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Bone marrow mesenchymal stromal cells induce nitric oxide synthase-dependent differentiation of CD11b⁺ cells that expedite hematopoietic recovery

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ABSTRACT

Bone marrow microenvironment is fundamental for hematopoietic homeostasis. Numerous efforts have been made to reproduce or manipulate its activity to facilitate engraftment after hematopoietic stem cell transplantation but clinical results remain unconvincing. This probably reflects the complexity of the hematopoietic niche. Recent data have demonstrated the fundamental role of stromal and myeloid cells in regulating hematopoietic stem cell self-renewal and mobilization in the bone marrow. In this study we unveil a novel interaction by which bone marrow mesenchymal stromal cells induce the rapid differentiation of CD11b⁺ myeloid cells from bone marrow progenitors. Such an activity requires the expression of nitric oxide synthase-2. Importantly, the administration of these mesenchymal stromal cell-educated CD11b⁺ cells accelerates hematopoietic reconstitution in bone marrow transplant recipients. We conclude that the liaison between mesenchymal stromal cells and myeloid cells is fundamental in hematopoietic homeostasis and suggests that it can be harnessed in clinical transplantation.

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Introduction

Mesenchymal stromal cells (MSC) play a crucial role in tissue homeostasis whereby they control inflammation and regulate stem cell renewal and differentiation. Their immunomodulatory properties, which target both adaptive and innate immune responses, have been extensively documented *in vitro* and *in vivo*.¹⁻⁷ Although the underlying mechanisms are only partially known, it is widely accepted that the combination of soluble factors and contact-dependent interactions plays a fundamental role. Upregulation of inducible nitric oxide synthase (iNOS or NOS2), one of the key MSC transcriptional responses resulting in the secretion of a short-lived molecule with potent immunomodulatory effects, nitric oxide (NO),^{8,9} is not sufficient in itself to explain MSC the immunosuppressive activities. In fact, one of the main MSC effector mechanisms is the recruitment and functional modulation of myeloid cells, such as inflammatory monocytes and tissue macrophages.^{6,10-14}

MSC also contribute to the hematopoietic stem cell (HSC) niche in which they regulate hematopoietic cell number and differentiation.^{15,16} These properties have been harnessed therapeutically to promote hematopoietic regeneration. However, early *in vivo* animal studies have not been unequivocally confirmed by clinical investigations.^{18,19} Although the mechanisms by which MSC regulate HSC are still unknown, it is arguable that, resembling what has been described for their immunosuppressive actions, MSC require other cells to execute their functions.²⁰ In particular, a few studies have described that the interaction between MSC and bone marrow (BM) macrophages contributes to the retention of HSC in the BM²¹ and prevents their

exhaustion.^{20,24} The nature of this interaction has not, however, been elucidated.

In this work, we have tested the hypothesis that MSC may skew the differentiation and expansion of BM myeloid progenitors with the ability to accelerate hematopoietic reconstitution. We have observed that MSC selectively promote the expansion and differentiation of CD11b⁺ cells from the BM and that this function is largely dependent on NOS2. *Ex-vivo* generated MSC-induced CD11b⁺ cells exhibit the ability to accelerate hematopoietic engraftment and reconstitution.

Methods

Cell cultures and media

Murine BM MSC were generated from crushed femora and tibiae of wild type (WT) C57Bl/6 or *Nos2*^{-/-} mice (for further information, see the *Online Supplementary Appendix*). Human BM MSC were generated from human BM mononuclear cells (MNC). All samples were collected after informed consent had been obtained in accordance with Ethics Committee approval from the Province of Monza-Brianza (Italy) (approval date: 16 Oct 2014, approval file name: BM-NICHE). Further details on the methods used to generate murine and human MSC, and on the generation of MSC-induced myeloid cells, are presented in the *Online Supplementary Appendix*.

Flow cytometry

Unspecific binding sites were blocked with phosphate-buffered saline supplemented with 1% fetal bovine serum and either Fc blocker (CD16/32, eBioscience) or whole mouse IgG (Sigma Aldrich) before cells were incubated with the respective monoclonal antibody at 4°C for 30 min. After incubation, cells were washed twice with phosphate-buffered saline and analyzed by flow cytometry with a BD FACSCalibur, BD FACS CantoII, BD

LSRII or BD Fortessa (BD Biosciences, NJ, USA). Data were subsequently analyzed using FlowJo software (Oregon, USA). A complete list of antibodies used is given in the *Online Supplementary Appendix*.

In vivo experiments

For the adoptive transfer of MSC, sublethally irradiated (split dose of 800 cGy) WT CD45.1 C57Bl/6 recipients were transplanted by tail vein injection with 2x10⁶ BM cells and 0.2x10⁶ WT or *Nos2*^{-/-} MSC 4 h after the second dose of irradiation. Mice were sacrificed after 13 days and spleens and BM analyzed by FACS.

For the adoptive transfer of CD11b⁺ cells, sublethally irradiated (split dose of 800 cGy) WT CD45.1 C57Bl/6 recipients were transplanted with 5x10⁴ BM cells alone or with 2x10⁶ CD45.2⁺ MSC-induced CD11b⁺ cells 4 h after the second dose of irradiation. Peripheral blood samples were taken every 2 weeks after the transplant up to 4 months. Peripheral blood was lysed with lysis buffer (150 mM NH₄Cl, 1 mM KHCO₃, 0.1 mM EDTA, pH 7.3), and cells were stained for flow cytometry.

Statistical analysis

Data were analyzed using GraphPad Prism Software. An unpaired two-tailed Student *t*-test was run with a confidence interval of 95%, and expressed as mean ± standard error of the mean (SEM) or standard deviation (SD). For analysis of three groups of data, one-way analysis of variance with Bonferroni's multiple comparison test was used, and expressed as mean ± SEM.

Results

Mesenchymal stromal cells induce the expansion and differentiation of CD11b⁺ cells from bone marrow myeloid progenitors

Unfractionated BM MNC were cultured in the presence

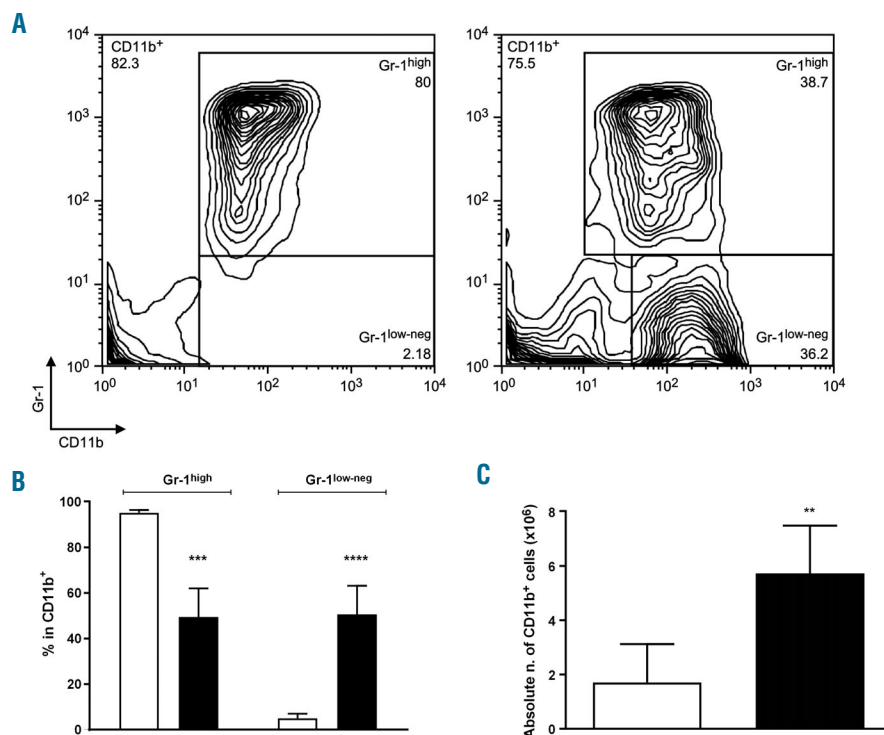


Figure 1. Mesenchymal stromal cells induce the differentiation of bone marrow mononuclear cells into myeloid cells. BM MNC were cultured alone or with MSC (ratio 5:1) for 4 days. (A) Proportion of CD11b⁺Gr-1^{high} and CD11b⁺Gr-1^{low-neg} cells in the live gate. A typical result of ten independently performed experiments is shown. (B) Percentage of Gr-1^{high} and Gr-1^{low-neg} cells in CD11b⁺ cells. Mean of ten independent experiments, ± SEM. *** *P*<0.001 **** *P*<0.0001 unpaired *t* test. (C) Absolute number of CD11b⁺ cells recovered from initial seeding from BM cultured alone (white bars) or with MSC (black bars) for 4 days. Mean of ten independent experiments, ± SEM ***P*<0.005, unpaired *t* test.

or absence of MSC. After 4 days, cultures were analyzed for the expression of the myeloid markers CD11b and Gr-1²⁵ (Figure 1A). Whilst in the control cultures the vast majority of BM cells consisted of CD11b⁺Gr-1^{high}, in those with MSC there was a marked skew towards the formation of a large proportion of CD11b⁺Gr-1^{low-neg} (50.6% ± 3.9%) (Figure 1B). Analysis of absolute cell numbers revealed that the presence of MSC could not only maintain the survival of total CD11b⁺ cells as compared to BM MNC cultured alone ($5.7 \times 10^6 \pm 1.8 \times 10^6$ compared to $1.7 \times 10^6 \pm 1.4 \times 10^6$) (Figure 1C), but also drive a selective retention and/or expansion of the Gr-1^{low-neg} population.

At morphological analysis the MSC-induced CD11b⁺ myeloid cells consisted of a fairly homogeneous population of large cells with reniform nuclei and abundant pale vacuolated cytoplasm with granules (Figure 2A). The immunophenotype of CD11b⁺ sorted cells revealed a 6-fold increase in F4/80⁺ (36.5% ± 10.3%), a 3-fold increase in IL4Rα⁺ (18.2% ± 7.5%), and a 2-fold increase in CD169⁺ (2.3% ± 0.6%) cells when compared to BM MNC cultured alone (Figure 2B, left panel). BM MNC cultured with MSC also expressed CD115 (48.6% ± 12.4%), CD206 (20.6% ± 2%) and CD68 (16.5% ± 4.9%) (Figure 2B, left panel). These macrophage markers were expressed only in the Gr-1^{low-neg} subset (Figure 2B, right panel), whilst CD115 was detected both in the Gr-1^{high} and the Gr-1^{low-neg} subsets.

To understand the target cells of MSC-induced myeloid differentiation, FACS-sorted HSC, common myeloid progenitors (CMP) or granulocyte/macrophage progenitors (GMP) were cultured with MSC. Megakaryocyte/erythroid progenitors (MEP) and unfractionated BM MNC were used as negative or positive control of differentiation, respectively. MSC induced the differentiation of only CMP and GMP into CD11b⁺Gr-1^{high} and CD11b⁺Gr-1^{low-neg} cells, with no effect on HSC or MEP (Figure 3A). The proportion of Gr-1^{low-neg} cells from CMP cultures was higher than in the cultures with unfractionated BM (Gr-1^{low-neg}: 60.1% ± 8.9% versus 35% ± 12.8% in unfractionated BM+MSC) (Figure 3B), and, accordingly, a 2-fold increase in the percentage of CD11b⁺F4/80⁺ cells (63.6% ± 9.8% versus 36.8% ± 13.7% in BM+MSC) and a higher percentage of CD11b⁺CD115⁺ cells (85.8% ± 1.3% versus 38.6% ± 18.9% in BM+MSC) (Figure 3C).

Mesenchymal stromal cell-induced CD11b⁺ Gr-1^{low-neg} formation is Nos2-dependent

Since NOS2 is one of the key effector molecules in MSC immunomodulatory properties,^{8,9} we tested the hypothesis that it could also be fundamental for the generation of Gr-1^{low-neg} cells. The differentiation of CD11b⁺Gr-1^{low-neg} cells was significantly impaired in cultures with *Nos2*^{-/-} MSC (16.6% ± 1.2% versus 30.8% ± 2.5% in BM+MSC WT), which correlated with an increased proportion in CD11b⁺Gr-1^{high} cells (82.3% ± 1.2% versus 68.2% ± 2.2% in BM+MSC WT) (Figure 4A,B). *Nos2*^{-/-} BM cells did not affect the ability of MSC to induce the generation of CD11b⁺Gr-1^{low-neg} cells (Figure 4A).

Mesenchymal stromal cells increase macrophage formation during hematopoietic reconstitution

The ability of MSC to drive the expansion and differentiation of CD11b⁺ cells was then studied *in vivo*. Sublethally irradiated mice were transplanted with BM cells from a congenic (CD45.1⁺) donor either alone or with WT or *Nos2*^{-/-} MSC. The proportion of the donor myeloid

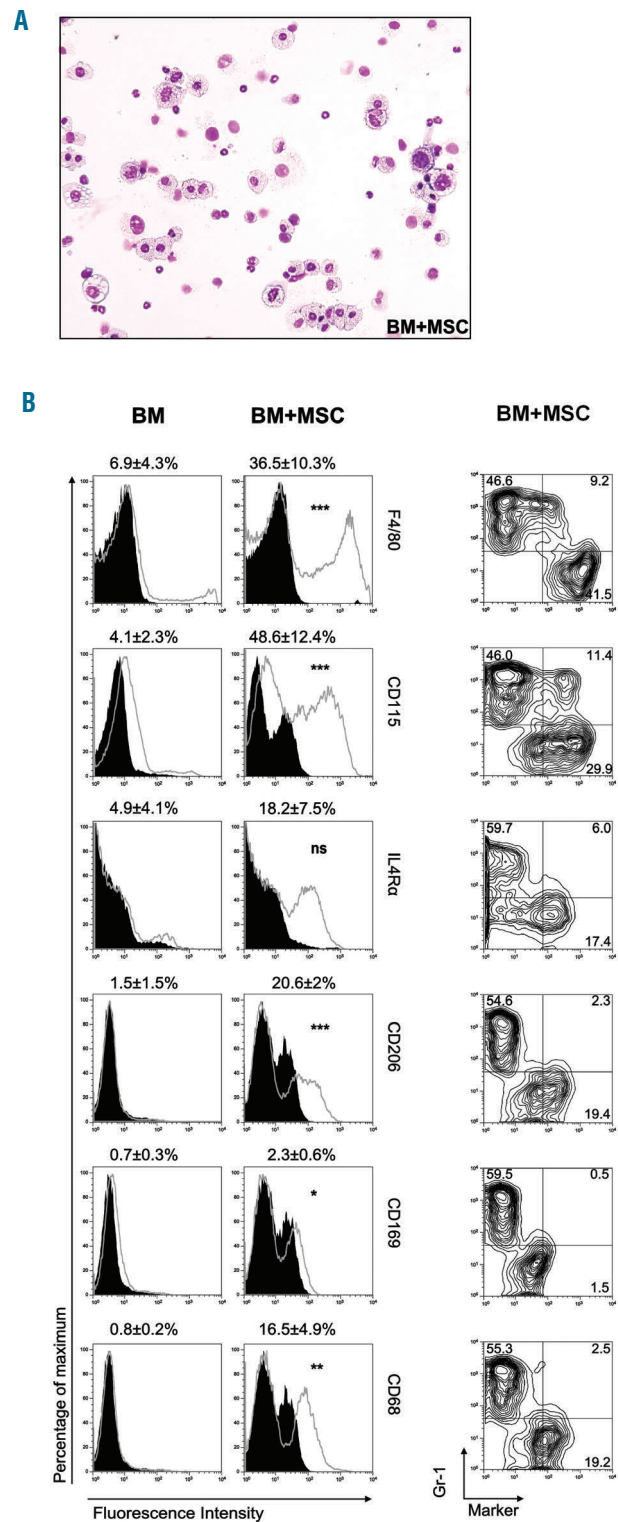


Figure 2. Mesenchymal stromal cell-induced CD11b⁺ cells consist of a large proportion of M0 macrophages. (A) May-Grünwald Giemsa staining of cytospin preparations of CD11b⁺ cells isolated from BM MNC cultured with MSC for 4 days. (B) BM MNC cultured alone or with MSC for 4 days were evaluated for the expression of macrophage surface markers within the CD11b⁺ gated population (open histograms) against their matched isotype controls (filled histograms). Contour plots within the CD11b⁺ gated population show the expression of each surface marker versus Gr-1 expression in BM MNC cultured with MSC. Contour and histogram plots from one out of six independent experiments, and mean fluorescence intensity values presented as mean ± SD of six independent experiments. **P* < 0.05, ***P* < 0.01, ****P* < 0.001, unpaired *t* test, all comparisons between 'BM' versus 'BM+MSC'.

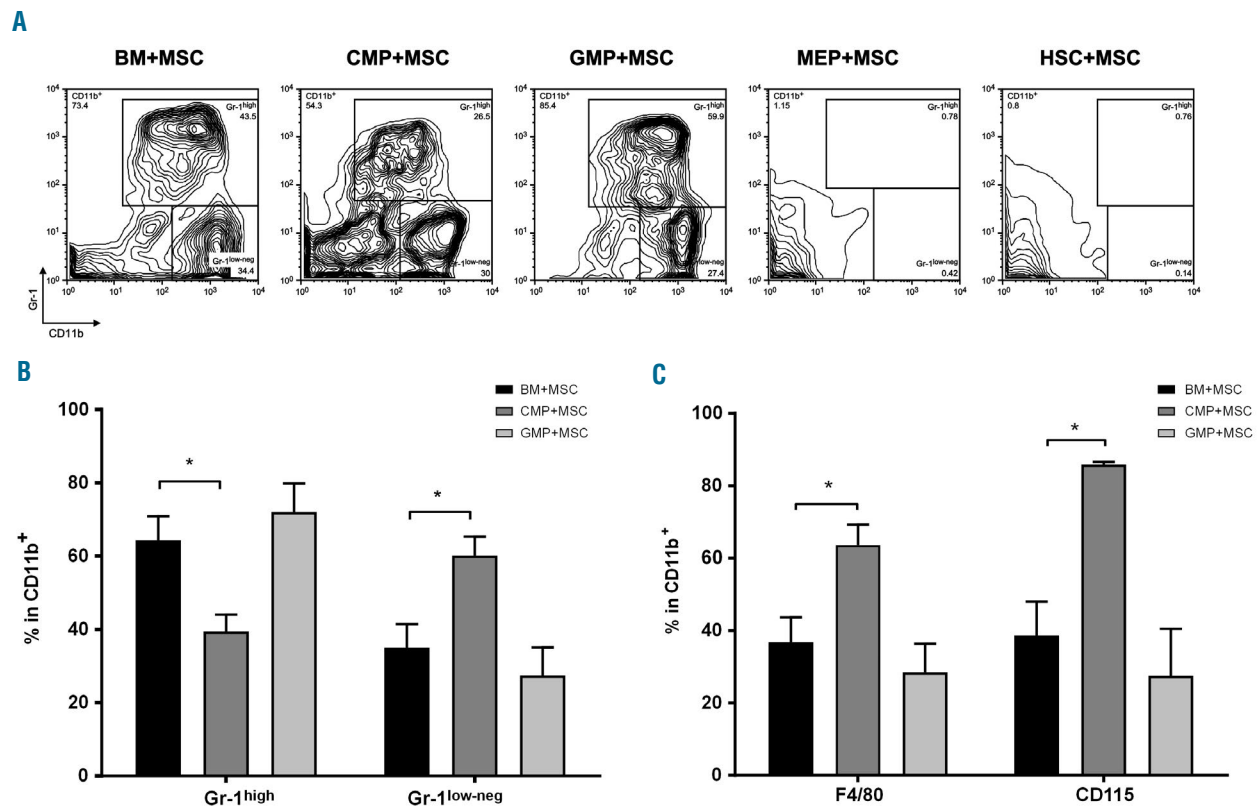


Figure 3. Mesenchymal stromal cell-induced CD11b⁺ differentiation targets committed myeloid progenitors but not hematopoietic stem cells. Unfractionated BM MNC (BM+MSC) or sorted CMP (CMP+MSC), GMP (GMP+MSC), MEP (MEP+MSC) and HSC (HSC+MSC) were cultured with MSC for 4 days. (A) Proportion of CD11b⁺Gr-1^{high} and CD11b⁺Gr-1^{low-neg} cells in the live gate. A typical result of four independently performed experiments is shown. (B) Percentage of Gr-1^{high} and Gr-1^{low-neg} cells within CD11b⁺ cells from BM, CMP or GMP cultures with MSC. Mean of four independent experiments, \pm SEM. * $P < 0.05$ unpaired t test. (C) Percentage of F4/80⁺ and CD115⁺ cells within the CD11b⁺ gate. Mean of four independent experiments, \pm SEM * $P < 0.05$, unpaired t test.

populations was analyzed in the BM and spleens of recipient mice 13 days after the transplant. At this time-point there was no difference in CD45.1⁺ engraftment (Figure 5A). However, the proportion and absolute numbers of Gr-1^{neg}F4/80⁺SSC^{low} macrophages were increased in mice receiving BM and WT MSC compared to the control group (3.2% \pm 0.3% versus 1.44% \pm 0.18%; absolute number: 0.46 \times 10⁶ \pm 0.12 \times 10⁶ versus 0.16 \times 10⁶ \pm 0.06 \times 10⁶) (Figure 5B,C). An increase in the proportion (Figure 5D) but not in the absolute number (Online Supplementary Figure S4E) of Ly6G⁺Ly6C⁺ monocytes was found in the WT MSC-treated group (23.5% \pm 1.9% versus 16.9% \pm 1.5%; absolute number: 1.06 \times 10⁶ \pm 0.14 \times 10⁶ versus 0.96 \times 10⁶ \pm 0.25 \times 10⁶).

Confirming *in vitro* results, the adoptive transfer of *Nos2*^{-/-} MSC failed to increase the proportion and absolute number of CD11b⁺Gr-1^{neg}F4/80⁺SSC^{low} macrophages and the proportion of Ly6G⁺Ly6C⁺ monocytes within the donor CD45.1⁺ engraftment (Gr-1^{neg}F4/80⁺SSC^{low} macrophages: 2.1% \pm 0.26% BM+*Nos2*^{-/-} MSC; absolute number: 0.32 \times 10⁶ \pm 0.07 \times 10⁶ BM+*Nos2*^{-/-} MSC. Ly6G⁺Ly6C⁺ monocytes: 18.1% \pm 1.3% BM+*Nos2*^{-/-} MSC) (Figure 5B-D).

There was no difference in the proportion or absolute number of neutrophils or eosinophils in the donor engraftment in all treated groups (Online Supplementary Figure S4C,D). The adoptive transfer of MSC and BM did not affect hematopoietic generation of donor myeloid cells in the spleen of recipient mice (*data not shown*).

Ex-vivo mesenchymal stromal cell-induced CD11b⁺ cells accelerate hematopoietic reconstitution

In order to further characterize the role of MSC-induced CD11b⁺ cells *in vivo*, we investigated their effect on hematopoietic reconstitution. For this purpose, sublethally irradiated mice received a BM transplant from a congenic donor with or without CD11b⁺ cells purified from the BM-MSC co-cultures. The engraftment of donor cells was monitored in the peripheral blood every 2 weeks. The group given CD11b⁺ cells showed a higher proportion (11.7% \pm 1.5% versus 20.9% \pm 2.1%) and absolute number (17.0 \pm 4.2 versus 43.3 \pm 7.5 CD45.1⁺ leukocytes/ μ L) of donor cells as compared to the control group already at 2 weeks. Such an increment was also observed in each leukocyte compartment (neutrophils: 37.4% \pm 3.4% versus 53.3% \pm 2.9%, and 8.4 \pm 2.4 versus 22.8 \pm 3.8 CD45.1⁺ neutrophils/ μ L; monocytes: 16.4% \pm 2.4 versus 21.2% \pm 2.9%, and 4.9 \pm 1.2 versus 12.1 \pm 2.9 CD45.1⁺ monocytes/ μ L, $P = 0.0195$; B lymphocytes: 10.7% \pm 1.5% versus 18.2% \pm 2.4%, and 3.9 \pm 0.7 versus 9.3 \pm 2.7 CD45.1⁺ B lymphocytes/ μ L; T lymphocytes: 0.06% \pm 0.03% versus 0.3% \pm 0.09%, and 0.02 \pm 0.01 versus 0.1 \pm 0.04 CD45.1⁺ T lymphocytes/ μ L) (Figure 6A,B).

Analysis of the absolute numbers of leukocyte populations in peripheral blood showed that CD11b⁺ cells not only expanded donor engraftment but they also enhanced the recovery of total hematopoiesis. At 2 weeks, the group

of mice given CD11b⁺ cells showed higher absolute counts of neutrophils (15.2±3.4 versus 42.1±5.9 neutrophils/μL), B lymphocytes (27.8±4.8 versus 51.5±7.7 B lymphocytes/μL) and monocytes (22.6±3.7 versus 37.9±4.1 monocytes/μL) (Online Supplementary Figure S5A).

Monitoring of the hematopoietic engraftment at later stages revealed that the CD11b⁺-driven hematopoietic reconstitution was still evident 4 weeks after the transplant (Online Supplementary Figures S5B and S6A,B). At 6 weeks the enhancement effect was selectively observed within the B and T lymphocytes compartments, whilst no

difference in the quality and quantity of engraftment was found from 8 weeks onwards (Online Supplementary Figures S5B and S6A,B).

Human mesenchymal stromal cells induce the differentiation of CD14⁺ cells from bone marrow mononuclear cells

Finally, we asked whether the newly discovered ability of murine MSC to induce macrophage differentiation is also a property of human MSC. For this purpose, human BM MNC were cultured alone or with human MSC for 7

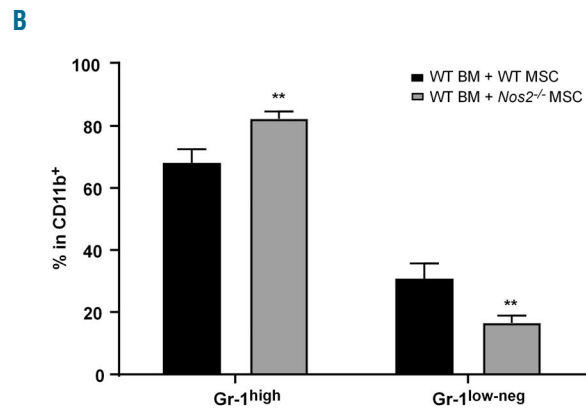
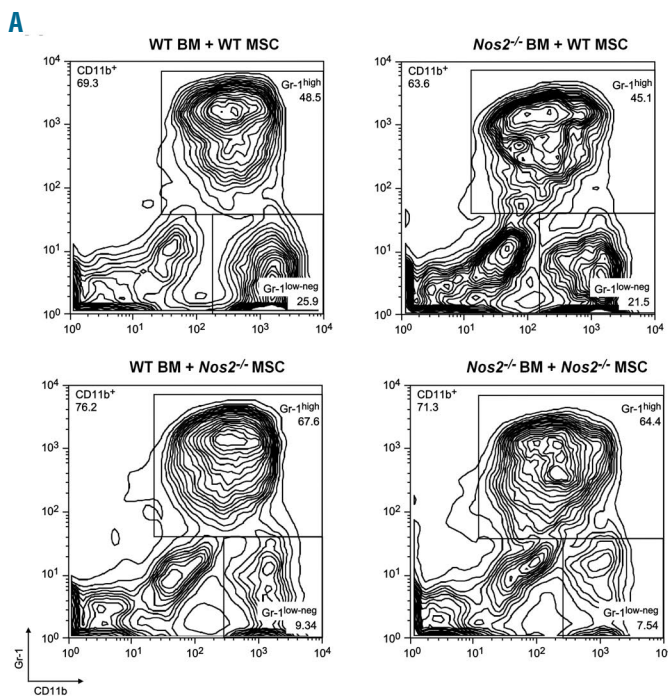


Figure 4. Mesenchymal stromal cell-induced myeloid cell formation is Nos2-dependent. (A, B) BM MNC isolated from WT or *Nos2*^{-/-} mice were cultured with MSC isolated from WT or *Nos2*^{-/-} mice for 4 days. (A) Contour plots represent CD11b⁺Gr-1^{high} and CD11b⁺Gr-1^{low-neg} proportions in the live gate. A typical result of three independently performed experiments is shown. (B) Percentages of Gr-1^{high} and Gr-1^{low-neg} within the CD11b⁺ gate. Mean of three independent experiments ±SEM, ***P*<0.01, unpaired t test.

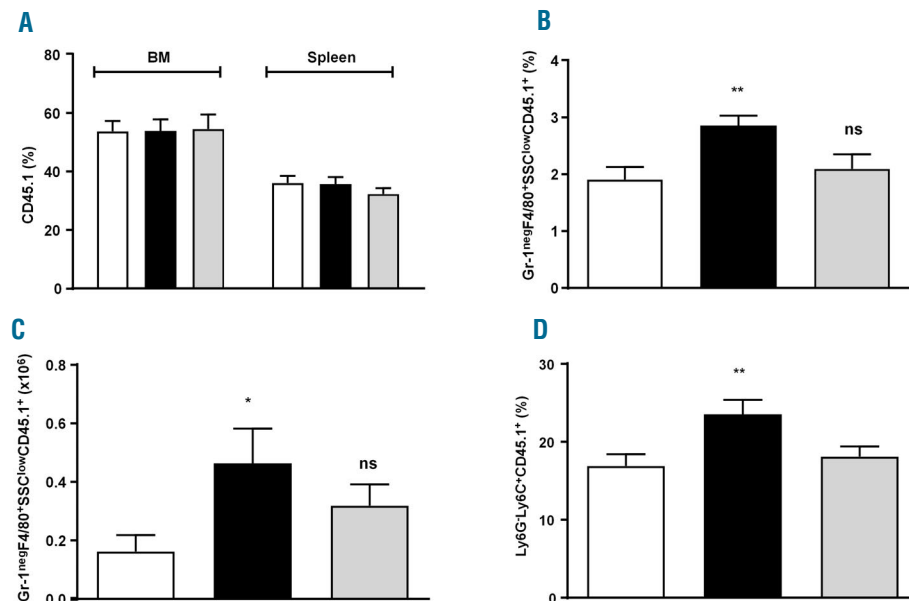


Figure 5. Adoptive transfer of mesenchymal stromal cells increases macrophage formation during hematopoietic reconstitution. Sublethally irradiated (800 cGy) CD45.2 WT recipients were injected with 2x10⁶ CD45.1 WT BM cells, either alone (BM - white bars) or in combination with 0.2x10⁶ CD45.2 WT MSC (BM+WT MSC - black bars) or 0.2x10⁶ CD45.2 *Nos2*^{-/-} MSC (BM+Nos2^{-/-} MSC - gray bars). 13 days after the transfer, BM and spleen were analyzed by FACS. (A) Percentage of donor engraftment in BM and spleen. (B, C) Percentage (B) and absolute number (C) of Gr-1^{high}F4/80⁺SSC^{low}CD45.1⁺ cells in the BM. Mean of three independent experiments ±SEM, **P*<0.05, ***P*<0.01 One-way analysis of variance (ANOVA) with the Bonferroni multiple comparison test. (D) Percentage of Ly6G⁺Ly6C⁺ monocytes within donor hematopoiesis (CD45.1⁺ cells) in the BM. Mean of three independent experiments ±SEM, ***P*<0.01 One-way ANOVA with the Bonferroni multiple comparison test.

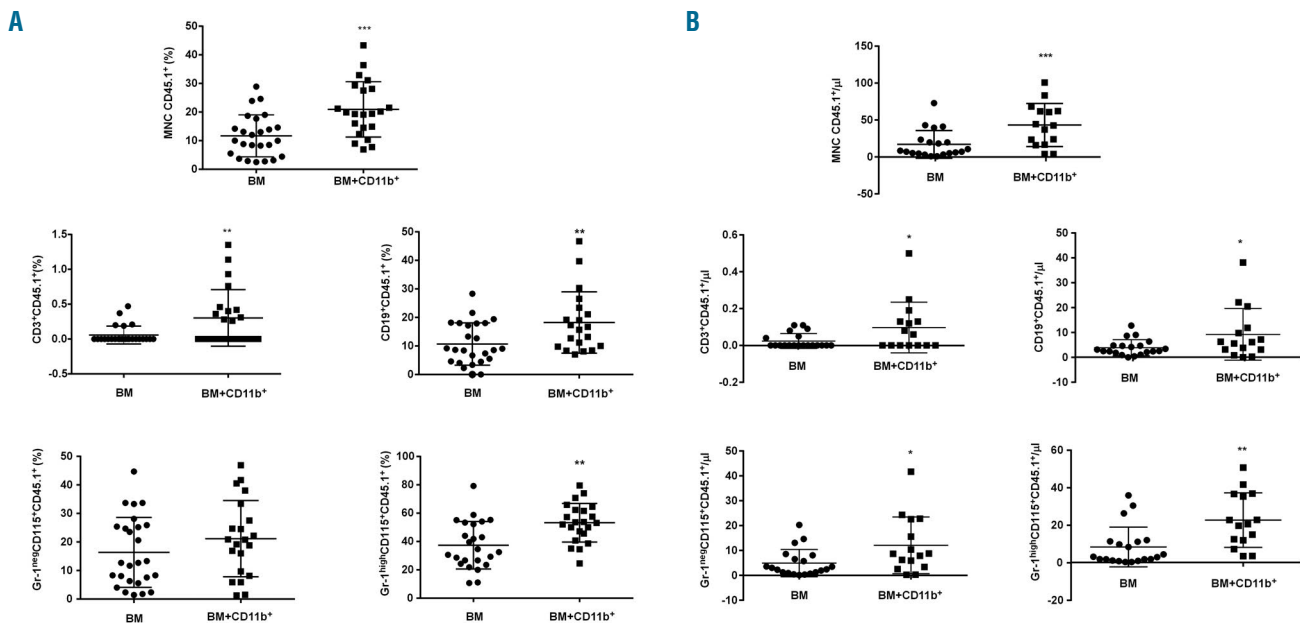


Figure 6. Adoptive transfer of mesenchymal stromal cell-induced CD11b⁺ cells accelerates engraftment. Sublethally irradiated (800 cGy) CD45.2 WT recipients were injected with 5×10^4 CD45.1 WT BM cells, either alone (BM) or in combination with 2×10^6 MSC-induced CD11b⁺ cells (BM+CD11b⁺). Peripheral blood samples were taken 14 days after the transplant, and the frequency of donor hematopoiesis analyzed by FACS as described in the *Methods*. (A) Frequency of donor hematopoiesis in MNC, T cells (CD3⁺), B cells (CD19⁺), monocytes (Gr-1^{hi}CD115⁺) and neutrophils (Gr-1^{hi}CD115⁻). Mean of four independent experiments \pm SEM, ** $P < 0.01$, *** $P < 0.001$, unpaired t test. (B) Absolute number of donor hematopoietic cells in MNC, T cells (CD3⁺), B cells (CD19⁺), monocytes (Gr-1^{hi}CD115⁺) and neutrophils (Gr-1^{hi}CD115⁻). Mean of four independent experiments \pm SEM, * $P < 0.05$, ** $P < 0.01$, unpaired t test.

days. We decided to analyze the proportions of CD14⁺, CD14⁺HLA-DR^{low-neg} and CD14⁺CD16⁺ cells, which best represent total myeloid population, macrophages and non-classical monocytes, respectively. The presence of human MSC induced an increment in CD14⁺ cells with a selective effect on non-classical monocytes (CD14⁺: $15.5\% \pm 2.2\%$ versus $25.8\% \pm 3.6\%$, in human BM and human BM cultured with human MSC, respectively; CD14⁺CD16⁺: $2.2\% \pm 0.6\%$ versus $7.5\% \pm 1.6\%$, in human BM and human BM cultured with human MSC, respectively) (Figure 7A). The analysis of the absolute numbers within the CD45⁺ population confirmed the statistically significant increase in non-classical monocytes (CD14⁺CD16⁺: $0.01 \times 10^6 \pm 0.003 \times 10^6$ versus $0.08 \times 10^6 \pm 0.03 \times 10^6$ in human BM and human BM cultured with human MSC, respectively) (Figure 7B). In contrast to what we observed in the murine system, the addition of the NOS2 inhibitor 1400W to cultures did not affect myeloid differentiation (*data not shown*).

Discussion

Our study unveils a previously unknown property of BM MSC consisting in the ability to differentiate and expand *in vitro* and *in vivo* myeloid cells from BM progenitors. Importantly, the administration of these myeloid cells accelerates hematopoietic reconstitution in BM transplant recipients.

The mechanism by which MSC promote myeloid expansion is largely dependent on NOS2. Although NOS2

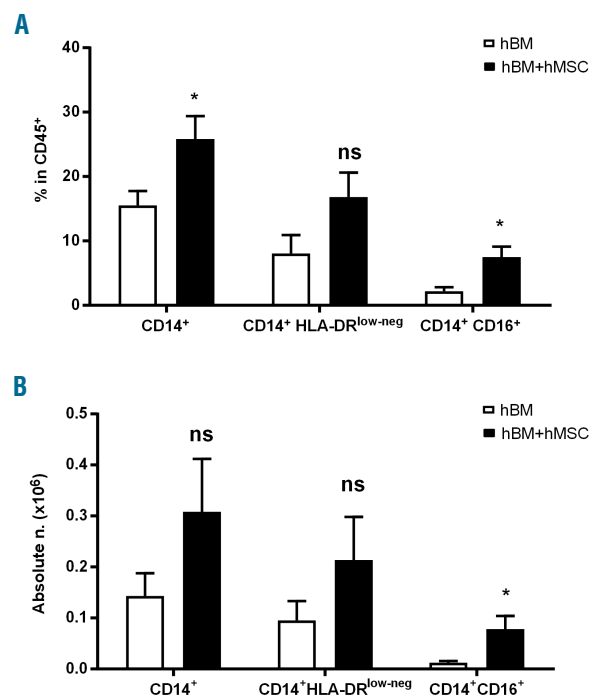


Figure 7. Human mesenchymal stromal cells induce the differentiation of human bone marrow mononuclear cells into myeloid cells. Human BM MNC were cultured alone or with MSC (ratio 10:1). After 7 days, the percentages (A) and absolute numbers (B) of CD14⁺, CD14⁺HLA-DR^{low-neg} and CD14⁺CD16⁺ in the CD45⁺ population were analyzed. Mean of six independent experiments, \pm SEM. * $P < 0.05$, unpaired t test.

has been extensively identified as the main effector mechanism accounting for MSC immunosuppressive activity and similarly implicated in myeloid immunobiology, this is the first study that attributes NOS2 the ability to differentiate and expand myeloid cells and macrophages. In contrast to the MSC-mediated immunosuppressive activity,⁸ we observed that the generation and expansion of myeloid cells was not influenced by inflammatory molecules such as tumor necrosis factor- α and interferon- γ (*data not shown*). This suggests that, despite using similar mechanisms, MSC-induced myeloid differentiation occurs independently of inflammation.

The potent ability of myeloid cells expanded *ex-vivo* by MSC to accelerate hematopoietic regeneration has at least two implications. The first is that the role of MSC in regulating HSC differentiation is not exclusively a direct one but can also be exerted by recruiting accessory cells. This has already been documented in the landscape of MSC immunomodulation.²⁶ Furthermore, the regenerative properties on the hematopoietic system may apply to other tissues. The molecular and functional profile of MSC-educated myeloid cells recapitulates that of the resident macrophages which have been described as involved in tissue repair activity in other organs.^{27,28} Therefore, our data might lend support to the notion that stroma/fibroblasts orchestrate tissue homeostasis.

The second set of implications affects the clinical approach to the use of MSC. MSC have long been studied for their ability to promote hematopoietic engraftment based on the evidence that they are a crucial component of the stem cell niche. After the initial investigation in xenogeneic models showing that MSC co-transplanted with umbilical cord blood CD34⁺ cells resulted in an increase in BM engraftment,¹⁷ subsequent studies indicated that such engraftment is only transient.^{29,30} Similar confounding factors can be found in clinical studies in pediatric^{31,32} and adult^{19,33-35} cohorts with small and heterogeneous groups of

patients. If MSC graft facilitating activity is the consequence of inducing BM progenitors to differentiate into regenerative myeloid cells, the variability in clinical outcome may result from the content of progenitors in the original donor inoculum rather than the actual direct activity of MSC on HSC differentiation.

Our data highlight the complexity of the MSC 'clinical niche effect'. Although the adoptive transfer of MSC at the time of BM transplantation significantly increased the percentage and number of donor macrophages in the BM (Figure 5B,C), we could not observe a consequent increment in the overall donor hematopoietic engraftment (Figure 5A). This discrepancy could be explained by the low number of macrophages produced by the MSC infusion. In fact, when BM cells were infused with a high number of *ex-vivo* MSC-induced myeloid cells, the hematopoietic engraftment was greatly enhanced (Figure 6).

The crucial role of myeloid cells in facilitating engraftment is also supported by recent clinical evidence in which MSC have been used to condition cord blood HSC *ex-vivo* in order to accelerate hematopoietic recovery. Our data shed light on the interpretation of these successful clinical results.³⁶ In that study, the MSC-conditioned cord blood unit did not engraft, indicating it mainly played a facilitating effect on the engraftment of the co-transplanted unmanipulated unit. Our study suggests that the graft facilitating effect might have been mediated by MSC-expanded monocytes, as confirmed by the phenotypic analysis of their MSC-educated cord blood HSC. Therefore, we propose the intriguing possibility that *ex-vivo* MSC induced myeloid cells could be harnessed in the clinical setting to expedite hematopoietic recovery and immune reconstitution in high-risk transplantation procedures.³⁷

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