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Novel potential therapeutics to modify iron metabolism and red cell synthesis in diseases associated with defective erythropoiesis

Short title: Erythropoiesis, iron metabolism and diseases

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Abstract
Under normal conditions, iron metabolism is carefully regulated to sustain normal cellular functions and the production of hemoglobin in erythroid cells. Perturbation to the erythropoiesis-iron metabolism axis can result in iron imbalances and cause anemias or organ toxicity. Several congenital and acquired diseases associated with abnormal red cell production are characterized by aberrant iron absorption. Several recent studies have shown that improvements in red blood cell production also ameliorate iron metabolism and vice versa. Many therapeutics are now under development with the potential to improve a variety of hematologic diseases, from beta-thalassemia and iron-refractory iron deficiency anemia (IRIDA) to anemia of inflammation and polycythemia vera. This review summarizes selected mechanisms related to red cell production and iron metabolism and describes potential therapeutics and their current uses. We also consider the potential application of the discussed therapeutics on various diseases, alone or in combination. The vast repertoire of drugs under development offers new opportunities to improve the clinical care of patients suffering from congenital or acquired red blood cell disorders with limited or no treatment options.

Erythropoiesis and iron metabolism

Erythropoiesis, the production of mature enucleated erythrocytes from early progenitor cells, is regulated by molecules that control proliferation and differentiation. For brevity, the descriptions of the molecules that regulate erythropoiesis and iron metabolism will be limited to those relevant to the diseases and potential therapies discussed in this review.

GATA1 is a key transcription factor essential for erythropoiesis. GATA1 promotes maturation and survival of erythroblasts, which develop into enucleated red cells. GATA1 stimulates several pathways, including those responsible for hemoglobin synthesis and stress erythropoiesis. The erythrocyte-specific isoform of pyruvate kinase, R-type pyruvate kinase, is regulated by GATA1 and is involved in the final steps of glycolysis, which leads to the formation of 2 adenosine triphosphate and 2 pyruvate molecules per glucose molecule. R-type pyruvate kinase is required for the survival of mature red cells and to sustain the tricarboxylic acid cycle in erythroid precursors.

Erythropoietin (EPO) binds to the EPO receptor (EPOR) to regulate early-stage erythropoiesis, primarily by phosphorylation of JAK2 kinase (Figure 1). Phosphorylated JAK2 kinase is responsible for the activation of downstream pathways that regulate expression of genes that control proliferation, survival, and iron metabolism, such as the hormone erythroferrone (ERFE) (Figure 1). A subset of transforming growth factor-beta (TGF-beta) superfamily receptors and ligands may play a major role in late-stage erythropoiesis by modulation of differentiation and red cell production.

Hepcidin (encoded by HAMP) is a liver-secreted hormone that plays a major role in iron homeostasis (Figure 1). It binds and inhibits the iron exporter ferroportin, preventing release of iron into the circulation from duodenal enterocytes, macrophages, and hepatocytes. Increased hepcidin expression causes anemia by limiting iron availability, whereas low hepcidin expression leads to iron overload. Activation of hepcidin transcription is mediated by the iron-sensor complex on liver parenchymal cells, which includes bone morphogenetic protein (BMP) receptors from the TGF-beta receptor superfamily as well as coreceptors, ligands, and modulators (Figure 1). Liver endothelial cells secrete the BMP ligands BMP2 and BMP6, which bind class I and II BMP receptors in hepatocytes and activate the SMAD1/5/8 complex and HAMP expression (Figure 1). Hemojuvelin (HJV) acts as a BMP
coreceptor, whereas TMPRSS6 (which encodes for matriptase-2, a transmembrane serine protease) downregulates hepcidin by cleaving HJV (Figure 1).^{17}

Additional proteins in the liver modulate the activity of the iron-sensor complex by sensing the level of iron bound to transferrin (TF), the main molecule responsible for the transportation of iron from the duodenum (Figure 1).^{16} Increasing amounts of diferric TF displace HFE from TF receptor 1. This not only enables iron uptake in the liver but also stabilizes TF receptor 2 (TFR2), potentiating signaling of the iron-sensor complex (Figure 1).^{16} Although TFR2 was thought to only modulate hepcidin synthesis in the liver, it was recently shown that TFR2 colocalizes with EPOR on erythroid precursor cells to modulate EPO sensitivity (Figure 1).^{16,18} The role of TFR2 in this setting is thought to fine-tune erythropoiesis based on the amount of iron bound to TF.^{19}

**Diseases associated with abnormal red cell synthesis and their pathophysiology**

Erythropoiesis can be altered by genetic mutations that impair red cell production, such as in congenital anemias or polycythemia vera (PV), or by conditions that perturb normal iron metabolism, such as iron-refractory iron deficiency anemia (IRIDA).^{20} Conditions associated with chronic inflammation, such as chronic kidney disease (CKD), are also characterized by reduced red cell production and survival.^{21} In this review, we focus on diseases that illustrate key pathophysiological mechanisms responsible for altered red cell production, the role of iron metabolism, and strategies targeting these mechanisms that may serve as potential novel treatment approaches.

Anemia is a consequence of inadequate functional red cells in circulation, which can result from reduced red cell production, increased red cell destruction, or a combination of these two processes. Diseases associated with genetic mutations that directly alter red cell production are generally characterized by defects at the erythroid cell-differentiation stage where the mutated genes exert their functions. These mutations may also affect the quality and lifespan of surviving mature red cells.

**Beta thalassemia**

Beta thalassemia (BT) and alpha thalassemia (AT) are two of the most widespread forms of congenital anemias.^{22} These diseases are triggered by mutations that reduce the synthesis of beta-globin or alpha-globin, respectively.^{22} Heterozygosity for BT null mutations (BT trait) is usually clinically silent but becomes symptomatic with coinheritance of extra alpha-globin genes, owing to a relatively increased amount of free alpha-globin protein.^{23,24} Hence, the globin-chain imbalance with an accumulation of free alpha-globin is a major determinant of BT pathophysiology and causes a variety of effects that impairs red cell maturation and survival (Figure 2).^{22}

Excess alpha-globin chains precipitate as hemichromes, which are insoluble aggregates of alpha-chains and heme molecules that form large inclusion bodies and cause iron accumulation and free radical production (Figure 2).^{25,26} These events lead to premature destruction by apoptosis of a portion of red cell precursors in the bone marrow as well as the production of abnormal enucleated erythrocytes with reduced lifespan.^{27} In addition, excess alpha-globin chains also interfere with erythroid differentiation. In normal erythropoiesis, chaperone heat shock protein 70 (HSP70) protects GATA1 from caspase-3 cleavage (Figure 2).^{28} However, in BT, HSP70 interacts directly with free alpha-globin chains, and therefore HSP70 is sequestered and unable to protect GATA1, resulting in end-stage maturation arrest (Figure 2).^{28} Altogether,
this mechanism limits the production of enucleated red cells and consequent oxygen delivery (Figure 2). The resulting hypoxia increases the production of EPO in the kidneys. EPO further stimulates the production and expansion of early erythroid progenitors without generating sufficient mature red cells (Figure 2).\textsuperscript{27} This leads to a net imbalance between nucleated and enucleated erythroid cells—the number of terminally differentiated red cells is disproportionately low compared with the number of progenitor red cells (Figure 2).\textsuperscript{27} This abnormal production of erythroid progenitors, combined with a hypoxic environment, profoundly affects iron metabolism. The production of ERFE, by the early progenitor red cells, increases as the number of erythroid precursor cells increases, leading to suppression of hepcidin and results in enhanced iron absorption (Figure 2).\textsuperscript{27} In addition, the hypoxic environment also triggers the synthesis of genes responsible for iron absorption in the duodenum, further exacerbating the process of iron overload in organs.\textsuperscript{27,29}

Iron accumulation results in free radical damage to cells and organs and contributes to the morbidity and mortality of iron overload (Figure 2).\textsuperscript{30} Although the main source of extra iron in patients with severe BT comes from blood transfusions, the main cause of iron overload in patients that only sporadically receive transfused blood is from increased iron absorption.\textsuperscript{31} When total body iron is significantly elevated the transferrin iron-binding ability is exceeded and circulating non-transferrin-bound iron (NTBI) appears in the plasma. NTBI rises markedly when the transferrin saturation reaches high levels, and a reactive Fe\textsuperscript{2+} subspecies of NTBI called labile plasma iron (LPI) increases concomitantly.\textsuperscript{32,33} It is important to underline that clinical observations suggest that NTBI was absent in NTDT patients without previous transfusions.\textsuperscript{32,33} Further, LPI was not detected in most NTDT patients, whereas other iron storage and turnover markers were elevated.\textsuperscript{32,34} Nevertheless, in both TDT and NTDT patients, iron chelation is the only treatment to eliminate excess iron and prevent organ damage.\textsuperscript{31} In both these scenarios, iron chelation is the only treatment to eliminate excess iron and prevent organ damage.\textsuperscript{31}

Owning to extensive studies in mouse models and humans, many novel drugs and genetic approaches are now in development to improve the quality of life or to provide a cure to patients with BT. These drugs can be loosely classified based on their mechanism of action, such as the improvement of red cell quality and production (eg, luspatercept, mitapivat, and drugs that target excess alpha-globin chains), improvement of anemia, suppression of erythropoiesis and improvement of splenomegaly (eg, JAK2 inhibitors), limitation of iron absorption and/or erythroid iron intake (eg, hepcidin agonists, hepcidin inducers, ferroportin or ERFE inhibitors, and TF therapy), all which lead to improvements of iron overload, and, potentially, improvement of red cell production over time (Table 1).\textsuperscript{35,36} In addition, novel gene therapy approaches and strategies aimed to pharmacologically reactivate fetal hemoglobin have the potential to significantly improve symptoms of BT or to cure BT, but these topics are beyond the scope of this review.

**Drugs that improve red cell quality and production**

Luspatercept (Reblozyl\textsuperscript{®}) is a ligand trap made using the extracellular domain of the activin receptor type IIB (Table 1).\textsuperscript{37,38} This drug binds to a subset of TGF-beta ligands to inhibit aberrant SMAD2/3 signaling and enhances late-stage erythropoiesis.\textsuperscript{37-39} Luspatercept has recently been approved in the United States and Europe for the treatment of adult patients with transfusion-dependent BT (TDBT).\textsuperscript{37} During this clinical trial, a significantly greater proportion of patients in the luspatercept group had a reduction in transfusion burden compared with the placebo group.\textsuperscript{37} Additional observations also suggest that patients with non-transfusion-dependent BT (NTDBT) may also benefit from this drug.\textsuperscript{38} Although luspatercept treatment
reduced transfusion burden and iron loading in patients affected by TDBT, NTDBT, or myelodysplastic syndrome (MDS), likely caused by improved erythropoiesis and iron consumption, luspatercept treatment also increased ERFE levels and, consequently, reduced hepcidin levels.\(^40\) Therefore, it is possible that luspatercept may negatively modulate hepcidin transcription by binding TGF-beta family ligands implicated in hepcidin synthesis.\(^41\) These observations suggest it could be beneficial to use luspatercept in combination with iron chelators or other therapeutics, as elaborated in the section “Anemias associated with impaired iron absorption.”

Mitapivat (AG-348; Pyrukynd\(^6\)) is a small-molecule activator of R-type pyruvate kinase that enhances the affinity of R-type pyruvate kinase for its substrate, phosphoenolpyruvate (Table 1).\(^42,43\) Mitapivat was initially developed to treat patients with pyruvate kinase deficiency, as it can improve pyruvate kinase activity, increase red cell deformability, and reduce transfusion burden in some patients that receive regular blood transfusions.\(^44\) Additional observations demonstrate that mitapivat induces a rapid and sustained increase in hemoglobin levels in approximately 50% of patients with pyruvate kinase deficiency who did not receive regular blood transfusions.\(^43\) In mouse models of BT, oral mitapivat administration ameliorated ineffective erythropoiesis and anemia by improving red cell metabolism.\(^9\) Likely caused by improved erythropoiesis, mitapivat administration resulted in reduced levels of soluble ERFE, increased liver hepcidin expression, and diminished liver iron overload.\(^9\) Mitapivat also reduced duodenal iron absorption, suggesting that this drug could further reduce iron overload independent of its indirect effect on hepcidin expression.\(^9,45\) Ongoing observations in patients with non-transfusion-dependent AT (NTDAT) and NTDBT treated with mitapivat suggest that this drug could ameliorate anemia in these patients.\(^45\)

Other studies aim to target pathways that enhance the removal of excess alpha-globin in BT erythroid precursor cells (Table 1).\(^24\) This type of detoxification strategies allow BT to be treated similar to other protein aggregation disorders.\(^46\) Pharmacological induction of protein-targeted quality control pathways and autophagy to eliminate relevant unstable proteins could be potentially beneficial in patients with BT.\(^47,48\) As mTORC1 inhibits autophagy, systemic treatment with the mTORC1 inhibitor rapamycin reduces alpha-globin precipitates and lessens pathologies in BT mice.\(^49\) These findings may allow investigation of additional pathways that can be regulated by drugs for amelioration of BT symptoms. However, we do not know if rapamycin or similar drugs also improve iron metabolism in patients with BT, although rapamycin has been shown to increase hepcidin expression in wild-type mice.\(^50\)

Selective deletion of Tfr2 in exclusively cells of the bone marrow in both NTDBT and TDBT mice significantly ameliorated anemia and highlight a major role for TFR2 in modulating the relationship between erythroid delivery of iron by TF, EPO sensitivity, and ineffective erythropoiesis (Table 1).\(^51,52\)

**Drugs that limit erythropoiesis to reduce splenomegaly**

In BT, in response to anemia and hypoxia, high levels of EPO are associated with increased activation of the JAK2 pathway and increased numbers of erythroid progenitors in both the bone marrow and the spleen.\(^53\) This extramedullary hematopoiesis leads to hepatosplenomegaly, which enhances the entrapment of red cells in the spleen and exacerbates the anemia, further worsening the hepatosplenomegaly.\(^27,53\) Additionally, erythropoiesis occurs at extramedullary sites, most commonly resulting in a paraspinal mass.\(^54\) Unfortunately, splenectomy considerably increases the rate of thromboembolic complications.\(^34\) Although administration of JAK2 inhibitors does not reduce transfusion
burden, the combination of blood transfusions with JAK2 inhibitors could be effective in reversing splenomegaly and therefore could be used as an alternative to splenectomy (Table 1).55,56

Drugs that limit iron absorption and heme synthesis

Identification of hepcidin and elucidation of the pathways that control its synthesis have been crucial to the development of pharmacological compounds that mimic the activity of hepcidin, increase hepcidin expression, decrease ferroportin or ERFE activity, which all lead to decreased iron absorption and TF saturation.36 In a similar fashion, administration of exogenous apo-TF can also decrease TF saturation levels.57 These potential therapeutics have reduced iron overload in animal models of BT.36 Intriguingly, in BT mice, the same drugs also lead to reduction in anemia.35,58 These drugs decreased not only iron absorption but also erythroid iron intake.35,59 In thalassemic erythroid progenitor cells, reduced iron uptake reduces the detrimental effects of oxidative stress triggered by the excess iron and heme that are not included in functional hemoglobin molecules.35,58 Overall, the consequences of iron-restriction to erythroid cells improve the quality and lifespan of mature red cells and increase circulating hemoglobin levels.35,58 An ERFE antibody is also a potential therapeutic that targets a component of the iron-regulation pathway, as it prevented hepcidin suppression and corrected the iron-loading phenotype in a mouse model of BT.60

These observations suggested that iron-restricted erythropoiesis could benefit BT patients in a similar fashion, and a series of clinical trials were designed using drugs that restrict iron absorption. The hepcidin mimetic rusfertide® (PTG-300) can reduce iron absorption by targeting ferroportin.61 Rusfertide treatment improved iron metabolism and erythropoiesis in BT mice and decreased serum iron levels in patients with TDBT (https://www.protagonist-inc.com/investors-media/press-releases/default.aspx) but failed to improve symptoms in patients that received regular blood transfusions.62 Similarly, the ferroportin inhibitor vamifeport® (VIT-2763) showed similar endpoints in BT mice. However, also in this case it failed to improve symptoms in NTDBT patients (https://thalassaemia.org.cy/fr/clinical-trial-updates/ferroportin-inhibitors/#:~:text=A%20randomized%2C%20controlled%2C%20multicentred%20phase%202%20proof-of-concept%20trial%2C%20serum%20iron%20for%20up%20to%2024%20hours%20post-dose).63 Since in these clinical trials, administration of these reagents was associated with decreased transferrin saturations levels, it is possible that the biology of human red cells may be less sensitive to the deleterious effect of hemichromes or that the effects on erythroid iron restriction on the viability of BT red cell may surpass the beneficial effects of reducing hemichrome formation. However, the ability of these reagents to modulate hemichromes formation in the red cells of BT patients has never been investigated. Although these drugs failed to improve anemia in BT patients, it is important to underline that they were efficacious in reducing dietary iron intake and transferrin saturation, which is a prerequisite to improving iron overload. This suggests that these drugs could be used to prevent iron overload or could be combined with other medications that improve erythropoiesis or reduce iron overload (eg Luspatercept, Mitapivat or iron chelators) to rapidly reduce iron overload, as shown in animal studies.35,64,65

Bitopertin is an oral inhibitor of glycine transporter type 1, a key membrane transporter required to supply red cells with sufficient glycine to support erythropoiesis.66 By limiting glycine uptake, bitopertin regulates downstream heme synthesis.66 For instance, in a mouse model of erythropoietic protoporphyria, bitopertin reduced the excessive production of protoporphyrin IX (a precursor of heme) and improved liver fibrosis.67 Based on these preclinical studies, bitopertin is currently being tested in patients with erythropoietic protoporphyria.67 Bitopertin also limited abnormal levels of heme and reduced anemia in BT
mice. However, a clinical trial in patients with NTDBT failed to reverse the anemia, with decreased levels of mean corpuscular hemoglobin and hemoglobin.

**Additional conditions that may benefit from modulation of erythropoiesis or suppression of iron absorption**

### Alpha thalassemia

Patients with severe forms of AT who require chronic blood transfusion therapy for survival (transfusion-dependent AT (TDAT)), almost exclusively make hemoglobin H, which is a tetramer made of 4 beta-globin chains that has extremely high oxygen affinity but has poor oxygen delivery to tissues. These patients share several pathophysiological manifestations with patients with TDBT, such as iron overload and the requirement for iron chelation therapy. However, patients with TDAT require a more aggressive transfusion regimen than patients with TDBT. If not, patients with TDAT continue to have a high proportion of circulating nonfunctional hemoglobin H and exhibit features of hypoxia and EPO-driven increased erythropoietic activity. These observations suggest that ineffective erythropoiesis in patients with TDAT may be qualitatively and quantitatively different from that observed in patients with TDBT, with potentially more enucleated but nonfunctional cells in circulation and increased iron absorption. If these predictions are confirmed by future studies, this will indicate that drugs that alter red cell quality may not be efficacious in patients with TDAT, although therapeutics that limit iron absorption or splenomegaly (such as JAK2 inhibitors) may be beneficial (Table 1).

In contrast, the less severe forms of AT (ie, NTDAT), where some level of functional hemoglobin is present, may benefit from the same therapeutics that are being tested for patients with NTDBT, such as luspatercept and mitapivat. In a preliminary study, patients with NTDAT seemed to benefit from administration of mitapivat. As is the case with severe forms of AT, other drugs that limit iron absorption or erythroid iron intake could also be effective in the less severe forms of AT associated with iron overload (Table 1).

### Polycythemia vera

There are conditions where modulation of iron metabolism could be beneficial to reduce excessive production of red cells. Mutations in JAK2 can lead to PV, which is characterized by erythrocytosis, bone marrow erythroid and megakaryocytic hyperplasia, fatigue, and splenomegaly. Patients with PV are frequently iron deficient at the time of diagnosis, which is further exacerbated by therapeutic phlebotomies for the maintenance of hematocrit below 45% to decrease thrombotic risk.

Studies in PV mice demonstrated that administration of exogenous hepcidin or induction of endogenous hepcidin using TMPRSS6-ASO, an antisense oligonucleotide to TMPRSS6 messenger RNA, was able to reverse erythrocytosis, decrease splenomegaly, and sequester iron in splenic macrophages, suggesting that the approaches that limit erythroid iron absorption and/or heme synthesis could be beneficial in patients with PV (Table 1). Additionally, preliminary results from clinical trials that aimed to evaluate the safety and efficacy of the hepcidin mimetic rusfertide in phlebotomy-requiring patients with PV suggest that approaches that limit erythroid iron absorption and/or heme synthesis could eliminate phlebotomy requirements, increase systemic iron stores, and decrease systemic symptoms.

### Other conditions
In brief, some conditions, such as congenital dyserythropoietic anemia type II (CDA-II), MDS, and Diamond-Blackfan anemia (DBA), may benefit from therapeutics that modulate erythropoiesis, erythroid iron intake, heme synthesis, and/or decreased dietary iron absorption. For instance, a subset of patients affected by MDS benefit from luspatercept treatment and may also benefit from iron restriction. Similar approaches may also benefit a subset of patients with CDA-II (mutations in \(SEC23B\)) who have anemia associated with increased ERFE expression, decreased hepcidin levels, and iron overload. Finally, erythroid cells in patients with DBA synthesize excess heme, which could be reduced by limiting erythroid iron intake and/or heme synthesis by treatment with a drug such as bitopertin.

Anemias associated with impaired iron absorption

Other forms of anemia can help us to further understand the relationship between ineffective erythropoiesis and iron metabolism. In this part of the review, we will discuss informative forms of anemia and speculate on novel potential drugs to treat them.

Anemia of inflammation (AI) and IRIDA are two of the clinical manifestations associated with inappropriately high hepcidin levels. AI is common in inflammatory diseases with complex pathophysiological features, such as CKD, autoimmune diseases, and some forms of cancer and myelofibrosis. Here, we review conditions in which inflammatory cytokines together with hepcidin contribute to iron-restricted erythropoiesis. The primary clinical goal for patients with AI is to treat the cause of inflammation; however, treatment for anemia is often managed by administration of intravenous iron and erythropoiesis-stimulating agents, which is ineffective for some patients and is associated with adverse effects.

Myelofibrosis is a myeloproliferative neoplasm characterized by splenomegaly, debilitating constitutional symptoms, and bone marrow failure. Disease-related anemia is common and associated with increased expression of hepcidin. IRIDA develops when mutations in \(TMPPRS6\) lead to high hepcidin levels. Patients with IRIDA have microcytic anemia and low plasma-iron levels and are refractory to oral iron therapy but are partially responsive to parenteral iron administration.

In all these conditions, high levels of hepcidin not only limit iron absorption but also sequester the iron provided to the patients by oral or intravenous administration to parenchymal cells and macrophages. Therefore, drugs that target proteins that interfere with the iron-sensor complex and decrease hepcidin levels or that can redistribute iron to TF may make iron more available for erythropoiesis. In addition, drugs that increase EPO sensitivity could also improve red cell production.

Activin-like receptor 2 (ALK2) belongs to the TGF-beta receptor superfamily and is part of the iron-sensor complex. ALK2 is the target for the drugs momelotinib and LJ000328. These drugs can downregulate hepcidin expression and increase availability of iron for erythropoiesis. In clinical testing, momelotinib improved hemoglobin levels and reduced transfusion burden in patients with myelofibrosis and baseline anemia, while also reducing spleen size and symptom burden. In mice, LJ000328 repressed hepcidin activity and significantly reduced the symptoms of IRIDA.
Mutations in \textit{HJV} are associated with a severe form of hemochromatosis.\textsuperscript{20,92} Therefore, pharmacological targeting of HJV could lead to hepcidin suppression. Anti-HJV antibodies, such as DISC-0974, have been effective in suppressing hepcidin expression in normal individuals and in preclinical models of AI and IRIDA, where the antibodies were shown to increase hemoglobin levels in both inflammatory and noninflammatory states.\textsuperscript{93-96}

It has been recently observed that luspatercept treatment in patients with BT or MDS resulted in the release of excess iron from stored organs into peripheral blood.\textsuperscript{40} This was associated with decreased hepcidin levels, suggesting that luspatercept (which targets a subset of TGF-beta ligands) may also inhibit the activity of the iron-sensor complex, possibly by targeting BMP or BMP-like molecules involved in hepcidin synthesis.\textsuperscript{41} These observations provide a mechanistic rationale to also evaluate luspatercept in diseases of elevated hepcidin synthesis, such as AI and IRIDA.

Deferiprone (Ferriprox\textsuperscript{8}) does not directly affect iron absorption, but it chelates iron and is used to treat iron overload in BT.\textsuperscript{97} Deferiprone readily reaches the major intracellular sites of iron accumulation, facilitating the extraction of iron deposited in organs.\textsuperscript{98} Deferiprone has a lower iron affinity than TF, so deferiprone could redistribute iron sequestered in organs to erythroid cells.\textsuperscript{99} Therefore, in patients with high hepcidin levels, deferiprone could shuttle the sequestered iron to unbound TF, paradoxically improving erythroid iron consumption and anemia in AI and IRIDA. This strategy may also require concurrent iron administration to provide the optimal amount of iron needed for efficient erythropoiesis.

Finally, it could be beneficial to patients with many of the above-mentioned conditions to target TFR2 on erythroid cells, as deletion of TFR2 in hematopoietic cells can improve EPO sensitivity and increase red cell production.\textsuperscript{100} Because deletion of TFR2 in the liver decreases hepcidin expression and leads to increased iron absorption, it is currently challenging to selectively target TFR2 to erythroid cells and improve anemic conditions associated with iron overload (eg, AT and BT). However, it may be beneficial to target TFR2 on both erythroid cells and liver cells in conditions in which an increase in both EPO sensitivity and iron absorption may be beneficial (ie, CKD).\textsuperscript{101}

\textit{Additional emerging technologies}

Current nucleoside-modified mRNA-lipid nanoparticle technology has successfully paved the way for next-generation vaccinations against SARS-CoV-2 during the COVID-19 pandemic.\textsuperscript{102} This emerging technology is also being used for additional applications, such as the ability to deliver small interfering RNA, messenger RNA, or plasmid DNA to treat most genetic diseases by silencing pathological genes, expressing therapeutic proteins, or gene editing. Using unmodified or targeted lipid nanoparticles encapsulating mRNA, several studies are underway to treat a variety of non-liver and liver diseases.\textsuperscript{103,104} As the pathways that control iron metabolism are primarily localized to the liver, it is only a question of time before this technology will be applied to modify the expression of iron-related genes and to improve iron overload or to directly correct genetic defects, as in IRIDA or HFE-related hemochromatosis.

\textit{Combinatorial therapies and conclusion}

Future clinical trials should clarify which drugs benefit all or a subset of patients with diseases associated with defective erythropoiesis. Combination therapies may emerge as the best approach, not only to improve anemia and iron overload but also to enhance treatment efficacy in a larger number of
In conclusion, a variety of promising novel drugs may expand the armory of therapeutics available to patients with diseases associated with defective erythropoiesis. If proven to be safe, selective, and effective, these drugs will increase the chance for treatment success, and competition between drug companies will likely diminish treatment costs.
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<td>Improve red cell quality/Pyruvate Kinase inducer</td>
<td>Pyruvate Kinase/Pyrulkynd (Mita pivat)</td>
<td>+</td>
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<td>+</td>
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<tr>
<td>Inhibition of autophagy</td>
<td>mTORC1/Rapamycin or similar</td>
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<tr>
<td>Decrease iron absorption</td>
<td>FPN/Rusfertide (PTG-300)</td>
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<tr>
<td>Decrease iron absorption</td>
<td>TMPRSS6 mRNA/TMPRSS6-ASO</td>
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<tr>
<td>Decrease iron absorption</td>
<td>Ferroportin/Vamifeport (VIT-2763)</td>
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<tr>
<td>Decrease iron absorption</td>
<td>ERFE</td>
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<tr>
<td>Decrease TF saturation</td>
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<td>Modulate heme synthesis</td>
<td>Glycine Transporter 1 Inhibitor/Btopertin</td>
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<tr>
<td>Increase iron absorption</td>
<td>HJV/DISC-0974</td>
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<tr>
<td>Increase iron absorption</td>
<td>ALK2/Momelotinib/LU000328</td>
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<tr>
<td>Increase TF saturation</td>
<td>DFO</td>
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DFO: deferiprone; Myel.: myelofibrosis.
Table 1
Potential use of drugs already approved or under development for diseases discussed in this review. The check mark √ indicates the use of already approved drugs. The plus sign (+) indicates drugs in use in preclinical studies or discussed in this review. The question mark (?) indicates the potential use of the drugs in other conditions, but these drugs have not been discussed in this review. The sign – indicates not applicable or not determined. The plus signs and question marks in this table are purely speculative. The use of these drugs under these conditions would need to be tested in animal models and preclinical trials to prove their efficacy and safety.

Figure legends

Figure 1
The pathways of a subset of the molecules involved in red cell production and iron metabolism. Only those molecules discussed in this review are included in this schematic.

Figure 2
Schematic representation of normal and beta thalassemic erythropoiesis. A few consequences associated with ineffective erythropoiesis in BT, such as iron overload and hepatosplenomegaly, are presented.