Homeostatic and pathogenic extramedullary hematopoiesis

Chang H Kim
Laboratory of Immunology and Hematopoiesis, Department of Comparative Pathobiology; Purdue Cancer Center, Purdue University, West Lafayette, IN, USA

Abstract: Extramedullary hematopoiesis (EH) is defined as hematopoiesis occurring in organs outside of the bone marrow; it occurs in diverse conditions, including fetal development, normal immune responses, and pathological circumstances. During fetal development, before formation of mature marrow, EH occurs in the yolk sac, fetal liver, and spleen. EH also occurs during active immune responses to pathogens. Most frequently, this response occurs in the spleen and liver for the production of antigen-presenting cells and phagocytes. EH also occurs when the marrow becomes inhabitable for stem and progenitor cells in certain pathological conditions, including myelofibrosis, where marrow cells are replaced with collagenous connective tissue fibers. Thus, EH occurs either actively or passively in response to diverse changes in the hematopoietic environment. This article reviews the key features and regulators of the major types of EH.

Keywords: myelopoiesis, cytokines, myeloid derived suppressor cells, T cells, infection, dendritic cells, FoxP3, myelofibrosis

Introduction

Extramedullary hematopoiesis (EH) refers to the hematopoiesis that occurs in organs other than bone marrow. A classic example of EH is the increased ectopic erythropoiesis in a liver or spleen in hypoxia due to increased erythropoietin production. EH can be further classified into active or passive categories. Normal hematopoiesis, which occurs in the fetal yolk sac, liver, and spleen, is an example of active EH; it is programmed as an essential process for routine fetal development. Another example of active EH is that which occurs in the spleen and liver during immune responses following infection. By contrast, EH also occurs as a result of failed marrow hematopoiesis in peripheral organs such as the liver and spleen and this is considered a passive form of EH. Both active and passive EH produce blood cells such as antigen-presenting cells, granulocytes, NK cells, red blood cells, and platelets for the growth or survival of the host. Inadequate EH leads to insufficient production or maturation of blood cells, while excessive EH leads to inflammatory diseases (Figure 1). This article will review the key aspects of major types of EH.

Factors required for hematopoiesis

Hematopoiesis is a complex process regulated by multiple factors. Hematopoiesis occurs in specialized tissue sites (eg, “hematopoietic niches”) conducive for the maintenance and differentiation of stem and progenitor cells. For example, hematopoietic stem cells, which are quiescent and self-renewing, are present in the marrow...
stem cell niche, which is low in oxygen and favorable to maintaining their identity as stem cells. The marrow stem cell niche is composed of various cells and their products which positively and negatively regulate the process. Key regulatory cells include osteoblasts, CXC chemokine ligand 12 (CXCL12)-expressing reticular cells, and vascular endothelium cells. Moreover, signals from the sympathetic nervous system and osteoclasts regulate the hematopoietic stem cell egress from the bone marrow through regulation of a critical stem cell homing and retention factor termed CXCL12. The requirement of a specialized niche is a limiting factor for EH in the periphery. As a consequence, EH is usually limited in the periphery at specific times and under certain conditions.

A number of molecular pathways including Wnt, calcium-sensing receptors, angiopoietin 1, Tie-2, and extracellular matrix components are involved in the process to finely control the stem cell niche. Although incompletely understood, these pathways are thought to be involved not only in the maintenance of the stem cell niche but also in renewal and differentiation of hematopoietic stem cells within the niche. Various cells produce hematopoietic cytokines such as stem cell factor (SCF), notch ligands, bone morphogenic proteins, transforming growth factor β, thrombopoietin (TPO), fibroblast growth factors, and insulin-like growth factor 2, which maintain and regulate the primitive hematopoietic stem cells. Other cytokines, such as granulocyte colony stimulating factor (G-CSF), interleukin-3 (IL-3), IL-7, erythropoietin (EPO), granulocyte macrophage (GM)-CSF, and macrophage (M)-CSF, play important roles in the differentiation of hematopoietic stem and progenitor cells to committed cell lineages.

In addition to the aforementioned cells and factors, hematopoiesis can be regulated by a number of other means,
such as Toll-like receptor ligands, metabolic/physiological products, various inflammatory mediators, and hormones. Many of the cytokines produced in inflammation act as myelopoietic factors, certain Toll-like receptor ligands promote myelopoiesis, and hypoxia is a well known inducer of erythropoiesis. Also, parathyroid hormone (PTH) and the insulin-like growth factors (IGF) control the hematopoietic stem cell niche; PTH can increase the number of bone marrow stem and progenitor cells and IGF can regulate the survival and expansion of hematopoietic stem and progenitor cells. Many of the cell types produced in the marrow, such as monocytes and B cells, require further maturation in the periphery to become fully functional immune cells. Monocytes will migrate into various tissue sites to become macrophages or dendritic cells and B cells must be activated in the periphery to become memory and plasma cells. Naive T cells are made in the thymus from progenitors that originated in the marrow and they undergo further differentiation in response to antigens proffered by antigen-presenting cells. Thus, the term “extramedullary hematopoiesis” refers to a broad range of hematopoietic activities from the early stages of lineage commitment to late stages of hematopoietic cell maturation.

**Early EH during fetal development**

Bone marrow becomes functional as a site of hematopoiesis in the fetus from 4–5 months in human pregnancy. In mice, the marrow hematopoiesis is delayed somewhat and becomes active following birth. The developing fetus needs hematopoietic cells for the supply of oxygen and other less obvious reasons. Therefore, nonmarrow tissues serve as sites of hematopoiesis before the bone marrow takes over the role as the major hematopoietic site. In early embryo, the yolk sac serves as the primary site of hematopoiesis. During mid-gestation, yolk sac cells colonize the umbilical cord, the aorta-gonad-mesonephros (AGM) region, and subsequently the embryonic liver. The yolk sac produces hematopoietic stem cells and red blood cells initially, but later produces myeloid cells as well. In this regard, the yolk sac is an important place, harboring primitive erythroblasts and essentially all definitive HPC. Later in embryonic development, some of these cells seed the fetal liver. As mentioned, the
next hematopoietic tissue site after the yolk sac is AGM and later, the fetal liver becomes a major hematopoietic site. Ultimately, the bone marrow becomes the predominant site of hematopoiesis. Following birth, the number of colony-forming hematopoietic progenitors in the spleen increases, peaking at two weeks of age in mice. In fact, the spleen remains a hematopoietic organ in mice throughout their lives, albeit at low levels.

It is thought that the liver ceases to act as a site of hematopoiesis after birth. However, the liver maintains hematopoietic stem cells, erythropoiesis, and myelopoiesis at low levels during adult life. Moreover, the liver is considered a maturation site for unconventional T cells including NKT cells, CD8αα T cells, CD4CD8- double-negative T cells, and γδ T cells.

**Induced EH occurs during infection and immune responses**

Following birth in mammals and rodents, extramedullary hematopoiesis occurs when immune responses occur in the periphery (Figure 1). The liver and spleen are the main sites of extramedullary hematopoiesis. Other organs such as the lungs, kidney, and the peritoneal cavity can also become the sites of hematopoiesis when in diseased states.

The increased extramedullary hematopoiesis in the spleen and liver of FAS (CD95)-deficient mice provides a useful insight into the cause of EH. In FAS-deficient mice, hematopoietic cells are resistant to apoptosis, allowing these cells to grow in organs which are normally not conducive for growth and differentiation of hematopoietic stem and progenitor cells. Also, immune cells are greatly expanded in these mice, producing large volumes of cytokines that can feed the growth of hematopoietic progenitor cells in the periphery. Another example of induced EH is the increased myelopoiesis found in many organs of FoxP3-null scurfy mice. Scurfy mice display increased CD11b+ myeloid cells in the spleen and the liver, which is consistent with increased expression of GM-CSF and IL-3. In scurfy mice, T cells producing GM-CSF and IL-3 are overly activated in the absence of FoxP3+ T cells, driving excessive EH in the spleen and liver. By contrast, the transfer of functional FoxP3+ T cells into newborn scurfy mice completely suppresses EH and scurfy-related inflammatory disease. An important function of the FoxP3+ T cells is to suppress the differentiation of naïve T cells into hematopoietic cytokine-producing T effector T cells (Figure 2). FoxP3+ T cells function mainly in the T-cell zone of the spleen rather than in the red pulps where EH occurs.

NK cells appear to play a negative role in the regulation of EH in the spleen. Depletion of NK cells with the anti-NK1.1 antibody in postnatal mice increased myeloid progenitor cells by 3–10 times in the spleen. This study, however, did not determine if the increase was the outcome of cell mobilization from the bone marrow or the product of the EH in the spleen. In vitro, NK cells can conversely decrease the numbers of progenitor cells. While the mechanism remains to be determined, NK cells could perhaps regulate the hematopoiesis with soluble inhibitory factors or through their cell-killing activity.

Toxic shock syndrome toxin-1 (TSST-1) is a superantigen produced by most *Staphylococcus aureus* strains. Superantigens can conjugate the MHC II molecules of antigen-presenting cells and certain β chains of T cell receptors for polyclonal activation of T cells. T cells are a good source of hematopoietic factors such as GM-CSF, IL-3, and oncostatin M. In this regard, TSST-1 can induce the production of hematopoietic factors. *Staphylococcus enterotoxin* B (SEB) is another superantigen produced by *Staphylococcus aureus*. SEB can activate naïve T cells and induce their differentiation into effector T cells that produce GM-CSF and IL-3.

Microbial components such as TLR ligands can affect hematopoiesis in both the marrow and spleen. TLR ligands such as lipopolysaccharides (LPS, a TLR2 ligand) and Pam3CSK4 (a TLR4 ligand) can directly activate hematopoietic progenitor cells through their receptors. Consequently, differentiation of progenitors into myeloid cells, such as macrophages and dendritic cells, is greatly enhanced in vitro while myeloid cells in marrow and spleen are increased following the injection of LPS *in vivo*. Interestingly, TLR ligands can turn even lymphoid progenitors into myeloid cells. Thus, TLR ligands are effective regulators of extramedullary hematopoiesis. This suggests that infection may significantly affect the hematopoiesis in the marrow and peripheral organs, such as the spleen.

The function of TLR ligands in regulation of hematopoiesis suggests that infection is an important element for EH to occur. As an example, it has been determined that Leishmania major infection increases colony-forming unit cells or myeloid progenitors in the spleen. Interestingly, the response in BALB/c mice was greater than that in C57BL/6 mice. This could be due to the lesser ability of BALB/c mice to clear the infection by intracellular pathogens. One of the mechanisms for the increased myelopoietic activity appears to be the production of GM-CSF and TNF-α, which can synergistically support the growth and maturation of myeloid progenitors.
Infection with *Plasmodium berghei* (ie, malaria) provided more insights into the kinetics of hematopoietic shift from the bone marrow to spleen. Infection with *Plasmodium berghei* increased numbers of CFU-GM in the bone marrow during the first week of infection, which was followed by a rise of CFU-GM in the peripheral blood and finally in the spleen around the end of the second week. The marrow progenitor numbers were normalized within two weeks, but those of the spleen remained elevated for a more prolonged time period. It is unclear how the hematopoiesis in the marrow and the spleen is sequentially regulated.

EH in an infection would generate sufficient numbers of mature myeloid cells to help clear pathogens, however, increased myelopoiesis without proper maturation of the myeloid progenitors could induce CD11b^+^ GR-1^+^ myeloid-derived suppressor cells. Myeloid-derived suppressor cells emerge in the spleen and other organs impacted by infection and cancer. These cells are highly heterogeneous and are considered normal constituents of myelopoiesis. Thus, incomplete EH could ultimately hamper immune responses to pathogens.

**EH as the result of chronic myeloproliferative syndrome**

Extramedullary hematopoiesis occurs if the bone marrow is no longer functional. Primary myelofibrosis is a form of Philadelphia-negative chronic myeloproliferative syndrome. In primary myelofibrosis, displacement and mobilization of stem and progenitor cells occur. As a consequence, hematopoietic stem and progenitor cells occupy the liver and spleen as alternative sites of hematopoiesis. At the same time, the bone marrow stem cell niche is altered so that it no longer supports normal hematopoiesis. Thus, in primary myelofibrosis, the site of hematopoiesis changes from the marrow to the spleen and liver. Primary myelofibrosis and related Philadelphia-negative chronic myeloproliferative syndrome are associated with a point mutation in the tyrosine kinase JAK2 (JAK2V617F). It is thought that this mutation makes hematopoietic stem and progenitor cells more sensitive to growth factors, alters the marrow stem cell niche, and causes the cells to mobilize to the spleen and liver. Another notable feature of primary myelofibrosis is the high production of inflammatory cytokines such as SDF-1, HGF, IL-6, IL-8, SCF, and VEGF. Likewise, the factors that promote fibrosis and angiogenesis (bFGF, TGF-β, PF4, VEGF, etc) are increased. These molecules are produced mainly by hematopoietic cells and contribute to the regulatory humoral changes occurring within the medullar and spleen niches.

In addition to the JAK2 mutation, activating mutations affecting the thrombopoietin receptor MPL (MPLW515L and MPLW515K) have been found in small numbers of patients with Philadelphia-negative chronic myeloproliferative syndrome. EH, particularly EH in myelofibrosis, can never fully replace the marrow hematopoiesis in production of necessary blood cells, however.

**Conclusion**

While the bone marrow is the major site of hematopoiesis, it can occur in many other tissues both during fetal development and after birth. Extramedullary hematopoiesis can occur as long as there are appropriate supporting cells, accommodation of hematopoietic progenitors, and local production of soluble and cell-bound hematopoietic factors that maintain and induce differentiation of the stem and progenitor cells. Extramedullary hematopoiesis occurs under several conditions, both actively and passively. It plays an essential role during fetal development, namely the survival of the fetus before formation of the functional marrow hematopoietic niche. As a normal response to infection and inflammation, myelopoiesis occurs in the spleen and liver to produce phagocytic cells and antigen-presenting cells. In malignant conditions, such as various forms of myelofibrosis, the marrow hematopoietic niche becomes inhabitable, and hematopoietic stem and progenitor cells move out to the periphery. EH cannot fully supplant the marrow hematopoiesis in terms of the production of necessary hematopoietic cells, rather it is imperative for the maturation of hematopoietic cells which are produced as immature cells in the marrow. While programmed extramedullary hematopoiesis is required to supplement the hematopoietic activity in the bone marrow, excessive and disease-associated extramedullary hematopoiesis can occur and mediate chronic inflammation. More research is required to discover useful strategies for controlling unwanted extramedullary hematopoiesis and chronic inflammation.

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References