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# Regulation of Hematopoietic Stem Cells by Bone Marrow Stromal Cells

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# Abstract

Hematopoietic stem cells (HSCs) reside in specialized microenvironments (niches) in the bone marrow. The stem cell niche is thought to provide signals that support key HSC properties, including self-renewal capacity and long-term multilineage repopulation ability. The stromal cells that comprise the stem cell niche and the signals that they generate that support HSC function are the subjects of intense investigation. Here we review the complex and diverse stromal cell populations that reside in the bone marrow and examine their contribution to HSC maintenance. We highlight recent data suggesting that perivascular CXCL12-expressing mesenchymal progenitors and endothelial cells are key cellular components of the stem cell niche in the bone marrow.

# Keywords

hematopoietic stem cells; HSCs; stem cell niche; mesenchymal stem cells; osteoblast; CXCL12

# The hematopoietic stem cell niche

Hematopoiesis is the process by which all mature blood cells are produced. It must balance enormous production needs (more than 500 billion blood cells are produced every day) with the need to precisely regulate the number of each blood cell type in the circulation. In vertebrates, the vast majority of hematopoiesis occurs in the bone marrow and is derived from a limited number of hematopoietic stem cells (HSCs) that are multipotent and capable of extensive self-renewal. In mammals, it is estimated that there are approximately 10,000 HSCs, of which, in humans, approximately 1,000 contribute hematopoiesis at any given time [1]. Prospective identification of HSCs using cell surface markers and flow cytometry is best described for murine HSCs. c-Kit<sup>+</sup> Sca-1<sup>+</sup> lineage<sup>-</sup> (KSL) CD150<sup>+</sup> CD48<sup>-</sup> cells [2] and CD34<sup>-</sup> Flk2<sup>-</sup> KSL cells [3] represent the two most commonly used murine HSCs phenotypes, each containing approximately 50% HSCs based on long-term repopulating assays.

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Key properties of HSCs are multipotency, self-renewal capacity, and quiescence. The bone marrow microenvironment appears to be uniquely adapted to support these, and other, HSC properties. As first proposed by Schofield in 1978 [4], the concept of a "stem cell niche" in the bone marrow has gained widespread popularity. In this model, specialized niche cells in the bone marrow are physically associated with HSCs and provide specific signals that help maintain their function. In this review, we highlight recent studies defining the bone marrow stromal cells that comprise the stem cell niche and the signals that they generate that contribute to HSC maintenance.

# Anatomy of the Bone Marrow

The bone marrow is the major site of hematopoiesis and bone formation in most vertebrates. Thus, in addition to containing hematopoietic cells, the bone marrow contains cells that contribute to bone homeostasis, including mesenchymal stem cells (also called skeletal stem cells), osteoprogenitors, osteoblasts, osteocytes, and chondrocytes (Figure 1). To add further complexity, other stromal cell populations that reside in the bone marrow may regulate hematopoiesis, including neuronal cells, glial cells, and adipocytes. Recent advances in imaging technologies have greatly advanced our understanding of the organization of the bone marrow. It is clear that the bone marrow in both long bones (e.g., femur) and flat bones (e.g., calvaria) is highly vascular [5, 6]. In long bones, central longitudinal arteries give rise to radial arteries that in turn branch into arterioles near the endosteum (Figure 2) [5]. The transition from Sca-1<sup>+</sup> arterioles to Sca-1<sup>-</sup> venous endothelium occurs in close proximity to the endosteum. Venous sinusoids extend back towards the central cavity where they coalesce into a large central sinus. Of note, the venous sinusoids, which contain numerous fenestra, are thought to be the major sites of leukocyte egress from the bone marrow into the circulation [7]. Osteoblasts and bone lining cells form a layer between mineralized bone and the bone marrow. Spindle-shaped N-cadherin<sup>+</sup> osteoblasts (SNO cells), which may represent an immature osteoblast, also are localized near the endosteum [8]. There is a rich network of stromal cells interspersed between islands of hematopoietic cells and include mesenchymal stem cells, CXCL12-abundant reticular cells, and adipocytes.

The location of HSCs in the bone marrow is controversial. Initial studies using labeled HSCenriched cell populations transplanted into recipients suggested a mostly endosteal location for HSCs [9–11]. This is consistent with data showing that long-term BrdU retaining cells (presumed to be HSCs) are preferentially localized next to SNO cells at the endosteum [8]. In contrast, using a rigorous definition of HSCs (CD150<sup>+</sup> CD48<sup>-</sup> CD41<sup>-</sup> lineage<sup>-</sup> cells), Kiel et al showed that the majority of HSCs are in contact with sinusoidal endothelium at bone-distant sites [2]. This is consistent with studies showing that most HSCs are in direct contact with perivascular CXCL12-abundant reticular (CAR) cells [12] and nestin-GFP<sup>+</sup> stromal cells [13]. Interestingly, a recent study suggested that both HSCs and lineagecommitted hematopoietic progenitor cells (HPCs) were localized near endothelium with a preference for the endosteal region [5]. Collectively, these data suggest that the majority of HCS are perivascular and enriched in the highly vascular endosteal region.

# Hypoxia and the stem cell niche

The prevailing view is that HSCs are maintained in a hypoxic niche in the bone marrow. As noted, HSCs are localized to the endosteal region, which previous studies characterized as having low perfusion and relative hypoxia [14–16]. Indeed, HSCs with lower cellular levels of reactive oxygen species have higher self-renewal potential [17]. Moreover, disruption of hypoxia-inducible factor-1 alpha (HIF-1 alpha) results in loss of HSC quiescence and repopulating activity [18], while stabilization of HIF-1 alpha induces HSC quiescence and enhances repopulating activity [18, 19]. However, as noted in the previous section, the

endosteal region is actually highly vascularized and most HSCs are perivascular [5]; thus, HSCs are likely to be relatively well oxygenated. Interestingly, hematopoietic stem/ progenitor cells (HSPCs) display a hypoxic profile (defined by strong retention of pimonidazole and expression of HIF-1 $\alpha$ ) regardless of their location in the bone marrow [5]. Indeed, even HSPCs in the peripheral circulation display a hypoxic profile. Thus, intrinsic differences in cellular metabolism rather than localization to a hypoxic microenvironment may define the hypoxic profile of HSCs.

# Stromal cells in the Bone Marrow

The bone marrow contains several stromal cell populations implicated in the regulation of hematopoiesis. There is emerging data that specific stromal cell populations regulate distinct hematopoietic progenitor populations; both positive and negative regulators of hematopoiesis have been identified. In the following sections, we review studies that focus on the contribution of specific stromal cell populations in the regulation of hematopoiesis. However, it is important to keep in mind, that it is the sum of the complex signals generated by these different stromal cell populations that are sensed by HSCs and other hematopoietic progenitor cell populations in the bone marrow.

#### Osteolineage cells

The localization of HSCs to the endosteum supported studies of the role of osteoblast lineage cells (which we will refer to as osteolineage cells) in the stem cell niche. Osteolineage cells produce a number of cytokines implicated in HSC regulation, including G-CSF [20], thrombopoietin [21, 22], and CXCL12 [23]. Expansion of osteolineage cells through enforced expression of parathyroid hormone receptor 1 [24] or through conditional inactivation of bone morphogenic protein receptor-1 [8] is associated with an increase in HSCs. Conversely, conditional ablation of osteolineage cells is associated with a loss of HSCs in the bone marrow and extramedullary hematopoiesis [25].

Despite these findings, the contribution of osteolineage cells, particularly mature osteoblasts, to HSC maintenance is controversial. A recent study suggests that parathyroid hormone treatment increases short-term HSCs, not through osteoblast expansion, but through Wnt ligand production by T cells [26]. Moreover, an increase in osteoblasts is not sufficient to expand HSCs. Treatment of mice with the bone anabolic agent, strontium, leads to mature osteoblast expansion but has no effect on HSC number or function [27]. Conversely, in mice with chronic inflammatory arthritis, resulting in osteoblast suppression, HSCs are normal [28]. Finally, as discussed in more detail in a later section, conditional deletion of *Cxcl12* [29, 30] or stem cell factor (SCF, *Kitl*) [31] from mature osteoblasts has no effect on HSCs. These apparently discrepant data may be due, at least in part, the heterogeneity of osteolineage cells. In particular, a recent study showed that more primitive osteolineage cells express higher levels of SCF and CXCL12 and support the long-term repopulating activity of HSCs better than more differentiated osteolineage cells [32]. Thus, while mature osteoblasts appear not to play a major role in vivo, immature osteolineage cells may contribute to HSC maintenance, a concept we will explore in a later section.

#### The role of N-cadherin in the stem cell niche

The contribution of N-cadherin to HSC maintenance remains unclear. Cadherins are calcium-dependent homotypic adhesion molecules that form adherens junctions. N-cadherin is expressed on a subset of osteolineage cells, with higher expression observed on immature osteolineage cells [32]. As discussed in the previous section, at least a subset of HSCs are localized near SNO cells [8, 10, 11], and N-cadherin has been implicated in the adhesion of HSCs to osteolineage cells [32]. Inhibition of N-cadherin signaling through knock-down of

N-cadherin or expression of a dominant-negative N-cadherin in HSCs inhibits their repopulating activity [33, 34]. On the other hand, conditional deletion of *Cdh2* (encoding for N-cadherin) in HSCs has no effect on HSC number or function [35]. Moreover, conditional deletion of *Cdh2* in osteolineage cells has no effect on HSC number, trafficking, cell cycle status, or repopulating activity [36, 37]. Thus, the preponderance of evidence, suggests that N-cadherin is not required for normal HSC function. It is also important to note that these results do not discount a role for SNO cells in the regulation of HSCs. SNO cells are suggested to be immature osteoblasts, and N-cadherin may simply mark an earlier developmental stage of osteoblasts important for niche maintenance.

#### Perivascular CXCL12-expressing stromal cells

The perivascular location of most HSCs has focused recent attention on the stromal cells that reside in the perivascular region as candidate niche cells. Besides endothelial cells, the perivascular region contains mesenchymal stem cells and a heterogeneous population of stromal cells characterized by very high CXCL12 expression. CXCL12 (stromal-derived factor-1, SDF-1) is a chemokine that plays a crucial role in maintaining HSC function. Three perivascular stromal cell populations that express high levels of CXCL12 have been identified: CXCL12-abundant reticular (CAR) cells, nestin-GFP<sup>+</sup> stromal cells, and leptin receptor<sup>+</sup> stromal cells. These stromal cell populations are defined by transgene expression using defined stromal-specific promoters, and it likely that there is considerable overlap.

CAR cells were identified using mice with GFP knocked into the *Cxcl12* locus as perivascular stromal cells with very high GFP expression [12, 38]. CAR cells are mesenchymal progenitors that have both adipogenic and osteogenic potential in vitro [39]. HSPCs and certain lymphoid progenitors directly contact CAR cells in the bone marrow [12, 38]. Conditional ablation of CAR cells using transgenic mice expressing the diphtheria toxin receptor (DTR) under control of *Cxcl12* regulatory elements (*Cxcl12-Dtr* mice) leads to a reduction in HSCs and HSC long-term repopulating activity but increased HSC quiescence [39]. CAR cells are the major source of SCF and CXCL12 in the bone marrow, and conditional ablation is associated with a marked loss of bone marrow SCF and CXCL12. Of note, although no obvious toxicity was observed in endothelial cells or osteoblasts, these cells express CXCL12, and it is possible that their function was altered after conditional ablation in *Cxcl12-Dtr* mice.

Nestin-GFP<sup>+</sup> cells are defined as perivascular stromal cells that express high levels of GFP under control of the nestin (*Nes*) promoter [13]. Nestin-GFP<sup>+</sup> stromal cells are enriched for mesenchymal stem cell activity with approximately 1% of cells having colony-forming units-fibroblast (CFU-F) activity. Nestin-GFP<sup>+</sup> cells express several genes implicated in HSC maintenance, including CXCL12, SCF, and angiopoietin. Transplanted HSCs preferentially home near nestin-GFP<sup>+</sup> cells. Finally, conditional ablation of nestin-expressing stromal cells results in a modest loss of HSCs.

Lineage mapping using Cre-recombinase expressed under the control of leptin receptor (*Lepr*) regulatory elements identified a perivascular stromal cell population in the bone marrow [31]. Similar to CAR cells, leptin receptor<sup>+</sup> stromal cells express high levels of CXCL12 and stem cell factor, suggesting that CAR cells and leptin-receptor<sup>+</sup> cells identify largely overlapping stromal cell populations. Indeed, we observed high level leptin receptor expression in sorted CAR cells [30]. This also is consistent with data showing that CAR cells have adipogenic (as well as osteogenic) capacity in vitro [39]. However, lineage mapping studies using *Lepr:Cre* showed no targeting of osteoblasts [31], raising the possibility that a subpopulation of leptin-receptor-negative CAR cells with osteogenic capacity exists. In any case, deletion of stem cell factor (*Kitl*) from leptin-receptor<sup>+</sup> stromal

#### Mesenchymal stem cells

A rigorous definition of MSCs is a cell with self-renewal capacity that is able to generate all mesenchymal cells within a skeletal element, including osteoblasts, chondrocytes, adipocytes, and fibroblasts [40]. Although there is currently no prospective way to identify MSCs on a per cell basis, several groups have reported methods to identify MSC-enriched cell populations. In mice, approximately 4% of CD45<sup>-</sup> lineage<sup>-</sup> PDGFRa<sup>+</sup> Sca<sup>+</sup> (PaS) cells have CFU-F activity [41] and approximately 1% of nestin-GFP+ stromal cells have CFU-F activity [13]. We recently showed that the Prx1-targeted subset of PaS is further enriched for MSC activity, with greater than 10% of these cells having CFU-F activity [30]. Surprisingly, we observed that neither *Prx1-Cre* targeted PaS cells nor CAR cells express nestin [30]. One potential explanation for the disparate results is that the nestin-GFP transgene results in aberrant expression of GFP that does not accurately reflect nestin expression. We suggest that nestin-GFP<sup>+</sup> expression identifies a heterogeneous stromal cell population that includes MSCs and CAR cells.

In human bone marrow, CD146-expressing stromal cells identifies an MSC-enriched cell population [42]. Recently, Pinho and colleagues showed that PDGFR $\alpha$  and CD51 expression define a bone marrow stromal cell population in both mice and humans that is highly enriched for MSCs and can support HSPC expansion in vitro [43].

#### Endothelial cells, adipocytes, neuronal, and glial cells

Hemogenic endothelium in the dorsal aorta gives rise to the first definitive HSCs during embryonic development [44, 45]. Bone marrow endothelial cells express several genes implicated in HSC maintenance, including CXCL12, SCF, and angiopoietin, and they support the proliferation of HSPCs in vitro [46]. Regeneration of sinusoidal endothelial cells is required for hematopoietic recovery from myeloablation [47, 48]. Moreover, deletion of the endothelial specific adhesion molecule, E-selectin, results in increased HSC quiescence, suggesting that endothelial cells regulate HSC proliferation [49]. Finally, endothelial cells contribute to HSC maintenance through production of SCF, as deletion of *Kitl* specifically from endothelial cells results in the loss of HSCs [31]. Collectively, these data show that endothelial cells are a key component of the stem cell niche.

Adipocyte number in the bone marrow increases with age (especially in humans). Naveiras et al. showed that the hematopoietic activity of distinct regions of the mouse skeleton correlated inversely with adiposity [50]. That is, adipocyte-rich bone marrow, such as the vertebrae of mice, has a decreased number of HSCs compared with adipocyte-poor bone marrow. In mice with impaired adipogenesis, hematopoietic activity is increased in bone marrow sites that are normally adipocyte-rich. Of potential clinical relevance, pharmacological inhibition of adipogenesis results in enhanced hematopoietic recovery following stem cell transplantation in mice [50]. These data suggest that adipocytes play an inhibitory role in HSC maintenance. Elegant studies by the Frenette group show that the sympathetic nervous system coordinate the circadian egress of HSCs into the circulation by regulating local production of CXCL12. Genetic or pharmacological ablation of adrenergic signaling inhibits G-CSF-induced HSPC mobilization [52].

A recent study also implicates glial cells, specifically nonmyelinating Schwann cells, in the regulation of HSCs [53]. Transforming growth factor  $\beta$  (TGF- $\beta$ ) is an important regulator of HSC function. It induces HSC quiescence in vitro [54] and loss of TGF- $\beta$  signaling in HSCs

results in impaired long-term repopulating activity [53]. While the latent form of TGF- $\beta$  is produced by many cell types, Yamazaki and colleagues provide data suggesting that nonmyelinating Schwann cells are the major source of active TGF- $\beta$  in the bone marrow. Importantly, surgical disruption of sympathetic nerves resulting in loss of Schwann cells is associated with decreased active TGF- $\beta$  expression and the loss of HSCs.

Distinct contributions of Cxcl12-producing niche cells to HSC maintenance CXCL12 plays a crucial role in maintaining HSC function, including retention in the bone marrow [55–58], quiescence [59, 60], and repopulating activity [60]. To test the hypothesis that CXCL12 production from different stromal cell populations would have distinct effects on HSCs, the Morrison group and our group independently deleted *Cxcl12* from candidate niche cells [29, 30]. Loss of CXCL12 from mature osteoblasts had no effect on HSC number or function. Loss of CXCL12 from CAR cells using osterix (Osx)-Cre results in HSPC mobilization, and a modest (50%) reduction in HSCs [30]. Deletion of Cxcl12 from leptinreceptor<sup>+</sup> stromal cells also results in constitutive HSPC mobilization, but no defect in HSC function was observed [31]. These observations show that CXCL12 expression from CAR cells or leptinreceptor<sup>+</sup> stromal cells is essential for efficient retention of HSPCs in the bone marrow. Both groups showed that deletion of Cxcl12 using Prx1-Cre resulted in a significant loss of HSCs, long-term repopulating activity, and HSC quiescence. Like Osx-Cre, Prx1-Cre targets CAR cells and osteoblasts but also targets PaS mesenchymal progenitors [30]. A modest decrease in long-term repopulating activity (but not HSC quiescence) also was observed in mice with *Cxcl12* deleted from endothelial cells. Together, these data suggest that CXCL12 expression from perivascular stromal cells is required for HSC maintenance. Whereas CAR cells and endothelial cells are minor contributors, mesenchymal progenitors are the major source of CXCL12 that supports HSCs.

# **Concluding remarks**

A model of the murine stem cell niche incorporating recent findings is shown in Figure 3. Current evidence supports a perivascular location for most HSCs, with a preference for the endosteal region. Key components of the perivascular niche include endothelial cells, mesenchymal progenitors, and CXCL12-expressing stromal cells (including CAR cells, leptin receptor<sup>+</sup> cells, and Nestin-GFP<sup>+</sup> cells). These perivascular stromal cells provide signals, including CXCL12, SCF, and angiopoietin that maintain HSC quiescence and self-renewal capacity. The sympathetic nervous system and associated glial cells and adipocytes also contribute to HSC maintenance. There are several important unresolved issues in field (Box 1). A better understanding of the interaction of HSCs with stromal cells in the bone marrow may provide potential therapeutic targets for manipulation to modulate HSC and their progeny. Moreover, since the stem cell niche may provide signals that support certain malignancies [61, 62], drugs that target the niche hold promise as a way to sensitize cancer cells to chemotherapy.

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## Outstanding areas for future study

- The relationship between CAR cells, nestin-GFP<sup>+</sup> stromal cells, and leptin receptor<sup>+</sup> stromal cells is not clear. Although they likely overlap considerably, it will be important to define precisely what cell types are included in each cell population.
- The genes expressed by niche cells that contribute to HSC maintenance have not been fully defined.
- The signals that regulate stromal cells in the niche are poorly understood. For example, the pro-inflammatory cytokine G-CSF induces HSC mobilization, at least in part, by disrupting the stem cell niche. Of particular interest are the signals that determine osteoblastic versus adipogenic differentiation of mesenchymal progenitors. Strategies that suppress adipogenic differentiation might reverse the age-associated loss of hematopoietic activity.



Figure 1. Distinct stromal cell populations in the bone marrow contribute to HSC maintenance A complex and diverse group of stromal cells in the bone marrow have been implicated in HSC maintenance. Endothelial cells, mesenchymal stem cells (MSCs), and CXCL12expressing mesenchymal progenitors (CEMP cells) are perivascular stromal cells that produce a number of factors that support HSCs, including CXCL12, angiopoietin, and stem cell factor (SCF). CEMP cells have been identified as CXCL12-abundant reticular (CAR) cells, leptin receptor<sup>+</sup> stromal (Lepr<sup>+</sup>) cells, and Nestin-GFP+ cells; these stromal cell populations likely overlap considerably. Osteoblasts and spindle-shaped N-cadherin<sup>+</sup> osteoblast (SNO cells) produce a number of factors that support HSCs, including thrombopoietin (TPO) and CXCL12. Sympathetic neurons indirectly regulate HSCs by targeting CXCL12 expression. Finally, glial cells, through production of active transforming growth factor- $\alpha$  TGF- $\alpha$  and adipocytes regulate HSCs.



#### Figure 2. The bone marrow is highly vascularized

Confocal microscopic images of the femoral diaphysis stained with the pan-vasculature marker laminin (green) and the arterial specific marker, Sca-1 (red). A Sca-1<sup>+</sup> central artery runs through the central marrow. The central artery branches off to smaller arterioles towards the endosteum. Reprinted by permission from Macmillan Publishers Ltd: Nature Cell Biology; 15 (5):533-43, copyright (2013).



#### Figure 3. Model of murine HSC niche

Conditional deletion of two key HSC maintenance genes, *Kitl* (stem cell factor, SCF) and *Cxcl12*, in candidate niche cells has emphasized the importance of stromal cells in the perivascular region. SCF production from endothelial cells and leptin receptor<sup>+</sup> stromal (Lepr<sup>+</sup>) cells but not osteoblasts (Ob) is required for HSC maintenance. CXCL12 production from mesenchymal stem cells (and to a lesser extent endothelial cells) is required for HSC maintenance, while CXCL12 production form CXCL12 abundant reticular (CAR) cells or Lepr<sup>+</sup> cells is required for efficient retention of hematopoietic progenitors in the bone marrow. Nestin-GFP<sup>+</sup> stromal cells likely overlap with CAR cells and Lepr<sup>+</sup> cells.