Umbilical Cord-derived Mesenchymal Stem Cells with Surfactant Protein B Alleviates Inflammatory Response in Acute Respiratory Distress Syndrome by Regulating Macrophage Polarization

Hongtao Xu, Guoming Nie, Taiyong Yin, Cui Shao, Dongsheng Ding, Minshu Zou

Department of Pediatrics, General Hospital of Central Theater Command of Chinese People’s Liberation Army, China

INTRODUCTION

Acute respiratory distress syndrome (ARDS) is a severe condition from acute lung injury (ALI) and is a respiratory failure disorder due to hypoxemia, with extremely high morbidity and mortality in critical care. It accounts for approximately 75,000 deaths annually in America. Pulmonary infection is the primary etiology of ARDS, in which, especially, lipopolysaccharide (LPS) in Gram-negative bacillus increases the permeability of the alveolar-capillary barrier, thereby leading to non-cardiogenic pulmonary edema. The inflammatory cascading reaction that is dominated by macrophages and neutrophils is the main mechanism in the pathological process of ARDS. Mass accumulation and infiltration of inflammatory cells in the lungs contribute to excessive inflammatory cytokine release and anti-inflammatory cytokine reduction with a consequence of inflammatory disequilibrium. Additionally, macrophages are considered indispensable mediators in ALI. In the lung microenvironment, the alveolar macrophages are polarized into two subpopulations, including M1 and M2 phenotype macrophages. M1-like polarization contributes to the pro-inflammatory reaction of host defense, whereas M2-like polarization exerts an important function in anti-inflammatory response and tissue remodeling. Therefore, alveolar macrophage polarization is critical for inflammatory equilibrium in the lungs.

Mesenchymal stem cells (MSCs) possess a strong therapeutic potential and exist in wide types of tissues. Promising features, including pluripotency and immune properties of hypo-immunogenicity, make MSCs superior candidates for cell therapies. Presently, MSC transplantation has been used in refractory pulmonary disease studies, including bronchopulmonary dysplasia, pulmonary fibrosis, and ARDS. Compared with other MSCs, umbilical cord-derived MSCs (UC-MSCs) show stronger abilities in tissue damage repair, immunoregulation, and differentiation, as well as...
higher gene transfection efficiency.\(^\text{17}\) Surfactant protein B (SP-B) is a key protein in pulmonary surfactant and functions in maintaining mammalian breathing\(^\text{18}\) and is required to maintain lung function, as genetic SP-B knockdown leads to respiratory failure in newborns\(^\text{19}\) and SP-B knockout mice.\(^\text{20}\) Animal-derived lung surfactants that contain native SP-B have been reported to improve outcomes in patients with RDS having surfactant deficiency.\(^\text{21}\)

We hypothesized that SP-B-transfected MSCs, which retain cell activity and various physiological effects, may yield stronger therapeutic effects on ARDS after transplantation into the lung. Therefore, this study aimed to transplant UC-MSCs transfected with SP-B into the lung of ARDS mice. By analyzing the pathological changes and inflammatory response of the lung tissue, the role of SP-B transfected UC-MSCs in ARDS was explored.

**MATERIALS AND METHODS**

**Isolation of UC-MSCs**

UC tissues (6–9 cm) were taken from pregnant women with full-term deliveries. This study was approved by the Ethics Committee of General Hospital of Central Theater Command of Chinese People’s Liberation Army (Hubei, China) (approval number: KY2019-023). Informed consent was signed by all participants. The tissues of 1 mm\(^3\) were washed with phosphate-buffered saline (PBS), digested in 0.1% collagenase II (Gibco, USA) that contain 30 U/ml DNase I for 1.5 h, and treated with 0.125% trypsin (Gibco) for 1 h. After terminating the trypsin activity, the mixture was filtered through a 100-μm cell strainer. Cells were then washed and cultured in Dulbecco’s Modified Eagle Medium/Nutrient Mixture F-12 (DMEM/F12) (1:1) (Gibco) that contain 100 μg/ml of penicillin, 10% fetal bovine serum (FBS) (Gibco), and 10 μg/ml of streptomycin (Sigma–Aldrich) at 37°C with 5% CO\(_2\). The adherent cells were incubated until confluence.

**Flow Cytometry**

Flow cytometry was applied to determine the presence of specific surface antigens of MSCs. Briefly, 1 x 10\(^5\) UC-MSCs that were suspended in 500 μl staining buffer with 1% FBS were cultured for 30 min at 4°C. Afterward, cells were stained with the phycoerythrin-conjugated monoclonal antibodies against CD44, CD29, CD34, and CD45 cell surface markers. All primary antibodies were obtained from Becton Dickinson (CA, USA). CytoFLEX Flow Cytometer (Beckman Coulter) that is equipped with CellQuest software (Becton Dickinson) was used to analyze the results.

**Osteogenic Differentiation**

UC-MSCs of passage 5–6 were incubated in the osteogenic differentiation culture medium containing DMEM/F12 (1:1), 100 μg/ml of penicillin, 10% FBS, 60 μM of ascorbic acid, 10 μg/ml of streptomycin, 10 nM of dexamethasone, and 10 mM of β-glycerophosphate disodium (Sigma)\(^\text{22}\) for 21 days. The medium was changed every 3 days. The staining was performed using Alizarin red S staining (Sigma). Mineralization during osteogenesis was monitored.

**Adipogenic Differentiation**

UC-MSCs of passage 5–6 were incubated in the adipogenic differentiation medium containing DMEM/F12 (1:1), 100 μg/ml of penicillin, 10% FBS, 50 mM of indomethacin, 10 μg/ml of streptomycin, 200 nM of insulin, 100 nM of dexamethasone, and 0.5 mM of isobutyl-methylxanthine. Cells were seeded in 6-well plates at 2 x 10\(^4\) cells/cm\(^2\). After 21 days, cells were stained with Oil Red O.

**Plasmid Construction and Transfection**

The whole coding region of SP-B was ligated into the pIRES2-enhanced green fluorescent protein (EGFP) plasmid at specific restriction sites (BamHI and HindIII). The target fragment sequence was tested by sequencing. UC-MSCs were cultured up to 70% confluence and then transfected with pIRES2-EGFP-SP-B recombinant plasmid using Lipofectamine 3000 (Invitrogen). Cells were cultured in a selection medium that is supplemented with 400 μg/ml of geneticin after 48 h. Stable clones were screened. Western blotting was used to confirm the efficiency.

**Western Blotting**

Radioimmunoprecipitation assay buffer (Invitrogen) was employed to extract the protein from the transfected UC-MSCs. Protein was separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and then transferred to polyvinylidene difluoride membranes. The membranes were incubated with primary antibodies SP-B (ab271345) and glyceraldehyde 3-phosphate dehydrogenase (ab181602) overnight at 4°C after blocking with 5% skim milk. Appropriate secondary antibodies were then added for 1 h of incubation at 4°C. Blot bands were determined with enhanced chemiluminescence reagent (Amersham, USA) and analyzed with Image J software.

**Experimental Animal**

Male BALB/c mice (8-week-old; Vital River Co. Ltd, Beijing, China) were raised under controlled conditions (22°C–26°C; 12 h light/dark cycle). Animal experiments were approved by the Animal Ethics Committee of General Hospital of Central Theater Command of Chinese People’s Liberation Army (Hubei, China) (approval number: DWLL2019-011).

Forty-two mice were evenly divided into six groups as follows: sham (intratracheally received normal saline); UC-MSCs (intratracheally received normal saline plus UC-MSCs); UC-MSCs-SP-B (intratracheally received normal saline plus UC-MSCs transfected with pIRES2-EGFP-SP-B); LPS (intratracheally received LPS); LPS + UC-MSCs-vector (intratracheally received LPS plus UC-MSCs transfected with empty plasmid), and LPS + UC-MSCs-SP-B (intratracheally received LPS plus UC-MSCs transfected with pIRES2-EGFP-SP-B).

**Experimental Model**

BALB/c mice were administrated with LPS.\(^\text{23}\) Animals were anesthetized via an intraperitoneal injection of pentobarbital sodium (60 mg/kg) before trachea exposure. After the anesthesia,
mice were intratracheally injected with LPS (O55:B5; 4 mg/kg; Sigma) in 100 μl of PBS or normal saline (0.1 ml/mouse) as control through a 20-gage catheter. After 6 h, 1 × 10⁶ UC-MSCs or UC-MSCs transfected with SP-B in 100 μl of PBS were intratracheally injected into the mice. After 72 h, the mice were euthanized by cervical dislocation under isoflurane. Lung tissues and bronchoalveolar lavage fluids (BALFs) were collected.

**Histology Analysis**

Lung tissues from each group were excised 72 h post-transplantation and fixed in 10% formalin under a pressure of 15 cm H₂O. After fixation for 24 h, tissues were embedded in paraffin. Next, 5-μm-thick specimens were deparaffinized and then stained with hematoxylin and eosin. Lung injury degree was observed with a microscope and semi-quantified according to a previously published lung injury scoring system.²⁴

**Wet/dry (W/D) Ratio**

Lung tissues were stored in a dry plastic plate, and the wet weight was immediately weighed with an electronic scale. The tissues were dried in an oven for 48 h at 60 °C and the dry weight was weighed. The W/D ratio indicated the ratio of the wet weight to the dry weight.²⁵

**Arterial Blood Gas**

As documented²⁶, mice were anesthetized with pentobarbital sodium 72 h post-transplantation. All mice maintained spontaneous breathing at the time of the experiment. Blood was taken from the celiac artery. An ABL90 FLEX analyzer (Radiometer, Denmark) was used to analyze oxygen partial pressure (pO₂) and saturation. The pO₂/fraction of inspired oxygen (FiO₂) ratio was calculated.

**Measurement of BALF**

Lung tissues were washed with 0.35 ml PBS for BALF sample collections, and BALF samples were centrifuged at 300 g for 10 min at 4 °C. Inflammatory cytokines were measured in the supernatants. Total protein concentration was assessed with bicinchoninic acid assays (Sigma). The precipitated cells were centrifuged at 220 g for 10 min at 4 °C. After washing, cells were then stained with CD11b (Cat.553310), CD11c (Cat. 558079), F4/80 (Cat. 565410), CD45 (Cat. 557659), and CD206 (Cat. 564996) for 1 h at 4 °C. Flow cytometry (Accuri C6, BD) was used to detect the single-cell suspension containing 10⁶–10⁷ cells. T-regulatory (Treg) (Foxp3⁺/CD25⁺) and Th17 (IL-17⁺/CD4⁺) were stained with antibodies, including anti-Foxp3-APC and anti-IL-17A-perCP, to detect T cell and T-helper 17 (Th17) cell differentiation.²⁷ All fluorescent-labeled antibodies, intracellular staining reagents, and FcR blockers for this assay were purchased from BioLegend (BioLegend, USA) or eBioscience (eBioscience, USA). Flow cytometry data were analyzed using FlowJo software (BD Biosciences, USA).

**Statistical Analysis**

Statistical analysis was performed using the Statistical Package for the Social Sciences version 21.0 (Chicago, IL, USA). Values are expressed as the means ± standard deviation. One-way analysis of variance followed by Tukey’s post-comparison test or t-test was applied to analyze the differences between groups. The normal distribution of data was tested using the Shapiro–Wilks test. The Mann–Whitney U test was used for non-normally distributed data. A p-value of <0.05 was statistically significant. A sample size of seven cases for each group was required to detect at least 50% difference between the control and test groups, with a power of 85% at the 1.25% significance level.

**RESULTS**

Identification of MSC Characteristics

First, the antigenic profile of isolated UC-MSCs was determined according to minimal criteria for defining pluripotent mesenchymal stromal/stem cells.²⁸ The population of UC-MSCs highly expressed the MSCs markers, CD44 and CD29, rather than the hematopoietic markers, CD34 and CD45 (Figure 1A-1D). Next, UC-MSCs were induced to differentiate into osteoblasts and adipocytes in a selective culture medium. As shown in Figure 1E, Alizarin Red Staining of UC-MSCs showed large calcium phosphate deposits, and blue staining was observed with hematoxylin and eosin. Lung injury degree was observed with a microscope and semi-quantified according to a previously published lung injury scoring system.²⁴

**Enzyme-linked Immunoassay (ELISA)**

The concentrations of interleukin (IL)-1β (Cat.EMC001b), IL-10 (Cat.EMC005), IL-13 (Cat.EMC124), tumor necrosis factor α (TNFα) (Cat.EMC102a), transforming growth factor (TGF-β) (Cat. F2686-B), and prostaglandin E2 (PGE2) (Cat.F9428-B) in BALF were measured using the ELISA kits (Neo-Bioscience, Shanghai; Hufeng Chemical, Shanghai) following the manufacturer’s instructions.

**Macrophage Phenotype Analysis**

Entire lung tissues were collected, chipped, and cultured on a shaker at 260 rpm for 1 h. The mixture was filtered through a 40-μm cell strainer to remove the nondigested fragment. Next, 1 ml of FBS was added to the cell suspension, and the mixture was centrifugated at 220 g for 10 min. The collected precipitates were resuspended, and red blood cells were discarded with lysis buffer. Afterward, the samples were washed with PBS containing 1% FBS. Cells were cultured with antibody CD16/CD32 (Cat.553141; Mouse BD Fc Block) for 15 min at 4 °C and then washed with PBS. Fixable viability stain 620 (Cat.564996; BD Pharmingen) was incubated with the cell suspension for 20 min at 18 °C. After washing, cells were then stained with CD11b (Cat.553310), CD11c (Cat. 558079), F4/80 (Cat. 565410), CD45 (Cat. 557659), and CD206 (Cat. 564996) for 1 h at 4 °C. Flow cytometry (Accuri C6, BD) was used to detect the single-cell suspension containing 10⁶–10⁷ cells. T-regulatory (Treg) (Foxp3⁺/CD25⁺) and Th17 (IL-17⁺/CD4⁺) were stained with antibodies, including anti-Foxp3-APC and anti-IL-17A-perCP, to detect T cell and T-helper 17 (Th17) cell differentiation.²⁷ All fluorescent-labeled antibodies, intracellular staining reagents, and FcR blockers for this assay were purchased from BioLegend (BioLegend, USA) or eBioscience (eBioscience, USA). Flow cytometry data were analyzed using FlowJo software (BD Biosciences, USA).
Xu et al. Role of Umbilical Cord-derived Mesenchymal Stem Cells in ARDS

SP-B Strengthens the Protection of UC-MSCs Against LPS-stimulated ARDS

The therapeutic effects of UC-MSCs were examined. Histological changes in the lung were observed by hematoxylin and eosin staining. No significant difference was found in the histologic change among sham, UC-MSCs, and UC-MSCs-SP-B groups (Figure 2A). Normal alveolar and interstitial tissue structures were destroyed, and the inflammatory infiltration was serious in the LPS group, whereas these changes were markedly attenuated after UC-MSCs transplantation together with SP-B transfection. LPS increased lung injury score in mice (vs. sham group: \( P = 0.000 \)). UC-MSCs-SP (vs. LPS group, \( P = 0.009 \)) and UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group, \( P = 0.017 \)) markedly reduced lung injury score (Figure 2B). The effect of UC-MSCs with SP-B transfection on pulmonary function was evaluated by measuring the W/D ratio and arterial blood gases. Data showed that the mice in LPS + UC-MSCs (vs. LPS group: \( P = 0.018 \)) and LPS + UC-MSCs-SP-B (group vs. LPS + UC-MSCs-vector group, \( P = 0.003 \)) groups had lower W/D ratio of 6.0 and 4.2, respectively, than 8.0 in LPS (Figure 2C), which suggest that UC-MSCs either alone or combined with SP-B attenuates the degree of LPS-induced lung edema. LPS also reduced the pO2/FiO2 ratio (vs. sham group: \( P = 0.000 \)), whereas UC-MSCs (vs. LPS group: \( P = 0.008 \)) and UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: \( P = 0.017 \)) markedly restored the pO2/FiO2 ratio (Figure 2D), which demonstrated lung function improvement. Overall, SP-B transfection into UC-MSCs strengthens the protective effects of UC-MSCs against LPS-stimulated ARDS.

UC-MSCs and UC-MSCs-SP-B Decreases the Degree of BALF Changes

Cell counting and protein level in BALF were measured for further analysis of lung damage and inflammation. Total protein level was elevated in LPS group (vs. sham group: \( P = 0.000 \)) but reduced in LPS + UC-MSCs-vector (vs. LPS group: \( P = 0.012 \)) and LPS + UC-MSCs-SP-B groups (vs. LPS + UC-MSCs-vector group: \( P = 0.029 \)) (Figure 3A). Cell number was increased in LPS group (vs. sham group: \( P = 0.000 \)) but reduced in LPS + UC-MSCs-vector (vs. LPS group: \( P = 0.015 \)) and LPS + UC-MSCs-SP-B groups (vs. LPS + UC-MSCs-vector group: \( P = 0.008 \)) (Figure 3B). Additionally, LPS increased the neutrophils in BALF (vs. sham group: \( P = 0.000 \)), whereas UC-MSCs (vs. LPS group: \( P = 0.002 \)) or UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: \( P = 0.002 \)) decreased this number (Figure 3C). These findings suggested that UC-MSCs either alone or combined with SP-B not only improve lung injury in ARDS but attenuate cellular infiltration.

**FIG. 1.** Identification of MSC characteristics. (a-d) For immunophenotypic analysis, cell surface markers (CD44, CD29, CD34, CD44, and CD45) in isolated UC-MSCs were detected using flow cytometry analysis. (e) Osteoblastic and adipogenic potential of UC-MSCs was examined by Alizarin Red and Oil Red O staining, respectively. (f) Transfection efficiency of pIREs2-EGFP-SP-B into UC-MSCs was verified by reverse transcription-quantitative polymerase chain reaction. \(^* P < 0.01.\)**
SP-B Strengthens the Promotive Effect of UC-MSCs on M2 Polarization

The impact of UC-MSCs on alveolar macrophage polarization in LPS mice was assessed using flow cytometry analysis. CD11c−CD206− indicates an M1-like polarization and CD11c−CD206+ indicates an M2-like polarization. As illustrated in Figure 4A-4C, percentages of both M1 and M2 macrophages were elevated in LPS mice compared to sham mice (vs. sham group: P = 0.000). However, LPS + UC-MSCs-vector (vs. LPS group: P = 0.006) and LPS + UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: P = 0.004) groups showed lower percentage of M1 and higher percentage of M2 than in LPS group. Changes in the M2/M1 ratio revealed a dominant M1 polarization compared to M2 polarization in the LPS group (vs. sham group: P = 0.017), whereas UC-MSCs (vs. LPS group: P = 0.001) or UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: P = 0.002) significantly promoted macrophage polarization toward M2, as shown by increased M2/M1 ratio (Fig. 4D). ELISA was used to measure the concentrations of pro-inflammatory cytokines (IL-1β and TNFα) and anti-inflammatory chemokines (IL-10, IL-13, TGF-β and PGE2) in BALF to analyze inflammatory regulation mediated by UC-MSCs. The increased levels of LPS-induced IL-1β and TNFα were significantly reversed by UC-MSCs (vs. LPS group: P = 0.006 and P = 0.006, respectively) or UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: P = 0.001 and P = 0.003, respectively) (Figure 4E, 4F). Additionally, in parallel, the levels of IL-10, IL-13, TGF-β, and PGE2 were increased in LPS (vs. sham group: P = 0.004, P = 0.001, P = 0.001, and P = 0.001, respectively), LPS + UC-MSCs-vector group (vs. LPS group: P = 0.010, P = 0.001, P = 0.001, and P = 0.001, respectively), and UC-MSCs-SP-B (vs. LPS + UC-MSCs-vector group: P = 0.013, P = 0.015, P = 0.010, and P = 0.008, respectively) (Figure 4G-4I). Overall, UC-
MSCs and UC-MSCs-Sp-B inhibit inflammation by promoting M2-like polarization in ARDS.

**UC-MSCs-Sp-B Reduces Th17 Differentiation and Promotes Treg Differentiation**

The effects of UC-MSCs-Sp-B on the regulation of T cell and Th17 were further examined. As expected, a markedly increased number of IL-17+CD4+ cells and reduced Foxp3+CD25+ after the addition of LPS was found (vs. sham group: \( P = 0.001 \) and \( P = 0.001 \), respectively). Notably, UC-MSCs (vs. LPS group: \( P = 0.013 \) and \( P = 0.001 \), respectively) or UC-MSCs-Sp-B (vs. LPS + UC-MSCs-vector group: \( P = 0.029 \) and \( P = 0.017 \), respectively) inhibited Th17 differentiation, whereas promoted Treg differentiation (Figure 5A, 5B). Therefore, UC-MSCs-Sp-B displays a key role in modulating the inflammatory differentiation of T cells.

**DISCUSSION**

ARDS is related to diffuse epithelial injury that is induced by an impaired alveolar-capillary barrier with consequently raised permeability, pulmonary edema, and neutrophil accumulation. During this process, persistent inflammation and plasma protein loss contribute to lung dysfunction. Insufficient effective treatment for ARDS is associated with the multifactorial etiology of this disorder.29,30 MSCs could be a promising candidate for cell therapy in inflammatory pulmonary diseases. Evidence has shown beneficial effects from the transplant of bone marrow or umbilical cord-derived MSCs in lung injury treatment.31 The effects of UC-MSCs in combination with SP-B in ARDS were preliminarily explored.

Inflammation is a critical pathophysiological event in ARDS, which is a complicated and highly coherent process.32 The transition from acute inflammation to chronic inflammation may result in tissue damage and ultimately multiple organ failure.33 In the present investigation, LPS, an important component of the gram-negative bacilli cell wall, induces a significant inflammatory response in the BALF of mice. Normal alveolar and interstitial tissue structure were destroyed accompanied by a high lung injury score after LPS instillation. Additionally, LPS aggravated pulmonary edema and neutrophil infiltration. UC-MSCs have more primitive, proliferative, and immunosuppressive characteristics.17 UC-MSCs effectively restore protein permeability in ALI and confer more practical and functional advantages than their adult counterparts.34 Studies have suggested that UC-MSCs infusion effectively decreases lung inflammation, prevents pulmonary fibrosis, increases blood counts, recovers hematopoiesis, and improves survival in shrews with acute radiation injury.35 Moreover, UC-MSC infusion was an effective therapeutic tool for patients with ARDS.36 This study revealed that intratracheal infusion of UC-MSCs in combination with SP-B attenuated inflammation and improved lung function in experimental ARDS models.

Resident inflammatory cells-alveolar macrophages are activated after stimulation and produce numerous cytokines, which promote the chemotaxis of neutrophils, monocytes, and lymphocytes, thereby aggravating the inflammatory lung responses.37 M1 macrophages potentiate the local inflammatory response by secreting inflammatory mediators, whereas M2 macrophages can inhibit inflammatory damage by releasing anti-inflammatory mediators. Once the pulmonary microenvironment is stimulated, alveolar macrophages are immediately transformed into the M1 phenotype.8,38 Our results showed that the M1 phenotype was dominant after LPS stimulation compared to the M2 phenotype. MSCs promote the differentiation of macrophages into M2 and exert an anti-inflammatory effect in neonatal lung injury.39 MSCs treatment alleviates bronchopulmonary dysplasia, improves lung function, and ameliorates pulmonary hypertension by pulmonary macrophage phenotype regulation.40 According to previous research, the ratio of pulmonary M1/M2 macrophages is increased.

**FIG. 3.** UC-MSCs and UC-MSCs-SP-B decreases the degree of BALF changes. (a) Total protein level in BALF. (b) Cell counting in BALF. (c) Neutrophil counting in BALF. \( P < 0.05 \), \( ** P < 0.01 \), *** \( P < 0.001 \). \( N = 7 \).
FIG. 4. SP-B strengthens the promotive effect of UC-MSCs on M2 polarization. (a) Polarization of alveolar macrophages using flow cytometry analysis. (b) The change in M1 (CD11c\(^+\)CD206\(^-\)) polarization. (c) The change in M2 (CD11c\(^-\)CD206\(^+\)) polarization. (d) The M2/M1 ratio. (e-j) The concentrations of IL-1\(\beta\), TNF\(\alpha\), IL-10, IL-13, TGF-\(\beta\), and PGE\(2\) in BALF were measured by ELISA. \(* P < 0.05, \quad ** P < 0.01, \quad *** P < 0.001. N = 7.\)
accompanied by reduced SP-B expression, which indicates lung injury initiation. Here, after LPS treatment, intratracheal transplantation of either UC-MSCs or UC-MSCs-SP-B suppressed the M1-like polarization and augmented the M2-like polarization in the BALF. This suggested that UC-MSCs transfected with SP-B regulate inflammation in ARDS through macrophage polarization modulation. Moreover, UC-MSCs-SP-B reduced Th17 differentiation and promoted Treg differentiation after ARDS.

SP-B preferentially binds to anionic surfactant lipids that potentiate the capacity of SP-B of facilitating lipid adsorption to the air-water interface. The SP-B dimer is oligomerized in a circular structure, which mediates the connection between surfactant membranes. SP-B knockdown contributes to lung inflammation, thereby supporting the models that surface force disruption activates pro-inflammatory signaling. After SP-B is lost, L-selectin is rapidly produced. L-selectin can mediate lung inflammation through increased leukocyte mobility into the lungs. STAT-3 activates many genes that are involved in lung protection. After doxycycline withdrawal, STAT-3 is phosphorylated and upregulated, which may indicate a compensatory response to SP-B deficiency. Additionally, phosphorylation of STAT-3 was reported crucial for repolarization of M1 to M2 type. Therefore, M1/M2 polarization mechanisms regulated by SP-B may involve multiple molecules and signals, which need further investigations in the future.

Overall, our findings showed that transplantation of UC-MSCs transfected with SP-B could potentiate M2 macrophage polarization and further relieve LPS-stimulated lung injury. This demonstrated that the combination of UC-MSCs with SP-B yields significant alleviative effects on ARDS, which may be a promising strategy for ARDS treatment.
REFERENCES


Xu et al. Role of Umbilical Cord-derived Mesenchymal Stem Cells in ARDS


37. Fan EKY, Fan J. Regulation of alveolar macrophage death in acute lung inflammation. Respir Res. 2018;19:50. [CrossRef]


