Expansion of human SCID-repopulating cells under hypoxic conditions

Guénahel H. Danet, …, Dominique A. Bonnet, M. Celeste Simon


It has been proposed that bone marrow (BM) hematopoietic stem and progenitor cells are distributed along an oxygen (O₂) gradient, where stem cells reside in the most hypoxic areas and proliferating progenitors are found in O₂-rich areas. However, the effects of hypoxia on human hematopoietic stem cells (HSCs) have not been characterized. Our objective was to evaluate the functional and molecular responses of human BM progenitors and stem cells to hypoxic conditions. BM lineage-negative (Lin−) CD34+CD38− cells were cultured in serum-free medium under 1.5% O₂ (hypoxia) or 20% O₂ (normoxia) for 4 days. Using limiting dilution analysis, we demonstrate that the absolute number of SCID-repopulating cells (SRCs) increased by 5.8-fold in hypoxic cultures compared with normoxia, and by 4.2-fold compared with freshly isolated Lin−CD34+CD38− cells. The observed increase in BM-repopulating activity was associated with a preferential expansion of Lin−CD34+CD38− cells. We also demonstrate that, in response to hypoxia, hypoxia-inducible factor-1α protein was stabilized, surface expression of angiogenic receptors was upregulated, and VEGF secretion increased in BM Lin−CD34+ cultures. The use of low O₂ levels to enhance the survival and/or self-renewal of human BM HSCs in vitro represents an important advance and could have valuable clinical implications.
Expansion of human SCID-repopulating cells under hypoxic conditions

Guénahel H. Danet,1 Yi Pan,2 Jennifer L. Luongo,1 Dominique A. Bonnet,3 and M. Celeste Simon2

1Division of Hematology-Oncology, University of Pennsylvania School of Medicine, Philadelphia, Pennsylvania, USA
2Howard Hughes Medical Institute and Abramson Family Cancer Research Institute, University of Pennsylvania Cancer Center, Philadelphia, Pennsylvania, USA
3Cancer Research UK, London Research Institute, London, United Kingdom

It has been proposed that bone marrow (BM) hematopoietic stem and progenitor cells are distributed along an oxygen (O2) gradient, where stem cells reside in the most hypoxic areas and proliferating progenitors are found in O2-rich areas. However, the effects of hypoxia on human hematopoietic stem cells (HSCs) have not been characterized. Our objective was to evaluate the functional and molecular responses of human BM progenitors and stem cells to hypoxic conditions. BM lineage–negative (Lin−) CD34+CD38− cells were cultured in serum-free medium under 1.5% O2 (hypoxia) or 20% O2 (normoxia) for 4 days. Using limiting dilution analysis, we demonstrate that the absolute number of SCID-repopulating cells (SRCs) increased by 5.8-fold in hypoxic cultures compared with normoxia, and by 4.2-fold compared with freshly isolated Lin−CD34+CD38− cells. The observed increase in BM-repopulating activity was associated with a preferential expansion of Lin−CD34+CD38− cells. We also demonstrate that, in response to hypoxia, hypoxia-inducible factor-1α protein was stabilized, surface expression of angiogenic receptors was upregulated, and VEGF secretion increased in BM Lin−CD34+ cultures. The use of low O2 levels to enhance the survival and/or self-renewal of human BM HSCs in vitro represents an important advance and could have valuable clinical implications.


Introduction
The effects of hypoxia on human hematopoietic progenitors and stem cells remain poorly understood. It has only recently been reported that, in normal volunteers, the partial pressure of oxygen (O2) and O2 saturation in human bone marrow (BM) are lower than in peripheral blood (1). The presence of hypoxic areas in the BM is determined by the architecture of medullary sinuses and the pattern of arterial blood flow in the marrow (2). It has been proposed that hematopoietic stem cells (HSCs) and progenitors are distributed along an O2 gradient, with stem cells residing in the most hypoxic areas and proliferating progenitors in O2-rich areas (3). The data in support of this model were generated using “closed systems” in which O2 consumption by cells results in a progressive decrease in O2 level below the initial 1%. In these conditions, Cipolleschi et al. (3) demonstrated that the ability of HSCs to repopulate the BM of lethally irradiated mice and give rise to myeloid colonies was better preserved in hypoxia (<1.5% O2 for 5 days) than in normoxic conditions. In contrast, committed progenitors (GM-CFU) could not be preserved under the same low O2 levels. When an O2 regulation device and growth factor stimulation were used, Ivanovic et al. (4) reported a smaller expansion of mouse progenitors and a better maintenance of the number of myeloid progenitors in the recipients’ BM after 8 days in hypoxia compared with normoxia. In addition, we have shown that the proliferation of embryonic hematopoietic progenitors is regulated by a hypoxia-mediated signaling pathway (5).

The available data on the effects of hypoxia on human hematopoietic cells is limited to the clonogenic activity of umbilical cord blood, mobilized peripheral blood, or BM progenitor cells. Cipolleschi et al. (6) reported an increase in burst-forming unit, erythroid (BFU-E) and a decrease in GM-CFU colonies when cord blood CD34+ cells were cultured in a 14-day clonogenic assay under severe hypoxia (<1% O2). An increase in both BFU-E and GM-CFU activity in BM CD34+ cells cultured in 1.5% O2 for 6 hours has also been reported (7). In contrast, the culture of mobilized peripheral blood CD34+ cells under severe hypoxia (<1% O2) for 7...
days resulted in a decrease in the total number of colony-forming cells (CFCs) (8). Although these in vitro assays provide clues on the effect of hypoxia on the clonogenic activity of human progenitors, they may not reflect its effects on human HSCs. It is necessary to use a xenotransplantation model, such as NOD/SCID mice (9), to evaluate the functional response of human BM-repopulating cells to culture under low O2 levels. Human HSCs capable of extensive proliferation and multilineage repopulation of the BM of NOD/SCID mice are defined as SCID-repopulating cells (SRCs) (10–12). SRCs are highly enriched in the BM lineage-negative (Lin-) CD34+CD38– fraction (10–12) and present at a higher frequency in umbilical cord blood than in adult BM (13, 14). We and others (15, 16–24) have examined various culture conditions for their ability to maintain or enhance SRC activity in vitro. From these studies, it has been shown that the ex vivo expansion of CFCs and long-term colony-initiating cells (LTC-ICs) is not necessarily associated with an increase in SRCs. Furthermore, limiting dilution analysis is required to quantitatively compare the effect of culture conditions on SRC activity (12, 17–20, 24).

The cellular mechanisms by which human hematopoietic progenitors and stem cells respond to hypoxia have not been characterized. However, the response to hypoxia has been investigated in a variety of other cell types and models (see refs. 25–29 for reviews). The hypoxia-inducible factor-1α (HIF-1α) protein, which is rapidly degraded under normoxic conditions, becomes stabilized (30) under low O2 levels (<5%) and forms a dimer with aryl hydrocarbon receptor nuclear translocator protein (ARNT). In contrast to HIF-1α, ARNT levels are not regulated by hypoxia. The heterodimer HIF-1α-ARNT is a transcriptional activator of genes encoding for a wide variety of genes including erythropoietin, VEGF, glucose transporters, and glycolytic enzymes (27, 31). HIF-2α (32–34) and HIF-3α (35) are closely related to HIF-1α but have a more restricted pattern of expression and partially overlapping functions (27).

The objectives of this study were to evaluate the functional and cellular responses of human BM hematopoietic progenitors and stem cells to hypoxia. We used normal adult BM as a source of human HSCs because these cells reside in a hypoxic environment (1) and therefore may represent a more relevant model than other sources of human HSCs such as cord blood or mobilized peripheral blood. In this study, we demonstrate quantitatively that adult BM SRCs can be expanded in vitro under hypoxic conditions. This increase in BM-repopulating activity was associated with the preferential expansion of Lin-CD34+CD38− cells, an upregulation of angiogenic receptors, and an increase in VEGF production. Our results suggest that hypoxia could play a critical role in regulating the self-renewal of human BM HSCs and that the increase in VEGF secretions induced by hypoxia could be involved in the maintenance of HSCs in vitro.

**Methods**

**BM cell isolation.** BM aspirates (n = 137) were obtained from healthy volunteers in accordance with the guidelines of the University of Pennsylvania Institutional Review Board for Human Subjects. At least three BM samples from three different donors were pooled for the isolation of Lin− cells. Light-density mononuclear cells were isolated using Ficoll-Paque Plus (Amersham Biosciences Corp., Piscataway, New Jersey, USA), erythrocytes were lysed using ammonium chloride, and Lin− cells were isolated using immunomagnetic beads (Stem-Cell Technologies Inc., Vancouver, British Columbia, Canada). Briefly, cells were incubated with a mixture of lineage-specific antibodies (CD2, CD3, CD14, CD16, CD19, CD24, CD56, CD66b, CD41, and glycoporphin A) followed by incubation with a secondary antibody conjugated to metal colloid. Cells were then eluted through a magnetized column to deplete the suspension from cells expressing lineage markers. Lin− cells were then stained with anti-human CD34-phycocerythrin (CD34-PE), anti-human CD38-allophycocyanine (both from Becton, Dickinson and Co., San Jose, California, USA), and propidium iodide (Molecular Probes Inc., Eugene, Oregon, USA) and sorted on a MoFlo cell sorter (Cytomation Inc., Fort Collins, Colorado, USA).

**Liquid cultures.** BM Lin-CD34+ or Lin-CD34+CD38− cells (10^6/ml) were cultured in serum-free medium consisting of StemSpan SFEM medium (StemCell Technologies Inc.) supplemented with IL-3 (10 ng/ml), IL-6 (10 ng/ml), stem cell factor (SCF) (300 ng/ml), Flt-3 ligand (300 ng/ml), G-CSF (50 ng/ml), 1% HEPES, gentamicin, and LDL (10 µg/ml). Cells were cultured for 4–9 days at 37°C in 5% CO2-humidified incubators in normoxic (20% O2) or hypoxic (1.5% O2) conditions. Hypoxic cultures were performed in a two-gas incubator (Jouan Inc., Winchester, Virginia, USA) equipped with an O2 probe to regulate N2 levels. In some experiments, normoxic cultures of BM Lin-CD34+ cells were supplemented with human VEGF (100 ng/ml), angiopoietin-1 (100 ng/ml), or angiopoietin-2 (100 ng/ml), all obtained from R&D Systems Inc. (Minneapolis, Minnesota, USA).

**Phenotypic analysis and cell division tracking.** Lin-CD34+ cells were harvested after 4 days of culture in hypoxic (1.5% O2) or normoxic conditions. Cell number and viability were evaluated using trypan blue exclusion. Cells were incubated with anti-human CD34 and the following antibodies: CD31, CD38, CD117 (c-kit), and HLA-DR (Becton, Dickinson and Co.), CD90 from Beckman Coulter (Miami, Florida, USA), CD133 from Miltenyi Biotec (Auburn, California, USA), CD135 and CD162 from Immunotech Inc. (Westbrook, Maine, USA), CXCR4, Tie-1, Tie-2, Flt-1, and VEGF receptor 2 (VEGFR2, also known as Kdr, Flk-1) from R&D Systems Inc., and vWF from Santa Cruz Biotechnology Inc. (Santa Cruz, California, USA). In some experiments, freshly isolated Lin-CD34+ cells were labeled with CFSE (Molecular Probes Inc., Eugene, Oregon, USA) as previously described (15). Cells were cultured
for 4 days in hypoxic (1.5% O₂) or normoxic conditions and stained for CD34-allophycocyanin and either the appropriate isotype control or one of the following anti-human antibodies: CD38, CD39, CD117, CD133, CD135, or HLA-DR. Cells were then analyzed on a FACSCalibur flow cytometer (Becton, Dickinson and Co.).

Cell cycle analysis. Lin−CD34+ cells cultured for 4 days in hypoxic (1.5% O₂) or normoxic conditions were harvested and cell cycle analysis was performed as previously described (36). Briefly, cells were incubated with CD34-PE, then fixed with a 0.4% formaldehyde-buffered solution, washed, and permeabilized with a 0.2% Triton X-100 solution on ice. After two washes in PBS plus 2% FBS, cells were labeled with Ki-67–FITC (Becton, Dickinson and Co.). Finally, each sample was washed twice and resuspended in a 10 µM solution of DAPI (Molecular Probes Inc.) in PBS plus 2% FBS. Cells were analyzed using a Becton-Dickinson LSR flow cytometer equipped with a 325-nm helium-cadmium UV laser and a 488-nm argon-ion laser.

CFC and LTC-IC assays. Human CFCs were assayed in semisolid methylcellulose medium in standard conditions (12). LTC-IC cultures were established on preformed M210B4 stroma layers using human myeloid long-term culture medium (MyeloCult S100; StemCell Technologies Inc.) according to described methods (37, 38). Limiting dilutions (at least three replicates/dilution) were performed in 96-well plates in which conditioned media (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three replicates/dilution) were performed in 96-well plates in which conditioned media (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three mice per cell dose per experiment). Lin−CD34+CD38− cells were coinjected with 0.5 million accessory cells consisting of irradiated (1,500 cGy) BM or cord blood mononuclear or lineage-positive (at least three mice per cell dose per experiment).

RT-PCR. RNA was isolated from freshly purified Lin−CD34+CD38− or Lin−CD34−CD38+ cells using the RNeasy isolation kit (Ambion Inc., Austin, Texas, USA). RNA extracted from K562 cells was used as a positive control for the PCR reactions. Standard reverse transcription was done using SuperScript II reverse transcriptase (Invitrogen Corp., San Diego, California, USA). The primers used were HIF-1α: 5′-AGATCCTCGGATT-GCAGCGCA and 5′-AGTTAGTCTAAAATCGAATTTACCC; HIF-2α: 5′-GCCCTGCTGTCTGTTCACTCA and 5′-ATCC-GTCTGGTTACTGCTGGTTC; and ARNT: 5′-AGGAAATGTTGGCCTACGCCCCCT and 5′-ATGGTTGAGCT-GTTGCTCTG. PCR was done using the Advantage- GC cDNA PCR kit (Clontech Laboratories Inc., Palo Alto, California, USA). The amplification cycles were: 94°C for 4 minutes followed by 35 cycles of 94°C for 10 minutes + 58°C for 30 seconds + 72°C for 2 minutes, and a final extension step at 72°C for 10 minutes was included before samples were cooled at 4°C.

Western blot analysis. Equal numbers of Lin−CD34+ cells were cultured overnight in serum-free conditions (10% /ml) at 1.5% O₂ in a hypoxic workstation (Invivox; Ruskinn Technology, Leeds, United Kingdom) or in normoxic conditions. Cells were directly lysed in 1× SDS loading buffer and boiled at 95°C for 10 minutes. The mouse anti-HIF-1α antibody used was from Becton, Dickinson and Co. (catalog no. H72320) and the rabbit anti-ARNT antibody was from Novus Biologicals Inc. (catalog no. NB 100-110; Littleton, Colorado, USA).

Human VEGF ELISA and lactate assay. Conditioned media from overnight cultures (n = 5) of Lin−CD34+ cells were harvested and human VEGF and erythropoietin concentrations were determined by ELISA (R&D Systems Inc.) according to the manufacturer’s procedure. Lactate concentrations in the conditioned media were also determined after overnight culture in hypoxic (1.5% O₂) and normoxic conditions using a colorimetric assay (Sigma-Aldrich, St. Louis, Missouri, USA) based on the enzymatic conversion of lactate to pyruvate and H₂O₂ by lactate oxidase.

Statistics. Data are presented as mean ± SD or mean ± SEM. Statistical differences were evaluated using the Student t test. In the limiting dilution analysis used to determine SRC frequencies, mice with at least 0.1% human cells were considered engrafted. The data from limiting dilution experiments were analyzed using the single-hit Poisson model, and SRC frequencies were determined using the maximum likelihood estimator as previously shown (12, 13, 16–20, 24). We used χ² analysis to verify the internal consistency of our data and validate the use of Poisson statistics.

Results

Preferential expansion of Lin−CD34+CD38− cells in hypoxia. Lin−CD34+ and Lin−CD34−CD38+ cells were cultured in serum-free conditions for 4 days under normoxia (20% O₂) or hypoxia (1.5% O₂). We used this level of O₂ based
on previous reports indicating that mouse BM-repopulating cells can be maintained when cultured under 1% O₂ (3, 4). As shown in Figure 1a, the level of expansion of Lin–CD34+ cells was decreased in hypoxia compared with normoxia. In contrast, the expansion of Lin–CD34–CD38– cells, a subpopulation enriched in primitive progenitors and stem cells, was greater in hypoxic conditions than in normoxia (2.4-fold vs. 1.5-fold, respectively). The differential effect of hypoxia on subsets of Lin–CD34+ cells was further investigated using CFSE to track the division history of each primitive subset (CD90+, HLA-DR–, CXCR4+, CD117low, CD133+, CD135+) during the 4-day culture (Figure 1b). Overall, CFSE profiles indicated that all examined subsets of Lin–CD34+ cells cultured in hypoxia divided at a slower rate than in normoxia. In particular, HLA-DR– cells divided only once or twice during the 4-day culture period. Despite this general slow proliferative rate, a small fraction (4–18%) of the cells in each subset was associated with a lower CFSE fluorescence, indicative of a rapid rate of division. These results show that a small subset of primitive BM CD34+ cells can rapidly proliferate under low O₂ levels while most CD34+ cells have a decreased rate of division. This suggests that O₂ levels can differentially regulate the proliferation of primitive subsets of human BM cells.

Hypoxia regulates the cell cycle of Lin–CD34+ cells. We evaluated the effect of hypoxia on the cell cycle of Lin–CD34+ cells after 4 days of culture in hypoxic or normoxic conditions. Before culture, less than 1% of sorted Lin–CD34+ cells was found in the S or G2/M fractions (data not shown). After 4 days under hypoxia, 34.2% ± 3.5% of CD34+ cells was found in S + G2/M phases compared with 46.3% ± 3.4% in normoxia (Table 1). A similar trend was observed for Lin–CD34– cells (data not shown). These results indicate that the slower rate of expansion of Lin–CD34+ cells under hypoxia is at least partially due to slower cell cycling and fewer cell divisions (as shown by DAPI and CFSE, respectively). We also examined the effect of hypoxia on the transition from G₀ to G₁ phase using Ki-67 to distinguish cells in G₀ phase (Ki-67–) from those in G₁ phase (Ki-67+). Table 1 shows that, in normoxic conditions, the G₀/G₁ ratio (% cells in G₀ to % cells in G₁) was 87 ± 12, indicating that most cells found in the G₀ + G₁ peak (as defined by DAPI staining) were resting. In hypoxic conditions, the G₀/G₁ ratio was only 13 ± 3 due to a 6.7-fold increase in cells in G₁ phase and fewer cells in G₀ phase. These results suggest that hypoxia preferentially promotes the transition of Lin–CD34+ cells from G₀ to G₁ and/or induces a cell cycle arrest in G₁ phase.

Effects of hypoxia on CFCs and LTC-ICs. We evaluated the clonogenic activity of Lin–CD34+ cells after 4 days of culture in hypoxic or normoxic conditions (Figure 2). After 4 days of culture, the total number of CFCs was increased by twofold compared with freshly isolated Lin–CD34+ cells (Figure 2a). The level of O₂ had no significant effect on lineage-committed progenitors, with the exception of granulocytic colonies (G-CFU), which decreased in number under hypoxic conditions. We also examined the effect of hypoxia on more primitive progenitors. LTC-ICs were equally expanded (1.4-fold) after 4 days of culture in hypoxia or normoxia (Figure 2b). In normoxia, only LTC-IC granulocyte-macrophage colonies significantly increased while, in hypoxia, both granulocyte-macrophage and erythroid colonies increased in number.
increased. This increase in BFU-E was the only hypoxia-specific effect on primitive progenitors we observed. These results indicate that a 4-day culture in hypoxia has a very limited effect on both committed and primitive progenitors.

**SCID-repopulating activity in cells cultured under hypoxic conditions.** First, we evaluated the effect of hypoxia on the SCID-repopulating ability of BM Lin–CD34+CD38– cells cultured for 4, 6, and 9 days in the presence of IL-3, IL-6, SCF, Flt-3 ligand, and G-CSF. In these conditions, SRCs could be maintained for only 4 days in culture, regardless of O2 levels. No human cells were detectable in the BM of mice (n = 16) transplanted with cells cultured for 6 or 9 days. These results indicate that in our conditions, hypoxia did not delay the loss of BM-repopulating activity observed after 4 days in normoxic cultures.

We used a limiting dilution analysis to quantitatively evaluate the effect of hypoxia on SCID-repopulating cells after 4 days. For this purpose, increasing numbers (ranging from 300 to 20,000) of fresh BM Lin–CD34+CD38– cells or cells cultured under hypoxia or normoxia for 4 days were transplanted at increasing doses into NOD/SCID mice (Figure 3). For each experimental condition (day 0, day 4 hypoxia, and day 4 normoxia), we estimated the number of Lin–CD34+CD38– cells required to statistically inject one SRC per mouse (12, 13, 20). Table 2 shows the frequencies of engrafted mice for each cell dose used in each experimental condition. We estimated that the frequency of SRCs in freshly purified Lin–CD34+CD38– cells (Table 2) was 1 in 1,009. After 4 days of culture, we determined that the frequency of SRCs in hypoxic conditions (1 in 577 Lin–CD34+CD38– cells) was significantly higher (P = 0.0017) than in normoxic cultures (1 SRC in 2,108 Lin–CD34+CD38– cells). Consequently, we observed a 3.6-fold increase in SRC frequency in hypoxic cultures compared with normoxia. In contrast, the apparent increase in SRC frequency between freshly isolated Lin–CD34+CD38– cells (1 SRC in 1,009 cells) and day 4 hypoxic cultures (1 in 577 cells) was not significant (P = 0.098). However, taking into account the expansion of Lin–CD34+CD38– CFCs after 4 days (Figure 1a), we estimate that in hypoxia, the absolute number of SRCs increased by 5.8-fold compared with the number in normoxia, and by 4.2-fold compared with freshly isolated Lin–CD34+CD38– cells. In contrast, in normoxic conditions, the absolute number of SRCs was just maintained in comparison with freshly isolated Lin–CD34+CD38– cells. In this study, we did not formally assess the effect of hypoxia on the self-renewal of human HSCs since no secondary transplants were performed. However, our results clearly demonstrate that low O2 levels can substantially increase the number of BM-repopulating cells compared with normoxic conditions.

**Molecular response of Lin–CD34+CD38– cells to hypoxia.** We examined whether human BM progenitors and stem cells express the critical components of the cellular response to hypoxia, such as HIF-1α and ARNT. Figure 4a shows that both HIF-1α and ARNT are constitutively expressed in purified Lin–CD34+CD38– cells. HIF-2α expression could not be detected in Lin–CD34+CD38– cells but was present at very low levels in Lin–CD34–CD38– cells, a subpopulation enriched in CD34– SRCs (data not shown). Since HIF-α subunits are regulated at the protein level,
we evaluated the effect of hypoxia and normoxia on HIF-1α protein. Figure 4b shows that HIF-1α protein can be detected in cells cultured in hypoxia but not under normal O2 levels. In contrast, ARNT protein levels were unaffected by O2 levels. These results demonstrate that HIF can mediate the hypoxic response of human BM progenitors and stem cells.

Hypoxia regulates expression of angiogenic factors in Lin–CD34+ cells. Angiogenic factors and receptors such as VEGF and Flt-1 are regulated by O2 levels (27). VEGF has been implicated in the regulation of HSC survival (40) and recruitment (41). To investigate whether the increase in SRC activity observed under hypoxia was associated with changes in the expression of angiogenic molecules, Lin CD34+ cells cultured for 4 days at 1.5% O2 or at normoxia were analyzed by flow cytometry for cell surface expression of VEGF receptor 1 (VEGFR1, also known as Flt-1), VEGFR2, Tie-1, Tie-2, and vWF. Levels of VEGF cell surface receptors (VEGFR1 and VEGFR2) increased slightly, whereas angiopoietin receptors (Tie-1 and Tie-2) were more clearly upregulated in cells cultured in hypoxic conditions (Figure 5a). In contrast, c-kit and Flt-3 ligand were unchanged, and vWF and MHC class II (HLA-DR) expression was downregulated under the same conditions (Figure 5a). It is unlikely that the upregulation of VEGFR1, VEGFR2, Tie-1, and Tie-2 was due to a nonspecific response to low O2 levels since other surface molecules were either unchanged or downregulated. These results suggest that human progenitors and stem cells can specifically enhance their responsiveness to angiogenic factors under hypoxic conditions.

Hypoxia-regulated genes include VEGF (42–44), erythropoietin (45), and glycolytic enzymes (46, 47) such as lactate dehydrogenase (48). To determine whether these HIF targets are upregulated in BM progenitors and stem cells, we cultured Lin–CD34+ cells for 18 hours under normoxic or hypoxic conditions and assayed VEGF, erythropoietin, and lactate concentrations in conditioned medium. Figure 5b shows that both VEGF and lactate secretions were increased by more than twofold, demonstrating that hypoxia upregulates VEGF production and glycolytic enzyme activity in human BM progenitors and stem cells. In contrast, erythropoietin was undetectable in the medium after 18 hours of culture regardless of O2 levels, suggesting that HIF targets may be differentially regulated in BM cells. Our results indicate that the hypoxic response of human progenitors and HSCs is characterized by a rapid increase in VEGF secretion and glycolytic activity. This rapid increase in VEGF secretion may play a critical role in the survival and expansion of human BM-repopulating cells under hypoxia.

To determine whether angiogenic cytokines play a role in the upregulation of angiogenic receptors observed under hypoxic conditions, Lin–CD34+ cells were cultured in normoxia in the presence of VEGF, angiopoietin-1, or angiopoietin-2 for 4 days. Interestingly, VEGFR2 surface expression was enhanced to levels comparable to those observed under hypoxia when cells were cultured in the presence of either VEGF,

<table>
<thead>
<tr>
<th>Table 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Frequency of NOD/SCID engraftment observed with increasing doses of fresh or cultured Lin–CD34+CD38+ cells after 4 days of culture in hypoxia or normoxia</td>
</tr>
<tr>
<td><strong>No. of NOD/SCID mice engrafted/transplanted</strong></td>
</tr>
<tr>
<td><strong>Day 0</strong></td>
</tr>
<tr>
<td><strong>No. injected cells</strong></td>
</tr>
<tr>
<td>300</td>
</tr>
<tr>
<td>600</td>
</tr>
<tr>
<td>1,000</td>
</tr>
<tr>
<td>1,800</td>
</tr>
<tr>
<td>2,500</td>
</tr>
<tr>
<td>≥5,000</td>
</tr>
</tbody>
</table>

NOD/SCID mice (n = 88) were injected with increasing doses of BM Lin CD34+CD38+ cells and analyzed 8–10 weeks after transplantation. Mice with BM containing at least 0.1% human myeloid (CD33+) and lymphoid (CD19+) cells were considered engrafted. *Significantly different from day 0 (P = 0.04). **Significantly different from day 4 normoxia (P = 0.0017).
In this report, we demonstrate for the first time that SRCs from human BM can be expanded in vitro under hypoxic conditions. We used BM Lin–CD34+CD38− cells, a population highly enriched in SRCs (12), and limiting dilution analysis to quantitatively evaluate the effect of hypoxia on SRCs. First, we determined that the frequency of SRCs in fresh adult BM is 1 in 1,009 Lin–CD34+CD38− cells. This result demonstrates that the frequency of SRCs in adult BM is only half of the SRC frequency that was reported previously for cord blood (1 in 617 Lin–CD34+CD38− cells) (12). This is consistent with other reports indicating that the frequency of SRC is higher in cord blood than in BM or mobilized peripheral blood (13, 14). To evaluate the effect of hypoxia on BM SRCs in vitro, we used culture conditions (IL-3, IL-6, SCF, Flt-3 ligand, and G-CSF in serum-free medium) capable of expanding SRCs in cord blood Lin–CD34+CD38− cells (16). Interestingly, after 4 days in the same culture conditions, SRCs in BM Lin–CD34+CD38− cells could not be expanded. The lack of SRC expansion after 4 days in culture under normoxic conditions suggests that human adult BM HSCs have different culture/cytokine requirements than cord blood HSCs. In contrast to normoxic conditions, we observed a significant increase in SRCs after 4 days in culture under 1.5% O2. Not only were SRCs increased relative to normoxia, but we also observed a 4.2-fold expansion of the number of SRCs in hypoxic cultures compared with freshly isolated BM Lin–CD34+CD38− cells. These results demonstrate that low levels of O2 can greatly enhance the survival and/or self-renewal of human BM-repopulating cells in vitro. This finding is consistent with a previous report indicating that mouse BM-repopulating cells were better maintained in culture under severe hypoxia than were more committed progenitors (3). We could no longer detect SRC activity after 6 or 9 days in culture under 1.5% or 3% (data not shown) or 20% O2. This finding indicates that, in the present culture conditions, which may not be optimal for BM stem cells, the positive effect of hypoxia on SRCs is short-lived.

The effect of hypoxia on clonogenic (CFC) and LTC-IC activity was limited to a reduction of granulocytic progenitors (G-CFU) and an increased number of LTC-ICs giving rise to BFU-E. The limited response of BM progenitors to hypoxia compared with the significant increase in SRCs observed in the same conditions demonstrates that the activity of human BM stem cells and progenitors is regulated differently by O2.

Discussion
In this report, we demonstrate for the first time that SRCs from human BM can be expanded in vitro under hypoxic conditions. We used BM Lin–CD34+CD38− cells, a population highly enriched in SRCs (12), and limiting dilution analysis to quantitatively evaluate the effect of hypoxia on SRCs. First, we determined that the frequency of SRCs in fresh adult BM is 1 in 1,009 Lin–CD34+CD38− cells. This result demonstrates that the frequency of SRCs in adult BM is only half of the SRC frequency that was reported previously for cord blood (1 in 617 Lin–CD34+CD38− cells) (12). This is consistent with other reports indicating that the frequency of SRC is higher in cord blood than in BM or mobilized peripheral blood (13, 14). To evaluate the effect of hypoxia on BM SRCs in vitro, we used culture conditions (IL-3, IL-6, SCF, Flt-3 ligand, and G-CSF in serum-free medium) capable of expanding SRCs in cord blood Lin–CD34+CD38− cells (16). Interestingly, after 4 days in the same culture conditions, SRCs in BM Lin–CD34+CD38− cells could not be expanded. The lack of SRC expansion after 4 days in culture under normoxic conditions suggests that human adult BM HSCs have different culture/cytokine requirements than cord blood HSCs. In contrast to normoxic conditions, we observed a significant increase in SRCs after 4 days in culture under 1.5% O2. Not only were SRCs increased relative to normoxia, but we also observed a 4.2-fold expansion of the number of SRCs in hypoxic cultures compared with freshly isolated BM Lin–CD34+CD38− cells. These results demonstrate that low levels of O2 can greatly enhance the survival and/or self-renewal of human BM-repopulating cells in vitro. This finding is consistent with a previous report indicating that mouse BM-repopulating cells were better maintained in culture under severe hypoxia than were more committed progenitors (3). We could no longer detect SRC activity after 6 or 9 days in culture under 1.5% or 3% (data not shown) or 20% O2. This finding indicates that, in the present culture conditions, which may not be optimal for BM stem cells, the positive effect of hypoxia on SRCs is short-lived.

The effect of hypoxia on clonogenic (CFC) and LTC-IC activity was limited to a reduction of granulocytic progenitors (G-CFU) and an increased number of LTC-ICs giving rise to BFU-E. The limited response of BM progenitors to hypoxia compared with the significant increase in SRCs observed in the same conditions demonstrates that the activity of human BM stem cells and progenitors is regulated differently by O2.

Figure 4
Expression of HIF in BM progenitors and stem cells. (a) RT-PCR analysis of ARNT and HIF-1α performed on RNA extracted from BM Lin CD34+CD38−, Lin CD34−CD38−, and K562 cells (as a positive control). (b) Western blot analysis of cell extracts from BM Lin CD34+ cells cultured overnight under hypoxic (H) or normoxic (N) conditions using antibodies against ARNT and HIF-1α. All gels are representative of at least three experiments.

Figure 5
Hypoxia regulates the expression of HIF target genes in BM progenitors and stem cells. (a) Representative histograms of cell surface expression of angiogenic and hematopoietic cytokine receptors in BM Lin CD34+ cells after 4 days in culture under normoxic (dashed lines) or hypoxic (1.5% O2, solid lines) conditions. (b) The conditioned medium from BM Lin CD34+ cells cultured overnight in normoxia or hypoxia (1.5% O2) was assayed for lactate and human VEGF. Bars represent the fold increase in lactate and VEGF concentrations measured in hypoxic cultures relative to normoxic conditions. Error bars show SD. (c) Effect of VEGF, angiopoietin-1 (Ang-1), and angiopoietin-2 (Ang-2) on cell surface expression of VEGFR2 in Lin−CD34+ cells after 4 days in culture under normoxic conditions (N).
To further characterize the hypoxic response, we evaluated the effects of low O2 tension on the proliferation and cell cycle distribution of human BM progenitors and stem cells in vitro. The expansion of Lin CD34+ cells was slightly reduced at 1.5% O2 compared with normoxia, indicating that most BM progenitors can survive and even proliferate under hypoxia. Interestingly, the expansion of purified Lin CD34+CD38- cells was greater in hypoxic than in normoxic conditions. This positive effect of hypoxia on Lin CD34+CD38- cells demonstrates that primitive progenitors and stem cells are particularly well adapted to proliferating in a low-O2 environment.

To further evaluate the effect of hypoxia on the proliferation of BM cells, we used CFSE to track the division history of primitive subsets (CD90+, CXCR4+, CD117low, CD135+, and HLA-DR-) of BM Lin CD34+ cells cultured under hypoxia and normoxia. We show that under hypoxic conditions, the overall proliferative rates of Lin CD34+ cells and subsets was slower in hypoxic conditions, resulting in fewer cells after 4 days in culture compared with normoxia. Hypoxia increased the percentage of the cells in G0 or G1 phase, while fewer cells were found in S or G2/M. Furthermore, hypoxic conditions increased the proportion of cells in G1 phase relative to resting (G0) cells, resulting in a dramatically decreased G0 to G1 ratio. This observation suggests that low O2 levels can either promote the transition from a resting state to G1 phase and/or inhibit the transition from G1 to S phase in BM progenitors and stem cells. It has been shown that hypoxia can induce G1 arrest in tumor (49, 50) and endothelial cells (51). Gardner et al. (52) reported that, in fibroblasts, severe hypoxia can inhibit the G1/S transition through regulation of p27 expression. Wilpshaar et al. (53) found no difference in SCID-repopulating ability between uncultured cord blood CD34+ cells in G0 and those in G1 phase. However, in cultured mobilized peripheral blood CD34+ cells, the level of human chimerism in NOD/SCID BM was lower in mice transplanted with cells in G1 phase compared with G0 (54). The expansion of SRCs we observed after 4 days in hypoxia and the increased number of cells in G1 phase suggests that SRCs may be found in both G0 and G1 phases under low oxygen levels.

In this study, we also investigated the cellular mechanisms involved in the hypoxic response of BM hematopoietic progenitor and stem cells. We found that HIF-1α and ARNT are constitutively expressed in Lin CD34+CD38- cells. However, HIF-2α transcripts could not be detected in Lin CD34+CD38- cells. Our Western blot analysis of Lin CD34+ cells shows that HIF-1α protein is present under hypoxic conditions and is undetectable in normoxia. In contrast, ARNT protein is detectable regardless of O2 levels. This is consistent with previous reports showing that, in normoxic conditions, HIF-1α is modified by a prolyl hydroxylase, an O2-dependent enzyme (55, 56). The modified form of HIF-1α can interact with the von Hippel-Lindau tumor suppressor protein (VHL). This interaction induces the ubiquitination of HIF-1α and its rapid degradation (57–60). Under hypoxic conditions, the hydroxylation no longer occurs, and HIF-1α remains stable and can upregulate expression of its target genes.

Among the genes induced by hypoxia (25–29), angiogenic factors and receptors appear to play a critical role in the regulation of HSC survival and self-renewal. In this report we show an increase in VEGF secretion and an upregulation of the cell surface expression of angiogenic receptors (Tie-2 and to a lesser extent VEGFR2 and Tie-1) under hypoxia. The hypoxia-induced increase in VEGF expression has been well characterized in other models (42–44). Recently, Gerber et al. (40) reported that VEGF regulates the survival of mouse HSCs through an internal autocrine loop. It has also been shown that human BM-repopulating cells express VEGF receptors (61, 62). It is thus tempting to speculate that a similar mechanism may be involved in regulating the survival of SRCs. Tie-2, the receptor for angiopoietin-1, has also been reported to be regulated by O2 levels (63) and can be found on HSCs (64). Thus, the combined effect of increased VEGF and enhanced responsiveness to angiogenic growth factors may play a role in the expansion of human BM-repopulating cells we observed. Future studies will be needed to further investigate the role of VEGF in the expansion of SRCs under hypoxia.

In conclusion, we show that the use of low O2 levels in vitro not only improves SRC survival compared with normoxia but also expands the number of human BM-repopulating cells by 4.2-fold compared with freshly isolated Lin CD34+CD38- cells. The present report also establishes that hypoxia increases both VEGF secretions and responsiveness to angiogenic factors in BM progenitors and stem cells. In addition, we show that stem cells and progenitors respond differently to hypoxic conditions. These findings suggest that the balance between the survival, self-renewal, and differentiation of human BM HSCs might be tightly regulated by O2 levels. The beneficial effect of hypoxia on human BM-repopulating cells may be mediated by angiogenic factors such as VEGF. Finally, the use of low O2 tension could represent an important advance in the ex vivo expansion of human HSCs and may have important translational and clinical implications.

Acknowledgments

We are grateful to Jennifer Sinibaldi, Matthew Lessie, Amy Cleck, and Martin Carroll from the Stem Cell and Leukemia Core at the University of Pennsylvania Cancer Center for providing the large number of normal BM samples necessary for this study. The authors also thank William Demuth at the Abramson Family Cancer Research Center (University of Pennsylvania) for cell sorting, Tammi Coleman and Derrick Dow for their help with the NOD/SCID mouse colony, and Diana
cells increases the SCID-repopulating capacity of adult human bone marrow. Blood. 100:4433–4439.

Ramirez-Bergeron, Andy Arsham, Brian Keith, and other members of the Simon Laboratory for their valuable input. M.C. Simon is an Investigator of the Howard Hughes Medical Institute.