Nogo-A expression dynamically varies after spinal cord injury

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Abstract
The mechanism involved in neural regeneration after spinal cord injury is unclear. The myelin-derived protein Nogo-A, which is specific to the central nervous system, has been identified to negatively affect the cytoskeleton and growth program of axotomized neurons. Studies have shown that Nogo-A exerts immediate and chronic inhibitory effects on neurite outgrowth. In vivo, inhibitors of Nogo-A have been shown to lead to a marked enhancement of regenerative axon extension. We established a spinal cord injury model in rats using a free-falling weight drop device to subsequently investigate Nogo-A expression. Nogo-A mRNA and protein expression and immunoreactivity were detected in spinal cord tissue using real-time quantitative PCR, immunohistochemistry and western blot analysis. At 24 hours after spinal cord injury, Nogo-A protein expression was low in the injured group compared with control and sham-operated groups. The levels then continued to drop further and were at their lowest at 3 days, rapidly rose to a peak after 7 days, and then gradually declined again after 14 days. These changes were observed at both the mRNA and protein level. The transient decrease observed early after injury followed by high levels for a few days indicates Nogo-A expression is time dependent. This may contribute to the lack of regeneration in the central nervous system after spinal cord injury. The dynamic variation of Nogo-A should be taken into account in the treatment of spinal cord injury.

Key Words: nerve regeneration; spinal cord injury; contusion; Nogo-A; axon growth; immunohistochemistry; fluorescent quantitative PCR; neural regeneration

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Introduction
The discovery of the Nogo family of proteins provides an opportunity to develop interventions to promote axonal regeneration in the central nervous system after injury. Basic research and clinical study of Nogo has increased our understanding of the mechanisms underlying spinal cord injury, multiple sclerosis, and neurodegenerative diseases. There are three major isoforms of Nogo: Nogo-A, Nogo-B and Nogo-C. The inhibitory actions of Nogo-A are dependent on two domains, Nogo-66 which is present in all three isoforms, and a sequence referred to as Nogo-delta 20, that is unique to the long N-terminal domain of Nogo-A. Nogo-A exerts its effects by binding to Nogo Receptor 1, via the Nogo-66 domain (Wu et al., 2014). Nogo-A specifically negatively regulates axonal regeneration in the central nervous system because Nogo receptor is only expressed by limited classes of neurons (Hunt et al., 2002b; Josephson et al., 2002). However, Nogo-A has additional inhibitory domains in its unique sequence (Prinjha et al., 2002) that do not require Nogo receptor for activity (Niederost et al., 2002). Growing evidence shows that lack of nerve regeneration after spinal cord injury is associated with a high expression of myelin-associated inhibitory molecules. Nogo-A is considered to be the major growth inhibitor, but its expression after spinal cord injury is not entirely clear (Schweigreiter et al., 2006; Cafferty et al., 2010; Lee et al., 2010; Fernet et al., 2012). Understanding its expression pattern after injury will allow the development of methods to effectively manipulate its levels and will be beneficial for the treatment of axonal regeneration after spinal cord injury (Walmsley et al., 2007; Müller et al., 2008; Montani et al., 2009; Kern et al., 2013). In this study, we used weight-drop-caused spinal cord injury rats to investigate the variation of Nogo-A expression in spinal cord tissue at different time points using immunohistochemistry, real-time quantitative PCR and western blot analysis. A comparative study using normal and sham-operated rat spinal cord tissue was carried out to clarify the role of Nogo-A protein in the nerve regeneration process.

Materials and Methods
Animals
A total of 108 adult female Sprague-Dawley rats weighing...
180–220 g (Experimental Animal Center of Nanjing University of Traditional Chinese Medicine, Nanjing, Jiangsu Province, China) were maintained separately and were given free access to food and water at the Animal Experimental Center of Nanjing University of Traditional Chinese Medicine in China (license No. SCXK (Zhe) 2008-0033). The protocol was performed in accordance with the Guidance Suggestions for the Care and Use of Laboratory Animals made by the Ministry of Science and Technology of China.

**Animal grouping and model establishment**

The rats were randomly and equally assigned to three groups: normal group, sham-operated group and model group (\(n = 36\)). The rats in the normal group were not operated. In the sham-operated and model groups, the vertebral column was exposed and a laminectomy was performed. The skin, subcutaneous tissue and deep fascia of rats in the sham-operated group were sutured and they received no spinal cord injury.

A contusion injury was performed on rats in the model group according to Allen’s (1911) advanced method. Under anesthesia of chloral hydrate (0.4 mL/100 g, intraperitoneal injection, Aoxin, Jiangsu Province, China), the vertebral column of the rats was exposed and a laminectomy was performed at the T\(_{9-11}\) level. A contusion injury was inflicted using an advanced self-made weight-drop device in rats. A weight of 10 g was dropped from a height of 25 mm on the exposed spinal cord. Spastic convulsions could be seen in the rats’ hindlimb several times which was followed by flaccid paralysis. Following contusion, the incision was closed with wound clips. Penicillin (80,000 units/day, intraperitoneal injection, Yakang, China) was used to protect from postoperative infection. During recovery, urine was expressed manually, twice daily, to assist in urination until intrinsic function returned to normal.

**Spinal cord extraction**

Rats were anesthetized and decapitated at 24 hours, 3 days, 7 days or 14 days postoperatively according to the studies of Josephson et al. (2001), Hunt et al. (2003) and Wang et al. (2008). At each time point, three rats in each group were randomly selected for immunohistochemistry, western blot analysis and real-time quantitative PCR test. Using aseptic procedures, rats were flushed with saline and fixed with 4% paraformaldehyde/PBS for 2 hours. The injured region was removed (about 1 cm) from the spinal cords and immediately frozen at −70°C. These were used for the detection of Nogo mRNA and protein expression. Samples for immunohistochemistry were fixed in 4% paraformaldehyde/PBS overnight and then frozen at 4°C.

**Immunohistochemistry**

Tissue samples of spinal cord were dehydrated and embedded in paraffin following routine methods. Ten serial sections were cut (4–5 μm thick) coronally through the lesion site. Five sections were randomly selected to detect Nogo-A using the EnVision Kit (Dako, Produktionsvej, Denmark) according to the manufacturer’s protocol (Primary antibody: rabbit anti-mouse polyclonal antibody, 1:400, Abcam, HK, China. Secondary antibody: goat anti-rabbit IgG-Horseradish Peroxidase, 1:2,000, Bioworld, MN, USA). 3,3′-Diaminobenzidine coloration was used with hematoxylin as a counterstain. Sections were observed using an inverted biological microscope, and images taken on a digital camera were analyzed using Image-Pro Plus 6.0 software (Media Cybernetics, MD, USA). Five fields of view were randomly chosen per section within each slice at 200× magnification. Nogo-A immunoreactivity, indicated by pale brown staining, was quantified using the mean value of optical density in five fields of view.

**Western blot analysis**

Spinal cords were homogenized in RIPA buffer for 30 minutes on ice, and the homogenate was centrifuged at 15,000 × g for 30 minutes at 4°C. Protein was extracted from the supernatant and concentrations were then determined using the Bradford method. Loading buffer was used for protein denaturation. Equal amounts of protein were then resolved on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (10% for β-actin, 6% for Nogo-A) and transferred onto polyvinylidene fluoride membrane with Trans-blot®SD Semi-dry Electroblotting Apparatus (25 V, 30 minutes for Nogo-A, 25 V, 22 minutes for β-actin). The membrane was blocked for 2 hours at room temperature with PBS containing 5% skimmed milk and 0.1% Tween-20, washed with 1 × TBST for five times and then incubated for 12 hours at 4°C with primary antibodies specific for β-actin (rabbit anti-mouse monoclonal antibody, 1:1,000, CST, Danvers, MA, USA) and Nogo-A (rabbit anti-mouse polyclonal antibody, 1:1,000, Abcam, HK, China) in the blocking buffer. The membrane was subsequently incubated with goat anti-rabbit IgG-horseradish peroxidase (for Nogo: 1:1,500, for β-actin: 1:3,000, Bioworld, MN, USA), then visualized using a chemiluminescence system (Image Quant LAS 4000 mini, GE, NY, USA). Fuji X-ray film (Fuji, Kanagawa, Japan) with cassette closure times was used to obtain adequate exposure and to visualize bands. Bands were analyzed using Image-Pro Plus 6.0 software (Media Cybernetics, MD, USA). β-Actin was taken as an internal control. The relative protein expression of Nogo-A was demonstrated by the ratio of gray scale between Nogo-A and β-actin.

**Real-time quantitative PCR**

RNA from tissue specimens in each group were purified using Trizol RNA isolation reagent according to the manufacturer’s protocol (Biyuntian, Nanjing, Jiangsu Province, China). A reverse transcription kit (Biyuntian) was used to make cDNA, and PCR amplification was performed. The upstream primer sequence of Nogo-A was 5′-GAC AGA AAT GGG CAG CAT AGT-3′; the downstream primer sequence was 5′-CAG AGA CAG CAG CAG GAA TAA-3′. The PCR program began with an initial 30 second denaturation step at 95°C. This was followed by 40 cycles of 5 seconds at 95°C for denaturation and 34 seconds at 60°C for annealing and extension. The relative expression of Nogo-A mRNA was estimated using the CT comparative method (2\(^{-\text{ΔΔCt}}\)) by Liebscher (2005). GAPDH was taken as the internal reference gene. The upstream prim-
er sequence of GAPDH was 5'-CTG AGC ACT CTC CCT CAC AAT TC-3'; the downstream primer sequence was 5'-GTG CAG CGA ATC TTA TTG ATG GT-3'.

**Statistical analysis**

All normally distributed values are presented as the mean ± SD. Statistical analysis was performed using SPSS version 19.0 software (SPSS, Chicago, IL, USA). The significance of any differences between three groups was evaluated using one-way analysis of variance and the least significant difference test. A probability of 95% was taken to indicate significant difference.

**Results**

**Immunoreactivity of Nogo-A in rat spinal cord**

In the sham-operated group, weak Nogo-A immunoreactivity was observed (compared with the normal group) at each time point (24 hours, 3 days, 7 days, 14 days post-injury).
The mRNA expression of Nogo-A in rat spinal cord

At each time point, Nogo-A mRNA expression in the sham-operated group was not significantly different compared with that in the normal group ($P > 0.05$). However, in the model group, the expression of Nogo-A mRNA was significantly lower at 24 hours post-injury compared with the sham-operated group. It dropped to its lowest level at 3 days, then rapidly rose to a peak at 7 days before declining slightly at 14 days post-injury. At both 7 and 14 days post-injury, the expression of Nogo-A mRNA in the model group was still significantly higher than that in the sham-operated group. The differences in expression were significantly different at every time point between sham-operated group and model group ($P < 0.01$; Figure 3).

**Discussion**

The microenvironment of the central nervous system is crucial to the survival and regeneration of damaged nerve tissue. Nogo protein is one of the myelin-associated growth inhibitory proteins and is thought to play an important role in spinal cord regeneration (Hunt et al., 2003; Baumann et al., 2009; Hånell et al., 2010; Hou et al., 2010; Raiker et al., 2010; Anja et al., 2012). Nogo-A is a myelin-related axonal growth inhibitory factor secreted by oligodendrocytes and stored in the white matter of the central nervous system, which is rich in myelinated membranes (Vitellearo-Zuccarello et al., 2007; Chong et al., 2012; Saha et al., 2014). Intracellular Nogo-A is released by oligodendrocytes and myelin into the extracellular matrix after spinal cord injury, and inhibits axonal regeneration. As an important nerve growth inhibitory factor, Nogo-A is currently at the forefront of biomedical research (Liebscher et al., 2005; Weinmann et al., 2006).

In this study, we found that a spinal cord injury model had low protein and mRNA expression at 24 hours after spinal cord injury, and dropped to the lowest level at 3 days, rapidly rose to a peak at 7 days, and gradually declined at 14 days. Compared with sham-operated group, the expression of Nogo-A protein and mRNA was significantly increased at 7 and 14 days post-injury. This confirms that spinal cord injury was responsible for the high expression of Nogo-A.

It is widely believed that Nogo-A launches a signaling cascade after binding with its receptors and induces transformation of actin-cytoskeleton in growth cones and contraction of both filopodia and lamellipodia. All of these change the shape and stability of growth cones and then cause degeneration, which inhibits axon regeneration (Broggini et al., 2010; Joset et al., 2010; Wu et al., 2010; Hui et al., 2013). As a result, the high expression of Nogo-A after spinal cord injury may be one of the main reasons why nerves do not regenerate in the central nervous system.

The dynamic variation of Nogo-A in the rat spinal cord has been previously reported (Wang et al., 2013). Huber’s team (2002) found expression levels of Nogo-A of postnatal Lewis rats were not changed significantly after traumatic lesions to the spinal cord. Hunt et al. (2003) showed that following injury to the spinal cord, Nogo-A mRNA was upregulated around the lesion and Nogo-A protein was strongly expressed in injured dorsal column fibres and their sprouts which entered the lesion site. Guo et al. (2005) showed high expression of Nogo-A appeared immediately after spinal cord injury in rats. The differences observed in these studies may be explained by the use of different breeds of animals, type of spinal cord injury surgery and the time points observed in the experiment. However, assessment of Nogo-A
immunohistochemistry, real-time quantitative PCR and western blot analysis and observation for 14 days had not previously been studied. Our study showed that after spinal cord injury, Nogo-A was first expressed at a lower level, dropped even further, and then rapidly rose to a peak, which is similar to the study by Wang et al. (2008). In contrast to Wang’s study, we used three different methods of Nogo-A measurement and found a slow decline in process in the last 7 days. We speculate that the initial low expression is relevant to acute injury and necrosis of spinal cord. The linearized expression of Nogo-A indicates 7 days post spinal cord injury as a time window for intervention.

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Conflicts of interest: None declared.

References


