in 24-well plates. In each well, 10 ng of phRL-TK renilla luciferase vector (Promega, USA) was co-transfected to normalize transfection efficiency. A concentration of 500 ng of calpain 1-UTR or calpain 1-UTR-Mut plasmids together with 10 nM miR-1124 mimics or negative control was also co-transfected. Transfection was done using Lipofectamine 2000 and Opti-MEM I-reduced serum medium (Life Technologies, California, USA). Briefly luciferase activity was measured using the Dual luciferase assay kit (Promega). Normalized relative luciferase activity (RLA) was calculated as the following formula: RLA = [firefly luciferase]/ [renilla luciferase].

## Cell transfection

The pre-miR-124 precursor molecule (miR-124 mimics), siRNA targeting calpain 1, activated

caspase 3 and negative control RNA-oligonucleotides were gained from Ambion corporation (Ambion, Austin, USA). The day before transfection, HEK 293T cells were seeded in antibiotic-free medium. Transfection of miRNAs was carried out using Lipofectamine 2000 in accordance with the manufacturer's procedure (Invitrogen, California, USA). The level of miR-124 mimics expression in the HEK 293T cells was assayed by real-time PCR.

## Basso-Beaie-Bresnehan scale (BBB) score

A large dataset was constructed by pooling BBB data from 236 subjects scored at Texas A&M University and 407 subjects scored at Ohio State University. The subjects scored at Texas A&M were male Sprague Dawley rats (Harlan, Houston, TX) with moderate (12.5 mm, MASCIS impactor) spinal contusion injuries. All contusion surgeries were performed by the same individual. BBB scores were assigned by three raters that had high inter-rater reliabilities (all r 0.9). Subjects were scored daily for the first 14 days after surgery, and then every other day until day 26. After day 26, subjects were scored every 4th day until day 42. The BBB data collected at Ohio State were generated by scoring 407 male and female Long Evans hooded rats (Simonsen Laboratories, Gilrov, CA), All subjects were contused using a MASCIS impactor with weight-drop heights of 12.5 (n = 223), 25 (n = 130), or 50 mm (n = 54). Data was collected by raters that had high inter-rater reliability (r > 0.88).

## Flow cytometry

To study the influence of expression changes of miR-124 on the apoptosis of colon cancer cells, flow cytometry was used. The colon cancer cell HEK 293T at logarithmic phase were selected and plated in a 96-well plate at a density of 2 × 10<sup>3</sup> cells/well in supplemented RPMI 1640 and incubated for 16 h before the cells were subjected to treatment in triplicate wells. After treatment, the cells were washed twice in phosphate-buffered saline (PBS) (2.68 mM KCl, 1.47 M KH<sub>2</sub>PO<sub>4</sub>, 8 mM Na<sub>2</sub>KPO<sub>4</sub>, 136.75 mM NaCl) and counted. Fifty to one hundred thousand cells were selected and centrifuged 5 min at 1000 r/min. Annexin V-FITC mixed liquor of 195 µL was added to resuspend cytotrophoblast cells and 5 µL was added to mix. Centrifugation at 1000 r/min for 5 min was per-



**Figure 1.** The level of miR-124 was decreased in spinal cord injury (SCI). A: The relative level of miR-124 in sham, agomir-NC and miR-124- agomir group. B: BBB score in sham, agomir-NC and miR-124- agomir group after SCI construction for 28 days. \*\*P < 0.01, compared with sham group, the value in agomir-NC or miR-124- agomir group had significant difference; \*P < 0.05 or \*\*P < 0.01, compared with agomir-NC group, the value in miR-124- agomir group had significant difference.

formed after cultivation 10 min. Sample was obtained after discarding supernatant and 10  $\mu$ L propidium iodide (PI) was added. Afterwards, the sample was stilled in dark for 30 min. Finally, the apoptosis was detected using flow cytometry (FCM) on the Moflo (Dako Cytomation, Glostrup, Denmark).

#### Statistical analysis

Statistical analysis was performed using SPSS 18.0. Data are expressed as the mean  $\pm$  standard error of the mean (SEM) of three independent experiments. Pearson's product-moment correlation coefficient was used to assess the correlation between different groups. A paired Student's *t*-test was used to evaluate differences between two groups. Multiple group comparisons were analyzed using ANOVA with a post hoc test. *P* < 0.05 was considered to indicate a statistically significant result.

## Results

## MiR-124 promoted the function of spinal cord

MiR-124 was found to inhibit the activation of microglial cells. To determine whether miR-124 play an essential role in the cholinergic antiinflammatory action, BALB/c mice were injected intravenously with the miR-124 mimic agomir. Therefore, we treated SD mice with 20 nmol miR-124 agomir or agomir-NC before bitting the spinal cord of rats 1 day. The mRNA level of miR-124 in T10 segment of spinal cord tissues in sham, agomir-NC and miR-124 agomir group was determined by qRT-PCR and results were shown in Figure 1A. As shown, the level of miR-124 in miR-124 agomir group was significantly higher than that in sham and agomir-NC group (p < 0.01). It indicated the level of miR-124 in miR-124 agomir group was markedly upregulated. Then spine function of mice after spinal cord was injured 28 days was estimated by BBB score method. As shown in Figure 1B, the BBB

score in agomir-NC group was evidently higher than that in sham group (P < 0.01), while BBB score in miR-124 agomir group was significantly higher than that in agomir-NC group (P < 0.01). It indicated miR-124 agomir could effectively protect mice from septic shock.

## MiR-124 inhibited apoptosis by targeting calpain

To directly assess the effect of miR-124 on the production of pro-inflammatory cytokines, miR-124 agomir was transfected into SD rats. The mRNA levels of miR-124 and calpain 1 in sham, agomir-NC and miR-124 agomir group before and after the SCI models were constructed 1, 3 and 7 days. As shown in Figure **2A**, the expression values of the miR-124 and calpain 1 in sham group before and after SCI 1, 3 and 7 days had no obvious changes. When the spinal cord of rats was injured by bitting, the level of miR-124 decreased with the time of SCI increased. The miR-124 levels in agomir-NC group after SCI 3 and 7 days were evidently downregulated compared with the value before SCI and after SCI 1 day (Figure 2B), while the level of calpain increased distinctly. However, if mice were treated with miR-124 agomir before SCI models were constructed, the mRNA levels of miR-124 and calpain 1 in sham, agomir-NC and miR-124 agomir group had no obvious dif-



**Figure 2.** A-C: Relative mRNA level of miR-124 and calpain 1 in sham, agomir-NC and miR-124- agomir group before and after SCI for 1, 3 and 7 days, respectively. \*P < 0.05 and \*\*P < 0.01, compared with sham group, the relative mRNA level of miR-124 or calpain 1 in agomir-NC group after SCI for 3 or 7 days had significant difference. D, E: The cell apoptosis situation in T10 segment of spinal cord tissues in sham, agomir-NC and miR-124- agomir group after SCI for 7 days. #P < 0.05, compared with sham group, average number in each view in agomir-NC group had significant difference. \*P < 0.05, compared with agomir-NC group had significant difference. \*P < 0.05, compared with agomir-NC group, average number in each view in miR-124-agomir group had significant difference.



Figure 3. A: The protein levels of activated caspase 3, calpain 1 and its specific substrate 68 kD-NFP in T10 segment of spinal cord tissues in sham, agomir-NC and miR-124 agomir group after SCI for 7 days by western blotting. B: Density analysis results in sham, agomir-NC and miR-124- agomir group after SCI for 7 days by Western blotting. \*P < 0.05 and \*\*P < 0.01,

compared with sham group, relative protein levels of activated caspase 3, calpain 1 and 68 kD-NFP in agomir-NC group had significant difference. \*P < 0.05 and \*\*P < 0.01, compared with agomir-NC group, relative protein levels of activated caspase 3, calpain 1 and 68 kD-NFP in miR-124 agomir group had significant difference.

ference (**Figure 2C**). Therefore, we speculated the downregulation of miR-124 may upregulate the level of calpain 1.

Calpains are a family of calcium-dependant cysteine-proteases involved in cytoskeleton remodelling and muscle differentiation. Previous study reported calpain-1 activation contribute to cardiomyocyte apoptosis [13]. In this study, we used flow cytometry to observe the cell apoptosis situation in T10 segment of spinal cord tissues. As Figure 2D showed SCI promoted cell apoptosis in spinal cord tissues and the increase of miR-124 inhibited the cell apopto-

sis. To further verify the effect, we observed the number of apoptotic cells in each view. As shown in **Figure 2E**, the average cell number in each view in agomir-NC group was much higher than that in sham group, while the value in miR-124 agomir group was evidently decreased compared with agomir-NC group and the number was almost the same with that in sham group.

#### MiR-124 inhibited the activation of caspase 3

As above results showed downregulation of miR-124 could upregulate the expression of calpain 1 and had significant inhibitory effect on cell apoptosis in spinal cord tissues. K1 keratin polypeptide (68 kD) is the main component of the stratum corneum. The level of 68-KD NFP was often used to indicate the activity of calpain. Therefore, we then determined the protein expression levels of calpain 1, 68 kD NFP and activated caspase 3 in the T10 segment of spinal cord tissues in SD rats after spinal cord was injured 7 days. As **Figure 3A** and **3B** 



**Figure 4.** A: The prediction on the targeting of miR-124 by online software www.targetscan.org. B: The normalized luciferase activity of pMIR-Ctrl, calpain 1-UTR and calpain 1-UTR-mut in negative control and miR-124 mimic group. \*\*P < 0.01, compared with negative control, normalized luciferase activity of calpain 1-UTR in miR-124 mimic group had significant difference. C: The cell growth situation of neuron cells HEK 293T, isolated from the spinal cord of newborn rats, in Entranster-R4000, negative control and miR-124 mimic group. D: The percentage of apoptotic cells HEK 293T in Entranster-R4000, negative control and miR-124 mimic group. ##P < 0.01, compared with negative control and miR-124 mimic group. The percentage of apoptotic cells in miR-124 mimic group had significant difference.

showed, the protein expression levels of calpain 1 and activated caspase 3 were increased with the downregulation of miR-124 in agomir-NC group, and the values were decreased with the increase of miR-124 in miR-124 agomir. The level of 68-KD NFP was decreased with the decrease of miR-124 in agomir-NC group and increase with the upregulation of miR-124 in miR-124 agomir group.

# MiR-124 downregulated calpain 1 by targeting the 3' UTR end

To study the molecular mechanism about the influence of miR-124 on SCI, we predicted the targets and result was shown in **Figure 4A**. Neuronal cell HEK 293T was isolated from the spinal cord of newborn rats and luciferase reporter assay was used to determine the levels of pMIR-Ctrl, calpain 1-UTR and calpain 1-UTR-mut in negative control and miR-124 mimic group. As **Figure 4B** showed the normalized luciferase activity of calpain 1-UTR was decreased significantly in miR-124 mimic group

compared with that in the negative control group. The normalized luciferase activity of pMIR-Ctrl and calpain 1-UTR-mut in negative control group or miR-124 mimic group had no obvious difference. Then after the original generation of neurons of spinal cord was transfected with miR-124 mimic for 24 h, 1 mM L-sodium glutamate was used to treat neurons for 24 h to induce SCI and flow cytometry was used to analyze the proportion of apoptotic neurons. Neurons after treating with DMSO and negative control were taken as comparison. As Figure 4C and 4D showed, the apoptosis rate of neurons was significantly decreased in miR-124 mimic group compared with that in the negative control group and Entranster<sup>™</sup>-R4000 group. Then, the protein expression levels of calpain 1, 68-kD NFP and activated caspase 3 group in DMSO, negative control and miR-124 mimic group were determined by Western blotting. As Figure 5A and 5B showed, the level of calpain 1 and activated caspase 3 is significantly reduced in miR-124 mimic group compared with nega-



**Figure 5.** A, B: The electrophoretogram and relative protein levels of activated caspase 3, calpain 1 and 68 kD-NFP in negative control and miR-124 mimic group. \*P < 0.05 and \*\*P < 0.01, compared with DMSO group, the relative protein level of activated caspase 3, calpain 1 and 68 kD-NFP in negative control group had significant difference. \*P < 0.05 and \*\*P < 0.01, compared with negative control group, the percentage of apoptotic cells in miR-124 mimic group had significant difference.

tive control group. It indicated miR-124 may induced the degradation of calpain 1 mRNA and downregulate the protein expression of calpain 1 by targeting the 3' UTR end of calpain 1.

#### Discussion

Spinal cord injury (SCI) is an insult to the spinal cord resulting in a change, either temporary or permanent, in its normal motor, sensory, or autonomic function. It is devastating and has profound influences on modern society from physical, psychosocial, and socioeconomic perspectives [1]. It can result in loss of oligodendrocytes demyelination of surviving axons and severe functional impairment [14]. SCI patients experience varying degrees of wrist and shoulder pain. MicroRNAs (miRNAs) play important roles during development and also in adult organisms by regulating the expression of multiple target genes. Yu et al reported miR-133b is essential for functional recovery after SCI in adult zebra fish [15]. MiR-124 is expressed specifically in muscle and the central nervous system, and is considered to play an important role in SCI [9]. Several studies demonstrated that the overexpression of miR-124 in neural stem cells (NSCs) could lead the NSCs to differentiate into neurons and astrocytes, which may be important for functional recovery in SCI [16]. MiR-124 had a modest role in neuronal differentiation and cell cycle [10]. Therefore, in this study we aimed to study the role of miR-124 in SCI.

SD rats were selected for research and divided into 3 groups: miR-124 agomir: 20 nmol miR-

124 agomir was injected into the injured spinal cord of rats before SCI models were constructed: Agomir-NC group: Rats were injected 20 nmol normal agomir at the injured spinal cord before SCI models were constructed; Sham group: the spinal cords of rats were not injured and the SCI models were not constructed successfully. The levels of miR-124, calpain 1, 68-kD NFP and caspase 3 were determined, motor function was evaluated by BBB score method and cell apoptosis situation in spinal cord tissues were determined. Results showed the level of miR-124 in injured spinal cord in agomir-NC group decreased with the increase of time, while the level of calpain 1 increased gradually. MiR-124 had evident inhibiting effect on cell apoptosis in spinal cord tissues. Moreover, when the overexpression of miR-124 was induced by miR-124 agomir, SCI could not result in the prominent decrease of miR-124 which might result in the efficient inhibition of the increase of calpain 1 level. The overexpression of miR-124 resulted in the evident inhibition on the level of activated caspase 3 and decreased the level of 68-kD NFP. Calpains are ubiquitous calcium-regulated cysteine proteases that have been implicated in cytoskeletal organization, cell proliferation, apoptosis, cell motility, and hemostasis [17]. Calpain 1 was one of them and was recently shown to bind tightly to the proximal end of the I-band titan segment in a calcium-dependent manner [18]. It was activated in neuronal apoptosis [19]. SCI evoked an increase in intracellular free Ca2+ level resulting in activation of calpain 1 [20]. The 68-kD protein forms the backbone of the

neurofilaments and the 200- and 150-kD proteins appear to be more peripherally located [21]. The loss of 68-kD NFP is found to be highest at the site of lesion, and decreases in neighboring regions in inverse relation to the distance from the lesion epicenter [22]. 68-KD NFP was often used to indicate the activity of calpain. Indicated prevention of NFP (68 kD) by estrogen treatment of SCI for 48 h in lesion and penumbra could inhibit the increased calpain activity and activation, a common finding after SCI [23]. Caspase-3 is a newly characterized mammalian cysteine protease that promotes cell death during brain development [24]. It has been identified as a key protease in the execution of apoptosis, whereas calpains have mainly been implicated in excitotoxic neuronal injury [25]. Caspase-1 and caspase-3 were often activated following SCI [26].

To explore the molecular mechanism, online software www.targetscan.org, showed 3' UTR end of calpain 1 was the target site of miR-124. Therefore, we speculated miR-124 may express its regulating role in neuronal apoptosis by directly targeting calpain 1. Luciferase reporter assay was used to determine the expression of pMR-Ctd, calpain 1-UTR and calpain 1-UTR-mut in negative control and miR-124 mimic groups. The firefly luciferase reporter activity is significantly reduced in calpain 1-UTR vector compared with calpain 1-UTR-mut. Flow cytometry assay showed the number of apoptotic cells decreased significantly in neuronal cells HEK 293T infected with miR-124 mimic. It indicated miR-124 could inhibit the apoptosis of spinal cord cells. miR-124 could promote neuronal differentiation of neural progenitor cells [27]. The down-regulation of miR-124 was observed in injured hypoglossal motor neurons after nerve injury [28]. Moreover, western blotting showed miR-124 mimic could decreased the levels of calpain 1 and activated caspase 3, which further proved miR-124 could induce the degradation of calpain 1 and downregulate the expression of calpain 1 by targeting the 3' UTR end of calpain.

## Disclosure of conflict of interest

None.

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