

Laboratory Investigation

Epigenetic Regulation in the Brain after Spinal Cord Injury : A Comparative Study

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Objective : After spinal cord injury (SCI), functional and structural reorganization occurs at multiple levels of brain including motor cortex. However, the underlying mechanism still remains unclear. The current study was performed to investigate the alterations in the expression of the main regulators of neuronal development, survival and death, in the brain following thoracic contusive SCI in a mouse model.

Methods : Eight-week-old female imprinting control region mice (n=60; 30-35 g) were used in this study. We analyzed the expression levels of regulators such as brain-derived neurotrophic factor (BDNF), glial cell line-derived neurotrophic factor (GDNF), nerve growth factor (NGF) and histone deacetylase (HDAC) 1 in the brain following thoracic contusive SCI.

Results : The expression of BDNF levels were elevated significantly compared with control group at 2 weeks after injury ($p < 0.05$). The expression of NGF levels were elevated at 2, 4 weeks compared with control group, but these difference were not significant ($p > 0.05$). The GDNF levels were elevated at 2 week compared with control group, but these differences were not significant ($p > 0.05$). The difference of HDAC1 levels were not significant at 2, 4 and 8 weeks compared with control group ($p > 0.05$).

Conclusion : These results demonstrate that the upregulation of BDNF may play an important role in brain reorganization after SCI.

Key Words : Spinal cord injury · Regulators · Brain · Brain derived neurotrophic factor · Epigenetic.

INTRODUCTION

Structural and functional changes following spinal cord injury (SCI) include both retrograde and antrograde cell degeneration^{2,24}. SCI removes supraspinal input to the spinal sensory and motor networks and thus results in severe and permanent sensory and motor function impairment after injury site. Some degree of functional recovery, however, can be observed without management⁴. Since regeneration of injured axons is limited in mature CNS, spontaneous recovery in motor function appears to be mediated by reorganization of spared neuronal system. This compensatory remodeling occurs at multiple level of the neuraxis including spinal motor centers, descending supraspinal tracts, brainstem and brain cortex¹⁹. The sensorimotor cortex in adults retains the capability to reorganize in response to altera-

tion in peripheral sensory input or behavioral manipulation^{6,16}. A much greater extent of structural and functional change can be observed after large-scale injuries such as SCI or limb amputation^{11,18}. Spinal lesions in primates reshape the sensory representational map in the cortex^{11,15}. EEG and PET in SCI patients has shown reorganization related to the recovery of limb functions and extensive changes in cortical and subcortical activation^{3,8,9}. On the other hand, transcranial magnetic stimulation in paraplegics disclosed an enlargement of the cortical representations of non-affected muscles in the primary motor cortex, together with an increased excitability⁷. A few reports have shown the alterations in the expression of the neurotrophins (NTs), NTs receptors, and prohormone convertase in the injured site following SCI¹⁰. Epigenetically, histone deacetylase (HDAC) family have been reported it can be important role in

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a variety of neurodegenerative diseases and neurological conditions, such as stroke or traumatic brain injury^{1,13}). Despite these ongoing studies, molecular mechanism in the brain after SCI remains unexplored. In this study, we have analyzed the changes in the expression of the main regulators of neuronal survival and death in the mouse model of SCI.

MATERIALS AND METHODS

Experimental animal

Eight-week-old female imprinting control region mice (n=60; 30-35 g) were used in this study. All animal care, interventions and euthanasia procedures, were in accordance with the National Institute of Health Guide for the Care and Use of Laboratory Animals and guideline approved by Animal Care and Use Committees of our institute. Data was analyzed and unblinded by the statistician at the end of the experiment. At the beginning of the experiment animals were randomly assigned to one of two groups: naive control (n=30), SCI (n=30).

Moderate spinal cord contusion

The prescribed animals underwent a contusive impact SCI induced by the Infinite Horizons (IH) device (Precision Systems and Instrumentation, Lexington, NY, USA). Animals were anesthetized with mixture of ketamine and xylazine (0.05 mL/kg) and absence of blink and withdrawal reflexes were ensured. Lacrilube ophthalmic ointment (Allergan Pharmaceuticals, Irvine, CA, USA) was applied to the eye to prevent drying. During surgery, the mice were kept on homeothermic blanket system (Harvard Apparatus, Ltd., Kent, UK) to maintain the body temperature at $37\pm 0.5^{\circ}\text{C}$ as measured by rectal probe. Following anesthesia, a vertical incision was made along the low thoracic vertebra and the superficial muscle and skin retracted. A laminectomy performed at thoracic vertebra T12 exposed the dorsal surface of the spinal cord without disrupting the duramater. Stabilization clamps were placed around the vertebrae at T9 and L2 to support the column during impact. The injury was induced by the tip of impactor probe, which transduces a mod-



Fig. 1. Schematic drawing illustrating operation of spinal cord injury using Infinite Horizons Impactor.

erate force of 50 kdyne. Feedback from the IH device, including the actual impact force, cord displacement, graph of time versus force, and graph of time versus displacement, were recorded for each animal (Fig. 1). After SCI, the muscles were sutured in layers with 5-0 chromic gut, and skin was closed with 7-mm wound clips. After surgery, mice were allowed to recover in a warmed cage with water and food easily accessible. Gentamicin (5 mg/kg, intramuscular; Abbott Laboratories, North Chicago, IL, USA) was administered immediately post-surgery. Animals were doubly housed with Mouse Houses for the duration of the study. Post-operative care consisted manual bladder expression twice of a day until mice recovered some bladder expression, then once a day for the duration of the study. The mice were maintained for 8 week after injury with all brains collected at 2, 4, 8 week for HDAC related protein, brain-derived neurotrophic factor (BDNF), glial cell line-derived neurotrophic factor (GDNF) and nerve growth factor (NGF) analysis. All animal care, interventions and euthanasia procedures, were in accordance with the National Institute of Health Guide for the Care and Use of Laboratory Animals and guideline approved by Animal Care and Use Committees of the Yonsei University Wonju College of Medicine. Data was analyzed and unblinded by the statistician at the end of the experiment.

Immuno blotting

To compare the expression levels of neurotrophic factors, mouse brains were lysed in 500 μL of cold RIPA buffer [50 mM Tris-HCl, pH 7.5, 1% Triton X-100, 150 mM NaCl, 0.1% sodium dodecyl sulfate (SDS), and 1% sodium deoxycholate] with a protease inhibitor cocktail (Sigma-Aldrich, St. Louis, MO, USA). Tissue lysate was centrifuged at $13000\times g$ for 15 minutes at 4°C . The supernatant was harvested, and protein concentration was analyzed using a Qunt-iT protein assay kit (Molecular Probes, Eugene, OR, USA). For electrophoresis, 50 μg of protein was dissolved in sample buffer (60 mM Tris-HCl, pH 6.8, 14.4 mM β -mercaptoethanol, 25% glycerol, 2% SDS, and 0.1% bromophenol blue), boiled for 10 minutes and separated on a 10% SDS reducing gel. Separated proteins were transferred onto polyvinylidene difluoride membranes (Invitrogen, Carlsbad, CA, USA) using a trans-blot system. Blots were blocked for 1 hour in Tris-buffered saline (TBS) (10 mM Tris-HCl, pH 7.5, 150 mM NaCl) containing 5% nonfat dry milk at room temperature, washed three times with TBS and incubated at 4°C overnight with an anti-mouse monoclonal BDNF (1 : 1000, Abcam, Cambridge, MA, USA), anti-mouse glial cell-derived neurotrophic factor (GDNF, 1 : 1000, Abcam) and anti-GAPDH (1 : 3000, Cell signaling, Boston, MA, USA) antibody in TBST (10 mM Tris, pH 7.5, 150 mM NaCl, and 0.02% Tween 20) containing 5% nonfat dry milk. On the next day, blots were washed three times with TBST and incubated for 1 hour with horseradish peroxidase-conjugated secondary antibodies (1 : 3000, SantaCruz Biotech, Santa Cruz, CA, USA) in TBST containing 3% nonfat dry milk at room temperature. After washing three times with TBST, pro-

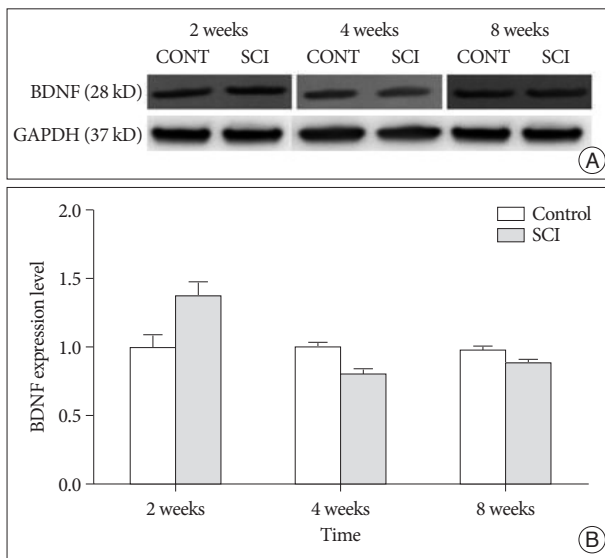


Fig. 2. Semi-quantitative immunoblot analysis of BDNF expression in brain after contusion of the spinal cord in mice (A and B). BDNF significantly elevated compared with those of control group at 2 week after SCI ($p < 0.05$). In this figure (B) data are plotted as mean \pm standard error of the mean. A sterisk indicate values significantly different from those of SCI and of control group by using a two-way repeated measures analysis of variance with repeated time factor. BDNF : brain-derived neurotrophic factor, SCI : spinal cord injury, CONT : control, GAPDH : glyceraldehyde 3-phosphate dehydrogenase.

teins were visualized with an ECL detection system (Amersham Pharmacia Biotech, Piscataway, NJ, USA).

Statistical analysis

Statistical analysis was performed using the Student's t-test and one-way ANOVA (Prism Graph Pad Software, San Diego, CA, USA). The data was expressed as mean \pm standard error of the mean. The significance level was assumed at $p < 0.05$, unless otherwise indicated.

RESULTS

To identify the neuromodulators associated with the neuronal cell development, growth, survival and apoptosis, the expression of four specific neurotrophic factors such as BDNF, NGF, GDNF and HDAC1 were investigated in the whole brain by Western blotting and multiplex ELISA assay.

Expression of BDNF was significantly elevated in the SCI group compared with those of control group at 2 weeks after SCI (1.38 ± 0.11 , $p < 0.05$). However, expressions of BDNF were not significant compared with those of control group at 4 weeks (0.80 ± 0.04) and 8 weeks (0.87 ± 0.04) after SCI (Fig. 2).

The level of NGF was elevated in the SCI group at 2 weeks (1.21 ± 0.10) and 4 weeks (1.17 ± 0.16), and decreased at 8 weeks (0.87 ± 0.03) compared with those of control group after SCI, however the differences were not significant ($p > 0.05$) (Fig. 3).

The level of GDNF was elevated in the SCI group at 2 weeks

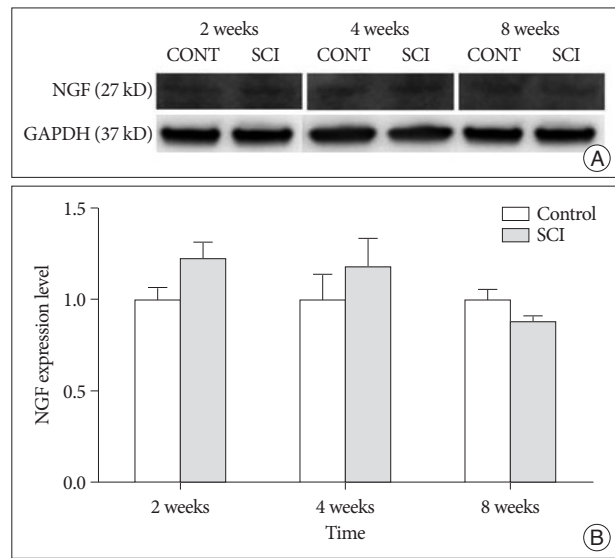


Fig. 3. Semi-quantitative immunoblot analysis of NGF expression in brain after contusion of the spinal cord in mice (A and B). NGF elevated compared with those of control group at 2 and 4 weeks, however the differences were not significant ($p > 0.05$). In this figure (B) data are plotted as mean \pm standard error of the mean. NGF : nerve growth factor, SCI : spinal cord injury, CONT : control, GAPDH : glyceraldehyde 3-phosphate dehydrogenase.

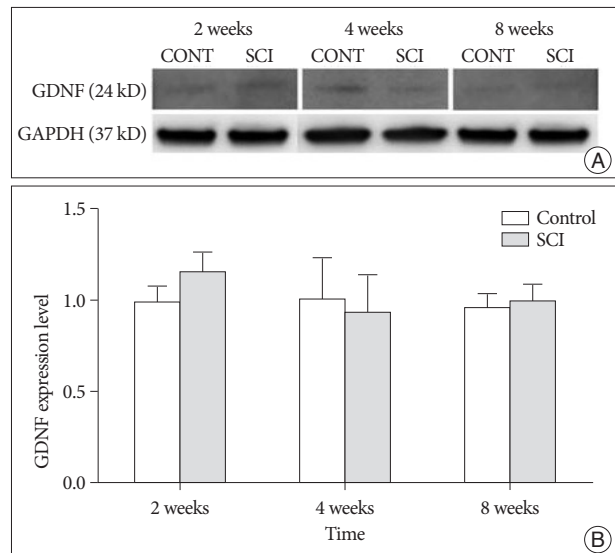


Fig. 4. Semi-quantitative immunoblot analysis of GDNF expression in brain after contusion of the spinal cord in mice (A and B). GDNF elevated compared with those of control group at 2 weeks, however the difference is not significant ($p > 0.05$). In this figure (B) data are plotted as mean \pm standard error of the mean. GDNF : glial cell line-derived neurotrophic factor, SCI : spinal cord injury, CONT : control, GAPDH : glyceraldehyde 3-phosphate dehydrogenase.

(1.16 ± 0.11) and 8 weeks (1.01 ± 0.09) compared with those of control group after SCI, however the differences were not significant ($p > 0.05$) (Fig. 4).

Among the regulators of gene expression, we investigated histone deacetylase-related protein. The expressions of HDAC1 were elevated at 2 weeks (1.11 ± 0.15), 4 weeks (1.15 ± 0.21) and 8

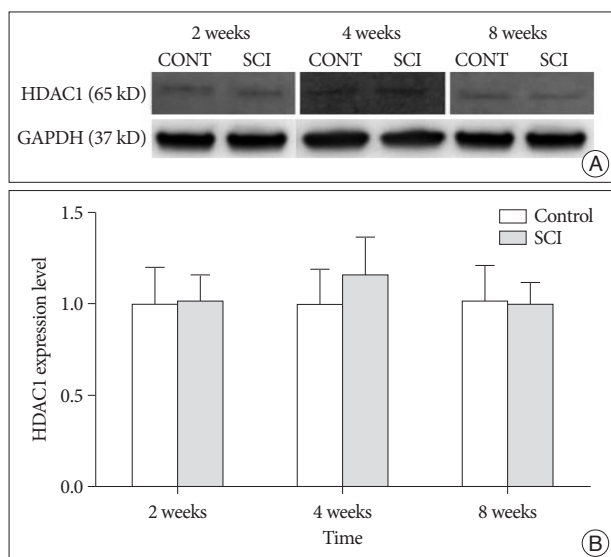


Fig. 5. Semi-quantitative immunoblot analysis of HDAC1 expression in brain after contusion of the spinal cord in mice (A and B). The difference of HDAC1 level in SCI and control group is not significant ($p > 0.05$). In this figure (B) data are plotted as mean \pm standard error of the mean. HDAC : histone deacetylase, SCI : spinal cord injury, CONT : control, GAPDH : glyceraldehyde 3-phosphate dehydrogenase.

weeks (1.09 ± 0.13) in SCI group compared with those of control group, but the differences were not significant ($p > 0.05$) (Fig. 5).

DISCUSSION

The purpose of this study was to investigate the alterations in the expression of the main regulators of neuronal development, survival and death related proteins. Here, we have examined the changes in expression of BDNF, NGF, GDNF and HDAC1 in the whole brain at specific time-points after SCI, in the mouse model of spinal cord contusion. BDNF protein showed significant increase at 2 weeks. NGF and GDNF proteins showed a tendency toward increase at 2 weeks compared with control, but it was not statistically significant. In addition, protein involved in gene regulation by relaxing the chromatin structure to a state that is transcriptionally active, HDAC1, showed elevated tendency, but the alteration was not significant. Over 100 years ago, it has been shown that following injury of a peripheral nervous system, axon distal to the injury undergo progressive retrograde degeneration, a process termed Wallerian degeneration.²²⁾ Also, recent studies suggest that Wallerian degeneration occur in the spinal cord of SCI patients, the topic of SCI results in anatomical changes of supraspinal central nervous system were clarified⁵⁾. For example, the recent study by Wrigley et al.²²⁾ found in living human that complete SCI results in anatomical changes in the human motor cortex and in the descending pathways from the motor cortex and anatomical changes in areas of the brain not directly involved in motor control, that is the medial prefrontal and anterior cingulate cortices. Furthermore, Kim et al.¹²⁾ observed that dynamic changes in density and

morphology of dendritic spines underlie reorganization of synaptic connectivity in the motor cortex following SCI. And PSD-95, a major postsynaptic density protein, and PSA-NCAM, a marker of active synaptic remodeling increase their expression. Some evidences observed that the alteration in the expression of enzymes in the spinal cord following SCI. Mature NTs induce cell survival upon binding to Trk receptors, their immature pro-protein counterparts promote apoptosis via activating P75NTR in the spinal cord¹³⁾. In accordance with this reports the members of PC enzymes functioning within regulated secretory pathway of neurons and neuroendocrine cells are significantly down-regulated in the spinal cord after SCI²¹⁾. In comparison, the expression of pro-NGF and pro-NT3 regulating enzymes are ubiquitous expressed in the cell of SCI. Pro-BDNF is primarily regulated secretory pathway of neuron in the spinal cord^{16,20)}. In the recent year, BDNF have emerged as important upstream regulators of long term potentiation in brain regions, including neocortex and hippocampus. These mean that BDNF appear to play an important role in neuronal cell development and survival. One of novel finding is significant elevation of BDNF level at 2 weeks. This is the first direct demonstration that BDNF signaling can be important regulation process in the brain reorganization following SCI. Although several HDACs are expressed at high level in the brain, their role in brain function has not been fully explored. The action of HDACs is opposed by histone acetyltransferases (HATs) such as CREB-binding protein and p300, which catalyze the transfer an acetyl moiety from acetyl-coenzyme A to specific lysine residues of histones¹⁴⁾. Acetylation of histones relaxes the chromatin structure to a state that is transcriptionally active, while histone deacetylation transforms chromatin to a transcriptionally repressed state. Hence, gene expression is regulated, in part, by the balance of HDAC and HAT activities¹⁴⁾. Recent study have reported that neuronal apoptosis by HDAC inhibitors involves stimulation of E2F-1, a transcription factor with established proapoptotic activity in neurons¹⁾. The expression of E2F target genes, such as those for cyclin E and Apaf-1, is induced in cerebellar granule neurons by pharmacological inhibition of HDACs¹⁾. Our data show elevated tendency of HDAC1 level at 2, 4 and 8 weeks but the elevation was not significant. Histone deacetylase-related protein expression is down regulated in apoptotic neurons, and this occurs prior to the time at which these neurons become irreversibly committed to death. Hence it is need to evaluate expression level of HDAC1 at the acute phase.

CONCLUSION

In summary, our studies reveal elevated expression of BDNF in the brain after SCI. Also it is need to explore neurorestorative effect might be induced by upregulation of BDNF, extending evidence that pharmacological means of BDNF may play an important role in brain reorganization following SCI. All together, the actions of NGF, GDNF and HDAC 1 on neuronal

changes in brain after SCI should be evaluated. Modulation of process such as extending its temporal window or histological analysis related alteration of proteins may be a future strategy to clarify main protein in the brain after SCI.

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