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Carbon Monoxide Improves Efficacy of Mesenchymal Stromal Cells During Sepsis by Production of Specialized Proresolving Lipid Mediators*

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*See also p. 2296.

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Objectives: Mesenchymal stromal cells are being investigated as a cell-based therapy for a number of disease processes, with promising results in animal models of systemic inflammation and sepsis. Studies are ongoing to determine ways to further improve the therapeutic potential of mesenchymal stromal cells. A gas molecule that improves outcome in experimental sepsis is carbon monoxide. We hypothesized that preconditioning of mesenchymal stromal cells with carbon monoxide ex vivo would promote further therapeutic benefit when cells are administered in vivo after the onset of polymicrobial sepsis in mice.

Design: Animal study and primary cell culture.

Setting: Laboratory investigation.

Subjects: BALB/c mice.

Interventions: Polymicrobial sepsis was induced by cecal ligation and puncture. Mesenchymal stromal cells, mesenchymal stromal cells-conditioned with carbon monoxide, fibroblasts, or fibroblasts-conditioned with carbon monoxide were delivered by tail vein injections to septic mice. The mice were assessed for survival, bacterial clearance, and the inflammatory response during sepsis in each of the groups. Mesenchymal stromal cells were also assessed for their ability to promote bacterial phagocytosis by neutrophils, the production of specialized proresolving lipid mediators, and their importance for mesenchymal stromal cells function using gene silencing.

Measurements and Main Results: Ex vivo preconditioning with carbon monoxide allowed mesenchymal stromal cells to be administered later after the onset of sepsis (6 hr), and yet maintain their therapeutic effect with increased survival. Carbon monoxide pre-conditioned mesenchymal stromal cells were also able to alleviate organ injury, improve bacterial clearance, and promote the resolution of inflammation. Mesenchymal stromal cells exposed to carbon monoxide, with docosahexaenoic acid substrate, produced specialized proresolving lipid mediators, particularly D-series resolvins, which promoted survival. Silencing of lipoxigenase pathways (5-lipoxygenase and 12/15-lipoxygenase), which are important enzymes for specialized proresolving lipid mediator biosynthesis,
Sepsis is a disease process initiated by an underlying severe infection that leads to a systemic inflammatory response (1–4). The desired outcome in sepsis is for the invading microorganism(s) to be cleared, followed by resolution of the inflammatory process (5), allowing homeostasis to be achieved and permanent organ dysfunction avoided. Although in the past the resolution of inflammation was felt to be a passive process, it has been demonstrated that specialized proresolving lipid mediators (SPMs) control this process in an orchestrated manner, stopping neutrophil influx and activating resolution pathways (6). These bioactive mediators are synthesized from omega-3 fatty acids (docosahexaenoic acid [DHA] and eicosapentaenoic acid). Omega-6 fatty acids also contribute to the production of SPMs; however, the arachidonic acid (AA)–derived mediators also include the proinflammatory eicosanoids (6).

Even with recent advances in standard of care, severe sepsis and septic shock remain associated with high rates of morbidity and mortality (7, 8). Investigators have recently begun to explore cell-based therapies for sepsis, including the use of mesenchymal stromal cells (MSCs) (9, 10). MSCs are considered to be a promising platform for cell-based therapy, and in preclinical animal models, the administration of MSCs has shown improved outcome in polymicrobial sepsis induced by cecal ligation and puncture (CLP) in mice (11–15). The benefit of treating with MSCs during CLP, or single organism bacterial peritonitis, appears to be better when given early (1–4 hr) after the onset of sepsis (11–13, 15), whereas later administration (6 hr) may require the addition of antibiotics for improved survival (14). Thus, timing of MSC administration in regard to the onset of sepsis is a very important factor determining therapeutic benefit.

Heme oxygenase (HO)-1 is an important cytoprotective molecule during experimental sepsis in mice (16–19). We previously demonstrated the benefits of MSCs when administered after the onset of polymicrobial sepsis in mice, although these effects were not dependent on endogenous HO-1 (12). A product of HO-1 degradation of heme, carbon monoxide (CO) has also been shown to be beneficial therapeutically in animal models of sepsis (16, 20, 21), and inhaled CO gas accelerates the resolution of inflammation in zymosan-induced peritonitis (22). Furthermore, ex vivo conditioning of cells (23, 24) or even tissues (25) with CO has shown protection from cell and organ injury. Thus, in the present study, we wanted to determine whether CO preconditioning would improve MSC function and therapeutic efficacy in experimental polymicrobial sepsis, and to determine potential mechanisms contributing to this effect.

**MATERIAL AND METHODS**

**Isolation of MSCs and Lung Fibroblasts**

Mouse MSCs were harvested from the compact bone of BALB/c mice (12 mice/harvest) as described (12), and lung fibroblasts were used as control mesenchymal cells. MSCs of passage 3–6 were used for experiments. Human MSCs were obtained from either Texas A&M Health Science Center, Institute for Regenerative Medicine, or Lonza. The cells were characterized using a BD fluorescent-activated cell sorting (FACS) Canto II (Becton Dickinson Biosciences, Bedford, MA) and analyzed using FlowJo software (Becton Dickinson Biosciences). Antibodies used for immunophenotyping of cells are listed in [Supplemental Table 1](#) (Supplemental Digital Content 1, http://links.lww.com/CCM/C8).

**Gene Silencing**

For the 5-lipoxygenase (LOX) and 12/15-LOX silencing experiments, lentiviral transfection was used. For detailed information, see [Supplemental Methods](#) (Supplemental Digital Content 2, http://links.lww.com/CCM/C9).

**CLP**

To induce polymicrobial sepsis, we used an established murine model of CLP (two-thirds cecum ligated, two 21-gauge needle punctures) as described (12, 16).

**Terminal Deoxynucleotidyl Transferase dUTP Nick-End Labeling Staining**

Terminal deoxynucleotidyl transferase dUTP nick-end labeling (TUNEL) was performed in the spleen, and the area of positive staining was assessed as described (27).

**Aspartate Aminotransferase, Alanine Aminotransferase, and Creatinine Assays**

Aspartate aminotransferase (AST), alanine aminotransferase, and creatinine plasma levels were measured by commercially available colorimetric assay kits (BioVision, Milpitas, CA).

**Assessment of Blood/Peritoneal Colony-Forming Units**

Serial dilutions of whole blood were performed, and aliquots were cultured on Luria-Bertani agar plates as described (12). Colony-forming units (CFUs) were counted following overnight incubation at 37°C. In addition, serial dilutions of peritoneal lavages were also performed, and CFUs were counted.
**Neutrophil Isolation**
Isolation of mouse and human neutrophils is described in the Supplemental Methods (Supplemental Digital Content 2, http://links.lww.com/CCM/C9).

**Phagocytosis Assay**
Neutrophil phagocytosis of green fluorescent protein–labeled bacteria, ± MSCs, is described in the Supplemental Methods (Supplemental Digital Content 2, http://links.lww.com/CCM/C9). Bacterial phagocytosis was measured by flow cytometry as described (12), and Supplemental Figure 1 (Supplemental Digital Content 3, http://links.lww.com/CCM/C10; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21) shows our gating strategy and a representative flow cytometry pseudocolor density plot for neutrophil phagocytosis.

**Efferocytosis Assay**
The measurement of macrophages phagocytizing dead neutrophils from peritoneal fluid after CLP injury, ± MSCs, is described in the Supplemental Methods (Supplemental Digital Content 2, http://links.lww.com/CCM/C9). Supplemental Figure 2 (Supplemental Digital Content 4, http://links.lww.com/CCM/C11; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21) demonstrates representative flow cytometry pseudocolor density plots of efferocytosis assays in permeabilized (A) and nonpermeabilized (B) cells.

**Lipid Mediator Profiling Using Liquid Chromatography-Tandem Mass Spectrometry-Based Metabololipidomics**
Measurement of lipid mediator profiles was performed as described (28). For detailed information, see the Supplemental Methods (Supplemental Digital Content 2, http://links.lww.com/CCM/C9).

**Quantitative Real-Time Polymerase Chain Reaction, Western Blotting, and 3-(4,5-Dimethylthiazol-2-yl)2,5-Diphenyl-Tetrazolium Bromide Viability Assay**
Quantitative real-time polymerase chain reaction (qRT-PCR) was performed as described (29). Primer sequences are shown in Supplemental Table 2 (Supplemental Digital Content 5, http://links.lww.com/CCM/C12). For further details regarding qRT-PCR, Western blotting, or 3-(4,5-dimethylthiazol-2-yl)2,5-diphenyl-tetrazolium bromide viability assays, see the Supplemental Methods (Supplemental Digital Content 2, http://links.lww.com/CCM/C9).

**Statistical Analysis**
Data are expressed as mean ± SEM. Comparisons of mortality were made by analyzing Kaplan-Meier survival curves, and then the log-rank test was performed to assess the differences in survival. For comparisons between two groups, we used Student unpaired t test. One-way analysis of variance was used for analysis of more than two groups. When data were not normally distributed, nonparametric analyses were performed using Mann-Whitney U or Kruskal-Wallis testing, respectively. The numbers of samples per group (n), or the numbers of experiments, are specified in the figure legends. Statistical significance is accepted at p value less than 0.05.

**Study Approval**
The care and use of animals for all experiments were approved by the Harvard Medical Area Standing Committee on Animals, Harvard Medical School. Experiments involving human blood donation by consented volunteers, for isolation of neutrophils, were approved by the Partners Healthcare Institutional Review Board.

**RESULTS**

**Preconditioning of MSCs With CO Sustains the Efficacy of the Cells for Survival Benefit in CLP-Induced Sepsis**
MSCs phenotypically adhered to plastic and expressed markers of mesenchymal origin (CD105, CD90.2, CD73, CD140b, CD29, CD44) and Sca1, but not markers of hematopoietic origin (such as CD45, CD11b, and bone marrow lineage markers) or major histocompatibility complex II (Supplemental Fig. 3A, Supplemental Digital Content 6, http://links.lww.com/CCM/C13; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21). To understand the effect of CO preconditioning on MSCs, we administered the cells in vivo to mice after the onset of CLP-induced sepsis. MSCs or control fibroblasts—exposed to ambient air or CO at 250 ppm × 4 hours—were initially IV administered 2 hours (5.0×10^5 cells), 24 hours (2.5×10^5 cells), and 48 hours (2.5×10^5 cells) after CLP surgery. The administration of MSCs increased survival (65%) compared with fibroblasts (22%) after the onset of sepsis (Fig. 1A). Preconditioning of MSCs with CO tended to further increase survival (88%) compared with nonconditioned MSCs, although this was not a statistically significant increase. However, mice receiving MSCs preconditioned with CO had a marked increase in survival compared with mice receiving fibroblasts or fibroblast preconditioned with CO (37%). To determine whether the effects of CO were dependent on the expression of HO-1, we preconditioned either wild-type or HO-1–deficient MSCs and administered the cells 2, 24, and 48 hours after CLP surgery. Supplemental Figure 4 (Supplemental Digital Content 7, http://links.lww.com/CCM/C14; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21) demonstrates that the mice had a nearly identical survival after the onset of sepsis, when receiving either wild-type or HO-1–deficient MSCs preconditioned with CO. Finally, we administered MSCs later after the onset of sepsis (6 hr), and in only one dose (5.0×10^5), to further investigate the effects of CO preconditioning on MSCs (Fig. 1B). In this setting, mice receiving MSCs preconditioned with CO had a significantly improved survival (76%) compared with mice treated with...
Preconditioning of MSCs With CO Promotes the Resolution of Inflammation During Polymicrobial Sepsis

Next we investigated whether MSCs would increase the clearance of apoptotic neutrophils by macrophages in the peritoneum (effe-rocytosis), which is an important process during the resolution of inflammation (33). Treatment with MSCs 6 hours after the onset of CLP-induced sepsis was able to increase peritoneal effe-rocytosis compared with fibroblasts; however, MSCs preconditioned with CO had a more marked increase in effe-rocytosis (Fig. 4A).

Coinciding with the process of effe-rocytosis, SPMs are produced to help orchestrate the resolution of inflammation (6, 33). Because culture conditions alone, in contrast to the in...
vein injection 6 hr after CLP. At 24 hr, mice were anesthetized, and blood and spleens were harvested.

**Figure 2.** Administration of mesenchymal stromal cells (MSCs) preconditioned with carbon monoxide (CO) decreases tissue injury in cecal ligation and puncture (CLP)-induced sepsis. Mice were subjected to sham or CLP surgery (CLP+Fibro, CLP+MSCs, CLP+MSCs/CO). A total of 5 × 10⁵ cells were administered by tail vein injection 6 hr after CLP. At 24 hr, mice were anesthetized, and blood and spleens were harvested. A, Spleen injury was evaluated by terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) staining (brown staining, right), and quantified (left). The bar in the sham represents a length of 50 μm. Data are presented as box plots, which show median values and interquartile ranges. Analysis was done by Kruskal-Wallis testing (also used in **B** and **C**). Sham, white bar, n = 3; CLP+Fibro, light gray bar, n = 7; CLP+MSCs, dark gray bar, n = 8; CLP+MSCs/CO, striped bar, n = 8, p = 0.0001, with significant comparisons * versus sham and † versus CLP+Fibro. B, Plasma aspartate aminotransferase (AST) and alanine aminotransferase (ALT) data are shown as box plots. Sham, white bar, n = 3; CLP+Fibro, light gray bar, n = 6; CLP+MSCs, dark gray bar, n = 8; CLP+MSCs/CO, striped bar, n = 9, p = 0.002 and p = 0.0008, respectively, with significant comparisons * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs. C, Plasma creatinine levels are shown as box plots. Sham, n = 3; CLP+Fibro, light gray bar, n = 6; CLP+MSCs, dark gray bar, n = 6; CLP+MSCs/CO, striped bar, n = 5, p = 0.018, and significant comparisons of specific groups * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs.

In the vivo setting, do not provide the necessary substrates for production of SPMs, we incubated CO preconditioned MSCs with either DHA or AA to help elucidate the mediators contributing to the beneficial MSC response. Figure 4B demonstrates that administration of CO preconditioned MSCs to mice 6 hours after the onset of CLP-induced sepsis led to a significantly improved survival when the cells were exposed to DHA (92% survival), compared with cells exposed to AA (58% survival). The cells exposed to DHA predominantly expressed SPMs, whereas the cells exposed to AA expressed eicosanoids (Fig. 4C).

To determine SPMs produced by MSCs preconditioned with CO, we performed liquid chromatography-tandem mass spectrometry. **Supplemental Figure 6** (Supplemental Digital Content 9, http://links.lww.com/CCM/C16; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21) shows representative multiple reaction monitoring traces (A) for the measured lipid mediators and accompanying tandem mass spectrometry spectra (B) used for identification. As seen in **Figure. 4D**, SPMs were greater in DHA exposed cells, compared with AA exposed cells, particularly D-series resolvins (Rv). An entire profile of SPMs is shown in **Supplemental Table 3** (Supplemental Digital Content 10, http://links.lww.com/CCM/C17), and a complete list of abbreviations is provided in **Supplemental Table 4** (Supplemental Digital Content 11, http://links.lww.com/CCM/C18). We next used representative D-series Rv (RvD1, RvD2, and aspirin triggered [AT]-RvD3) to assess their role in the interaction of MSCs with neutrophils to promote bacterial phagocytosis. To confirm this response was applicable in human cells, we used human MSCs and neutrophils. MSCs increased the phagocytosis of E. coli by neutrophils, and MSCs preconditioned with CO produced an enhanced phagocytic response (**Fig. 4E**). Incubation of neutrophils with RvD1, RvD2, and AT-RvD3 alone showed evidence of neutrophil phagocytosis of E. coli. However, when MSCs were exposed to either RvD1 or RvD2, and then incubated with neutrophils, phagocytosis was robustly increased compared with neutrophils exposed to RvDs alone or MSCs alone, to a level analogous to neutrophils incubated with MSCs preconditioned with CO (Fig. 4E). Exposure of MSCs to AT-RvD3 did not promote neutrophil phagocytosis to a level significantly greater than neutrophils exposed to MSCs alone.

**Figure 2.** Administration of mesenchymal stromal cells (MSCs) preconditioned with carbon monoxide (CO) decreases tissue injury in cecal ligation and puncture (CLP)-induced sepsis. Mice were subjected to sham or CLP surgery (CLP+Fibro, CLP+MSCs, CLP+MSCs/CO). A total of 5 × 10⁵ cells were administered by tail vein injection 6 hr after CLP. At 24 hr, mice were anesthetized, and blood and spleens were harvested. A, Spleen injury was evaluated by terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) staining (brown staining, right), and quantitated (left). The bar in the sham represents a length of 50 μm. Data are presented as box plots, which show median values and interquartile ranges. Analysis was done by Kruskal-Wallis testing (also used in **B** and **C**). Sham, white bar, n = 3; CLP+Fibro, light gray bar, n = 7; CLP+MSCs, dark gray bar, n = 8; CLP+MSCs/CO, striped bar, n = 8, p = 0.0001, with significant comparisons * versus sham and † versus CLP+Fibro. B, Plasma aspartate aminotransferase (AST) and alanine aminotransferase (ALT) data are shown as box plots. Sham, white bar, n = 3; CLP+Fibro, light gray bar, n = 6; CLP+MSCs, dark gray bar, n = 8; CLP+MSCs/CO, striped bar, n = 9, p = 0.002 and p = 0.0008, respectively, with significant comparisons * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs. C, Plasma creatinine levels are shown as box plots. Sham, n = 3; CLP+Fibro, light gray bar, n = 6; CLP+MSCs, dark gray bar, n = 6; CLP+MSCs/CO, striped bar, n = 5, p = 0.018, and significant comparisons of specific groups * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs.

**5-LOX and 12/15-LOX Are Critical Enzymes for CO-Induced MSC Function and Benefit in Sepsis**

LOXs are the enzymes responsible for the biosynthesis of SPMs, with both 5-LOX and 12/15-LOX pathways contributing to the production of Rv from DHA (34, 35). To further assess the role of SPMs in promoting neutrophil phagocytosis by MSCs, we pretreated MSCs with the LOX inhibitor baicalein, and then incubated ambient air- or CO-exposed MSCs with neutrophils. Baicalein blunted the ability of ambient air-exposed MSCs to promote neutrophil phagocytosis, and the LOX inhibitor abolished the enhanced phagocytic response of neutrophils exposed to CO preconditioned MSCs (Fig. 5A). This lack of a CO response...
A total of $5 \times 10^5$ cells were administered by tail vein injection 6 hr after CLP. At 24 hr, peritoneal fluid (left) and blood samples (right) were collected and analyzed. Data are presented as box plots, which show median values and interquartile ranges. Analysis was done by Kruskal-Wallis testing (also used in B and C). Left, sham, $n = 11$; CLP+Fibro, white box, $n = 6$; CLP+MSCs, light gray box, $n = 9$; CLP+MSCs/CO, dark gray box, $n = 12$. Right, sham, $n = 7$; CLP+Fibro, white box, $n = 6$; CLP+MSCs, light gray box, $n = 10$; CLP+MSCs/CO, dark gray box, $n = 12$. $p < 0.0001$ left and $p = 0.001$ right, with significant comparisons * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs. B, Isolated neutrophils were incubated with green fluorescent protein (GFP)-labeled E. coli or E. faecalis in the presence of MSCs (light gray bars), MSCs+CO (dark gray bars) or in the absence of MSCs (white bars). Data are presented as box plots, $n = 5$–7 per groups from three independent experiments. $p < 0.0008$ for E. coli and $p = 0.0022$ for E. faecalis, with significant comparisons * versus no MSCs, and † versus MSCs. C, Isolated neutrophils were incubated with GFP-labeled E. coli or E. faecalis with addition of conditioned medium (CM) from MSCs (light gray bars), MSCs+CO (dark gray bars), or basal medium (white bars). Data are presented as box plots, $n = 4$–5 per group from three independent experiments. $p < 0.0064$ for E. coli and $p < 0.0047$ for E. faecalis, with significant comparisons * versus basal medium, and † versus CM of MSCs.

was evident in both human (Fig. 5A, left) and mouse (Fig. 5A, right) MSCs. To delineate the response of LOX enzymes to CO, we exposed either MSCs or control fibroblasts to ambient air or CO (250 ppm) and then assessed the level of 5-LOX or 12/15-LOX messenger RNA (mRNA) by qRT-PCR. The basal levels of both 5-LOX and 12/15-LOX were low in fibroblasts and not inducible by CO. Interestingly, the basal level of 5-LOX was significantly greater in MSCs compared with fibroblasts, and CO caused a further increase in 5-LOX expression (Fig. 5B, top). In contrast, the basal level of 12/15-LOX was not significantly different between fibroblasts and MSCs; however, the exposure of CO caused a marked increase in the level of 12/15-LOX (Fig. 5B, bottom). Thus, in MSCs, both the 5-LOX and 12/15-LOX enzymes are induced by exposure to CO.

Finally, to determine whether the 5-LOX and 12/15-LOX enzymes are critical for MSC function, we silenced these enzymes in mouse MSCs (silencing of 5-LOX and 12/15 LOX [shLOX] [as described in the narrative]) or introduced a scrambled shRNA sequence (shSCR). Supplemental Figure 7 (Supplemental Digital Content 12, http://links.lww.com/CCM/C19; legend, Supplemental Digital Content 14, http://links.lww.com/CCM/C21) demonstrates a reduction in 5-LOX and 12/15-LOX mRNA, protein, and a reduction in RvD1 and RvD2 production in shLOX cells. Although shSCR MSCs produced an increase in neutrophil phagocytosis, there was no significant increase in neutrophil phagocytosis by shLOX MSCs (targeting both 5-LOX and 12/15-LOX enzymes). Even more impressive, shLOX MSCs preconditioned with CO had a
Analysis was done by one-way analysis of variance (e1242 Tsoyi et al. CLP+MSCs, CLP+MSCs/CO). A total of 5 × 10^5 cells were administered by tail vein injection 6 hr after CLP. At 24 hr, Carbon monoxide (CO) preconditioning induces production of specialized proresolving lipid mediators

Figure 4.

Figure 4. Carbon monoxide (CO) preconditioning induces production of specialized proresolving lipid mediators (SPMs) in mesenchymal stromal cells (MSCs), which mimic the CO effect by enhancing neutrophil phagocytosis of bacteria. A. Mice were subjected to sham or cecal ligation and puncture (CLP) surgery (CLP+fibroblasts [Fibro], CLP+MSCs, CLP+MSCs/CO). A total of 5 × 10^5 cells were administered by tail vein injection 6 hr after CLP. A total of 5 × 10^5 cells were administered by tail vein injection 6 hr after CLP. Peritoneal fluid was collected and efferocytosis assays were performed. Data are presented as box plots, which show median values and interquartile ranges. Sham, n = 3; CLP+Fibro, light gray bar, n = 6; CLP+MSCs, dark gray bar, n = 6; CLP+MSCs/CO, striped bar, n = 7. Analysis was done by Kruskal-Wallis testing (p = 0.0004), with significant comparisons * versus sham, † versus CLP+Fibro, and ‡ versus CLP+MSCs. B. Mice were subjected to CLP, and 6 hr later, the mice received 5 × 10^5 MSCs treated with CO in the presence of docosahexaenoic acid (DHA) (10 μM), solid black line, n = 12 or MSCs-treated with CO in the presence of arachidonic acid (AA) (10 μM), dashed black line, n = 12. Survival of mice was monitored for 7 d and data are presented as a Kaplan-Meier survival curve, and analyzed by log-rank test. * versus MSCs+CO (AA), p = 0.040. C. Human MSCs were exposed to CO in the presence of DHA or AA, followed by lipid mediator profiling. Total SPMs are shown left, and total eicosanoids right. Data are presented as box plots, n = 4 samples per group from two independent experiments. Analysis was done by Mann-Whitney U testing. * versus MSCs+CO (AA), p = 0.03 for DHA or AA, p = 0.03 for eicosanoids. D. Representative pie charts of SPM levels in MSCs conditioned with CO, in the presence of DHA or AA. E. Human neutrophils were incubated with green fluorescent protein-labeled E. coli alone (white bars, n = 16), in the presence of MSCs (gray bar, n = 16), or in the presence of CO-conditioned MSCs (black bar, n = 16) or MSCs conditioned with resolvin D1 (RvD1, 10 nM), RvD2 (10 nM), or aspirin-triggered RvD3 (AT-RvD3, 10 nM) (black bars, n = 8). Data are presented as fold increase in mean fluorescence intensity, mean ± SEM, from three independent experiments. Analysis was done by one-way analysis of variance (p < 0.0001), with significant comparisons * versus no MSCs, † versus MSCs, and ‡ versus its RvD control. HETE = hydroxyeicosatetraenoic acid, Lx = lipoxin, PD1 = Protectin D1.

DISCUSSION

Gas molecules have been proposed as therapeutic agents for a variety of disease processes, including lung and even systemic illnesses (36, 37). One such gas is CO, which has been shown to be beneficial therapeutically in animal models of sepsis (16, 20, 21). Although preclinical studies and clinical trials are ongoing to more fully evaluate the safety and effectiveness of administering...
Figure 5. Lipoxygenase pathways contribute to the carbon monoxide (CO)-mediated effects of mesenchymal stromal cells (MSCs) on neutrophil phagocytosis, and silencing of 5-lipoxygenase (LOX) and 12/15-LOX results in a loss of improved MSC efficacy by CO conditioning. **A**. MSCs were pretreated with baicaicin (10 μM) or vehicle, and then cocultured with human (left) or mouse (right) neutrophils in the presence or absence of green fluorescent protein (GFP)-labeled *E. coli*. Data are presented as box plots, which show median values and interquartile ranges. Analysis was done by Kruskal-Wallis testing (also used in **B**) for 5-LOX and 12/15-LOX, with significant comparisons * versus Fibro, † versus Fibro+CO, and ‡ versus MSCs. **B**. MSCs were silenced with scrambled (SCR), or both 5-LOX and 12/15-LOX = small hairpin RNAs (shRNAs) (method used to silence 5-LOX and 12/15 LOX (shLOX) [defined in the narrative of the article]. The MSCs were exposed to CO or ambient air, and then cocultured with neutrophils in the presence of GFP-labeled *E. coli*. shLOX MSCs exposed to CO were also exposed to resolvin D1 (RvD1, 10 nM) or RvD2 (10 nM). Data are presented as fold increase in mean fluorescence intensity, mean ± SEM, n = 6–12 per group from three independent experiments. Analysis was done by one-way analysis of variance (p < 0.0001) with significant comparisons * versus no MSCs, † versus MSCs (SCR shRNA sequence [shSCR]), and ‡ between designated groups. **C**. BAECs (Fibro) and MSCs were incubated with CO or ambient air for 4 hr, and then RNA was extracted and quantitative real-time polymerase chain reaction performed for 5-LOX (upper) and 12/15-LOX (lower). Data are presented as box plots, n = 4–6 per group from three independent experiments. p = 0.0008 for 5-LOX and 12/15-LOX, with significant comparisons * versus Fibro, † versus Fibro+CO, and ‡ versus MSCs. **C**. MSCs were silenced with scrambled (SCR), or both 5-LOX and 12/15-LOX = small hairpin RNAs (shRNAs) (method used to silence 5-LOX and 12/15 LOX (shLOX) [defined in the narrative of the article]. The MSCs were exposed to CO or ambient air, and then cocultured with neutrophils in the presence of GFP-labeled *E. coli*. shLOX MSCs exposed to CO were also exposed to resolvin D1 (RvD1, 10 nM) or RvD2 (10 nM). Data are presented as fold increase in mean fluorescence intensity, mean ± SEM, n = 6–12 per group from three independent experiments. Analysis was done by one-way analysis of variance (p < 0.0001) with significant comparisons * versus no MSCs, † versus MSCs (SCR shRNA sequence [shSCR]), and ‡ between designated groups. **D**, BALB/c mice were randomly separated into two groups, shSCR MSCs + CO (solid black line, n = 14) or shLOX MSCs + CO (dashed black line, n = 14). Six hours after cecal ligation and puncture (CLP), the mice were treated with 5 × 10⁶ cells by tail vein injection. Animal survival was monitored for 7 d, and data are presented as a Kaplan-Meier survival curve, and analyzed by log-rank test. * versus shSCR MSCs+CO, p = 0.049. GAPDH = glyceraldehyde-3-phosphate dehydrogenase, mRNA = messenger RNA, NS = not significant.
of LXA4 (50). Conversely, in our study, LxA4 was expressed by CO
coli able to promote the resolution of acute lung injury induced by
organ injury (Fig. 2), and improved bacterial clearance (Fig. 3)
during experimental sepsis.

Once the invading bacteria are cleared, the resolution of inflammation is necessary to regain homeostasis and prevent organ injury during sepsis. SPMs are produced to coordinate the resolution response (6, 33, 35). In the presence of DHA substrate, MSCs exposed to CO predominantly produced SPMs and improved survival when administered after the onset of sepsis, whereas MSCs exposed to AA substrate resulted in eicosanoid production and reduced survival (Fig. 4). When assessing the differences in SPMs, the production of Rv family members was greater in cells exposed to DHA versus AA (Fig. 4). Interestingly, it was previously shown by Spite et al (47) that RvD2 improved survival in CLP-induced polymicrobial sepsis by regulating the inflammatory response, phagocytosis, and bacterial clearance. Taken together, these data suggest that production of SPMs (particularly D-series Rv) may contribute to the improved therapeutic response of MSCs exposed to CO ex vivo.

To confirm the importance of SPM production in MSCs preconditioned with CO, we exposed MSCs to a LOX inhibitor (baicalein) or silenced both 5-LOX and 12/15-LOX enzymes in MSCs. Either maneuver resulted in a loss of enhanced neutrophil phagocytosis induced by CO preconditioned MSCs (Fig. 5, A and C). In addition, silencing of both 5-LOX and 12/15-LOX resulted in a loss of survival benefit for MSCs preconditioned with CO, and a loss of MSCs efficacy when administered 6 hours after CLP-induced sepsis (Fig. 5D).

It has been suggested that exposure to CO promotes a beneficial response, in part, through a positive feedback loop increasing the production of its parent enzyme HO-1 (22, 48). In addition, the proresolving lipid mediator lipoxin A₄ (LxA₄) is known to increase HO-1 expression (49). However, we found that endogenous HO-1 was not required for the beneficial CO preconditioning response of MSCs during sepsis because HO-1–deficient cells were as effective as wild-type MSCs in protecting mice from CLP-induced mortality (Supplemental Fig. 4, Supplemental Digital Content 7, http://links.lww.com/ CCM/C14; legend, Supplemental Digital Content 14, http:// links.lww.com/CCM/C21).

Recently it was shown that human MSCs cocultured with alveolar epithelial type II cells have increased production of LXA₄ in the presence of proinflammatory cytokines (50). MSCs were also able to promote the resolution of acute lung injury induced by E. coli lipopolysaccharide, and this occurred in part by production of LXA₄ (50). Conversely, in our study, LxA₄ was expressed by CO preconditioned MSCs only in the presence of AA (Fig. 4D; and Supplemental Table 3, Supplemental Digital Content 10, http:// links.lww.com/CCM/C17), which resulted in the production of predominantly eicosanoids (Fig. 4C) and led to decreased survival in CLP-induced sepsis (Fig. 4B). The imbalance of increased production of AA-derived proinflammatory eicosanoids, and less production of SPMs (i.e., LxA4), likely contributed to a worse outcome. Taken together, these data suggest that LXA₄ is not playing as significant of a role in the MSC response to CO exposure, and its effect on outcome during polymicrobial sepsis.

Although the present study advanced our understanding of the therapeutic effects of MSCs in sepsis, we are presently limited to using a mouse model of disease. Furthermore, MSCs were not used in conjunction with antibiotics and hemodynamic resuscitation, standards of care for human patients, and they will need to be tested in the future. We believe that conditioning of MSCs with CO has the potential for future application; however, in this early stage of investigation, CO-conditioned MSCs are not immediately translatable to clinical application.

CONCLUSION
In the present study, we demonstrate that preconditioning of MSCs with CO gas improved MSC function and therapeutic efficacy during sepsis. Data including inhibition or silencing of LOX enzymes (5-LOX and 12/15-LOX) support the concept that the increased beneficial response of MSCs exposed to CO occurred in part through the production of SPMs, which contributed to the interaction of MSCs with neutrophils to promote bacterial phagocytosis, resolution of inflammation, and increased survival. We propose that MSCs preconditioned with CO may have future implications for the treatment of sepsis, allowing a beneficial response even with a delayed initiation of therapy.

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REFERENCES


