

TLR2/NFκB signalling regulates endogenous IL-6 release from marrow-derived mesenchymal stromal cells to suppress the apoptosis of PC12 cells injured by oxygen and glucose deprivation

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Abstract. Two previous studies published by our group identified that mesenchymal stromal cells (MSCs) conferred neuroprotection in a rat model of hypoxic-ischaemic brain damage (HIBD), and that MSCs secreted abundant interleukin-6 (IL-6) when co-cultured with oxygen and glucose deprivation (OGD)-injured PC12 cells. The present study has further investigated the role of IL-6, and explored potential signalling pathways *in vitro*. *In vitro* models were established by co-culturing OGD-injured PC12 cells with MSCs. Subsequently, the expression levels of the signalling molecules, Toll-like receptor 2 (TLR2)/nuclear factor κB (NFκB), and IL-6 were altered separately in this *in vitro* model by treatment with an agonist, antagonist, siRNA or overexpression adenovirus. The expression levels of B cell lymphoma-associated X (Bax), TLR2, NFκB and IL-6 were detected by western blot analysis, real-time polymerase chain reaction or ELISA. The resting membrane potential (RMP) of the PC12 cells was analysed by whole-cell patch-clamp recordings. Compared with controls or the PC12 co-culture

group, the MSC co-cultured group induced less expression of Bax, but more IL-6 secretion. Up- or down-regulation of the TLR2/NFκB signalling pathway resulted in a corresponding increase or decrease in the IL-6 expression level in the MSCs. Co-culture with siIL-6-MSCs increased the expression levels of Bax and increased the RMP in the OGD PC12 cells. In conclusion, the release of IL-6 from MSCs was regulated via the TLR2/NFκB signalling pathway. Endogenous IL-6 reduced apoptosis and protected OGD-injured PC12 cells when they were co-cultured with MSCs. The present study has reported a novel immunomodulatory effect of the microenvironment of neural damage during MSC cytotherapy.

Introduction

Cerebral hypoxic-ischaemic insult during the perinatal period leads to neonatal brain injury, including encephalopathy, motor and mental deficits, learning disabilities and epilepsy (1). Hypoxic-ischaemic brain damage (HIBD) can be caused by brain lesions and the induction of hippocampal long-term potentiation through a series of pathophysiological changes in cell metabolic dysfunction, cerebral blood flow abnormalities, excitatory amino acid neurotoxicity, and intracellular calcium overload, free radical (e.g., nitric oxide) accumulation and apoptosis (2,3). The majority of drug therapies in the clinical arena are based on maintaining cell activities and delaying neuron death by preventing apoptosis to improve brain function. However, currently, no specific treatments are available for HIBD. An oxygen and glucose deprivation (OGD) *in vitro* model was used to simplify the complex brain interaction network in HIBD. Previous studies demonstrated that mesenchymal stromal cells (MSCs) could effectively restore learning and memory function in neonatal rats submitted to hypoxic-ischaemic insult *in vivo* (4-6), and these results demonstrated that transplanted MSCs exert immunomodulatory effects in the injury microenvironment, and that the MSCs secreted abundant interleukin-6 (IL-6) when co-cultured with OGD-injured PC12 cells (7). These results suggested that the high level of IL-6 expression may exert a neuroprotective role in neuronal injury. IL-6 is a pleiotropic cytokine that is able to regulate a variety of cell functions, including cell

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Abbreviations: MSCs, mesenchymal stromal cells; HIBD, hypoxic-ischaemic brain damage; IL-6, interleukin-6; PC12 cells, pheochromocytoma cells; OGD, oxygen-glucose deprivation; TLR2, Toll-like receptor 2; NFκB, nuclear factor κB; Bax, B-cell lymphoma-associated X; RMP, resting membrane potential; PDTTC, pyrrolidine dithiocarbamate; PGN-SA, peptidoglycan from *Staphylococcus aureus*

Key words: bone marrow-derived mesenchymal stromal cells, interleukin-6, Toll-like receptor 2/nuclear factor-κB pathway, PC12 cells, oxygen and glucose deprivation

proliferation, cell differentiation, immune defence and the inhibition of apoptosis. Previous evidence has revealed that IL-6 acts a double-edged sword, having dual proinflammatory and anti-inflammatory effects. Several studies have revealed that the expression of IL-6 is markedly increased when the central nervous system (CNS) is injured or is disease-afflicted, causing brain injury aggravation (8-10). However, a previous study (11) identified that astrocyte and macrophage activation were markedly decreased in the brains of IL-6 knockout mice following brain injury. This decrease in immunological reactivity suggested that IL-6 may have a neuroprotective role in brain injury. Based on these findings, the present study aimed to identify how MSCs secrete IL-6, and the physiological role of IL-6 in OGD-injured PC12 cells.

Materials and methods

Animals. Ten Wistar rats (age, 21 days), including 5 female and 5 male rats, were purchased from the Animal Experiment Centre of Daping Hospital affiliated with the Third Military Medical University (Chongqing, China). The animals were housed under a 12 h light/dark cycle with food and water freely available [SYXK (Yu) 2012-0015]. The experimental animal procedures were approved by the Ethics Committee of Chongqing Medical University (Chongqing, China).

Isolation and culture of MSCs. MSCs were isolated from the bone marrow of 21-day-old Wistar rats. The bone marrow was washed repeatedly with Dulbecco's modified Eagle's medium (DMEM/F12; Gibco, Thermo Fisher Scientific, Inc., Waltham, MA, USA) in a laminar flow cabinet until the bone had become bleached and the bone marrow was flushed out. Subsequently, the fluid containing bone marrow was centrifuged at 10,000 x g at room temperature for 3 min, and the supernatant was discarded. Then, the bone marrow cells were scattered and plated in a 60 mm Petri dish. The DMEM/F12 culture medium with 10% foetal bovine serum (FBS; Gibco, Thermo Fisher Scientific, Inc.) was changed every 24 h for the first three days, and subsequently the medium was exchanged every other day for three additional days. When the MSCs reached 70-80% confluence, the cells were digested with TrypLE (Gibco; Thermo Fisher Scientific, Inc.) and passaged at a ratio of 1:2 or 1:3.

Culturing PC12 cells. PC12 cells were obtained from the Cell Bank of Shanghai Institute of Cell Biology, Chinese Academy of Science (Shanghai, China). The cells were cultured in DMEM containing 10% horse serum, 5% FBS, 100 U/ml penicillin and 100 µg/ml streptomycin (all reagents purchased from Gibco; Thermo Fisher Scientific, Inc.) at 37°C in a 5% CO₂ incubator. When the PC12 cells reached 90% confluence, they were digested with TrypLE (Gibco; Thermo Fisher Scientific, Inc.) and passaged at a ratio of 1:2 every three to four days.

In vitro OGD model construction and co-culture configurations. The PC12 cells were seeded and grown to ~90% confluence in a six-well plate. To establish the OGD model, the PC12 cell culture medium was changed to Earle's balanced salt solution (EBSS; GE Healthcare Life Sciences, Logan, Utah, USA) after the cells had been washed twice with D-Hank's solution

(GE Healthcare Life Sciences). Subsequently, the cells were placed in an incubator (Thermo Forma 3111; Thermo Fisher Scientific, Inc.) with 5% O₂ and 95% N₂ at 37°C for 4 h, after which the EBSS medium was replaced with normal cell medium. The OGD-injured PC12 cells were co-cultured with MSCs (2x10⁶; called the MSC co-culture group) or with normal PC12 cells (2x10⁶; called the PC12 co-culture group) for a further 24 h. OGD-injured PC12 cells without co-culture served as controls. The co-culture methods were performed as previously described (12), and the PC12 cell co-culture group without injury was used as a co-culture negative control.

MSC treatments. To up-regulate Toll-like receptor 2 (TLR2) or down-regulate NFκB, the MSCs were cultured in the presence of the TLR2 agonist, peptidoglycan from *Staphylococcus aureus* (PGN-SA; 8 µg/ml; InvivoGen, Hong Kong, China) or the NFκB inhibitor, pyrrolidine dithiocarbamate (PDTC; 10 µg/ml; Sigma-Aldrich, St. Louis, MO, USA) for 24 h. Subsequently, the MSCs were treated with a TLR2- or IL-6-targeting small interfering (si)RNA, siTLR2 and siIL-6, or recombinant Ad-IL-6 adenovirus to knock down the expression of TLR2 or to regulate the expression level of IL-6, respectively, according to our previous studies (13,14). MSCs treated with red fluorescent protein (RFG) served as the control. Stable siIL-6-MSCs were used to investigate the biological function of IL-6, as reported previously (15). Green fluorescent protein (GFP)-labelled MSCs were used as a control.

RNA isolation and real-time quantitative polymerase chain reaction (RT-qPCR) analysis. Total RNA was isolated using an RNA extraction kit (BioTek Corp., Beijing, China) and reverse-transcribed into cDNA using a PrimeScript[®] RT Reagent kit (DRR037A; Takara Bio, Inc., Shiga, Japan) and a Bio-Rad My Cycler (Bio-Rad Laboratories, Inc., Hercules, CA, USA), according to the manufacturer's protocol. The single-stranded cDNA was diluted 10-fold, and used as the template. qPCR was performed on a StepOne[™] v2.1 Real-Time PCR instrument (Applied Biosystems; Redwood City, CA, USA) using Real Master mix [SYBR[®] Green; Tiangen Biotech (Beijing) Co., Ltd, Beijing, China]. The thermocycling conditions were as follows: Initial denaturation at 95°C for 30 sec and denaturation at 95°C for 5 sec, followed by 40 cycles at 60°C for 30 sec. Melting curve analysis and gel electrophoresis were performed to ensure that a single PCR product was amplified in each reaction. The PCR primers were designed using Primer Premier 5.0 software (Premier Biosoft International, Palo Alto, CA, USA). The ratio of the relative quantity of the target gene to glyceraldehyde-3-phosphate dehydrogenase (GAPDH), as a reference gene, was calculated using the 2^{-ΔΔC_q} method (16). The primer sequences for GAPDH, TLR2, NFκB and IL-6 are shown in Table I.

Western blot analysis. The cells were lysed in radioimmuno-precipitation assay buffer containing phenylmethanesulfonyl fluoride (Biotek Co., Ltd., Taiwan, China). The cell lysates were collected, and the protein concentration was determined using a bicinchoninic acid protein concentration determination kit (Biotek Co., Ltd., Beijing, China). The cell lysates were purified by 10% sodium dodecyl sulphate-polyacrylamide gel

Table I. Primer sequences for polymerase chain reaction.

Gene	Primer sequence
GAPDH	F: 5'-CCTGGAGAAACCTGCCAAG-3' R: 5'-CACAGGAGACAACCTGGTCC-3'
TLR2	F: 5'-TCTCGGCAACTATGAGTCCC-3' R: 5'-ATCGGTGAGATCTGCATTCC-3'
NFκB	F: 5'-AGGACTGCCGGGATGGCTTCTAT-3' R: 5'-GGTCTGGATGCGCTGGCTAATGG-3'
IL-6	F: 5'-ACAGCCACTGCCTTCCCTAC-3' R: 5'-TTGCCATTGCACAACCTCTTTTC-3'

F, forward; R, reverse; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; TLR2, Toll-like receptor 2; NFκB, nuclear factor κB; IL-6, interleukin-6.

electrophoresis (Beyotime Institute of Biotechnology, Beijing, China), with 30–40 μg total protein loaded per lane. After the proteins were electrophoresed, they were transferred to polyvinylidene fluoride membranes (Millipore Corp., Billerica, MA, USA). Following blocking of the membranes with 5% bovine serum antigen in Tris-buffered saline/Tween-20 (TBST) buffer at room temperature for 1 h, they were probed overnight with primary antibodies, including mouse monoclonal anti-B-cell lymphoma-associated X (anti-Bax; cat. no. sc-7480; 1:1,000; Santa Cruz Biotechnology, Inc., CA, USA), rabbit polyclonal anti-TLR2 (cat. no. SAB2102440; 1:1,000; Sigma-Aldrich), rabbit polyclonal anti-NFκB (cat. no. ab12146; 1:500 to 1:1,000; Abcam, Cambridge, UK), mouse monoclonal anti-IL-6 (cat. no. MAB5061; 1:500; R&D Systems China Co., Ltd., Shanghai, China), and mouse monoclonal β-actin (cat. no. sc-47778; 1:100 to 1:1,000; Santa Cruz Biotechnology, Inc.) at 4°C, and subsequently incubated with the appropriate secondary antibody conjugated with horseradish peroxidase (HRP; Santa Cruz Biotechnology, Inc.) at room temperature for 1 h. The proteins were detected using an enhanced chemiluminescent (ECL) substrate kit containing chemiluminescent HRP substrate (Millipore Corp.), and protein levels were recorded using an ECL Imaging System (G:BOX; Syngene UK, Cambridge, UK).

Enzyme-linked immunosorbent assay (ELISA). The levels of IL-6 cytokine released into the culture media of the different treatment groups were measured with an ELISA kit (Beijing 4A Biotech Co., Ltd., Beijing, China), according to the manufacturer's protocol. Background values were also analysed as a control. The optical density absorbance was measured at a wavelength of 450 nm. The values were calculated based on a constructed standard curve, and the assay was performed in triplicate.

Whole-cell patch-clamp recordings. The resting membrane potential (RMP) of cells was measured with a whole-cell patch clamp, as previously reported (4). Briefly, 30 mm glass coverslips containing PC12 cells, following co-culture with GFP-labelled MSCs or siIL-6-MSCs, were placed in an acrylic chamber under a TE-2000U fluorescence inverted microscope (Nikon, Tokyo, Japan). The membrane potential was amplified

using a Multiclamp 700B amplifier, and a Digidata 1322 interface (Axon Instruments, Foster City, CA, USA) was used for acquisition and off-line analysis. Between seven and ten intact cells in each group were selected to record the RMP, and all the recordings were performed at room temperature and completed within 1 h following the removal of the cells from the incubator.

Statistical analysis. All the data are expressed as the mean ± standard error of the mean. Significant differences between samples were examined by repeated measurements of analysis of variance (ANOVA) with one-way ANOVA, followed by Duncan's multiple-range test. $P < 0.05$ was considered to indicate a statistically significant value.

Results

MSCs down-regulate the expression level of the apoptosis factor, Bax, and up-regulate the expression level of IL-6 in OGD-injured PC12 cells. To evaluate the effects of MSCs on the OGD-injured PC12 cells, an *in vitro* co-culture system of MSCs and OGD-injured PC12 cells was established. Bax protein expression in the OGD-injured PC12 cells, which were co-cultured with the same quantity of uninjured PC12 cells or MSCs, was analysed by western blot analysis. As shown in Fig. 1A, the protein expression level of Bax in the MSC co-culture group was the lowest, that in the uninjured PC12 cell co-culture group was at an intermediate level, and that in the OGD-injured PC12 cell group was the highest. These data demonstrated that co-culture with MSCs was able to promote anti-apoptotic effects on the OGD-injured PC12 cells. The supernatants of the three above-mentioned groups were collected to assess IL-6 expression by ELISA; as shown in Fig. 1B, IL-6 secretion in the MSC co-culture group was significantly higher compared with that in the uninjured PC12 cell co-culture group ($P < 0.001$), whereas no significant difference in IL-6 secretion was observed between the uninjured PC12 cell co-culture group and the OGD-injured PC12 cell group. These data demonstrated that IL-6 was secreted primarily by MSCs, which may regulate the anti-apoptotic effect on the OGD-injured PC12 cells.

Protein expression levels of TLR2, NFκB and IL-6 are altered following PGN-SA and siTLR2 treatment of MSCs. To investigate whether IL-6 expression was regulated via the TLR2/NFκB signalling pathway, the MSCs were treated with PGN-SA or siTLR2. The changes in the expression levels of TLR2, NFκB and IL-6 were detected by RT-qPCR and western blot analysis. As shown in Fig. 2A, when compared with the control group, the mRNA expression levels of TLR2 ($P < 0.001$), NFκB ($P < 0.01$) and IL-6 ($P < 0.05$) were significantly increased following PGN-SA treatment. The changes in TLR2, NFκB and IL-6 protein expression detected by western blot analysis were identical with those changes in mRNA expression (Fig. 2B), suggesting that treatment with 8 μg/ml PGN-SA not only increased the expression level of TLR2 efficiently, but also up-regulated the expression of NFκB and IL-6. Subsequently, the MSCs were infected with siTLR2 adenovirus; compared with the RFP group, the mRNA and protein expression levels of TLR2 ($P < 0.01$), NFκB ($P < 0.05$) and IL-6 ($P < 0.001$) were significantly decreased in

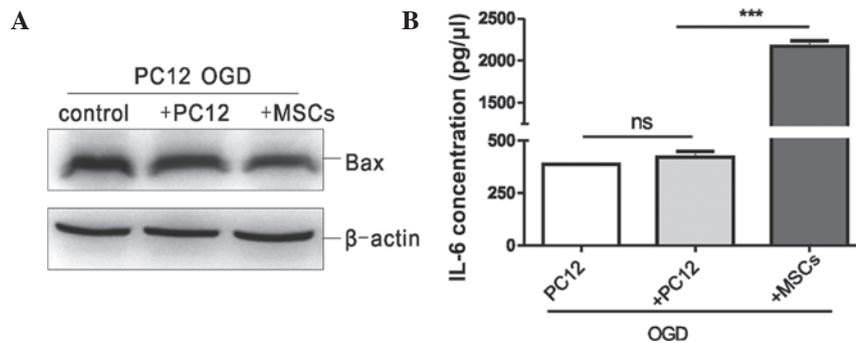


Figure 1. OGD-injured PC12 cells show an increased secretion of IL-6 and decreased protein levels of Bax when co-cultured with MSCs. (A) Detection of Bax protein levels in OGD-injured PC12 cells cultured alone and in OGD-injured PC12 cells co-cultured with uninjured PC12 cells or MSCs. A western blot representative of three independent experiments is shown. (B) Concentration of IL-6 in the supernatants of OGD-injured PC12 cells cultured alone, and of OGD-injured PC12 cells co-cultured with uninjured PC12 cells or MSCs, as detected by ELISA. Values are expressed as the mean \pm standard error of the mean ($n=3$) *** $P<0.001$, vs. the uninjured PC12 co-culture group. MSCs, mesenchymal stromal cells; Bax, Bcl-2-associated X; IL-6, interleukin 6; OGD, oxygen-glucose deprivation; ELISA, enzyme-linked immunosorbent assay; ns, not significant.

the siTLR2 group (Fig. 2C and D). The above results demonstrated that the down-regulation of NF κ B and IL-6 expression levels may be associated with reduced levels of TLR2, and further suggested that MSCs may regulate the secretion of IL-6 via the TLR2/NF κ B/p65 signalling pathway.

The levels of NF κ B and IL-6 protein expression are down-regulated by the NF κ B inhibitor, PDTC. To evaluate whether IL-6 is released from the MSCs via the TLR2/NF κ B pathway, MSCs were also treated with the NF κ B inhibitor, PDTC. Following treatment of the MSCs with 10 μ g/ml PDTC for 48 h, the protein levels of NF κ B, IL-6 and TLR2 were assessed by western blot analysis, and the quantification of the results is shown in Fig. 3. Significant decreases in the expression levels of NF κ B ($P<0.001$) and IL-6 ($P<0.005$) were observed following PDTC treatment compared with the control group. However, the expression levels of TLR2 were not significantly different between the PDTC group and the control group (Fig. 3A and B). This finding suggested that the inhibition of NF κ B expression was able to decrease IL-6 expression, although it did not affect TLR2 expression.

siIL-6 and Ad-IL-6 alter IL-6 expression, but do not affect protein expression in the TLR2/NF κ B pathway. To investigate whether IL-6 may regulate via a feedback process the TLR2/NF κ B signalling pathway, MSCs were treated with Ad-IL-6 or siIL-6, and the expression levels of IL-6, TLR2 and NF κ B were analysed by RT-qPCR and western blot analysis. The data in Fig. 4A and B demonstrate that IL-6 expression was markedly increased in the MSCs treated with Ad-IL-6 ($P<0.001$) compared with that of the RFP group, whereas the expression of TLR2 and NF κ B did not change. Subsequently, the MSCs were infected with siIL-6, and as expected, the IL-6 expression level correspondingly decreased ($P<0.001$); however, siIL-6 infection did not reduce the expression levels of TLR2 or NF κ B (Fig. 4C and D). These findings suggested that siIL-6 and Ad-IL-6 were able to change the expression level of IL-6, but did not affect the expression levels of TLR2 or NF κ B in the MSCs.

siIL-6-MSCs induce Bax expression and affect the RMP in the OGD-injured PC12 cells. To evaluate the biological function of IL-6 secreted from MSCs co-cultured with the OGD-injured

PC12 cells, the MSC cell line was screened for stable siIL-6 expression (15). The protein expression level of Bax was markedly increased in the OGD-injured PC12 cells co-cultured with the siIL-6-MSCs compared with the GFP-MSC group (Fig. 5A). The whole-cell patch-clamp recordings revealed that the RMP of the siIL-6-MSC group experienced a greater increase compared with that of the GFP-MSC group ($P<0.05$; Fig. 5B). These findings indicated that endogenous IL-6 from MSCs improved the restorative function of the OGD-injured PC12 cells.

Discussion

A previous study (11) has demonstrated that transplanted MSCs are not only able to ameliorate newly sustained brain damage, but are also able to improve the long-term prognosis *in vivo*. Transplanted MSCs have neuroprotective effects on the treatment of CNS injury, including traumatic brain injury and stroke, as well as on animal models of spinal cord injury. Previous studies (4,5,17) by our group have shown that MSCs can be induced and differentiated into neurons to promote the recovery of nerve function in HIBD rats. A further study (7) indicated that MSCs may secrete large quantities of IL-6 and IL-10, and that the levels of these cytokines may change the injury microenvironment and reduce H₂O₂-induced apoptosis. These results suggested that transplanted MSCs may have direct or indirect immunomodulatory effects on the injury microenvironment. In the present study of MSCs co-cultured with OGD-injured PC12 cells, it was demonstrated that the expression level of Bax in the MSC co-cultured group was significantly lower compared with that in the PC12 co-cultured and control groups, suggesting that MSCs have neuroprotective effects on OGD-injured PC12 cells.

An ELISA assay of the supernatants revealed that IL-6 secretion in the MSC co-culture group was significantly higher compared with that in the PC12 cell co-culture group, indicating that IL-6 be involved in the neuroprotective effects of MSCs and in microenvironment regulation. IL-6 is a secreted protein that consists of 184 amino acids, which is widely present in the human body, specifically in T cells, B cells, glial cells, fibroblasts, epithelial cells and certain tumour cells (18,19). IL-6 is able to generate an immune response, and has a role in macrophage and astrocyte activation during CNS injury. Under normal physiological condi-

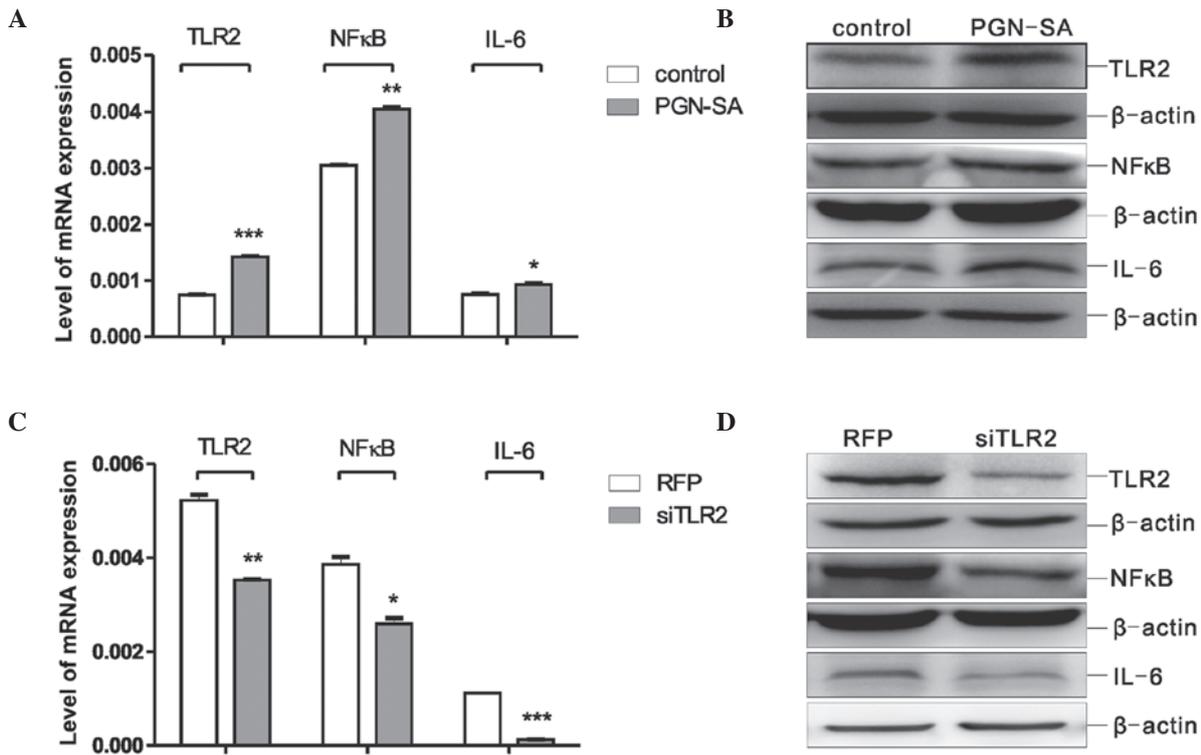


Figure 2. PGN-SA and siTLR2 caused changes in the TLR2/NFκB pathway and IL-6, respectively, in MSCs. (A) RT-qPCR was used to assess mRNA levels of TLR2, NFκB and IL-6 expression in MSCs after being treated with PGN-SA. The values are expressed as the mean ± standard error of the mean (*P<0.05, **P<0.01, ***P<0.001 vs. the respective control group cultured without PGN-SA). (B) Western blot analysis of the protein levels of TLR2, NFκB and IL-6 in MSCs following treatment with PGN-SA. (C) Real-time PCR was used to assess mRNA levels of TLR2, NFκB and IL-6 expression in MSCs after being treated with siTLR2. The values are expressed as the mean ± standard error of the mean (*P<0.05, **P<0.01, ***P<0.001 vs. the respective control RFP group). (D) Western blot analysis of the protein levels of TLR2, NFκB and IL-6 in MSCs following treatment with siTLR2. For the western blot analyses (B and D), the data were obtained from at least three independent experiments, analysed and normalised against β-actin. MSCs, mesenchymal stromal cells; RT-qPCR, real-time quantitative polymerase chain reaction; TLR2, Toll-like receptor 2; IL-6, interleukin-6; NFκB, nuclear factor κB; PGN-SA, peptidoglycan from *Staphylococcus aureus*; siTLR2, Toll-like receptor 2 small interfering RNA; RFP, red fluorescent protein.

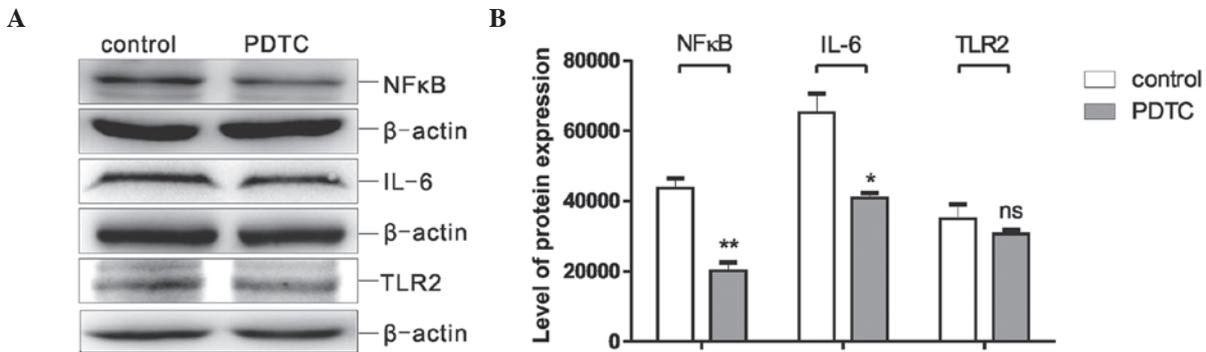


Figure 3. Protein expression levels of NFκB, IL-6 and TLR2 were assessed in MSCs following treatment with the NFκB inhibitor, PDTC. (A) Western blot analysis of NFκB, IL-6 and TLR2 protein expression in MSCs following treatment with 10 μg/ml PDTC. (B) Quantification (grey intensity analysis) of the western blot results. The data were obtained from at least three independent experiments, analysed and normalised against β-actin. The values are expressed as the mean ± standard error of the mean (*P<0.05, **P<0.01 vs. the respective control group). MSCs, mesenchymal stromal cells; TLR2, Toll-like receptor 2; NFκB, nuclear factor κB; IL-6, interleukin-6; PDTC, pyrrolidine dithiocarbamate.

tions, the expression level of IL-6 in the brain is extremely low. However, when the CNS is injured or becomes disease-afflicted, the level of IL-6 is significantly increased to induce the release of tumour necrosis factor-α (TNF-α) and IL-1β, leading brain injury aggravation. However, Erta *et al* (11), in studying IL-6 knockout mice following brain injury, demonstrated that astrocyte and macrophage activation markedly decreased in the injured brain, and that the immunological reaction was reduced,

suggesting that IL-6 is a key regulatory cytokine during brain injury. By contrast, another study (20) revealed that a high expression of IL-6 could reduce brain ischaemic injury, demonstrating that IL-6 also exerted neuroprotective effects. These results demonstrated that IL-6 exerts dual proinflammatory and anti-inflammatory effects. To evaluate the biological function of IL-6 secreted from MSCs co-cultured with OGD-injured PC12 cells, an MSC cell line was screened in the present study

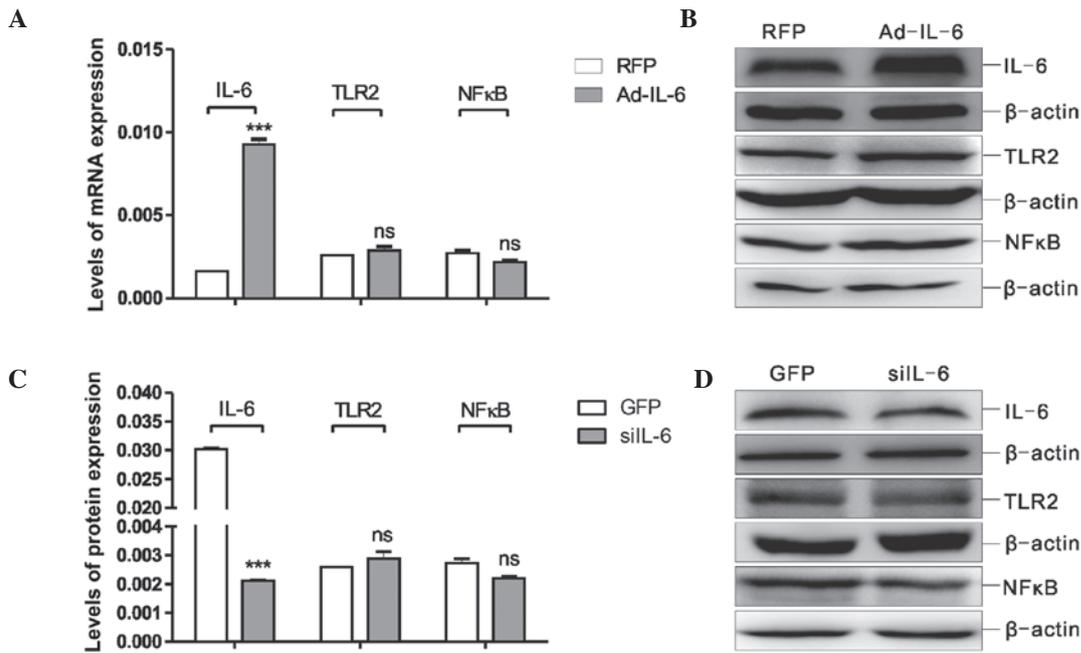


Figure 4. Determination of the changes in the TLR2/NFκB /IL-6 pathway in MSCs following treatment with Ad-IL-6 or siIL-6. (A) RT-qPCR analysis of the mRNA expression levels of IL-6, TLR2 and NFκB in MSCs following treatment with Ad-IL-6. The values are expressed as the mean \pm standard error of the mean. *** $P < 0.001$ vs. the RFP group. (B) Western blot analysis of IL-6, TLR2 and NFκB protein levels in MSCs following treatment with Ad-IL-6. (C) RT-qPCR analysis of the mRNA expression levels of IL-6, TLR2 and NFκB in MSCs following treatment with siIL-6. The values are expressed as the mean \pm standard error of the mean. *** $P < 0.001$ vs. the RFP group. (D) Western blot analysis of IL-6, TLR2 and NFκB protein levels in MSCs following treatment with siIL-6. For the western blot analyses (B and D), the data were obtained from at least three independent experiments, analysed and normalised against β -actin. MSCs, mesenchymal stromal cells; NFκB, nuclear factor κ B; IL-6, interleukin-6; Ad-IL-6, IL-6 adenovirus; RFP, red fluorescent protein; siIL-6, IL-6 small interfering RNA; GFP, green fluorescent protein; TLR2, Toll-like receptor 2.

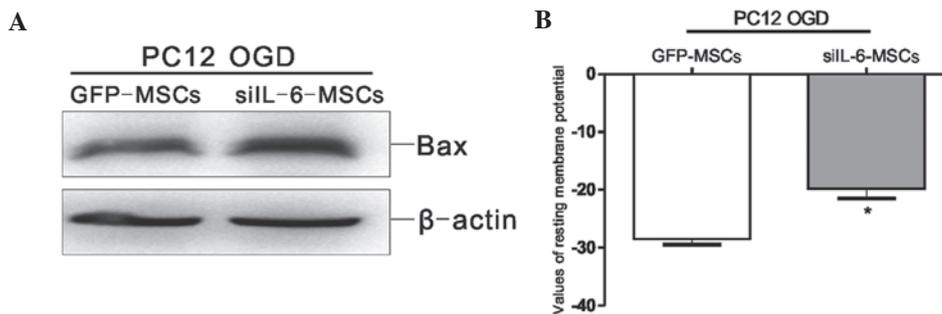


Figure 5. Effect of siIL-6-MSCs on the (A) expression level of Bax and (B) RMP in OGD-injured PC12 cells. (A) Changes in Bax protein expression in OGD-injured PC12 cells co-cultured with GFP-MSCs or siIL-6-MSCs. (B) Changes in the RMP of OGD-injured PC12 cells following co-culture with GFP-MSCs and siIL-6-MSCs, as determined by whole-cell patch-clamp recordings. The values are presented as the mean \pm standard error of the mean for data from 6-8 samples (* $P < 0.05$ vs. the GFP-MSC co-culture group). GFP, green fluorescent protein; MSC, mesenchymal stromal cells; RMP, resting membrane potential; siIL-6, interleukin-6 small interfering RNA; Bax, B-cell lymphoma-associated X; OGD, oxygen and glucose deprivation.

with stable siIL-6 expression. When the expression level of IL-6 decreased, the expression level of Bax increased and the RMP was induced. Bax is an apoptotic factor, and the increase in Bax expression levels indicated that apoptosis of the OGD-injured PC12 cells was aggravated. The RMP is the membrane potential of a nerve cell in an unstimulated state, and is known as the resting potential or transmembrane resting potential. The RMP threshold of neurons represents their viability and transmitting function. The whole-cell patch-clamp recordings revealed that the RMP threshold of the OGD-injured PC12 cells following co-culture with siIL-6-MSCs experienced a greater increase compared with that of the GFP-MSC co-culture group. These findings indicated that the decrease in the expression levels of

IL-6 in the MSCs reduced the capability of the OGD-injured PC12 cells to recover, thereby suggesting that endogenous IL-6 secreted from MSCs improved the functional restoration of OGD-injured PC12 cells.

The present study suggested that endogenous IL-6 secreted from MSCs had neuroprotective effects; however, the mechanism by which the MSCs secreted IL-6, and the signalling pathway(s) that are involved, have yet to be fully elucidated. In this preliminary study, the expression levels of TLR2 and IL-6 were shown to be increased when the PC12 cells were subjected to OGD injury. In the CNS, TLRs are not only key components of the innate immune system, but they also are associated with nerve degeneration and tissue damage (21,22). TLR2 is a member of

the TLR family that is highly expressed in the microglia of the CNS. NF κ B is activated by TLR2 via the myeloid differentiation primary response gene 88 (MyD88) pathway in the presence of interleukin-1 receptor-associated kinase (IRAK)-1 and IRAK-4, inducing the production of cytokines and chemical factors, including IL-1, IL-6 and TNF (23-25). In the current study, to investigate whether endogenous IL-6 secretion from MSCs was also regulated via the TLR2/NF κ B signalling pathway, the MSCs were treated with the TLR2 agonist, PGN-SA. Compared with the control group, the expression levels of TLR2, NF κ B and IL-6 increased following PGN-SA treatment. After the MSCs had been treated with siTLR2 adenovirus, the expression levels of TLR2, NF κ B and IL-6 markedly decreased. These results demonstrated that TLR2 expression in MSCs directly affected the expression levels of NF κ B and IL-6. NF κ B is an important factor in the TLR2 signalling pathway, and exerts a key role in the induction of trans-shipment in the innate immune system (26-28). The NF κ B inhibitor, PDTC, was also used in the present study to treat MSCs. Following treatment of the MSCs with 10 μ g/ml PDTC for 48 h, a significant decrease in the expression level of IL-6 in the MSCs was observed. However, the expression level of TLR2 was not significantly different compared with untreated MSCs. These data further demonstrated that endogenous IL-6 secreted from MSCs was regulated via the TLR2/NF κ B signalling pathway.

At this stage, it had not been established whether IL-6 feedback regulated the expression levels of TLR2 and NF κ B. To investigate this hypothesis, MSCs were treated with Ad-IL-6 adenovirus and siIL-6, and Ad-IL-6 and siIL-6 were revealed to change the expression level of IL-6, but not affect the expression levels of TLR2 or NF κ B in the MSCs. These data suggested that endogenous IL-6 secreted from MSCs was regulated by the TLR2/NF κ B signalling pathway, although IL-6 did not regulate the expression levels of TLR2 and NF κ B via a feedback mechanism. Therefore, further studies should focus on the biological effects of endogenous IL-6 from MSCs on the process of HIBD treatment.

In conclusion, the release of IL-6 from MSCs was regulated via the TLR2/NF κ B signalling pathway. Endogenous IL-6 secreted from MSCs was able to reduce the levels of apoptosis and to improve the functional restoration of OGD-injured PC12 cells following co-culture with MSCs. These findings represent a novel immunomodulatory effect of the neural injury microenvironment during MSC cytotераpy.

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