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# In Vivo Effects of Mesenchymal Stromal Cells in Two Patients With Severe Acute Respiratory Distress Syndrome

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#### Abstract

Mesenchymal stromal cells (MSCs) have been investigated as a treatment for various inflammatory diseases because of their immunomodulatory and reparative properties. However, many basic questions concerning their mechanisms of action after systemic infusion remain unanswered. We performed a detailed analysis of the immunomodulatory properties and proteomic profile of MSCs systemically administered to two patients with severe refractory acute respiratory distress syndrome (ARDS) on a compassionate use basis and attempted to correlate these with in vivo anti-inflammatory actions. Both patients received  $2 \times 10^{6}$  cells per kilogram, and each subsequently improved with resolution of respiratory, hemodynamic, and multiorgan failure. In parallel, a decrease was seen in multiple pulmonary and systemic markers of inflammation, including epithelial apoptosis, alveolar-capillary fluid leakage, and proinflammatory cytokines, microRNAs, and chemokines. In vitro studies of the MSCs demonstrated a broad antiinflammatory capacity, including suppression of T-cell responses and induction of regulatory phenotypes in T cells, monocytes, and neutrophils. Some of these in vitro potency assessments correlated with, and were relevant to, the observed in vivo actions. These experiences highlight both the mechanistic information that can be gained from clinical experience and the value of correlating in vitro potency assessments with clinical effects. The findings also suggest, but do not prove, a beneficial effect of lung protective strategies using adoptively transferred MSCs in ARDS. Appropriate randomized clinical trials are required to further assess any potential clinical efficacy and investigate the effects on in vivo inflammation. Stem Cells Translational Medicine 2015;4:1199–1213

# SIGNIFICANCE

This article describes the cases of two patients with severe refractory adult respiratory syndrome (ARDS) who failed to improve after both standard life support measures, including mechanical ventilation, and additional measures, including extracorporeal ventilation (i.e., in a heart-lung machine). Unlike acute forms of ARDS (such in the current NIH-sponsored study of mesenchymal stromal cells in ARDS), recovery does not generally occur in such patients.

#### INTRODUCTION

Acute respiratory distress syndrome (ARDS) is a severe, life-threatening medical condition characterized by widespread uncontrolled inflammation in the lungs associated with the loss of surfactant and impaired pulmonary capillary endothelial barriers, resulting in fluid accumulation in the distal airspaces. A number of etiologies can trigger ARDS, including primary respiratory infections, sepsis, and significant trauma. ARDS continues to result in high mortality and high costs related to often prolonged intensive care unit (ICU) and hospital stays [1–4]. Furthermore, ARDS survivors often have long-term pulmonary, neuromuscular, and cognitive symptoms and a diminished quality of life [2–4]. A number of different pharmacologic therapies have failed to demonstrate benefit, and treatment is currently limited to supportive care and appropriate use of mechanical ventilation and fluid management [5, 6]. The addition of extracorporeal membrane oxygenation (ECMO) has been used in patients with severe ARDS; however, in a recent systematic review and meta-analysis of current evidence, no association with improved outcomes could be demonstrated in adult patients [4]. As such, new therapeutic approaches are needed.

Adoptive transfer of mesenchymal stromal cells (MSCs) ameliorates experimentally induced acute lung injury in preclinical animal models and in ex vivo perfused human lungs [7-15]. Although not completely understood, the mechanisms of MSC actions in acute models of ARDS include the release of paracrine anti-inflammatory and antibacterial peptides and mitochondrial transfer through cellcell contact with damaged alveolar epithelial cells in the absence of permanent cell engraftment [10, 16-18]. Furthermore, MSCs release extracellular vesicles (EVs), which have reduced inflammation and promote tissue regeneration in different preclinical models [7, 19-21]. Because administration of non-human leukocyte antigen (HLA)matched allogeneic MSCs has also been demonstrated to be safe and potentially effective in a widening range of clinical applications in both lung diseases and other diseases [6, 8, 22-24], this approach could be beneficial in patients with ARDS. A recent phase I dose-escalation safety study demonstrated the safety of a single i.v. administration of 1-10 million cells per kilogram of MSCs in 9 patients with moderate to severe ARDS, and a phase II efficacy trial is currently under way [25].

However, despite abundant descriptions of in vitro and in vivo MSC actions in preclinical models, correlation of the in vitro MSC release of anti-inflammatory mediators and effects on immune effector cells in clinical applications remains poorly understood [26]. Furthermore, no animal models resembling patients with refractory ARDS who require additional support measures are available. Our incomplete understanding of the mechanisms of action of MSCs has limited the development of relevant in vitro potency assays that can define the MSC populations optimal for clinical use, including the selection of potent cell batches relevant to specific disease indications. More importantly, it has limited the selection of patients likely to benefit from treatment. The treatment of two patients with severe refractory ARDS using MSCs was studied in an attempt to correlate the in vivo actions of the MSCs on lung and systemic inflammation with a detailed investigation of in vitro potency assays, including the effects on inflammatory and immune modulatory cells, and proteomic assessments of the MSCs and EVs released by the MSCs.

#### MATERIALS AND METHODS

# **Ethical Considerations**

The production of clinical-grade MSCs was accredited by the Swedish Board of Health and Welfare (approval no. 9.1-57237/2012) and the Medical Products Agency (approval no. 5.9.2-2013-047346). ECMO is a last resort supportive therapy for ARDS. The condition of the two patients in our report had deteriorated with ECMO support; also, because the patients were sedated before MSC administration, informed consent could not be obtained. At a multidisciplinary conference, we decided to use MSCs on a compassionate use basis under the approval of the chief medical officer of the hospital, the hospital ethics committee, and the patients' relatives.

## Isolation and Expansion of Mesenchymal Stromal Cells

A total of 28 ml of bone marrow was aspirated from a 49-year-old HLA-mismatched, third-party, healthy male volunteer. Clinicalgrade MSCs were generated under good manufacturing practice (GMP) conditions according to a common protocol elaborated by the European Group for Bone and Marrow Transplantation Developmental Committee, accredited by the Swedish National Board of Health and Welfare [27]. Bone marrow mononuclear cells  $(146 imes 10^6)$  were seeded into 175 cm<sup>2</sup> flasks (Falcon, Franklin Lakes, NJ, http://www.BD.com) in Dulbecco's modified Eagle's mediumlow glucose (Life Technologies, Gaithersburg, MD, http://www. lifetechnologies.com) supplemented with lysed human platelets (final concentration, 10<sup>8</sup> per milliliter). When the cultures were near confluence (>80%), the cells were detached once by treatment with trypsin and EDTA (Invitrogen, Grand Island, NY, http://www.invitrogen. com) and replated once at a density of 4,000 cells per cm<sup>2</sup>. The MSCs were harvested and cryopreserved in 10% dimethyl sulfoxide (DMSO; WAK-Chemie Medical GmbH, Steinbach, Germany, http://www.wakchemie.net). After thawing, the cells were washed 3 times in phosphate-buffered saline (PBS) and resuspended in 0.9% saline solution with the addition of 10% AB plasma, to a final concentration of 2  $\times$  10<sup>6</sup> cells per milliliter. The MSC release criteria for clinical use included the absence of visible clumps, spindle shape morphology, the absence of contamination by pathogens (bacteria and mycoplasma), and viability >95%. The MSCs expressed CD73, CD90, CD105, and HLA-ABC and were negative for CD14, CD31, CD34, CD45, and HLA-DR (supplemental online Fig. 1A).

The total dose administered to the patients was  $2 \times 10^6$  cells per kilogram of recipient body weight and was determined from previous experience of MSC treatment of graft-vs.-host disease (GVHD) [27]. The dose was divided into 4 syringes of 50 ml, and each syringe was infused over 30 seconds. No visible clumps or cell debris were observed during the infusions.

#### **Clinical Outcome Measures**

Blood was collected as standard practice during routine care in the intensive care unit. Bronchoalveolar lavage (BAL) fluid was obtained from lavage performed at intervals as part of the routine clinical care provided to each patient. Blood samples for cell purification were collected in heparinized tubes. Serum and plasma samples for cytokines and extracellular vesicles were collected and frozen immediately after sample preparation and stored at  $-80^{\circ}$ C until analyzed. Chest x-rays (CXRs) and chest computed tomography scans were performed as part of the routine clinical care provided to each patient. Respiratory measurements, including lung volumes and pressure, were obtained from the mechanical ventilation record of each patient. Other standard laboratory measurements were taken as a part of the routine care for each patient and obtained from the patient's medical records.

#### **BAL Fluid and Serum Analyses**

The levels of pro- and anti-inflammatory cytokines in serial samples of BAL fluid and serum were assessed using a multiplex cytokine assay (EMD Millipore, Billerica, MA, http://www.emdmillipore.com) on a Luminex machine (EMD Millipore). In BAL fluid, the surfactant protein B concentration was determined by enzyme-linked immunosorbent assay (ELISA; USCN Life Science Inc., Houston, TX, http:// www.uscnk.com). Caspase-cleaved cytokeratin-18 (ccK18) and total K18 were measured using M30-Apoptosense ELISA (Peviva AB, Bromma, Sweden, http://www.peviva.com) and M65 EpiDeath ELISA (Peviva AB), respectively.

#### Assessment of Circulating MicroRNAs

Inhibitory microRNAs (miRNAs) were isolated from circulating EVs purified from the routine blood samples collected. EV purification

and miRNA isolation was performed by Exosome Diagnostic Inc. (New York, NY, http://www.exosomedx.com). In brief, 1.5-2 ml of the blood samples from each time point was purified using the exoRNeasy serum/plasma kit (Qiagen, Hilden, Germany, http://www.qiagen.com). Eluted RNA was processed for microRNA analysis using the low sample input protocol for the TagMan Open-Array Human MicroRNA Panel (Life Technologies). Megaplex RT Primers, Human Pool A and Human Pool B (Life Technologies), were used for reverse transcription followed by a preamplification step according to the manufacturer's protocol (95°C for 10 minutes, 55°C for 2 minutes, 72°C for 2 minutes, followed by 16 cycles of 95°C for 15 seconds, 60°C for 4 minutes, followed by 99°C for 10 minutes, followed by 4°C). The preamplified material was diluted 1:20 in Tris-EDTA (pH 8.0) for analysis. The samples were loaded onto OpenArray plates using the standard Accufill protocol (Life Technologies). Amplification was performed according to the protocol established for the TaqMan OpenArray Human MicroRNA panel downloaded for each plate (OpenArray Plate product page; http://www.lifetechnologies. com). A total of 758 miRNAs were assayed, and 200-300 miRNAs gave a detectable Ct value for each time point.

U6 small nuclear RNA (snRNA) is a noncoding small nuclear RNA commonly used for normalization of miRNA [28]. U6 was present in every sample and every subarray and was used to normalize for subarray-to-subarray variations and differences between two Megaplex pools.  $\Delta$ Ct values were calculated for each miRNA by subtracting the U6 Ct value located on the same subarray. The values were corrected for variation of the total RNA amount by subtracting each Ct value by the median Ct value of 128 miRNAs present in all samples in the first patient and by the median Ct of 79 miRNAs present in all samples in the second patient. This procedure was performed for all arrays and time points. Missing values were entered as the highest observed normalized Ct value per patient. Finally, the fold change for every miRNA relative to the normal healthy control sample was calculated.

# Mononuclear Cell Collection, Cell Purification, and In Vitro Mixed Lymphocyte Studies

The peripheral blood mononuclear cells (PBMCs) used in the in vitro studies were retrieved from the patients and healthy donors, isolated by density gradient-based centrifugation, and stored in 10% DMSO in liquid nitrogen until additional analysis. For further purification of T cells, a paramagnetic bead-based selection was used (Miltenyi Biotec, Bergisch Gladbach, Germany, http://www. miltenyibiotec.com). The same bone marrow-MSCs used in the patient were cocultured with allogeneic T cells (MSC/T-cell ratio 1:5 and 1:10) for 5 days. The T cells were stimulated with activating anti-CD2/CD3/CD28 antibodies (Miltenyi Biotec) in a 0.5 bead per cell ratio [29]. PBMCs were cultured for 5 days in the presence of MSCs (MSC/PBMC ratio of 1:5 and 1:10), and the monocytic compartment was subsequently analyzed. Polymorphonuclear leukocytes (PMNs) from buffy coats of healthy donors were ultra-purified under endotoxin-free conditions, as previously described [30]. PMNs (>95% purity) and MSCs were cocultured for up to 40 hours in the presence or absence of 100 ng/ml lipopolysaccharide (LPS; Invitrogen, Carlsbad, CA, http://www.invitrogen. com). In all cases, MSCs were plated for 72 hours before the start of cocultures. In selected experiments, MSCs were pretreated with recombinant human interferon- $\gamma$  (IFN- $\gamma$ ; 10 ng/ml; PeproTech, Rocky Hill, NJ, http://www.peprotech.com) and tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ; 15 ng/ml, R&D Systems, Minneapolis, MN, http://www.rndsystems.com) for 48 hours and then cocultured with freshly isolated PMNs. MSC and PMN cocultures were stained with May-Grünwald-Giemsa dye to observe the cell morphology after reciprocal interaction.

## In Vitro PMN Assessments

PMNs were stained according to the manufacturer's instructions with fluorochrome-coupled antibodies, as detailed in supplemental online Table 2. The LIVE/DEAD Fixable Aqua Dead Cell Stain Kit (Life Technologies) was used for the exclusion of dead cells for PBMC analysis and propidium iodide (Invitrogen) staining to test the viability of the MSCs. To perform intracellular staining, the cells were treated with the Cytofix/Cytoperm Fixation/Permeabilization Kit (BD Biosciences, San Jose, CA, http://www.bdbiosciences.com). After coculture with MSCs, PMNs were identified on the basis of their typical morphological parameters (forward scatter/side scatter) and their CD45 expression. An annexin-V-FITC staining kit (Miltenyi Biotec) was used to assess the levels of PMN apoptosis. The cells were analyzed using a FACS Canto II cytometer (BD Biosciences) and FlowJo, version 9.5 software (TreeStar, Ashland, OR, http://www.treestar.com).

# RNA Preparation and Quantitative Reverse Transcriptase-Polymerase Chain Reaction

Total RNA was extracted (RNeasy Mini Kit; Qiagen) and cDNA prepared (Superscript First Strand Synthesis System; Life Technologies) using a Mastercycler nexus (Eppendorf, Hamburg, Germany, http:// www.eppendorf.com). For cDNA transcription, a two-step procedure was performed in line with the manufacturer's recommendations. In brief, the final reaction was incubated for 10 minutes at 25°C, followed by 60 minutes at 60°C, and terminated by heating at 70°C for 15 minutes. Messenger RNA levels were quantified using quantitative polymerase chain reaction (PCR; Quantitect SYBR Green PCR Kit; Qiagen) on a Rotor Gene Q (Qiagen). Relative gene expression was determined by normalizing the expression to  $\beta_2$ -microglobulin using the  $2^{-\Delta\Delta Ct}$  method. The following gene-specific primers were used (forward and reverse sequence): indoleamine-2,3-dioxygenase (IDO), 5'-GCATTTTTCAGTGTTCTTCGCATA-3' and 5'-TCATACACCAGACC-GTCTGATAGC-3';  $\beta_2$ -microglobulin, 5'- TGCTGTCTCCATGTTTGATGTA-TCT-3' and 5'-TCTCTGCTCCCCACCTCTAAGT-3'.

#### In Vitro MSC Responses to Inflammatory Stimuli

The MSCs were tested in vitro for their responsiveness to inflammatory stimuli ("MSC licensing"). Culture in the presence of TNF- $\alpha$ (15 ng/ml) or IFN- $\gamma$  (10 ng/ml) for 48 hours induced the expression of CD54 (ICAM-1), CD106 (VCAM-1), and HLA-ABC and HLA-DR, as assessed by flow cytometry, and IDO mRNA transcripts was measured using quantitative PCR (supplemental online Fig. 1B).

#### **MSC EV Purification**

MSCs were cultured in serum-reduced medium (OptiMem; Gibco, Grand Island, NY, http://www.lifetechnologies.com) for 48 hours. The conditioned medium was harvested and spun at 300*g* for 5 minutes and filtered through 0.2- $\mu$ m sterile syringe filters to remove cell debris. The EVs were subsequently pelleted by ultracentrifugation at 110,000*g* for 90 minutes at +4°C. The pellets were resuspended in PBS, pooled, and ultracentrifuged again at 110,000*g* for 90 minutes at +4°C. The pellets were resuspended in PBS. Nanoparticle tracking analysis was performed using the NS500 (NanoSight, Wiltshire, U.K., http://www.malvern.com) to

measure the size distribution of the particles, which is based on the motion and light scatter of nanometer-size particles (Brownian motion) [31]. The number of particles and their movement were recorded using a camera level of 15 and automatic functions for all postacquisition settings except for the detection threshold, which was fixed at 6. The samples were diluted in PBS between 1:500 and 1:2,000 to achieve a particle count between 2  $\times$  10<sup>8</sup> and 2  $\times$  10<sup>9</sup> per milliliter. The camera focus was adjusted to make the particles appear as sharp dots. Using the script control function, 5  $\times$  30-second videos were recorded, incorporating a sample advance and a 5-second delay between each recording and analyzed using the NS500 software (NanoSight). The EVs were frozen at  $-80^\circ$ C until further analysis.

#### **EV Preparation for Proteomic Assessment**

Donor cell pellet and EVs were lysed with 4% SDS, 25 mM HEPES, 1 mM dithiothreitol (DTT). Lysates were heated to 95°C for 5 minutes followed by sonication for 1 minute and centrifugation at 14,000g for 15 minutes. The supernatant was mixed with 1 mM DTT, 8 M urea, and 25 mM HEPES (pH 7.6) and transferred to a centrifugation filtering unit, with a 10-kDa cutoff (Nanosep; Pall Corp., Port Washington, NY, http://www.pall.com), and centrifuged for 15 minutes at 14,000g, followed by another addition of the 8 M urea buffer and centrifugation. Proteins were alkylated by 50 mM indole-3-acetic acid, in 8 M urea and 25 mM HEPES for 10 minutes, centrifuged for 15 minutes at 14,000g, followed by 2 more additions and centrifugation with 8 M urea and 25 mM HEPES. Trypsin (Promega, Madison, WI, http://www. promega.com), at a 1:50 trypsin/protein ratio, was added to the cell lysate in 250 mM urea and 50 mM HEPES and incubated overnight at 37°C. The filter units were centrifuged for 15 minutes at 14,000g, followed by another centrifugation with Milli-Q water (EMD Millipore), and the flow through was collected. The peptides were cleaned using a strata-X-C-cartridge (Phenomenex, Torrance, CA, http://www.phenomenex.com).

# Nano Reverse Phase Chromatography Gradient Before Mass Spectrometry Analysis

Before analysis using the Q Exactive mass spectrometer (Thermo Fisher Scientific, San Jose, CA, http://www.thermoscientific.com), the peptides were separated using an Agilent 1200 nano-LC system (Agilent Technologies, Santa Clara, CA, http://www.agilent.com). The samples were trapped on a Zorbax 300SB-C18 (Agilent Technologies) and separated on a NTCC-360/100-5-153 (Nikkyo Technos, Ltd., Tokyo, Japan, http://www.nikkyo-tec.co.jp) column using a gradient of A (5% DMSO, 0.1% formic acid) and B (90% acetonitrile, 5% DMSO, 0.1% formic acid), ranging from 5% to 37% B in 240 minutes with a flow of 0.4  $\mu$ l/minute. The Q Exactive mass spectrometer (Thermo Scientific, Wilmington, DE, http://www.thermoscientific. com) was operated in a data-dependent manner, selecting the top five precursors for fragmentation by higher energy C-trap dissociation (HCD). The survey scan was performed at 70,000 resolutions from 300 to 1,700 mass-to-charge ratio (m/z), with a maximum injection time of 100 milliseconds and target of  $1 \times 10^{6}$  ions. For generation of HCD fragmentation spectra, a maximum ion injection time of 500 milliseconds and automatic gain control of  $1 \times 10^5$  were used before fragmentation at 30% normalized collision energy, 35,000 resolutions. Precursors were isolated with a width of 2 m/z and put on the exclusion list for 70 seconds. Single and unassigned charge states were rejected from precursor selection.

#### Peptide, Protein Identification, and Data Analyses

Proteome Discoverer, version 1.4 (Thermo Scientific), with SEQUEST-Percolator was used for protein identification. The precursor mass tolerance was set to 10 ppm and for fragments to 0.02 Da. Oxidized methionine was set as dynamic modification and carbamidomethylation as static modification. Spectra were matched to a combined database of UniProt human (140203) combined with the 250 most abundant proteins from 4-hour nLC-MS/MS analysis of FBS (Bos Taurus, UniProt 140203). The results were filtered to 1% false discovery rate (FDR). Identifications in Bos taurus were considered to originate from FBS and were removed. Gene ontology term enrichment analysis was done using the protein analysis through evolutionary relationships (PANTHER) classification system [32].

#### **Statistical Analysis**

The Wilcoxon paired test was used to compare the differences between two different groups. One-way analysis of variance was used to statistically evaluate the difference of sample means among multiple groups. The significance level was set at  $p \leq .05$ .

#### RESULTS

#### **Clinical Presentation**

#### Patient 1

A 58-year-old man with a history of hypertension was hospitalized 8 days after the onset of high fever, cough, and dyspnea. The CXR demonstrated diffuse bilateral infiltrates and reverse transcription (RT)-PCR of BAL fluid was positive for influenza A H1N1. Oseltamivir (Tamiflu; Roche, Basel, Switzerland, http://www.roche.com) was initiated; however, the patient's condition deteriorated over the next 2 days, with increased bilateral opacities on the CXR and progressive respiratory failure requiring mechanical ventilation, followed initially by venovenous (VV), and later by venoarterial, ECMO (CARDIOHELP System; Maquet, Rastatt, Germany, http://www.maquet.com) for refractory hypoxemia. He was diagnosed with severe ARDS according to the Berlin criteria (Table 1) [33]. Acute kidney failure developed, requiring continuous renal replacement therapy (CRRT).

Over the next 4 days, the patient's condition worsened, with persistent hypoxemia, progressive bilateral infiltrates, and liver failure. At this juncture, after extensive discussions with the family, hospital administration, and ethics board, HLA-mismatched allogeneic bone marrow-derived MSCs, obtained from a healthy volunteer, were systemically infused through a central venous catheter positioned in the right atrium of the heart on a compassionate use basis. The total dose ( $2 \times 10^6$  cells per kilogram of recipient weight) was divided into aliquots of 5 ml of cell suspension given at 6 time points over 20 minutes. To avoid infusion of MSCs into the ECMO circuit and to maximize the delivery of cells to the pulmonary circulation, the ECMO outflow cannula was clamped during each infusion and opened after six heartbeats. ECMO support was maintained for 3 minutes between each MSC infusion. At the time of infusion, the patient had no detectable influenza A H1N1 particles, as assessed by RT-PCR, and had no other detectable active infection.

#### Patient 2

A 40-year-old man with no previous medical history was admitted to the hospital because of malaise, weight loss, night sweats, gingival hyperplasia, and fever. Acute myeloid leukemia was diagnosed in the bone marrow aspirate, and induction chemotherapy was

**Table 1.** ARDS diagnostics according to the Berlin criteria at admission to the ICU

ARDS criteria according		
to the Berlin criteria	Patient 1	Patient 2
Lung injury of acute onset, within 1 week of an apparent clinical insult and with progression of respiratory symptoms	Yes	Yes
Bilateral opacities on chest imaging not explained by other pulmonary pathology	Yes	Yes
Respiratory failure not explained by heart failure or volume overload	Yes	Yes
PaO <sub>2</sub> /FiO <sub>2</sub> ratio <sup>a</sup>	55	67.5
Kidney failure	Yes	No
Liver failure	Yes	No
Body weight, kg	100	110

<sup>a</sup>Mild ARDS: A ratio of 201–300 mmHg indicates mild ARDS; 101–200 mmHg, moderate ARDS; and  $\leq$  100 mmHg, severe ARDS, with a positive end-expiratory pressure of  $\geq$ 5 cmH<sub>2</sub>O.

Abbreviations: ARDS, acute respiratory distress syndrome; FiO<sub>2</sub>, fraction of inspired oxygen; ICU, intensive care unit;  $PaO_2$ , partial pressure of arterial oxygen.

initiated. Because of caries and dental root abscesses, several teeth were extracted at the initiation of cytotoxic therapy. The following day, the patient became progressively hypoxemic and developed acute respiratory failure requiring intubation and mechanical ventilation. Blood and sputum cultures were negative, and CXR demonstrated bilateral infiltrates. Despite treatment with broad-spectrum antibiotics in the now neutropenic patient, progressive hypercapnia and hypoxia developed over the next 2 days, and VV ECMO was initiated. An echocardiogram demonstrated normal cardiac function. The patient was initially stable with ECMO; however, the clinical course was complicated by severe transfusion-dependent cytotoxic chemotherapy-induced thrombocytopenia. Given the large fluid volume from the transfusions, CRRT was initiated to maintain an appropriate fluid balance and reduce interstitial edema. Fourteen days after initiation of ECMO, the patient's condition continued to deteriorate, with increasing hypoxia and progressive bilateral infiltrates on CXR. Possible contributing factors included progressive severe ARDS according to the Berlin criteria [33] (Table 1), possibly secondary to an initial infection, although the findings from blood and sputum cultures and viral screens had remained negative. Transfusion-related acute lung injury could also have played a role in the patient's deterioration. After extensive discussion with the family, hospital administration, and ethics board, HLAmismatched allogeneic bone marrow-derived MSCs (from the same donor and passage number used for patient 1;  $2 \times 10^{\circ}$  cells per kilogram recipient weight) were administered in the same manner as for the first patient 28 days after the initiation of ECMO.

#### **Clinical Course After Allogeneic MSC Infusion**

No obvious adverse events or changes in clinical status were observed in either patient during the MSC infusions, including no alterations in hemodynamic parameters or in oxygen saturation. Serial CXRs demonstrated progressive decreases in pulmonary infiltrates that were evident 24 hours after infusion in both patients (Figs. 1A, 2A). In accordance with local guidelines for patients receiving ECMO support, the positive end-expiratory pressure (PEEP) and peak inspiratory pressure were set and maintained within a defined range while the patient was receiving ECMO support. Concomitant clinical improvements were observed in the tidal volumes and compliances, thus, implying improvement in respiratory dynamics and recovery of the injured lung without manipulation of the ventilator settings. Oxygenation and pulmonary compliance improved in both patients only a few days after MSC administration in the setting of unchanged ECMO settings and PEEP (supplemental online Fig. 2). In patient 1, pulmonary compliance increased from 6 ml/cmH<sub>2</sub>O before infusion to 20 ml/cmH<sub>2</sub>O at 2 days, increasing to 44 ml/cmH<sub>2</sub>O at day 8 (Fig. 1B). In patient 2, pulmonary compliance increased from 20 ml/cmH<sub>2</sub>O before infusion to 35 ml/cmH<sub>2</sub>O at day 1, increasing to 73 ml/cmH<sub>2</sub>O at day 12 (Fig. 2B). The tidal volumes similarly increased; increasing in patient 1 from 100 to 420 ml over a 2-day period, reaching 720 ml at 5 days (Fig. 1C). In parallel with the improvements in respiratory status, patient 1, who was hemodynamically unstable at the time of infusion, stabilized over the first days and was weaned off intravenous pressor agents.

At 5 days after MSC administration, patient 1 developed nosocomial pneumonia with fever, an increased white blood cell count (Fig. 1D), and new infiltrates on CXR (Fig. 1A). The blood cultures were positive for coagulase-negative staphylococci. Vancomycin was added to the antibiotic regimen at day 6, with clinical resolution of the pneumonia by day 8. In patient 2, the tidal volumes increased from 221 to 617 ml over a 7-day period (Fig. 2C). The improvements in respiratory status persisted, allowing discontinuation of ECMO support and, subsequently, a progressive decrease in the need for mechanical ventilation (Figs. 1A, 2A). Liver function in patient 1 improved during the first week, with normalization of transaminase and bilirubin levels (Fig. 1E). Kidney function also steadily improved in patient 1, allowing cessation of CRRT. Patient 2 did not develop renal or hepatic failure and the bone marrow function continued to be reconstituted after MSC infusion, with resolution of leukopenia (Fig. 2D) and thrombocytopenia (Fig. 2E). Patient 1 was discharged from the ICU after 46 days and left the hospital at day 69 (Fig. 1A). Two months later, the patient had returned to work and had recovered both physically and mentally, scoring 38.4 on the physical component and 67.4 on the mental component of the SF-36 quality of life assessment (norm-based scale, with an average of 50 for the physical and mental components both). Patient 2 was weaned off ECMO after 8 days, was removed from mechanical ventilation after 12 days, and returned to the general ward 15 days after the MSC infusion (Fig. 2A). The patient then underwent consolidation chemotherapy for acute myeloid leukemia and subsequently could not perform the SF-36 test.

# Changes in Pulmonary and Systemic Inflammatory Markers After MSC Infusion

Detailed assessments of lung and systemic inflammation before the MSC infusions demonstrated elevations in some, but not all inflammatory, markers in both patients, suggestive of some degree of ongoing inflammation. In patient 1, MSC administration was subsequently followed by a decrease in a range of inflammatory markers in both BAL fluid and serum. In patient 1, the BAL fluid albumin levels, a marker of alveolar-capillary barrier integrity and permeability, decreased from 2 g/l at the time of MSC infusion to undetectable levels the next day. Interleukin-6 (IL-6) in BAL fluid, together with IL-8 and IFN- $\gamma$  levels in plasma,



**Figure 1.** Clinical characteristics of patient 1 before and after MSC administration. (**A**): Disease progression and concomitant chest x-rays (CXRs) from the onset of influenza A H1N1 until hospital discharge. As demonstrated, a progression of bilateral opacifications of the lungs occurred after the onset of the influenza infection, and the patient required both mechanical ventilation and ECMO support. By 24 hours after MSC infusion, a decrease had occurred in the pulmonary infiltrates on CXR, followed by progressive improvement until the time of weaning of ECMO and extubation 4 weeks after MSC administration. In parallel with the improvement on the CXRs, both the pulmonary compliance (**B**) and tidal volumes (**C**) improved to normal levels within the first 2 days. However, 5 days after MSC administration, the patient eveloped nosocomial pneumonia with an increased WBC count (**D**) and new infiltrates on the CXR (**A**), which resolved within 3 days. The patient's liver function also improved during the first week, with normalization of serum ALT, AST, and bilirubin levels (**E**). Abbreviations: ALT, alanine transaminase; AST, aspartate transaminase; ECMO, extracorporeal membrane oxygenation; MSC, mesenchymal stromal cell; WBC, white blood cell.

similarly decreased during the first days after MSC administration (Fig. 3A, 3B). The plasma IL-6 and IFN- $\gamma$  levels subsequently increased transiently in conjunction with the hospital-acquired pneumonia, and IL-8 demonstrated a similar response pattern to that in the BAL fluid (Fig. 3A, 3B). Plasma and BAL fluid granulocyte macrophage colony-stimulating factor levels increased in response to both MSC administration and pneumonia (Fig. 3A, 3B). The proinflammatory chemokine inducible protein 10 (IP10) responded similarly to IL-8 in the BAL fluid with an initial rapid decrease after MSC infusion, followed by a small increase as a response to the pneumonia and a decline again at day 7. No significant change in the plasma IL-8 levels was observed (Fig. 3A).

In patient 2, BAL was only performed twice because of concern of provoking significant bleeding secondary to severe thrombocytopenia. Changes over time in pro- or anti-inflammatory cytokines, such as were seen in patient 1, could not, therefore, be evaluated in the BAL fluid of patient 2 (Fig. 3A). Nonetheless, no significant increase in proinflammatory cytokines was observed in either BAL fluid or plasma after MSC administration (Fig. 3A, 3B).



**Figure 2.** Clinical characteristics of patient 2 before and after MSC administration. **(A)**: Progression of acute respiratory distress syndrome both on chest x-rays (CXRs) and serial computed tomographic scans. As demonstrated, a progressive decrease occurred in pulmonary infiltrates starting the first day after MSC infusion. By day 7, the CXR findings had improved back to normal, and the patient was weaned off ECMO the next day and extubated on day 12. Pulmonary compliance **(B)** and tidal volumes **(C)** improved to normal levels during the first week. Bone marrow function continued to be reconstituted with resolution of leukopenia **(D)** and thrombocytopenia **(E)**. Abbreviations: ECMO, extracorporeal membrane oxygenation; MSC, mesenchymal stromal cell; WBC, white blood cell.

# Improvement in Markers of Lung Epithelial Cell Recovery

BAL fluid levels of total K18, indicative of total epithelial cell death, and ccK18, indicative of epithelial apoptosis [34], were elevated in both patients. ccK18 was elevated in patient 1 before MSC transfer and decreased, starting a few hours after infusion (Fig. 3C). Despite a transient elevation of ccK18 during the episode of nosocomial pneumonia in patient 1, a progressive decrease in both ccK18 and K18 occurred, suggesting no clear shift between epithelial apoptosis and necrosis. In both patients' BAL fluids, surfactant protein B levels increased during the first 4 days after MSC administration, further suggesting recovery of alveolar epithelial integrity and function (Fig. 3D).

# In Vivo MicroRNA Profiling of Blood Extracellular Vesicles

EVs, small membrane vesicles with a lipid bilayer containing a variety of substances, including inhibitory miRNAs, are secreted by nearly all cells and have been increasingly recognized for roles in cell-cell communication, particularly in the transfer of miRNAs [35]. In both patients, circulating levels of several proinflammatory miRNAs in circulating EVs (miR-409-3P, 886-5P, 324-3P, 222, 125A-5P, 339-3P, 155) that were elevated before MSC administration, demonstrated a significant decline within 24 hours after infusion (Fig. 3E). A transient increase in these miRNA levels was observed by day 7 in patient 1 and had returned to normal by day 28 (Fig. 3E). This coincided with the occurrence of nosocomial pneumonia. Although the proinflammatory miRNA trend was similar between the two patients, the absolute value of each miRNA was much lower in patient 2 than in patient 1. Other miRNAs related to inflammation (miR-638, 328, 26B, 29B, 30D, 27B) demonstrated a dichotomous pattern with an initial increase within the first hours to first day, followed by a decline and another peak by day 27 in patient 1 and day 14 in patient 2, returning to normal by day 28 in both patients (Fig. 3E).

# MSC-Mediated Promotion of Regulatory Leukocytes In Vitro

One of the key features of ARDS is the accumulation of neutrophils (PMNs) in the lung microvasculature, interstitium, and alveolar space. In both patients, MSC administration was temporally associated with changes in circulating neutrophil counts. Although the BAL fluid total and differential cell counts could not be assessed in these two patients, we nonetheless believed it of value to assess the MSC effects on neutrophil behavior in vitro, including both survival and activation state. To exclude the indirect effects of other leukocyte populations on both MSC and PMN function, highly purified circulating third-party human PMN preparations were used, obtained as previously described [30].

To mimic in part the local inflammatory lung milieu, ex vivoexpanded MSCs were either unprimed (MSCs) or pretreated (primed MSCs, pMSCs) with 10 ng/ml IFN- $\gamma$  and 15 ng/ml



**Figure 3.** Inflammatory marker response, lung epithelial recovery, and miRNA response to MSC administration. The cytokine response in BAL **(A)** and plasma **(B)** after MSC infusion. The orange line represents the mean values measured in healthy controls (*n* = 102–106). The area between the 5th and 95th percentile is shown in gray. Red lines, patient 1; black lines, patient 2. **(C)**: Levels of caspase-cleaved cytokeratin-18, indicative of epithelial apoptosis, and uncleaved cytokeratin-18, indicative of total epithelial death in BAL fluid, decreased in both patients within a few hours after MSC administration. **(D)**: In parallel, surfactant protein B increased during the first 3 days after MSC infusion, suggesting recovery of alveolar epithelial function. Several miRNAs related to inflammation demonstrated a significant decline within the first 24 hours after MSC administration (**[E]**, top row). Proinflammatory: purple, miR-409-3P; green, miR-886-5P; orange, miR-324-3P; red, miR-222; blue, miR-125A-5P; violet, miR-339-3P; brown, miR-155. A transient increase in these miRNA levels was observed by day 7 and had returned to normal by day 28. Other inflammation-related miRNAs demonstrated a dichotomous pattern with an initial increase within the first hours to the first day, followed by a decline and another peak by day 27 in patient 1 (red) and day 14 in patient 2 (black), returning to normal by day 28 in both patients (**[E]**, bottom row). Abbreviations: BAL, interferon-γ; IL-6, interleukin-6; IP10, inducible protein 10; K18, cytokeratin-18; m, month; miR, microRNA; MSC, mesenchymal stromal cell.

TNF- $\alpha$  for 48 hours and subsequently used in cocultures with PMNs with or without activation by endotoxin (100 ng/ml LPS) [36]. Compared with unprimed MSCs, primed MSCs upregulated cell surface expression of CD54 (ICAM-1), CD106 (VCAM-1), and HLA-ABC and HLA-DR (supplemental online Fig. 3) and the expression of IDO, a potent mediator of many MSC immune regulatory functions [37] (Fig. 4A). Subsequently, the viability and expression levels of surface markers by control or LPS-stimulated PMNs were investigated after 40 hours of direct coculture with either MSCs or pMSCs. As previously reported, CD16 (Fc $\gamma R$ -III) expression can be reliably used as a surrogate marker of PMN viability [30]. As shown in Figure 4B, in the absence of LPS, PMN survival was enhanced only in presence of pMSCs. When LPS was added to the coculture, both resting and pMSCs protected PMNs from apoptosis. Accordingly, the percentage of CD16-positive PMNs matched the percentage of viable PMNs in all culture conditions (Fig. 4C). In parallel, the expression of CD11b and CD54 is typically associated with PMN activation status [38]. As expected, the percentage of CD11b-positive PMNs was higher in the presence of MSCs and further enhanced by LPS treatment (Fig. 4D), although the CD11b relative mean fluorescence intensity (rMFI) did not change significantly, suggesting that more PMNs are becoming activated after MSC or pMSC exposure. In contrast, CD54 rMFI was upregulated by LPS treatment, and this effect was enhanced by coculture with MSCs (Fig. 4E). Overall, the higher PMN survival and activation triggered by MSCs suggests that MSCs might influence the PMNdependent innate response through functional modifications rather than proapoptotic effects.

Emerging evidence has suggested that a heterogeneous subpopulation of PMNs and monocytes, termed myeloid-derived suppressor cells (MDSCs), exert immune regulatory effects and are often increased in inflammatory or inflammatory-like conditions such as cancer, where they are postulated to act as an intrinsic negative feedback mechanism [39–41]. Whether they play a role in ARDS has not yet been ascertained; however, preclinical data have pointed toward beneficial effects [42]. Coculturing PMNs with MSCs led to a marked increase in mature CD11b<sup>bright</sup>/CD16<sup>bright</sup>/CD62L<sup>dim</sup> (N2-type) cells with hypersegmented nuclei, consistent with granulocytic MDSCs [41] (Fig. 4F, 4G). Similarly, coculturing MSCs with healthy control human PBMCs at different ratios promoted CD14<sup>+</sup>HLA-DR<sup>low</sup> monocytes, resembling monocytic MDSCs (Fig. 4H; supplemental online Fig. 3A), as shown previously [43, 44].

The proportion of CD4<sup>+</sup>CD25<sup>high</sup>CD127<sup>low</sup> regulatory T cells (T<sub>Regs</sub>), a key immune regulatory cell population [45], was also expanded in coculture experiments with PBMCs (Fig. 41; supplemental online Fig. 3B). In parallel, increased levels of circulating CD4<sup>+</sup>CD25<sup>high</sup>CD127<sup>low</sup>  $T_{Regs}$  were observed in both patients' peripheral blood for up to 20 days after MSC administration ( $T_{Regs}$  among CD4<sup>+</sup> T cells in healthy controls [n = 11], 3.84%  $\pm$  1.60%;  $T_{Reg}$  range in both patients, 3.34%–17.8%; Fig. 4J), similar to previous findings in MSC-treated patients with graft-vs.-host disease [46]. This finding could have resulted from an elevated thymic output, as indicated by the increased proportion of CD31<sup>+</sup> recent thymic emigrants among CD45RA<sup>+</sup> naïve T<sub>Regs</sub> in both patients (Fig. 4K) and/or the MSC-stimulated conversion of conventional T cells in the periphery (Fig. 4I; supplemental online Fig. 3B). The findings from animal models have suggested that recovery from acute lung injury involves upregulation of  $T_{Regs}$  [47].

# Proteome Characterization of Donor MSCs and Derived Extracellular Vesicles

The ability of the MSCs to ameliorate the diffuse inflammatory processes in ARDS likely results, at least in part, from the release of soluble mediators, including a variety of growth factors, anti-inflammatory cytokines and chemokines, and miRNAs, either directly secreted or delivered by EVs [48-50]. To determine the proteomic profile of the infused MSCs and corresponding EVs, cells were grown in serum-free media for 48 hours. Conditioned medium was collected, and the EVs were isolated by ultracentrifugation according to established protocols with minor changes [51]. Nanoparticle tracking analysis showed a typical EV distribution pattern with particles of a mode size of around 60 nm (supplemental online Fig. 4). Total protein lysates were generated on harvested cells and EVs followed by tryptic digestion into peptides. Digestion was analyzed using nano-reverse-phase chromatography gradient before mass spectrometry (nanoLC-MS/MS) analyses. The nanoLC-MS/MS analysis identified 3,040 proteins in the MSCs and 937 proteins in their EVs (1% FDR; supplemental online Table 1), with a total overlap of 754 proteins (Fig. 5A). The median number of peptide spectrum matches per protein was approximately 10 for cells and 8 for EVs (supplemental online Fig. 5A). Several proteins were only detected in EVs and not in the corresponding cells. This might reflect gradient separation limitations of the cell fraction, resulting in identification of the most abundant part of the proteome by nanoLC-MS/MS, with the potentially less abundant proteins that are packed into EVs missed in cell analyses. A relatively poor correlation of protein abundance was observed between EVs and cells. This indicates that EVs are not merely formed by lysed cells but rather result from selective protein packing into vesicles, corroborating previous reports on EVs derived from other cell types [48] (Fig. 5B; supplemental online Fig. 5B).

To delineate the potential functions of the identified proteins, GO enrichment analysis was performed using the on-line PAN-THER classification system [52]. An overview of the percentage of GO biological processes in donor cells and EVs is illustrated in Figure 5C. Examples of significantly enriched GO categories in EVs and/or cells (p < .001) include metabolic processes such as the tricarboxylic acid cycle and glycolysis pathway and biological processes and pathways such as adhesion and integrin signaling (Fig. 5D). Additional GO enrichment data on biological processes, molecular function, cellular component, protein class, and pathways can be found in the supplemental online data (supplemental online Fig. 6A–6E).

#### DISCUSSION

We have described the administration of non-HLA-matched allogeneic bone marrow-derived MSCs to 2 consecutive patients with severe refractory ARDS of different underlying etiologies under compassionate use. In both instances, clinical improvement was observed, with both patients achieving discharge from the ICU and hospital. Although causality can only be speculated in this small case series, the clinical outcomes paralleled those observed after MSC administration in a number of preclinical models of acute lung injury [7, 12–15, 19–21]. After MSC infusion, both patients demonstrated improved lung function, including tidal volumes and compliance, and CXRs with



**Figure 4.** MSC-mediated immunomodulation of leukocytes. **(A):** In selected experiments, MSCs were exposed to IFN-*γ* and TNF-*α* for 48 hours (pMSCs). After priming, IDO expression was substantially upregulated as assessed by quantitative polymerase chain reaction compared with untreated MSCs. **(B):** Unstimulated MSCs and pMSCs affected viability of control or LPS-stimulated PMNs. **(C):** CD16 ( $Fc\gamma R$ -III) expression was used as a marker of PMN viability and matched with the percentage of viable PMNs in all culture conditions. Similarly, CD11b and CD54 expressions were associated with PMN activation status. The percentage of CD11b-positive PMNs was higher in the presence of MSCs and further enhanced by LPS treatment **(D)**, and CD54 relative MFI was upregulated by LPS treatment, and this effect was enhanced by coculture with MSCs **(E)**. The capacity of MSCs to promote the generation of myeloid-derived suppressor cells (MDSCs) was tested within our coculture model with PMNs. Coculture with MSCs led to a marked increase of mature CD11<sup>bright</sup>/CD16<sup>bright</sup>/CD62L<sup>dim</sup> (N2-type) PMNs resembling granulocytic MDSCs **(F)**. **(G)**: MSC and PMN cocultures were stained with May-Grünwald-Giemsa dye to observe cell morphology after reciprocal interaction for 24 hours. According to the marked increase of mature CD11<sup>bright</sup>/CD16<sup>bright</sup>/CD62L<sup>dim</sup> (N2-type) PMNs shown at this time by flow cytometry, PMNs displayed a hypersegmented nuclear morphology after coculture with both resting (PMN MSCs at 24 hours) and pMSCs (PMN pMSCs at 24 hours). Similarly, coculturing MSCs with healthy control T cells **(I)**. **(J)**: Increased levels of circulating CD4<sup>+</sup>CD25<sup>brigh</sup>CD127<sup>low</sup> T<sub>Regs</sub>, when cocultured in different ratios with purified healthy control T cells **(I)**. **(J)**: Increased levels of circulating CD4<sup>+</sup>CD25<sup>brigh</sup>CD127<sup>low</sup> T<sub>Regs</sub>, were also observed up to 20 days after MSC administration (red, patient 1; black, patient 2). **(K)**: This could in part have resulted from increased thymic output, indicated by an increased propor

concomitant overall progressive clinical improvement. These are provocative clinical observations in patients with severe refractory ARDS, providing additional impetus for larger formal clinical investigations such as the recently published NIH-sponsored phase I trial that demonstrated the safety of systemic administration of allogeneic bone marrow-derived MSCs in 9 patients with moderate to severe ARDS and that has since been expanded to a phase II clinical trial [53, 54].



**Figure 5.** Proteome characterization of donor MSCs and donor-derived EVs. **(A)**: Overlap in protein identification between MSC donor cells and EVs. Proteins were considered identified if they had a quantifiable protein area. **(B)**: Correlation of protein abundance between MSC donor cells and EVs. Protein abundance was estimated by label-free quantification using protein signal intensity (area under peak of identified peptides/ protein). **(C)**: Pie chart of Gene Ontology (GO) major biological process categories in MSC donor EVs and donor cells. Percentages were based on GO assignment. **(D)**: Selected GO enrichment terms for donor cells (WCs) and EVs. All enriched GO terms can be found in supplemental online Figure 6A–6E. Abbreviations: EVs, extracellular vesicles; MSCs, mesenchymal stromal cells; WCs, whole cells.

At the time of MSC infusion, the first patient was critically ill with hemodynamic and respiratory instability and multiorgan failure. Within several hours after MSC administration, with no other supportive measures changed, including no changes in the ECMO or PEEP settings as per the standard clinical protocol, a number of clinical and laboratory parameters improved. This trend continued, with

subsequent discontinuation of supportive therapies, resolution of multiorgan failure, discharge from the ICU and, eventually, from the hospital, and a return to excellent functional status. A transient setback with nosocomial (ventilator-associated) pneumonia occurred several days after MSC administration but responded to appropriate antimicrobial treatment. Whether MSC administration increases the risk of infectious complications in ARDS must be assessed in future investigations; however, no such increase has been reported in clinical trials of MSCs given to severely immunoincompetent recipients [22, 27]. In contrast, the findings from a number of recent preclinical studies suggest that MSCs have antimicrobial effects [55–57]. The second patient mainly had severe respiratory failure but nonetheless demonstrated an improvement in clinical status after MSC administration, with subsequent discontinuation of supportive therapies and discharge from the ICU. Similar trends in inflammatory indices were observed, although the chemotherapy-induced bone marrow suppression likely altered the inflammatory pathways, obscuring interpretations of possible in vivo MSC effects.

The range of improvements in the in vivo inflammatory indices was broad, with decreases in multiple markers of inflammation, including markers of epithelial apoptosis, alveolar-capillary fluid leakage, and proinflammatory cytokines, miRNAs, and chemokines in plasma and BAL fluid. These results demonstrated that ongoing inflammation was still present long after the initial inciting events that had provoked the development of ARDS. Each of these inflammatory outcomes reflects altered inflammatory and immune pathways that play a role in the pathogenesis or perpetuation of ARDS. Furthermore, as a sign of recovery of alveolar epithelial function, the surfactant protein B levels increased in both patients' BAL fluid after MSC administration. This was more pronounced in patient 1 and could potentially be explained by the difference in clinical status at the time of MSC administration. The validity and relevance of the changes in inflammatory indices are suggested by the subsequent development of nosocomial pneumonia in patient 1, with correlating increases in the inflammatory markers that had decreased after MSC administration. Nonetheless, because this was a descriptive hypothesis-generating study, it would be difficult to know what is relevant with respect to measures of inflammatory mediators. Only further clinical and preclinical studies investigating the roles of specific mediators potentially involved in the pathogenesis and/or resolution of ARDS can answer this question.

The dose used in these patients was based on previous use of MSCs in GVHD patients [27]. Until the recently published phase I trial, a paucity of information was available concerning the appropriate dosing in ARDS patients; thus, the dosing used in these 3 patients was an empiric application. The study of systemic MSC administration in patients with chronic obstructive pulmonary disease also used an empiric approach for multiple dosing [24]. However, those were stable outpatients, not critically and acutely ill patients. Moreover, the preclinical studies investigating use of MSCs in models of acute lung injury have virtually all shown beneficial effects after a single dose or administration. As such, multiple dosing of the administration of MSCs was not considered.

The lack of understanding of MSCs' actions in vivo [26] and the lack of corresponding methods for evaluating their functional quality before application have limited the clinical application of MSCs. Previous investigations of patients receiving allogeneic MSC treatment have suggested that directly on contact with whole blood, such as by i.v. infusion, the cells elicit activation of complement and coagulation cascades (a reaction termed "instant blood-mediated reaction" [IBMIR]) that results in destruction of most of the infused cells and virtually no cell engraftment [58-60]. However, skewing toward a tolerogenic Th2 T-cell repertoire occurs and is most evident in patients as late as 20-30 days after infusion [46, 61]. The events subsequent to the instant IBMIR are unknown in humans; however, previous clinical studies have indicated that in vitro assays can correlate with ex vivo observations and potentially explain MSCs' clinical efficacy [46]. Accordingly, parallel in vitro studies of the MSCs used in our two patients demonstrated anti-inflammatory actions on relevant inflammatory and immune cells, in particular, monocytes and neutrophils, which play significant roles in ARDS pathogenesis or resolution. Although we did not specifically assess this in vivo in either BAL fluid or serum, promotion of the anti-inflammatory M2 monocyte (macrophage) phenotype is relevant for the treatment of ARDS. The role of  $T_{\mbox{\scriptsize Regs}}$  is less clear in ARDS compared with monocytes/macrophages and PMNs, although some data have suggested a role in experimental models of acute lung injury [62]. Nonetheless, the correlation between the induction of T<sub>Regs</sub> in vitro and increased levels of circulating  $T_{\text{Regs}}$  in vivo is striking. The role of PMNs in ARDS is complex, and it is unclear at present whether the promotion of PMN survival, activation, and/or increase in MDSC subsets will indicate beneficial clinical effects. Nevertheless, the marked increase in mature CD11b<sup>bright</sup>/CD16<sup>bright</sup>/ CD62L<sup>dim</sup> (N2-type) cells with hypersegmented nuclei found by coculturing PMNs with MSCs is in agreement with published data showing that the onset of granulocytic MDSCs is associated with such an immunological phenotype, which might be considered a surrogate pattern of PMN functional polarization.

As such, the in vitro findings are hypothesis generating and need to be extensively explored in future studies of a larger number of patients with ARDS of diverse etiologies. This also raises an overall consideration regarding whether the decrease in any latent inflammation present in severe persistent ARDS would be beneficial or harmful. A growing body of evidence suggests that failure of resolution of inflammation is an important pathogenic mechanism underlying ARDS [63]. It is perhaps in this regard in which the antiinflammatory actions of MSCs would be beneficial. Further preclinical and clinical studies of MSCs are needed to fully comprehend the potential role of the timing of MSC administration in ARDS.

In the in vitro experiments, we mimicked in part an inflammatory milieu by pretreating the ex vivo expanded MSCs with IFN- $\gamma$  and TNF- $\alpha$ . This is an established technique that can be performed to assess MSCs' immune modulatory potency. To this aim, the MSC Committee of the International Society for Cellular Therapy recently published a working proposal to highlight the need for standardized in vitro methods to support the various MSC-based clinical approaches used to treat immunological diseases [64]. The standard immune plasticity assay based on IFN- $\gamma$ , with or without TNF- $\alpha$ , is a good model of in vitro priming to assess the immune regulatory properties of MSCs. The use of serum from patients does not meet these criteria for in vitro studies owing to the high variability of the content of biological active molecules secondary to a large number of factors (e.g., comorbidities, treatments, sample collection methods).

None of the standardized in vitro tests, when used alone, can be considered the perfect surrogate and biomarker of in vivo behavior of MSCs. Nevertheless, the combination of different in vitro standardized tests can be useful to decipher the clinical results. Furthermore, the range of biologic mediators present in the inflammatory lung/ARDS environment that might alter or license MSC biology is not yet well understood. A recent study has demonstrated that exposure of MSCs to serum from ARDS patients altered the expression of a range of potential antiinflammatory mediators and improved efficacy of MSC actions when administered to mice with endotoxin-induced acute lung injury [65]. This suggests that licensing or priming of the MSCs with a standardized set of inflammatory cytokines or perhaps even specific patient serum or BAL fluid before administration might be beneficial in ARDS. This offers new opportunities to investigate and correlate disease-specific in vitro potency measures with clinical actions.

Many of the therapeutic effects of MSCs in ARDS are likely mediated by the release of soluble proteins and EVs containing a range of bioactive macromolecules. Some of these proteins and pathways were identified in the MSCs and EVs that have been suggested as potentially associated with the therapeutic effects in other disease models [66]. This includes pathways associated with platelet-derived growth factor receptor and epidermal growth factor receptor activation, which have been implicated in MSC recruitment [66-68], cell adhesion molecules [69-72], and glycolysis and tricarboxic acid cycle-related proteins and proteins involved in translation and ribosomal components. Although the relevance of these pathways to ARDS is not clear, enrichment of the latter proteins could conceivably complement an energy deficit in the ARDS-injured lung, critical in the restoration of normal surfactant production [73]. Just as with the in vitro MSC effects on the inflammatory and immune cells, these initial findings and potential correlations with the in vivo observations are hypothesis generating and need to be further explored in larger numbers of patients. This baseline characterization of successfully used MSCs and their vesicle components is also important for understanding the molecular determinants of functional cell batches and, in the future, to define biomarkers to select MSCs for clinical use in ARDS.

Our findings contrast with those observed in a recent study of the administration of non-HLA-matched allogeneic adipose-derived MSCs in ARDS patients for whom, although short-term improvement in oxygenation occurred after MSC administration, no improvement in ventilator-free days, ICU-free days, or length of hospital stay was observed [74]. Limited measures of inflammatory outcomes were obtained but demonstrated no significant changes in the circulating levels of IL-6, IL-8, or surfactant protein D (SP-D) after MSC administration. Whether this reflects differences in the adipose vs. bone marrow origin of the MSCs or differences in other aspects of the respective protocols used remains to be determined. To investigate the cytokine response after MSCs infusion in ARDS patients, larger cohorts are needed, as well as an expanded range of inflammatory mediators studied. For example, circulating C-reactive protein was not assessed in the present study. The MSCs used in our study were directly infused after thawing, and the adipose MSCs were thawed and cultured for 24-48 hours in each respective patient's own serum before infusion. The use of freshly thawed vs. cultured MSCs has been shown to result in different effects in the in vitro mixed lymphocyte reactions [75]: however. it is as yet unclear how these different approaches will affect the potential use of MSCs in ARDS.

#### CONCLUSION

Our overall results are congruent with those observed in preclinical models and suggest that MSCs might have clinical efficacy in severe refractory ARDS and, perhaps, ARDS resulting from a wider spectrum of etiologies than those in the two patients in our study. These case studies also highlight the wealth of mechanistic and hypothesis-generating information that can be obtained, even from a few patients, and the critical need to incorporate mechanistic studies and in vitro potency assessments as a part of any clinical study of MSCs in ARDS, as well as in other diseases.

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#### **AUTHOR CONTRIBUTIONS**

O.E.S., K.-H.G., K.L.B.: conception and design, collection and/or assembly of data, data analysis and interpretation, manuscript writing, final approval of manuscript, clinical care of patients; D.M., N.H., M.K., D.J.W.: conception and design, collection and/or assembly of data, data analysis and interpretation, manuscript writing, final approval of manuscript; M.D., L.H.-J., P.S., P.-O.J., H.H., M.M.: collection and/or assembly of data, data analysis and interpretation, manuscript writing, final approval of manuscript, clinical care of patients; G.B., H.J.J., R.J., S.R., M.C., S.E.A., O.P.B.W., J.Z.N., J.S., C.R., T.K., J.L., O.R.F., R.S., O.K.: collection and/or assembly of data, data analysis and interpretation, manuscript writing, final approval of manuscript.

## DISCLOSURE OF POTENTIAL CONFLICTS OF INTEREST

O.E.S., M.C., and K.-H.G. are cofounders and owners of Islet One AB, have uncompensated employment, and have compensated intellectual property rights and ownership interest. S.R. is a minority coowner of Biolamina and has compensated intellectual property rights and ownership interest. J.S. is founder and co-owner of Exosome Diagnostics and has compensated employment, intellectual property rights, and ownership interest. C.R. and T.K. are compensated employees of Exosome Diagnostics. D.J.W. has compensated research funding from Athersys Inc. and United Therapuetics. The other authors indicated no potential conflicts of interest.

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