Erythropoietin

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During the past century, few proteins have matched erythropoietin (Epo) in capturing the imagination of physiologists, molecular biologists, and, more recently, physicians and patients. Its appeal rests on its commanding role as the premier erythroid cytokine, the elegant mechanism underlying the regulation of its gene, and its remarkable impact as a therapeutic agent, arguably the most successful drug spawned by the revolution in recombinant DNA technology. This concise review will begin with a synopsis of the colorful history of this protein, culminating in its purification and molecular cloning. It then covers in more detail the contemporary understanding of Epo's physiology as well as its structure and interaction with its receptor. A major part of this article focuses on the regulation of the *Epo* gene and the discovery of HIF, a transcription factor that plays a cardinal role in molecular adaptation to hypoxia. In the concluding section, a synopsis of Epo's role in disorders of red blood cell production will be followed by an assessment of the remarkable impact of Epo therapy in the treatment of anemias, as well as concerns that provide a strong impetus for the development of even safer and more effective treatment.

In 1890, Viault (1890) observed that 2 weeks after traveling in Peru from sea level (Lima) to the mountain area of Morococha (4200 m) his red blood cell count went from 5.0 million to 7.1 million/mm³. Values of five other sojourners in his party ranged from 7.1 to 8.0 million. These simple observations provided the first convincing demonstration of the robust burst of erythropoiesis in man soon after exposure to high altitude hypoxia. During the transition into the next millennium the mechanism underlying this phenomenon became a topic of heated debate. Friedrich Miescher (1893), well known for his discovery of DNA, proposed that a decrease in oxygen tension within the bone marrow provided a direct stimulus to erythroid cells. A

half-century elapsed before this theory was disproven by carefully executed measurements of oxygen saturation in bone marrow specimens of patients with erythrocytosis, both primary (Berk et al. 1948) and secondary (Stohlman et al. 1954).

In 1906, Carnot and Deflandre (1906) proposed an alternate mechanism for hypoxic induction of erythropoiesis. They observed an increase in red blood cell counts following infusion of normal rabbits with serum from anemic animals and concluded that erythropoiesis is regulated by a humoral "factor" in the plasma. Attempts to reproduce this experiment over the ensuing decades yielded equivocal or negative results, thus casting doubt on this hypothesis.

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However, in the middle of the twentieth century, Krumdieck (1943) and Erslev (1953) modified the experimental design of Carnot and Deflandre by the addition of accurate measurements of reticulocytes and convincingly showed in rabbits the induction of new red cell production within 3–6 days following injection of anemic serum injected.

The notion that hypoxic stimulation of erythropoiesis involved an indirect humoral mechanism was strongly buttressed by experiments of Reissmann (1950) and Ruhenstroth-Bauer (1950). They used parabiotic pairs of rats whose circulations were connected at the capillary level by overlapping flaps of skin and elegantly showed that when one rat was exposed to low oxygen tension, whereas the other breathed room air and remained normoxic, both animals developed a surge of new red cell production and erythrocytosis.

Taken together, these studies led to the conception of a circulating erythroid-stimulating hormone, “erythropoietin” (Epo). Organ ablation studies in rats (Jacobson et al. 1957) and man (Nathan et al. 1964) firmly established that the kidney was the major site but not the sole site of Epo production. These findings led Eugene Goldwasser and his colleagues to undertake an intense and prolonged effort to isolate Epo. Initial attempts to obtain Epo from kidneys were unsuccessful owing to the release of proteolytic enzymes during tissue homogenization. In the search for a more tractable source of Epo, Goldwasser first turned to plasma of anemia sheep, then to urine from Argentinians with severe iron deficiency owing to hookworm infestation, and finally to urine from Japanese patients with aplastic anemia. This 15-year endeavor was greatly facilitated by the development of a sensitive and specific assay using incorporation of radio-labeled iron into newly produced red cells (Fried et al. 1956). By 1977, Goldwasser and his team were able to prepare 8 mg of highly purified human Epo (Miyake et al. 1977). Amino-terminal amino acid sequencing of this preparation enabled the synthesis of semidegenerate oligonucleotide probes that could then be used for the molecular cloning of the *Epo* gene (Jacobs et al. 1985; Lin et al.

1985). This advance opened up a new era in the exploration of the physiology and molecular biology of Epo and was exploited in the development of recombinant human Epo as a therapeutic agent for patients with various types of anemia.

For more detailed information on the history of Epo, see reviews by Grant and Root (1952), Erslev (1980, 1993), and Goldwasser (1996).

PHYSIOLOGY¹

In humans and other mammals, erythropoiesis normally proceeds at a low basal rate, replacing senescent red blood cells with young reticulocytes. In humans, red cell production can be enhanced as much as eightfold the baseline rate in a variety of clinical settings including hemorrhage, hemolysis, and other types of stress that impair the oxygenation of arterial blood or the delivery of oxygen to the tissues. Epo is the primary, and probably the sole mediator of hypoxic induction of erythropoiesis. During fetal development, Epo is produced mainly in the liver. However, in keeping with the above-mentioned report of Jacobson et al. (1957), following birth, the kidney accounts for ~80% of Epo production. Realizing that renal cortical blood flow closely matches oxygen consumption, Erslev et al. (1985) proposed that the proximal tubule is the ideal location for Epo production. In situ hybridization studies have indeed shown that *Epo* mRNA expression in the kidney is localized to a subset of peritubular fibroblasts in the cortex close to the boundary with the medulla (Koury et al. 1988, 1989; Lacombe et al. 1988; Bachmann et al. 1993; Maxwell et al. 1993a). In the liver, Epo is produced both in hepatocytes and in interstitial cells (Koury et al. 1991; Schuster et al. 1992). The latter have been shown to be Ito presinusoidal cells (Maxwell et al. 1994). In both Ito cells and the above-mentioned renal fibroblasts, in situ expression of Epo correlates with that of ecto-

¹For more information on all aspects of erythropoietin, see the recent detailed and comprehensive review by Wenger and Kurtz (2011).

5'-nucleotidase, probably because this enzyme is also controlled by local oxygen tension (Schmid et al. 1994).

The notion that Epo production is markedly up-regulated by hypoxia and that it stimulates erythropoiesis in a dose-dependent manner led

to the now well accepted paradigm of a negative feedback loop, similar to those underlying the regulation of blood glucose by insulin and of peripheral endocrine hormones by pituitary tropins. As shown in Figure 1A, hypoxia induces an increase in Epo hormone production in the

Figure 1. Regulation of red cell production by Epo. (A) Decreased oxygen delivery to specialized cells in the kidney results in increased expression and secretion of Epo, which circulates in the plasma and stimulates marrow progenitors, thereby increasing red cell production. If the increase in red cell mass relieves the hypoxic signal, Epo expression is down-regulated. (B) Plasma Epo levels (milliunits/mL) in patients with different types of and degrees of anemia and in those with primary erythrocytosis and secondary erythrocytosis. HIF, hypoxia inducible factor; PCV, polycythemia vera.

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kidney, which then circulates in the plasma and binds to receptors abundantly expressed on erythroid progenitor cells, thereby promoting the viability, proliferation, and terminal differentiation of erythroid precursors, and causing an increase in red blood cell mass. The oxygen-carrying capacity of the blood is thereby enhanced, increasing tissue oxygen tension, thus completing the feedback loop and suppressing further expression of Epo.

Further insight into the physiology of Epo production has come from measurements of plasma levels in normal individuals as well as in those with different types of anemia and erythrocytosis (Erslev et al. 1987). These studies became more conclusive with the development of accurate immune assays. Figure 1B depicts plasma Epo levels in normal individuals as well as in those with different types of anemia of varying degree. Note that the plasma Epo levels are plotted on a logarithmic scale. In patients with very severe anemia, Epo may be increased up to 1000-fold. The striking inverse relationship between plasma Epo levels and hematocrit or hemoglobin is seen in many different types of anemia, including blood loss, impaired red cell production, and hemolysis.

BIOCHEMISTRY AND CELL BIOLOGY²

Structure

The molecular cloning of the *Epo* gene led straightaway to high-level production of recombinant human erythropoietin (rhEpo) in cell culture in sufficient purity and quantity for both definitive scientific pursuits and the development of therapy (discussed at the end of this article). Human *Epo* messenger RNA (mRNA) encodes a 193-residue polypeptide. Following cleavage of a canonical leader sequence in the endoplasmic reticulum and glycosylation in the Golgi, a 166-residue polypeptide is released (Jacobs et al. 1985; Lin et al. 1985). The primary structure of rhEpo was shown to be identical to

that of the endogenous hormone except for the in vivo posttranslational cleavage of an arginine at the carboxyl terminus. Endogenous Epo and rhEpo have about 40% carbohydrate (w/w) of very similar structure, consisting of three *N*-linked polysaccharide groups and one *O*-linked group. Glycosylation of Epo slightly impedes its biological activity but is essential for ensuring prolonged circulation in the plasma (Goldwasser et al. 1974).

From its primary structure it was predicted that Epo, like other hematopoietic cytokines, folds into a globular three-dimensional structure consisting of a bundle of four amphipathic α helices connected by loops lacking secondary structure and stabilized by a critical disulfide bridge between the amino-terminal and carboxy-terminal helices (Bazan 1990; Boissel et al. 1993). This prediction was borne out by nuclear magnetic resonance spectroscopy of Epo in solution (Cheetham et al. 1998) as well as by X-ray crystallographic analysis of Epo in a complex with two extracellular domains of the Epo receptor (Syed et al. 1998).

Interaction with EpoR

Epo circulates in plasma with a plasma half-life of ~ 7 – 8 h and binds to high-affinity (~ 100 pM) receptors present in relatively small numbers (~ 1000 /cell) on the surface of erythroid progenitor cells (CFUe) in the bone marrow. The receptor-binding domains on the surface of Epo that were predicted from the properties of site-directed mutants (Wen et al. 1994; Elliott et al. 1996, 1997) have been confirmed and extended by structural analysis (Syed et al. 1998; Remy et al. 1999). EpoR is present on the surface of erythroid progenitors as a homodimer, even in the absence of ligand (Livnah et al. 1999). As shown in Figure 2, on binding to Epo, the receptor undergoes a conformational change that brings its intracellular domains into close apposition (Livnah et al. 1999; Remy et al. 1999), enabling cross phosphorylation via the binding of Jak2 kinase and the initiation of the signal transduction cascade.

The growth of CFUe in vitro was shown to be Epo dependent (McLeod et al. 1974). Ligand

²For more detailed information on Epo structure, interaction with receptor, and signal transduction, see review by Jelkmann and Wagner (2004).

Figure 2. Epo-dependent signaling. When Epo binds to its dimeric receptor (EpoR) on erythroid progenitor cells, the two receptor monomers are pulled together allowing phosphorylation of JAK2 kinase, which initiates the signal transduction cascade.

binding results in salvaging CFUe from apoptosis (Koury and Bondurant 1990) thereby enabling proliferation of erythroid cells as well as induction of terminal differentiation. The finding that both *Epo* mRNA and Epo protein are expressed in erythroid progenitors (Hermine et al. 1991; Stopka et al. 1998) has raised the intriguing possibility that tonic low-level erythropoiesis may be supported by autocrine stimulation, whereas circulating (hormonal) Epo provides a more robust stimulus to erythropoiesis during hypoxic stress.

In man, Epo's hematopoietic role appears to be restricted to the erythron, whereas in rodents, Epo stimulates megakaryocyte proliferation and maturation as well (Clark and Dessypris 1986; McDonald et al. 1987; Berridge et al. 1988).

NONHEMATOPOIETIC EFFECTS OF Epo³

As stressed in the sections above, by far the most abundant sites of Epo production are the kidney and liver, whereas expression of EpoR is vastly more robust in marrow erythroid progenitors

than in other cells. However, it has become increasingly evident that Epo has significant biological effects apart from regulation of red cell production. Attention has focused primarily on the central nervous system along with the heart, kidney, liver, and vasculature.

Within the brain, *Epo* and *Epo*-receptor mRNA are expressed widely (Tan et al. 1991; Masuda et al. 1994; Digicaylioglu et al. 1995; Sakanaka et al. 1998) and throughout development (Dame et al. 2000) in neurons, astrocytes, and endothelial cells (Yamaji et al. 1996). In contrast, demonstration of Epo and EpoR protein expression in situ has been considerably more challenging and less convincing. Oxygen-regulated expression of *Epo* has been observed in astrocytes both in vitro (Masuda et al. 1994; Marti et al. 1996) and in vivo (Tan et al. 1991; Marti et al. 1996), suggesting the possibility of a paracrine function for Epo in neural tissue.

The most compelling evidence of the importance of Epo signaling in the central nervous system comes from two independent sources. Targeted knockout of EpoR expression in the mouse brain has shown reduced neural cell proliferation and impaired poststroke neurogenesis (Tsai et al. 2006). Although global deletion of EpoR is lethal to mouse embryos, rescue by

³For more information on this topic, see review by Noguchi et al. (2008).

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expression of erythroid-specific EpoR results in survival to adulthood (Suzuki et al. 2002) but impaired neural cell proliferation and viability (Chen et al. 2007). In addition, administration of Epo protects against experimental brain injury in vivo (Sakanaka et al. 1998; Brines et al. 2000). Moreover, structural derivatives of Epo that lack erythroid stimulating activity have been shown to confer neuroprotection in animal models of stroke, spinal cord compression, diabetic neuropathy, and autoimmune encephalomyelitis (Leist et al. 2004). In a preliminary trial in patients with acute stroke, recombinant human erythropoietin (rhEpo) therapy was well tolerated and readily penetrated the blood–brain barrier (Ehrenreich et al. 2002). The authors' observations of reduction in infarct size and clinical improvement have not been confirmed in a more extensive clinical trial (Ehrenreich et al. 2009).⁴

High doses of rhEpo (Parsa et al. 2004; Nishiya et al. 2006) and the above-mentioned Epo derivatives (Fiordaliso et al. 2005; Brines et al. 2008) confer protection in animal models of cardiac ischemia/reperfusion and myocardial infarction. Moreover rhEpo is also effective in protecting the kidney and liver from ischemia/reperfusion injury (Yang et al. 2003; Patel et al. 2004; Sharples et al. 2004; Sepodes et al. 2006; Schmeding et al. 2009) as well as other experimental threats to kidney (Vaziri et al. 1994; Bahlmann et al. 2004) or liver (Le Minh et al. 2007; Schmeding et al. 2008, 2009) integrity.

The salutary effects of rhEpo and derivatives thereof on experimental models of ischemia and other types of injury may be due in part to their effect on the vascular endothelium. In vitro rhEpo stimulates the proliferation, mobilization, and differentiation of endothelial progenitor and precursor cells (Bahlmann et al. 2003; Heeschen et al. 2003) and also enhances endothelial cell viability (Chen et al. 2008) and survival by blocking apoptosis (Bahlmann et al. 2004). Thus, exogenous Epo can significantly enhance neovascularization induced by inflammation and ischemia (Heeschen et al. 2003).

⁴For more information on Epo and the brain, see reviews by Jelkmann (2005) and Noguchi et al. (2007).

Perhaps the most puzzling aspect of non-erythroid effects of Epo is the nature of the receptor. In erythroid cells, EpoR mRNA is expressed at sufficiently high levels for abundant display of EpoR protein on cell surface and dimerization that is required for high-affinity Epo binding. In contrast, the mRNA expression in nonerythroid cells is so low that convincing demonstration of surface EpoR protein by immunological means has been problematic at best. It is very likely that such low-level expression would not be sufficient for the mass action needed for EpoR dimerization to occur. Accordingly, the low-affinity binding site for Epo may be a heterodimer composed of EpoR and another partner, perhaps CD131, the β common cytokine receptor (Brines et al. 2008; Brines 2010). This may explain the requirement of high levels of exogenous Epo for biological effects in nonerythroid cells and possibly how an Epo derivative could have activity only in nonerythroid cells.

REGULATION OF THE *Epo* GENE⁵

Initial investigation of the molecular basis of *Epo* gene regulation was greatly facilitated by the discovery that the human hepatoma cell lines Hep3B and HepG2 produce significant amounts of Epo constitutively, with marked induction in response to hypoxia (Goldberg et al. 1987). The magnitude and time course of the induction of *Epo* mRNA paralleled Epo protein production. More recently, a human renal cell line has also been shown to display oxygen-dependent regulation of Epo production (Frede et al. 2011).

Tissue-specific expression of the *Epo* gene and its induction by hypoxia are dependent on far upstream *cis* elements and an enhancer element downstream from the polyadenylation signal. Transgenic mice experiments indicate the presence of element(s) between 0.4 and 6 kb 5' of the promoter, which suppress promiscuous expression, and element(s) 9.5 and 14 kb upstream, which are necessary for kidney-specific

⁵For more information on this topic, see review by Fandrey (2004).

expression (Semenza et al. 1989, 1990, 1991a; Madan et al. 1995).

The *Epo* gene has a weak promoter, owing in part to suppression of transcription by a GATA motif (Imagawa et al. 1994, 1997). Mutation of this response element in a 180-kb *Epo* transgene resulted in constitutive in vivo expression in renal peritubular cells (Obara et al. 2008). Enhanced binding of GATA-2 to this site may contribute to the inhibition of *Epo* production by inflammatory cytokines (La Ferla et al. 2002). GATA-4 is expressed exclusively in hepatocytes and binds strongly to this GATA motif, enhancing *Epo* expression (Dame et al. 2004). The marked reduction in expression of hepatic GATA-4 following birth is likely an important contributor to the switch in *Epo* production from liver to kidney (Dame et al. 2004).

A crucial 3' enhancer (Beck et al. 1991; Pugh et al. 1991; Semenza et al. 1991b; Semenza and Wang 1992) binds to two transcription factors, hypoxia-inducible factor 1 (HIF) and the nuclear receptor HNF-4. As shown in Figure 3, these two DNA-binding proteins interact with the transcriptional coactivator p300/CBP, triggering transcriptional activation.

HIF

Hypoxic induction of *Epo* depends in large part on the transcription factor HIF. HIF is activated in virtually all cells (Maxwell et al. 1993b; Wang and Semenza 1993b) by exposure to hypoxia, as well as cobalt or iron chelators. The activation of HIF enables it to bind to a consensus sequence (5'-TACGTGCT-3') first identified in the *Epo* 3' enhancer (Fig. 3) (Semenza et al. 1991b). Sub-

sequently, this hypoxia response element has been identified in more than a hundred genes and shown to be required for hypoxic induction of transcription. In addition to *Epo*, an impressive number of other physiologically relevant genes are regulated by HIF (Wenger et al. 2005) including those involved in angiogenesis, intermediary metabolism, and, of particular interest to readers of this collection, iron homeostasis (Peyssonnaud et al. 2007).

Wang and Semenza (1995) purified HIF by its affinity for the *Epo* 3' enhancer and showed that it is a heterodimer composed of a 120 kD α subunit and a 91–94 kD β subunit. Molecular cloning revealed that both subunits are basic helix-loop-helix proteins in the PAS family of transcription factors (Wang et al. 1995a). HIF- α , or HIF-1 α , is a novel protein, whereas HIF-1 β is the previously cloned and characterized aryl hydrocarbon receptor nuclear translocator (ARNT) (Hoffman et al. 1991).

The HIF- α subunit is widespread among animals, including not only mammals and fish but also invertebrates such as flies and worms. In man and other mammals, three homologous HIF- α genes have been identified: HIF-1 α , HIF-2 α (Tian et al. 1997), and HIF-3 α (Gu et al. 1998). HIF-1 α is the most abundant and present in most organs and tissues. In contrast, HIF-2 α has more restricted expression and was originally thought to be limited to the vascular endothelium (Tian et al. 1997). Knockout of HIF-1 α in mice is embryonic lethal (Iyer et al. 1998; Ryan et al. 1998), whereas HIF-2 α null mice survive gestation but have impaired survival owing to phenotypic abnormalities that are strain-dependent. Most genes that are

Figure 3. Diagram of the *Epo* gene. The five exons of the *Epo* gene are shown as rectangles with the coding regions in black. Far upstream of the *Epo* promoter is a kidney-inducible element (KIE) that is required for high-level up-regulation of *Epo* mRNA in the kidney. Just downstream from exon 5 is a critical enhancer that binds to HNF-4 and also to HIF when the *Epo*-producing cell is hypoxic. These two transcription factors bind to the transcriptional activator p300. This enhanceosome is a powerful inducer of *Epo* transcription.

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induced by hypoxia in an HIF-dependent manner require HIF-1 α (Warnecke et al. 2004). In contrast, Epo production and its up-regulation by hypoxia require HIF-2 α (Gruber et al. 2007) in the kidney (Scortegagna et al. 2005; Kapitsinou et al. 2010; Paliege et al. 2010) and an Epo-producing renal cell line (Frede et al. 2011), as well as in the fetal and adult liver (Rankin et al. 2007).

Oxygen Sensing and HIF Activation

To understand the mechanism by which HIF-1 is activated, it is necessary to determine how hypoxia impacts on the expression of its two subunits. The steady-state levels of HIF- α mRNA and ARNT mRNA are not significantly affected by oxygen tension (Gradin et al. 1996; Huang et al. 1996; Wood et al. 1996). At the protein level, the ARNT subunit remains abundant, irrespective of the oxygenation of the cell. In contrast, the HIF- α subunit cannot be detected in oxygenated cells (Wang et al. 1995b; Huang et al. 1996). HIF- α protein can only be detected in deoxygenated cells or in those exposed to iron chelators or certain transition-metal ions, most notably cobalt, all of which induce HIF-dependent gene transcription (Semenza and Wang 1992; Wang and Semenza 1993a). These observations suggest that the activation of HIF-1 depends on an increase in the amount of HIF- α protein in deoxygenated cells.

In cells exposed to 21% O₂, HIF- α is remarkably unstable (Huang et al. 1996). It forms a complex with von Hippel–Lindau protein (vHL) (Maxwell et al. 1999), enabling it to undergo ubiquitination and subsequent degradation in proteasomes (Salceda and Caro 1997; Huang et al. 1998). Low oxygen tension abrogates this process, thereby allowing the HIF- α subunit to accumulate so that it can enter the nucleus and form a stable heterodimer that can participate in transcriptional regulation. Oxygen-dependent degradation of HIF- α depends on a large (~200-residue) interior segment (Huang et al. 1998). When this domain is deleted, the protein is stable and capable of *trans*-activation, even in oxygenated cells. When this

domain is fused to a foreign protein, it confers rapid oxygen-dependent degradation.

The oxygen-dependent degradation domain of HIF-1 α contains two highly conserved LXXLAP sequences, which are recognition sites for an oxygen- and iron-dependent prolyl hydroxylase that converts the proline into a hydroxyproline (Ivan et al. 2001; Jaakkola et al. 2001; Masson et al. 2001; Yu et al. 2001). This posttranslational modification is necessary and sufficient for the binding of vHL to HIF- α . As shown in Figure 4, this interaction enables the ubiquitin E3 ligase to dock at this complex, thereby triggering the covalent addition of a number of small ubiquitin molecules (Cockman et al. 2000; Ohh et al. 2000). The polyubiquitinated HIF- α is then recognized by the proteasome where it is rapidly degraded. Ratcliffe and his colleagues (Epstein et al. 2001) showed that the hydroxylation of these two specific proline residues is mediated by a dioxygenase that is homologous to classic collagen prolyl hydroxylase. Shortly thereafter, a human homolog of HIF prolyl hydroxylase was biochemically purified and characterized (Ivan et al. 2002). These enzymes have an active center where an iron atom and a molecule of α -ketoglutarate interact with a conserved core of two histidines, an aspartate and an arginine residue. The two LXXLAP motifs within HIF- α each bind at this site, along with oxygen, catalyzing hydroxylation of the proline residue. This oxidative modification is inhibited by iron chelators as well as by cobaltous ion, which presumably displaces iron from the active site. The effect of the iron chelation and cobalt on the activity of HIF prolyl hydroxylases provides a satisfying explanation for how these agents mimic hypoxia in activating HIF. These enzymes are the proximate oxygen sensor for regulating HIF. Among the three HIF- α prolyl hydroxylases, PHD2 plays the predominant role at most sites (Berra et al. 2003) including renal production of Epo (Minamishima et al. 2008; Takeda et al. 2008).

Hemoglobin is encapsulated in circulating erythrocytes not only in man and other mammals but also in all other vertebrates. Oxygen homeostasis in these complex organisms is

Figure 4. Pathway by which the hypoxia-inducible transcription factor is up-regulated by low intracellular oxygen tension. In normally oxygenated cells, the α subunit (HIF- α) undergoes hydroxylation of two proline residues, one of which is shown here. In the presence of iron (Fe) and α -ketoglutarate, this oxygen-dependent posttranslational modification is catalyzed by an HIF- α -specific prolyl hydroxylase (PH). The von Hippel-Lindau protein (pVHL) binds to hydroxylated HIF- α . Subsequent docking of a ubiquitin ligase (UL) enables HIF- α to be polyubiquitinated.

well served by HIF-dependent regulation of a cytokine that is required for erythropoiesis. In contrast, in many invertebrate phyla, oxygen transport depends on the presence of free soluble polymeric hemoglobins in circulating hemolymph, and in some, hemoglobin production is enhanced during hypoxic stress. One of the most dramatic observations in marine biology is the change in the fresh-water crustacean *Daphnia magna* from pale gray to bright red soon after transfer from normoxic to hypoxic water (Fox 1947, 1949). Two of their four tandem globin genes have been shown to have functional HIF response elements within their promoters (Gorr et al. 2004). Thus, in keeping with the relative simplicity of this organism, the regulation of oxygen-carrying capacity of the blood is direct, rather than mediated indirectly via a cytokine.

HNF-4

Downstream from the HIF-1 binding site, the *Epo* 3' enhancer contains two tandem consensus steroid hormone response elements separated

by 2 bp (Fig. 1). Mutations at these sites either abolish or markedly inhibit hypoxic induction of reporter genes (Blanchard et al. 1992; Semenza and Wang 1992; Pugh et al. 1994; Hu et al. 1997). These elements in the *Epo* enhancer interact with an orphan nuclear receptor, i.e., a DNA-binding protein that shares structural homology with hormone-binding nuclear receptors but lacks a known ligand (Blanchard et al. 1992). The orphan receptor HNF-4 plays a critical role in *Epo* gene regulation (Galson et al. 1995; Makita et al. 2001). The expression of HNF-4 is limited to renal cortex and liver, sites of *Epo* production, and also intestine. Thus the binding of HNF-4 to the *Epo* enhancer is likely to contribute to both the high-level induction of the *Epo* gene as well as its tissue specificity.

p300

As shown in Figure 1, the carboxy-terminal portion of HIF-1 α binds specifically to p300 (Arany et al. 1996), a general transcriptional activator that participates in a number of biological functions such as induction of tissue-specific

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expression, regulation of cell cycle, and stimulation of differentiation pathways. This very large protein, which is closely homologous to CREB-binding protein (CBP), does not bind to DNA but does interact with a number of other proteins including HNF-4 and other nuclear receptors. Thus, as depicted in Figure 1, the HIF-1 heterodimer, activated by hypoxia, participates in a macromolecular assembly with p300 (or a related family member) and with HNF-4 to transduce a signal to the Epo promoter, enabling activation of transcription. It is very likely that such a combinatorial process applies not only to *Epo* but also to other genes that are induced by hypoxia (Ebert and Bunn 1998). Most, if not all, cells contain a protein factor called FIH (factor-inhibiting HIF) (Mahon et al. 2001) that hydroxylates an asparagine at the carboxyl terminus of HIF- α in an oxygen- and iron-dependent manner, thereby blocking binding to p300 and inhibiting transcription (Lando et al. 2002a,b).

IMPACT OF *Epo* PRODUCTION ON CLINICAL DISORDERS

In a variety of clinical settings, anemia can be caused by underproduction of *Epo*, whereas erythrocytosis can result from overproduction.

Underproduction of *Epo*

Patients with renal failure generally develop severe anemia, owing in part to suppression of erythropoiesis from a buildup of metabolic wastes and in part to a moderate reduction in red cell life span. However, the most important contributor to the anemia of uremia is insufficient *Epo* production (Fig. 1B) (Caro et al. 1979; Cotes 1989). The degree of anemia correlates roughly with the extent of renal functional impairment. In renal failure owing to a wide variety of etiologies, insufficient *Epo* production results either from direct damage to *Epo*-producing cells in the kidney or from the suppression of *Epo* production by inflammatory cytokines.

Patients with inflammatory disorders have a less marked impairment of *Epo* production. Examples include rheumatoid arthritis (Means

1994), cancer (Miller et al. 1990; Spivak 1994), and AIDS (Fischl et al. 1990). It is likely that inflammatory cytokines suppress *Epo* gene expression in these disorders.

Exposure to certain metals may result in disordered structure and function of the renal proximal tubule, resulting in suppression of *Epo* production, out of proportion to impairment of global renal function. Such a pathogenetic process is likely the basis for the marked anemia often encountered in cancer patients who have been treated with *cis*-platinum or individuals suffering from chronic cadmium intoxication (Horiguchi et al. 2000). *Epo* production is also inhibited in patients with increased plasma viscosity owing to monoclonal dysproteinemias (Singh et al. 1993).

Overproduction of *Epo*

Measurement of plasma *Epo* is useful in distinguishing primary from secondary erythrocytosis (Erslev and Caro 1984; Messinezy et al. 2002). Individuals with autonomous red cell overproduction due to polycythemia vera (or rarely to mutations at the carboxyl terminus of *EpoR*) have low plasma *Epo* levels, whereas individuals with various forms of chronic arterial hypoxemia have erythrocytosis owing to increased plasma *Epo* levels. The highest documented hematocrit levels have been observed in patients with right to left cardiac shunts. The erythropoietic response in patients with hypoxemia owing to chronic obstructive pulmonary disease is variable, depending in part on whether there is coexisting infection, which, as explained above, can suppress erythropoietin expression.

Increased oxygen affinity of red cells causes impaired delivery of oxygen to tissues and therefore hypoxia at the cellular level, which triggers increased *Epo* expression and therefore secondary erythrocytosis (Prchal and Prchal 1999). Such a “shift to the left” in the oxyhemoglobin dissociation curve can be caused by mutations in α - or β -globin subunits or in two red cell enzymes. Deficiency in cytochrome b5 reductase causes congenital methemoglobinemia and mild erythrocytosis. A more pronounced increase in red cell mass has been reported in

very rare families with deficiency of bisphosphoglycerate mutase that causes a deficit in red cell 2,3-BPG, a critical modulator of intracellular hemoglobin function (Rosa et al. 1978). These individuals with left-shifted O₂-binding curves are generally asymptomatic because impaired oxygen delivery is balanced by increased oxygen-carrying capacity.

Occasional individuals have chronic elevation of plasma Epo that is inappropriate and autonomous, i.e., not due to arterial hypoxemia or cellular hypoxia. These individuals usually have either sporadic or germline mutation in genes responsible for oxygen sensing and HIF regulation.

Erythrocytosis due to inappropriate overproduction of Epo can be encountered in various neoplasms, particularly those arising in kidney, liver, and cerebellum, organs that are physiologic sites of erythropoietin expression. In particular, secondary erythrocytosis occurs in patients with renal carcinomas, Wilms tumor, hepatomas, and cerebellar hemangioblastomas. In some cases, the tumor cells secrete Epo (Jelkmann 1992). In other cases, the surrounding normal kidney or liver tissue secretes Epo, presumably because of local ischemia.

Highly vascular renal and central nervous system tumors can arise in families and sporadically owing to mutations in the von Hippel-Lindau (VHL) gene. As explained earlier in this article, this protein is required for the oxygen-dependent degradation of HIF- α . Inactivating mutations lead to constitutive activation of HIF- α and therefore overexpression of HIF responsive genes (Gnarra et al. 1996; Iliopoulos et al. 1996), particularly vascular endothelial growth factor (VEGF), which causes the tumors to have enhanced vascularity. In a subset of affected individuals, overexpression of Epo leads to erythrocytosis.

A number of families with congenital erythrocytosis have been encountered in Chuvashia, a circumscribed region in the Russian Federation in which there may have been inbreeding and a high likelihood of a founder effect (Sergeyeva et al. 1997; Gordeuk and Prchal 2006). The erythrocytosis (mean hemoglobin levels of 23 g/dL and hematocrit of 67%) follows an au-

tosomal recessive pattern of inheritance. Affected individuals have elevated erythropoietin levels, and increased incidence of thrombotic and hemorrhagic complications. These individuals are homozygous for a missense mutation at the carboxyl terminus of the VHL gene (R200W) leading to HIF activation (Ang et al. 2002). The clinical phenotype is similar in a R200W “knock-in” mouse model (Hickey et al. 2007). Isolated families with congenital erythrocytosis have been encountered elsewhere that harbor the R200W mutation (Pastore et al. 2003; Cario et al. 2005; Perrotta et al. 2006). Affected individuals are usually homozygotes but some are compound heterozygotes, having R200W along with another VHL mutation (Pastore et al. 2003; Cario et al. 2005).

Families as well as isolated individuals have been encountered in which erythrocytosis owing to elevated plasma Epo levels is explained by heterozygosity for an activating mutation in HIF-2 α (Gale et al. 2008; Martini et al. 2008; Percy et al. 2008a,b, 2012; Furlow et al. 2009; van Wijk et al. 2010). These mutations generally impair the ability of HIF-2 α to bind to VHL or PHD2. In other families, affected individuals are heterozygous for loss-of-function mutations in PHD2 (Percy et al. 2006, 2007; Ladrone et al. 2008). Genome-wide association studies comparing lowland Tibetans to highlanders with erythrocytosis have shown highly significant polymorphisms in the vicinity of HIF-2 α (Beall et al. 2010; Yi et al. 2010) and PHD2 (Simonson 2010). Both these selected “experiments of nature” and the broader population studies, attest to the primacy of HIF-2 α and PHD2 in regulating the *Epo* gene.

THERAPY WITH rhEpo AND ERYTHROID-STIMULATING AGENTS⁶

By far the most common use of rhEpo has been in patients with chronic renal failure. Over one million patients worldwide have responded effectively to either rhEpo or a derivative, darbepoietin alfa (Aranesp), which has an enhanced

⁶For more information on this subject, see reviews by Bunn (2007), Elliott (2008), and Jelkmann (2008).

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plasma half-life (~ 22 h) owing to the introduction of two additional *N*-linked glycosylation sites.

The importance of erythropoietin in the pathogenesis of the anemia of renal failure has been affirmed by the dramatic efficacy of rhEpo therapy. Figure 5 shows one of the first cases treated with rhEpo. Within a few days after initiation of rhEpo therapy, the hematocrit approached normal, necessitating a reduction in dose. The marked increase in red cell mass following treatment was accompanied by enhanced utilization of iron stores, as reflected in a decline in serum iron and serum ferritin. Other patients who have normal or low iron stores before rhEpo therapy need concomitant administration of iron to achieve an optimal erythropoietic response.

The other common use of rhEpo has been in treating anemia in cancer patients with chemo-

therapy-associated anemia. In addition, rhEpo is effective in AIDs patients who have developed severe anemia as a result of antiviral therapy. A minority of patients with myelodysplastic anemia also responds to rhEpo. In some clinics, rhEpo is used in the management of the anemia of prematurity as well in surgical patients, in both the preoperative and postoperative periods.

The impressive efficacy and safety of rhEpo therapy notwithstanding, a number of problems warrant careful consideration. The most evident are the high cost of treatment and the need for parenteral administration. Of equal concern are outcome studies on large numbers of renal dialysis patients showing that doses of rhEpo and darbepoietin alfa that raise the hemoglobin level >12 g/dl are associated with thrombosis of arteriovenous access, impaired control of hypertension, and a small but sig-

Figure 5. Response of an anephric patient to recombinant human erythropoietin (rhEpo) therapy. Note that, before therapy, the patient was severely anemic and transfusion dependent. Treatment with rhEpo resulted in a reticulocytosis followed by a progressive increase in hemoglobin. The dose of rhEpo had to be lowered to prevent the hemoglobin from rising too high. Before rhEpo therapy, the patient was severely iron overloaded. The marked increase in red cell mass following therapy was accompanied by a significant reduction in iron stores. RBC, red blood cells; Fe, iron; TIBC, total iron binding capacity; Sat., saturation (Data from Eschbach et al. 1987.)

nificant increase in mortality (Besarab et al. 1998; Singh et al. 2006; Phrommintikul et al. 2007). Cancer patients are inherently at higher risk to develop thrombosis, but this risk is significantly enhanced in those receiving rhEpo therapy (Bohlius et al. 2006; Bennett et al. 2008).

It is likely that the prothrombotic effects of rhEpo dose are due to the high doses given rather than to the increase in hemoglobin level. As mentioned earlier in this article, nonerythropoietic effects are widespread and complex. Early reports suggest that rhEpo may also promote enhanced tumor growth in vivo (Henke et al. 2003; Leyland-Jones 2003) and in vitro (Mohyeldin et al. 2005). However, EpoR expression is low in the great majority of tumors and though expressed at the mRNA level, it cannot be detected on the cell surface (Sinclair et al. 2008). Moreover, more recent meta-analyses mentioned above do not support the notion that rhEpo therapy promotes tumor progression in vivo (Bohlius et al. 2006; Bennett et al. 2008).

RECENT DEVELOPMENTS

In view of the above-mentioned concerns regarding long-term rhEpo therapy, there has been increasing interest in developing erythroid-stimulating agents that offer advantages over rhEpo and darbepoietin alfa. One approach entailed the production of a large phage library expressing short (~15-residue) random peptides that are then screened for sequences that bind with relatively high affinity and specificity to the Epo receptor. This strategy led to the identification of peptides that lack any homology with Epo but are active both in vitro and in vivo in stimulating erythropoiesis through the same signaling pathway that is triggered by Epo (Livnah et al. 1996; Wrighton et al. 1996). However, in contrast to recent progress with thrombopoietin mimetic agents, little progress has been made to date in developing these Epo mimetic peptides into safe and effective drugs.

Another approach involves structural modification of Epo in the interest of prolonging its longevity in the circulation. Continuous erythropoietin receptor activator (CERA) is a de-

rivative in which a 30 kDa methoxy-polyethylene glycol polymer has been covalently linked to Epo. It has a half-life in the circulation of ~135 h, about sixfold greater than that of darbepoietin alfa (Macdougall 2005; Macdougall et al. 2006; Provenzano et al. 2007). This attribute greatly reduces the frequency of parenteral administration. CERA has been through extensive clinical trials and has passed muster in terms of both safety and efficacy. As a result, it received approval from the European Commission and the American Food and Drug Administration in 2008. However, it has not yet gained wide use in clinical practice.

The most novel strategy for pharmacologic stimulation of erythropoiesis involves the activation of HIF by inhibition of HIF- α prolylhydroxylases. As shown in Figure 4, small molecule analogs of α -ketoglutarate could prevent HIF- α hydroxylation, even in well-oxygenated cells and thereby induce expression not only of endogenous Epo but also of other HIF-dependent genes that are required for robust erythropoiesis. Several companies have developed inhibitors that induce Epo and erythropoiesis in mice (Safran et al. 2006; Yan et al. 2010) and in humans (Yan et al. 2010). Because these agents suppress expression of hepcidin (Volke et al. 2009), they may be particularly effective in the treatment of anemia of chronic inflammation. HIF prolylhydroxylase inhibitors offer for the first time the possibility of inexpensive oral therapy for stimulation of erythropoiesis. Of course, a major concern is specificity because such a large number of genes are regulated by HIF. The lead compounds that are currently undergoing clinical trials appear to inhibit HIF-2 α specifically. If so, they may also enhance duodenal iron absorption because the divalent metal transporter 1 (DMT1), like Epo, is up-regulated by HIF-2 α .

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