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Enigmatic niche brings challenge for HSC expansion

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The holy grail of hematopoietic stem cell (HSC) biology has been to harness self-renewal to expand stem cells for greater availability of transplantation procedures in people with cancer or hematologic diseases. Despite the advances in transplantation biology, many individuals do not have a suitable allogeneic donor. In these cases, unrelated cord blood-derived HSC has shown promise as a source of cells for transplantation, but the number of HSC recovered in a single cord is generally too low for adult transplantation, and infusion of two separate cords still led to delayed hematopoietic recovery¹. Thus, the development of methods to readily expand HSC *in vitro* would be a major leap forward.

In-depth understanding of the identities and function of each component of the hematopoietic microenvironment at the cellular and molecular levels will be crucial to learn how to properly expand HSCs. This knowledge gap has closed substantially after the concept of a stem cell niche was enunciated decades ago². It has been assumed by many, perhaps by wishful thinking, that a 'do-it-all' niche cell exists in the mammalian bone marrow. This idea comes from the original concept by Schofield², and also from strong experimental work in the germline of invertebrates, where specific niche cells are thought to provide the necessary 'goods' to maintain germline stem cells³. In the mammalian bone marrow, candidate niche cells including osteoblasts⁴, Cxcl12-abundant reticular cells⁵ and Nestin-positive mesenchymal stem cells⁶ have been reported, suggesting a key role for skeletal precursor cells. Two recent studies have added further complexity to the cellular make-up of putative HSC niches in bone marrow^{7,8} (Fig. 1).

Yamazaki *et al.*⁷ were searching for microenvironments enabling HSC dormancy in the bone marrow. Cell cycle quiescence is a hallmark of somatic stem cells and a key behavior to protect them from hematopoietic stresses and exhaustion⁹. Transforming growth factor- β (TGF- β) is a powerful molecule capable of inducing HSC quiescence through inhibition of lipid raft clustering that assembles growth factor signaling microdomains¹⁰. As it is secreted as an inactive form, active TGF- β is generated locally through local protease activity. If TGF- β was crucial in promoting HSC quiescence, we would expect that the putative niche would provide a specific environment for its activation. Yamazaki *et al.*⁷ have found that glial fibrillary acidic protein (GFAP)-positive Schwann cells (that ensheath sympathetic nerve fibers) express β 8 integrin, which promotes metalloproteinase-mediated TGF- β activation. Afferent sympathetic nerves form a complex with Nestin-positive cells within the

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bone marrow, in which the nerve itself may regulate HSC quiescence (Fig. 1). Unexpectedly, a reduction of HSC frequency in the bone marrow occurred when mice were surgically sympathectomized by ligation of the sympathetic trunk. The authors have suggested that the loss of Schwann cells led to HSC differentiation. Other less invasive, but also less specific, methods of sympathectomy (such as femoral and sciatic neurectomy), however, have not led to such reductions in HSC activity¹¹, suggesting the presence of other signals that may collaborate with those provided by the Schwann cells of sympathetic nerves. Although the latter issue still needs to be clarified, the findings support the crucial function of local **sympathetic** nerves in the regulation of hematopoiesis.

Another study in mice evaluated the cellular source of stem cell factor (SCF), also called kit ligand, a growth factor required for HSC maintenance in a non-cell autonomous manner¹⁰. SCF exists in two forms, membrane-bound and soluble, due to differential splicing and proteolytic cleavage; however, the membrane-bound form appears to play a prominent role in HSC maintenance ¹². When expressed under the Scf locus, green fluorescence protein (GFP) expression was mostly localized in cells associated with sinusoids, the fenestrated blood vessels within the bone marrow⁷. The authors made tissue-restricted deletions of Scfor knock-in expression of reporter genes including GFP in osteoblasts, perivascular cells expressing Nestin or Leptin receptor, and endothelial cells. A reduction of HSC frequency was only found in Lepr-expressing perivascular cells and endothelial cells lacking Scf, suggesting that SCF was provided by both cell types, although the contribution from the former type may be much greater. This study proposed that Lepr-expressing and Nestinexpressing cells are distinct as the deletion of Scf using transgenic lines producing Cremediated deletion under the Nestin promoter did not elicit the same phenotype. It is important to note, however, that Lepr was among the top 1% most highly expressed gene in sorted Nestin-positive cells by microarray analysis⁶, suggesting a potential overlap between the two niche cells, which will require further investigation. Thus, cells associated with the vasculature are the major source of SCF in the bone marrow under steady state. It will be important to evaluate whether the same stromal cells produce the cytokine in situation of regenerative stress, which may shed light on the requirements for building an artificial niche dedicated for HSC expansion.

These two studies bring new exciting angles that add further complexity to bone marrow niche(s); at the same time, they soberly remind us about the significant hurdles ahead of us to provide expanded HSC for clinical use. HSCs require an array of secreted and contact factors—most of which still remain unknown—for their maintenance and proliferation. If these factors are indeed provided by distinct stromal cell types, the challenge to concoct the right ratios of niche constituents to support HSC for clinical use becomes very significant. By the same token, one can imagine the translational difficulty to obtain a renewable source of good manufacturing practice (GMP) grade cellular cocktail that would meet the safety thresholds for clinical cell therapy. Further studies will aim at characterizing stromal cell types and, most important, the molecular constituents and signals that allow HSC proliferation while maintaining self-renewal.

A major obstacle for future translational HSC-based therapeutics, highlighted by the study of Yamazaki *et al.*⁷, concerns the propensity of adult HSC to remain quiescent. This natural

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preference was repeatedly shown in several studies where genetic mutations leading to HSC proliferation invariably produce HSC exhaustion⁹. How can we unlock HSC from their quiescent behavior for the purpose of expansion without reaching exhaustion? It is likely that the coupling of self-renewal and quiescence is not absolute given that it is not the case for all stem cells; for example, embryonic stem cells can both proliferate and self-renew indefinitely. The ultimate solution, through a greater understanding of the molecular basis of the niche, may lie in some sort of direct or indirect 'reprogramming' of adult (quiescent) HSCs into fetal-like (proliferative) stem cells, a feat that will have to be achieved without increasing the risk of malignancies. Every scientific hurdle, almost by definition, can be overcome. We are just one breakthrough away from surmounting this one.

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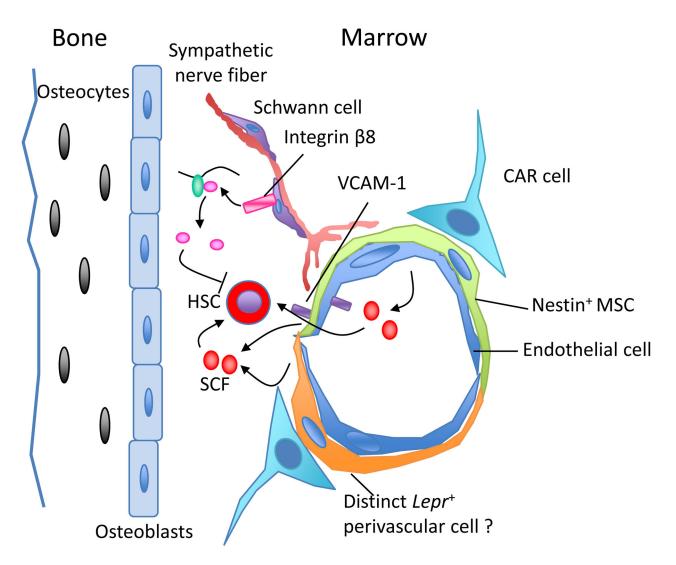


Figure 1.

The complex cellular and molecular make up of the HSC niche in the bone marrow. A variety of cells, including osteoblasts, Cxcl12-abundant reticular (CAR) cells, Nestinpositive mesenchymal stem cells (MSC), *Lepr*-expressing perivascular cells and endothelial cells have been shown as possible components of the niche. Cells from the neural system, nonmyelinating Schwann cells wrapping sympathetic nerve fibers, promote HSC quiescence through activation of TGF- β . The structural composition of these cells provides a specialized microenvironment that regulates HSC self-renewal and differentiation, either though contact-dependent signals such as vascular cell adhesion molecule-1 (VCAM-1)⁶, or via soluble factors such as SCF.