

Enhanced connexin 43 expression following neural stem cell transplantation in a rat model of traumatic brain injury

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Abstract

Introduction: Reestablishment of functional networks after traumatic brain injury (TBI) has been proffered as one of the goals of neural stem cell (NSC) transplantation therapeutics. Gap junctions provide essential means for direct cellular communication by transferring small molecules and ions, which may provide insights into the interplay between grafted NSCs and host cells.

Material and methods: Thirty-six adult male Wistar rats were used in this study. The controlled cortical impact (CCI) model of brain injury has been performed. Seventy-two hours after CCI injury, animals were randomly assigned to two groups: PBS- and NSC- transplanted group. NSCs-transplanted group received delivery of the NSCs suspension to the cortex below the injury cavity in the ipsilateral hemisphere. At 1, 2, and 4 weeks post-transplantation, we investigated the expression patterns of gap junction-associated connexin 43 (Cx43) in the transplant site and the border of CCI by immunohistochemistry, Western blot and RT-PCR.

Results: Our findings showed that Cx43 staining was significantly greater in the transplant site and the border of CCI in the NSCs-transplanted rats compared to the control rats at different time points ($p < 0.01$ at 1 week, $p < 0.05$ at 2 and 4 weeks). Significantly higher gene and protein expression of Cx43 was found in NSCs-transplanted rats compared to the control rats in the period of 4 weeks post-transplantation ($p < 0.01$), and remained at a higher level until 2 weeks with or without NSC transplantation.

Conclusions: It is proposed that gap junction-associated Cx43 might participate in NSCs' beneficial effects via gap-junctional coupling by which grafted NSCs integrate into host neural tissue following transplantation after TBI.

Key words: neural stem cell, traumatic brain injury, transplantation, connexin43.

Introduction

Traumatic brain injury (TBI) results in immediate neuronal and glial cell loss and associated neurological deficits. Recent knowledge on neural stem cells has brought novel approaches to the use of stem cells in the treatment of some neurodegenerative disorders, as well as in the management of central neural system (CNS) injuries [1]. Reestablishment of functional networks after TBI has been proffered as one of the goals of stem

cell-mediated therapeutics. Although it is becoming recognized that grafted neural stem cells (NSCs) interact with endogenous imperiled neurons *in vivo*, the underlying mechanisms are not clear. Our previous study showed that grafted NSCs could improve expression of synaptic protein in the core of transplantation and improve functional recovery following cell transplantation after TBI [2]. Others have attributed this rescue and protection to the release of a protective factor or specific neurotransmitters from NSCs and signals present in the injured brain via cell–cell mediated repair [3, 4]. However, this mechanism alone is inadequate to account for the beneficial effects of NSCs.

Gap junctions provide essential means for direct cellular communication by transferring small molecules and ions, which may provide insights into the interplay between grafted NSCs and host cells. The role played by this type of direct intercellular communication during the progression of the injuries is of some debate. Far from being a simple matter of judgment, the contribution of gap junctions to cell injury is a complicated phenomenon that depends on the specific insult and network in which it operates [5]. The alterations in the gap junction activities after brain injury have been proposed to be involved in ‘bystander death’, which is delayed neuronal death, by allowing the spread of death messengers from injured cells to otherwise healthy neighboring cells through the gap junction [6]. However, an early and essential way in which grafted NSCs integrate functionally into the host neural circuitry and affect host cells is via gap-junctional coupling, even before (and perhaps establishing a template for) mature electrochemical synaptic communication. The exogenous NSCs also protect host neurons from death and reduce such signs of secondary injury as reactive astrogliosis. Both *in vitro* and *in vivo*, these beneficial effects of NSCs were abrogated when gap junction formation or function was suppressed [7].

Connexin proteins form hemichannels with their typical hexamer conformation, and association of two hemichannels of adjacent cells compose a functional gap junction. The expressions of some connexins in the nervous system have been shown to be influenced by neuronal injury [8]. Of these, connexin 43 (Cx43) is the most ubiquitously expressed and is predominantly found in astrocytes, activated microglia, and developing neurons. Furthermore, Cx43 appeared to be most closely associated with immature NSCs and/or non-neuronal cells [7]. Here we report that NSC transplantation significantly increased the gene and protein expression of gap junction-associated Cx43 in the transplant core and border of the CCI injury in NSCs-transplanted rats compared to the saline control rats at 1, 2 and 4 weeks post-transplantation. It is proposed that gap junctions might

participate in the iterative developmental process by which grafted NSCs integrate into host neural tissue after TBI.

Material and methods

Culturing of NSCs

All procedures were approved by the Institutional Animal Care and Use Committee of the China Medical University, Shenyang, China. Primary NSCs were isolated and cultured as previously described [9, 10]. In brief, the forebrain tissue of embryonic 14-day-old Wistar rats (Experimental Animal Center of Dalian Medical University, China) was harvested. Accutase (Sigma, St. Louis, MO, USA) was added to the shredded tissue and incubated for 10 min at room temperature. Gentle pipetting was used to generate a single cell suspension. The suspension was centrifuged for 5 min, resuspended with culture medium, and then seeded into 25 cm² T-flasks at a density of 0.8×10^5 cells/ml as a suspension. Cells were maintained at 37°C in a humidified atmosphere with 5% CO₂. Expansion medium (DMEM and F12, 1 : 1; L-glutamine 2.92 g/100 ml, HEPES 5 mM, NaHCO₃ 7.5%, glucose 0.915 g/100 ml, and heparin 50 mg/100 ml, all from Sigma, USA) contained N₂ supplement (1%; Gibco, USA), epidermal growth factor (EGF), 10 ng/ml and basic fibroblast growth factor (bFGF), 10 ng/ml (Invitrogen, USA). As shown previously, their pluripotent stem cell fate was confirmed [11].

Surgical procedures and transplantation

Thirty-six adult male Wistar rats (weight, 200–220 g, Experimental Animal Laboratory of Dalian Medical University, China) were used in this study. All procedures were approved by the Institutional Animal Care and Use Committee of the Dalian Medical University, Dalian, China. The controlled cortical impact (CCI) model of brain injury has been described previously [2]. Briefly, on the day of surgery the animals were deeply anesthetized with sodium pentobarbital (60 mg/kg intraperitoneally) and placed in a stereotactic frame (Stoelting Co., USA). After the skull was exposed, a 3-mm craniotomy was performed over the left parietotemporal cortex between lambda and bregma, with the dura matter left intact. The CCI brain injury was induced by the use of a 2.0 mm diameter pneumatic impactor (Air-Power, Inc., High Point, NC) (tissue deformation 1 mm, 4.5 m/s tip velocity). Seventy-two hours after CCI injury, animals were randomly assigned to two groups. One group received delivery of the NSC suspension at a concentration of 1×10^6 cells/ μ l (3 μ l per animal) to the cortex below the injury cavity in the ipsilateral hemisphere: anteroposterior – 3.0 mm from bregma, mediolateral 1.0 mm, dorsoventral 1.1 mm ($n = 18$). The second (control) group

received an equivalent volume of phosphate-buffered saline in the same location ($n = 18$). A 10- μ l Hamilton syringe was slowly advanced through the dura and cortex until the desired depth was reached. With controlled use of the Stoelting quintessential injector (Stoelting Co., USA), the cell suspension was injected over a period of 3 min (1 μ l/min). The needle was left in place for an additional 5 min and then withdrawn slowly.

Histological analysis

At 1, 2 and 4 weeks, animals ($n = 3$ in each group at different time points) were given a lethal dose of sodium pentobarbital (200 mg/kg intraperitoneally) and transcardially perfused with cold phosphate-buffered saline and freshly hydrolyzed 4% paraformaldehyde (pH 7.4 in 0.1 mol/l phosphate buffer). Brains were harvested, postfixed overnight, cryoprotected in increasing sucrose concentrations, and then cryosectioned at a 16- μ m thickness along the coronal plane for immunohistochemistry. In brief, sections were rinsed in 0.3% Triton X-100 in phosphate-buffered saline (T-PBS) and pre-incubated in 10% normal goat serum in T-PBS, rinsed with PBS again, immersed in 3% H₂O₂ for 10 min, washed with PBS, blocked by normal goat serum and incubated at 4°C overnight with the primary antibodies rabbit anti-Cx43 (1 : 500; Sigma, USA). Then, biotinylated secondary antibody and ABC complex (Boster, Wuhan, China) were added in sequence and developed with DAB reaction. Images were taken with an Olympus optical microscope.

Western blotting

Another 3 animals per group were decapitated at each time point. Brain tissue encompassing the transplant site and the border of the lesion was obtained immediately and dissected into two parts on a bed of ice along the center of the injury cavity for Western blot and RT-PCR detection (described below). Western blot analysis was performed as described previously [12]. Each sample was sonicated in 0.0625 M Tris-HCl (pH 6.8) and then centrifuged at 12,000 g for 15 min at 4°C. The supernatant was assayed for protein using a Bradford assay. Aliquots (50 μ g) of protein were boiled in denaturing sample buffer (62.5 mmol/l Tris (pH 6.8), 2% SDS, 5 mmol/l EDTA, 10% glycerol, 0.25% 2-mercaptoethanol, and 0.01% bromophenol blue). Brain samples were loaded onto SDS-PAGE mini-gels, electrophoresed, and transferred to nitrocellulose membranes. After transferring, membranes were blocked with 5% milk in TBST (0.1% Tween 20 in 20 mM Tris-HCl, pH 7.4, and 410 mM NaCl) for 2 h at room temperature. Blots were incubated overnight at 4°C with rabbit anti-Cx43 (1 : 1000; Sigma, USA). Following primary antibody incubations, blots were washed three times for 5 min each with TBST, incubated for

1 h with horseradish peroxidase (HRP)-linked secondary antibodies, washed four times for 10 min each with TBST, and developed with an enhanced chemiluminescence Western blotting detection system kit (ECL Plus, Amersham Biosciences, NJ, USA).

RT-PCR analysis

Total mRNA was extracted from the tissue using an RNA isolation reagent (TRIZOL, TaKaRa, Dalian, China) according to the manufacturer's instructions and treated with RNase-free DNase (TaKaRa, Dalian, China). PCR was performed using Taq Dynazyme (TaKaRa, Dalian, China) under standard conditions. β -actin was co-amplified as an internal control. The products were observed after electrophoresis on 1.2% agarose gel, and the density of each band was analyzed on the Gel Image Analysis System. The level of Cx43 mRNA was determined by calculating the density ratio of Cx43 mRNA/ β -actin mRNA. All primer pairs were obtained from Shenggong Biotech Company (Shanghai, China) and were as follows: Cx43 (5'-AAAGGCGTTAAGGATCGCGTG/GTCATCAG-GCCGAGGCT-3') and β -actin (5'-GTCACCCACTGTGCCCATCT/ACAGAGTACTTGCGCTCAGGAG-3').

Statistical analysis

The positive products of Cx43 were analyzed with the Motic Image Advanced 3.0 image analysis system. We immunostained the 3 sections in the transplant core and boundary of the CCI injury with or without transplantation of NSCs per animal ($n = 3$). Under the microscope (400 \times), 3 regions per section were counted for Cx43 positive cells, and the average integral optical density (IOD) was determined. The Western blot and RT-PCR products were semi-quantitatively analyzed by Gel-Pro Analyzer Version 4.0 software (Media Cybernetics, Bethesda, MD) [13]. The ratio of Cx43/GAPDH or Cx43/ β -actin represented the level of protein or mRNA expression [14]. Independent sample *t*-test was performed to detect the difference between mean values for the NSCs-transplanted rats and the control rats at the same time points. The data were expressed as means \pm standard deviation and the difference was considered significant at $p < 0.05$.

Results

Cx43 immunohistochemical staining

We detected positive products of Cx43 in brain sections from NSCs-transplanted rats and the control rats at different time points post-transplantation. Representative Cx43 (Figures 1A-B) staining was found in the cell membrane, in the cytoplasm adjacent to the cell membrane, and in the processes in injured brain with or without NSC transplantation at 1, 2 and 4 weeks. Our study showed that Cx43 staining was significantly greater in the NSCs-transplanted rats compared to the control rats at different time points (Figure 1C).

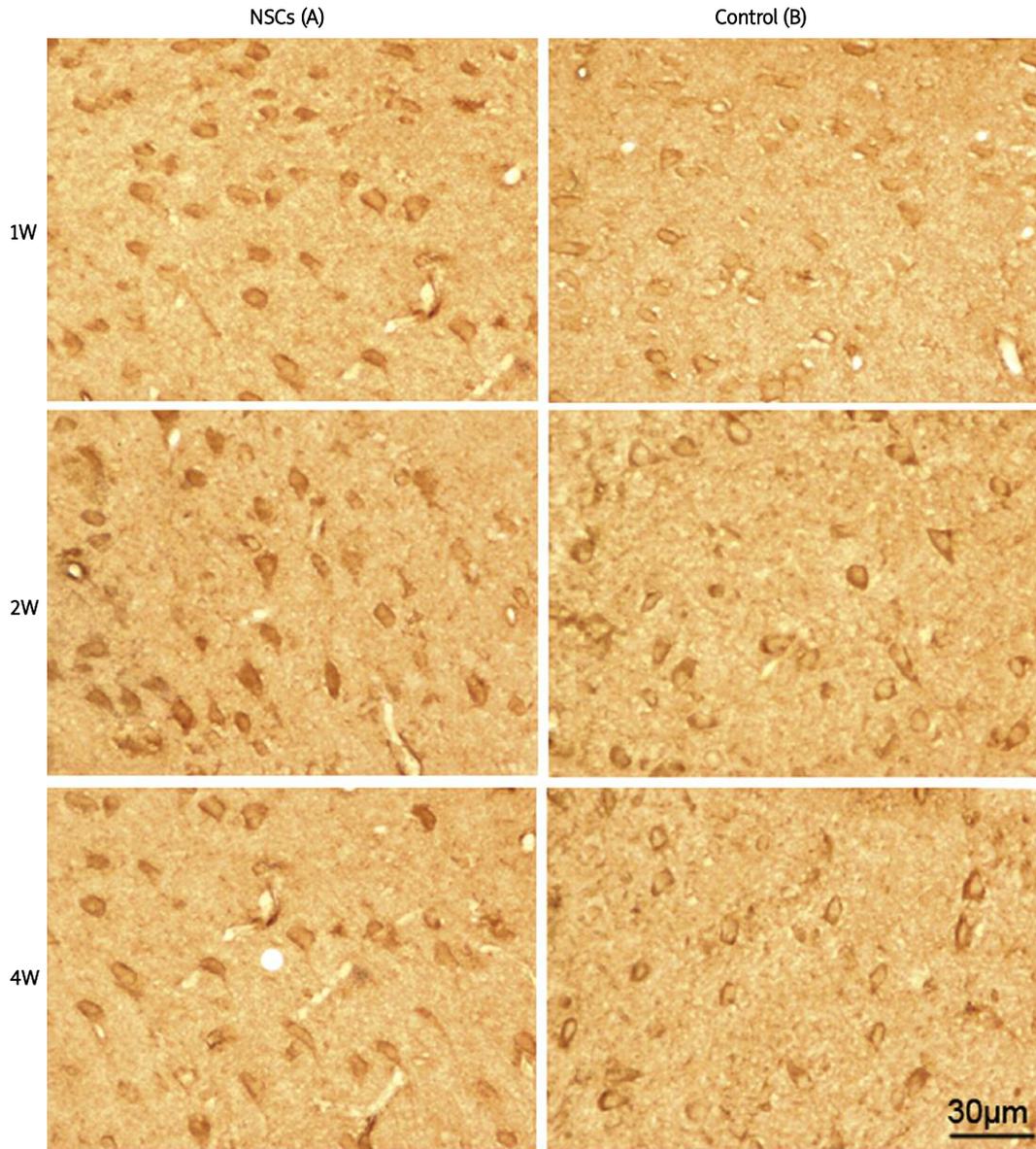
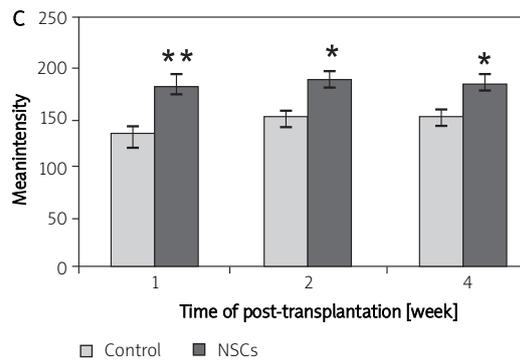


Figure 1. The immunoreactivity of Cx43 in the transplant core and the border of CCI at 1, 2 and 4 weeks post-transplantation. (A and B) Cx43-positive staining in 16-µm sections from the brains of the NSCs-transplanted and the control rats at 1, 2 and 4 weeks. Immunoreactivity of Cx43 was significantly increased in the brains of the NSCs-transplanted rats compared to the control rats at different time points (C). The scale bar represents 30 µm
* and ** indicate $p < 0.05$ and $p < 0.01$ vs. the control group at different time points



Expression of Cx43 protein and mRNA

We also performed Western blots and RT-PCR to detect protein and gene expression of Cx43 in the transplant core and border of the CCI injury at va-

rious time points post-transplantation. Figure 2 shows the expression of Cx43 at both the gene (Figures 2 A, B) and protein level (Figures 2 C, D) in the NSCs-transplanted groups and the control groups at various time points post-transplantation.

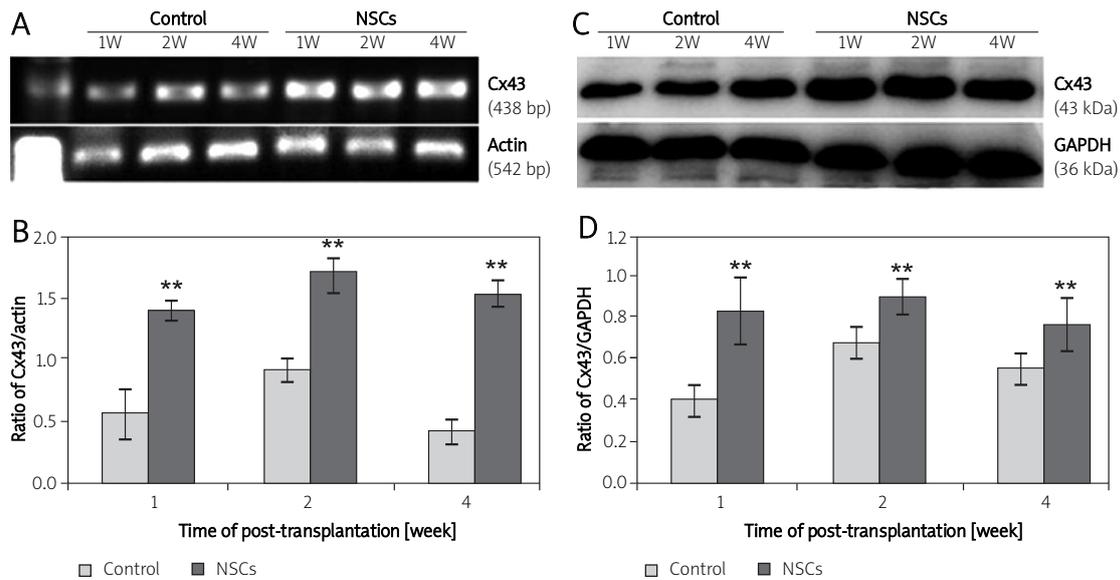


Figure 2. Gene and protein expression of Cx43 in the injured brain. In (A) and (C), lanes 1, 2 and 3 are from control rats and lanes 4, 5 and 6 are from NSCs-transplanted rats. Lanes 1 and 4, 2 and 5, and 3 and 6 are derived from rats at 1, 2 and 4 weeks post-transplantation, respectively. Representative results from semi-quantitative RT-PCR for Cx43 (A) and densitometric analysis of these bands (B). The intensity of each band was standardized to the band intensity of β -actin. Representative Western blots for Cx43 (C) and densitometric analysis of protein bands (D). The intensity of each band was standardized to the band intensity of GAPDH. Data are expressed as the mean \pm SD ($n = 3$ in each group) * and ** indicate $p < 0.05$ and $p < 0.01$ vs. the control group at different time points

As shown in Figure 2, the PCR product of Cx43 mRNA was 438 bp. The gene expression of Cx43 was 1.38-fold, 1.69-fold, and 1.53-fold at 1, 2, and 4 weeks, respectively, after transplantation in NSCs-transplanted rats, and 0.55-fold, 0.91-fold, and 0.42-fold, respectively, in the control rats (Figures 2 A, B). Significantly higher gene expression of Cx43 was found in NSCs-transplanted rats compared to the control rats in the period of 4 weeks post-transplantation ($p < 0.01$), and remained at a higher level until 2 weeks with or without NSC transplantation. Cx43 protein in the transplant core and border of the CCI injury predominantly migrated as a 43 kDa band as detected by Western blot. The results of the protein and gene expression of Cx43 were in agreement (Figures 2 C, D).

Discussion

Our previous study indicated that NSCs were capable of surviving and differentiating and significantly improved neurological function in a rat model of TBI. In this study, we performed the same model of TBI and cell transplantation strategy. Our data showed that NSC transplantation significantly increased the gene and protein expression of gap junction-associated Cx43 in the transplant core and border of the CCI injury in the NSCs-transplanted rats compared to the saline control rats at 1, 2 and 4 weeks post-transplantation. Enhanced gap junction-associated Cx43 might participate in the NSCs' beneficial effects in mild TBI.

The role of Cx43 gap junctions and hemichannels in CNS injury can be conceived to be either detrimental or beneficial. A process known as the "good Samaritan" was put forward by Shenton, who proposed a number of possible mechanisms by which gap junction communication might be beneficial *in vivo* following injury [15]. First of all, cellular injury results in accumulation of deleterious metabolites which are key players in excitotoxicity, such as calcium ions and glutamate in the cytoplasm. Although gap junction communication can also harm healthy neighboring cells by distribution of these toxic materials, a process known as the "bystander effect", gap junction communication promotes cell survival by allowing passage of toxic molecules out of the injured cell and neuroprotective substances, including neurotrophic factors and cytokines secreted by astrocytes, into the cell. In mild injury, increased Cx43 expression might be involved in diffusion or spreading of these deleterious substances, either buffered through gap junctions by healthy cells or released into the extracellular matrix via hemichannels [16], which might otherwise reach toxic levels. Thus, injured cells may be prevented from dying. On the other hand, astrocytes may mediate protection by buffering neurons from excitotoxicity and hypoxic depolarization by taking up extracellular glutamate and potassium, which might stimulate the survival of neighboring neurons and protect neurons from excitotoxic and oxidative insults [17]. Therefore, increased Cx43 is neuroprotective in mild injury. There is some evi-

dence of increased neurodestruction following global gap junction blockade *in vivo*. Cx43 null mice exhibited continuous neurodestruction or lesion spread [18, 19]. In addition, Cx43 might act directly at a genetic level, independent of gap junction or hemichannel activity [20].

Exogenous NSCs can protect or rescue endogenous neurons at risk for degeneration. Apart from diffusible factors that have typically been implicated [21, 22], an important feature that has been recognized is direct contact of NSCs with host cells. Connexin-associated gap junction formation and function are pivotal for ensuring host cell well-being. *In vivo*, engrafted NSCs-mediated rescue of imperiled host neurons did not occur when gap junction formation was suppressed by inhibiting gap junction function. It has been reported that functional gap junctions between grafted NSCs and host cells could be established within 2-18 h after NSCs grafting. Gap-junctional intercellular networks are an early form of communication that precedes and sets the stage for later electrochemical synapses and "traditional" electrophysiological communication between grafted and host cells. It is the earlier connexin, Cx43, that seems most pivotal in the NSCs-mediated rescue actions [7, 23].

It has also been reported that the role of Cx43 appears to vary with time following CNS injury. Cx43 mRNA and protein were up-regulated within hours in gray matter rostral to the lesion and reached over 3 times normal levels at 4 weeks following rodent spinal cord transection injury [24]. Another study found that the injured cortex showed an initial reduction in Cx43-ir up to 6 h after injury, followed by a subsequent increase in Cx43-ir from 24 h to 72 h in the model of TBI compared to controls [25]. It was proposed that it was a critical time in which gap junction coupling and hemichannel activation resulted in the mediation of injury. Initially after a CNS insult Cx43 hemichannels may be involved in the swelling of cells and spread of neurotoxic molecules from injured cells into the extracellular milieu. At later stages spatial buffering through the action of astrocytic gap junctions appears to be required for long-term neuroprotection [15]. What is more important, immunohistochemical staining and dye-transfer studies revealed that the NSCs already formed functional gap junctions prior to engraftment, thereby creating a substrate for subsequent graft and host communication. In an organotypic *ex vivo* model system for striatal tissue, at 0 and 7 d 74-94% of grafted NSCs and NSCs-derived cells expressed Cx43, suggesting a window of opportunity for successful host cell rescue by NSC transplantation [26].

In the present study, we investigated the expression patterns of Cx43 genes and proteins in the transplant core and the border of CCI, which is regarded as an important site as a therapeutic target because delayed responses take place mainly

in the penumbra, and the prevention of delayed responses may be required for successful treatment of injured brain [27]. Our results showed that the gene and protein expression of Cx43 were significantly increased at 1, 2 and 4 weeks post-transplantation compared to the control. We assumed that enhanced NSCs-mediated Cx43 may contribute to NSCs' beneficial effects via gap-junctional coupling in TBI rats treated with NSC transplantation. The mechanisms underlying how gap-junctional coupling facilitates NSCs' beneficial effects *in vivo* require further study. It is likely that initial NSC-astrocyte interactions and subsequent NSC-neuron couplings could be important especially during early stages in donor-host interactions. Further functional investigation of gap junction expression in graft and host cells and the associated variations in intercellular communication may help to understand and control the "good Samaritan" effects and "bystander effect" evidently mediated by gap junctional coupling following NSC transplantation after injury and thereby develop a novel therapeutic strategy to alleviate neuronal damage and/or to promote neuronal regeneration in future clinical NSC transplantation therapies.

In conclusion, the gene and protein expression of Cx43 in the transplant core and the border of CCI were significantly increased at 1, 2 and 4 weeks following NSC transplantation compared to the control. We assume that enhanced NSCs-mediated Cx43 might contribute to NSCs' beneficial effects via gap-junctional coupling in TBI rats treated with NSC transplantation. Further functional investigation of gap junction expression in exogenous and host cells and the associated changes in intercellular communication may help to understand and control the "good Samaritan" effects and "bystander effect" mediated by gap junctional coupling and thereby develop a novel therapeutic strategy to alleviate neuronal damage and/or to improve neuronal regeneration in future clinical NSC transplantation therapies.

Acknowledgments

Bo Yu and Haiying Ma contributed equally to this study.

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