The pathophysiology of transfusional iron overload

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ABSTRACT (247)

The pathophysiological consequences of transfusional iron overload (TIO) as well as the benefits of iron chelation are best described in Thalassemia Major (TM), although TIO is increasingly seen in other clinical settings. These consequences broadly reflect the levels and distribution of excess storage iron in the heart, endocrine tissues and liver. MRI-visible storage iron does not directly damage cells, but its intracellular turnover contributes to labile iron pools that generate harmful free radicals. TIO also increases the risk of infection, due to increased availability of labile iron to microorganisms. Although storage iron accumulates firstly and predominantly in the liver, heart failure from myocardial iron loading typically precedes the hepatic complications of cirrhosis and hepatocellular carcinoma by at least two With improved chelation, decreased cardiomyopathy and decades. increasing survival, hepatic complications are more commonly encountered. Storage iron distribution reflects the pattern of transferrin-independent iron uptake (NTBI), which in animal models has been linked to L-type calcium channels. The propensity of iron overload to distribute extra-hepatically differs between underlying clinical conditions. Thus Sickle Cell Disease (SCD) patients have a lower risk of myocardial and endocrine iron deposition than TM and also have disproportionately low NTBI levels. Conversely, Diamond-Blackfan Anemia (DBA) patients are prone to extra-hepatic iron deposition, and have high levels of NTBI. consistent with low transferrin iron utilization. We suggest that extra-hepatic iron distribution, and hence toxicity, is influenced by balance between generation of NTBI from red cell catabolism and the utilization of transferrin iron by the erythron.

IRON HOMEOSTATIC MECHANISMS (5004 WORDS)

Iron homeostatic mechanisms are key to the pathophysiology of TIO. In humans, these mechanisms are best adapted to increasing iron acquisition in conditions of iron deficiency or anemia, or to limiting iron distribution from the macrophage system during inflammation. They are not well adapted, however, to controlling the distribution of TIO or to eliminating excess iron. This is in marked contrast to rodents, where most studies on iron overload and iron metabolism have been performed and where iron overload is eliminated efficiently by the biliary route. Iron homeostasis is adapted to supplying only that which is essential for the functioning of proteins involved in oxygen transport, oxidative energy production, mitochondrial respiration and DNA synthesis, while minimizing the potential for iron toxicity from its redox cycling. These homeostatic mechanisms work at two levels: firstly at the level of the whole body through interactions of plasma hepcidin with membrane ferroportin and secondly at a cellular level through interaction of IRE-binding proteins (IRPs) with iron responsive elements (IREs) present on mRNAs of key iron metabolism-related proteins.

Body iron homeostasis

A healthy human contains 40-50mg/kg of iron, mainly as hemoglobin (30mg/kg). About 4mg/kg is present in muscle myoglobin, with 2mg/kg in cells as iron-containing enzymes. Storage iron, present as ferritin and its compact, partially degraded form hemosiderin, ranges from 0 to 2000mg.¹ This is mainly present in liver, spleen and bone marrow (BM) macrophages, formerly referred to as the reticulo-endothelial system (RES) and in hepatocytes². Liver iron concentration (LIC) rarely exceeds 1.8mg/g dry weight (dw) in healthy individuals in the absence of liver disease, hemochromatosis genes, inappropriate dietary supplementation or blood transfusion.

A healthy individual absorbs only about 10% of dietary iron or about 1-2mg/day, usually balanced by iron loss from skin, gut, menstruation or pregnancy. Anemia, hypoxia, ineffective erythropoiesis (IE), and the presence of variant HFE genes increase iron absorption, the common factor being low, or inappropriately low, plasma hepcidin levels³. The latter permit higher enterocyte ferroportin expression, allowing Fe(II) flux and hence increasing dietary iron absorption. Iron absorption is also increased through hypoxia-inducible factor-1-mediated signaling, by duodenal up-regulation of DcytB and DMT1 expression.⁴ Thus, in principle, any anemia will tend to increase the efficiency of iron absorption. The majority of body iron turnover however, is not directed through iron absorption, but through the plasma transferrin, which, while binding only 1-2mg of iron at any moment, in a healthy adult delivers about 20-30mg/day *via* transferrin receptors on the erythron for hemoglobin synthesis.

Hepcidin regulation is important both to iron absorption from diet and to iron egress from erythrophagocytic macrophages. Hepcidin controls iron egress from both macrophages and enterocytes by binding to and degrading

ferroportin, through which Fe(II) exits these cells^{5,6}. Hepatic hepcidin synthesis is controlled by at least 3 distinct regulatory mechanisms responsive to levels of iron, erythropoiesis or inflammation.

- Extracellular iron sensing involves the binding of diferric transferrin to transferrin-receptor-1 (TfR1), initiating the translocation of HFE from TfR1 to TfR2 and its subsequent signaling via ERK1/ERK2 and p38 MAP kinase to induce hepcidin expression. Storage iron sensing is effected by BMP6 signaling via BMP receptor (and SMADs pathway) whose sensitivity is markedly increased by its interaction with hemojuvelin, HFE and TfR2 in holotransferrin dependent manner, thus enhancing hepcidin transcription^{7,8}.
- Erythropoiesis sensing involves BM-derived factors that suppress hepcidin synthesis; conditions with high levels of IE will have high levels of these factors, the nature of which has been debated. These include GDF15⁹, twisted gastrulation factor-1¹⁰, and most recently erythoferrone¹¹ which has been identified as a key factor in mice, although its relevance in humans has yet to be demonstrated. Another separate erythropoiesis sensing mechanism likely involves desaturation of transferrin by the erythron as disruption of transferrin iron uptake into erythron in *hbd* mouse increases hepcidin despite ongoing anemia,^{8,12} presumably overriding the erythropoietic depressors of hepcidin (or demonstrating that depressors have an effect only with concomitant transferrin desaturation).
- Inflammation sensing mediated through IL-6/STATs pathway and other cytokines up-regulates hepcidin synthesis,¹³ being the key mechanism in hypoferremia of acute inflammation through the action of ferroportin degradation in macrophages⁵.

Erythropoietic drive overrides both iron sensing¹⁴⁻¹⁶ and inflammation sensing mechanisms¹⁷, but the extent to which these potentially opposing regulators of hepcidin synthesis play out in the context of iron metabolism in TIO are difficult to predict from murine studies alone and require careful clinical observations. These are discussed below under the clinical condition in question

Cellular iron homeostasis

Intracellular iron homeostasis is controlled not only by the synthesis of ferritin but also by the regulation of iron uptake through regulation of membrane transferrin receptors (Tfr). Both of these are regulated by IREs, stem-loop structures present in untranslated regions (UTR) of mRNA e.g. in the 5'UTR of H-ferritin or the 3'UTR of TfR mRNA, respectively. These are both sensitive to the magnitude of labile intracellular iron pools (LIP) through interaction with cellular IRPs; the conformation of IRPs and their binding to IREs are sensitive to LIP concentrations.¹⁸ IRE binding of both IRP1 and IRP2 increases in iron-deficient conditions, but both are rapidly degraded by iron and heme. IRP2 has predominant control overall,¹⁹ while IRP1 can switch from aconitase activity form in iron repletion (dependent on iron-sulphur cluster assembly, 4Fe-4S) to IRE-binding form in iron deficiency (losing iron: 3Fe-4S).²⁰ Therefore both their cellular level and the position of IRE on mRNA regulate in concert the onset and degree of translation events (5' UTR governing

access to matrices, 3'UTR governing stability of matrices by regulating the binding of nucleases). High levels of LIP thus increase ferritin synthesis while decreasing the membrane expression of TfR1. However, in the erythron such feedback is absent; instead *transcriptional* control permits high TfR despite high cellular iron or heme, consistent with hemoglobin synthesis requirements. Most ferroportin transcripts also contain IRE at 5'UTR, and therefore the amount of mRNA is increased in iron overload. However, the effective regulation of ferroportin happens post-translationally through hepcidin-dependent down-regulation⁶ or lack of Fe(II) acceptor.²¹

The availability of iron for the synthesis of iron containing molecules at a cellular level is directed through a transient low molecular weight iron pool, LIP, which in turn determines the levels and action of IRPs. Although LIP iron has been proposed to be coordinated mainly by glutathione from a thermodynamic perspective²², its exact nature still remains unclear, but it can potentially redox cycle between Fe(II) and Fe(III) with consequent generation of harmful free radicals. In order to minimize these risks, elegant homeostatic mechanisms carefully coordinate the distribution of body iron so as to provide iron pools for efficient synthesis of these proteins, while minimizing iron-mediated free radical generation.

IMPACT OF BLOOD TRANSFUSION ON IRON BALANCE

The rates and nature of blood transfusion regimens affect iron accumulation and its distribution in the body. This is key to the pathophysiology of iron overload and varies with the underlying clinical condition.

Thalassemia Major

In thalassemia major (TM), blood transfusion typically begins in the first year of life. Current transfusion recommendations in TM aim to keep the pretransfusion hemoglobin level at approximately 9.5g/dl and to maintain an average hemoglobin of 12g/dl,²³ which usually amounts to an iron load rate (ILR) of 0.3-0.5mg/kg/d.²³ This regimen has been arrived at so as to balance the beneficial effects of suppression of IE and dietary iron absorption with the iron accumulated from transfusion. The transfusional suppression of the endogenous BM activity can be assessed by monitoring circulating transferrin receptors, which show more suppression when the pre-transfusion hemoglobin level exceeds 10g/dL.²⁴ Maintenance of a mean pre-transfusion hemoglobin level of 9.4g/dl versus 11.3g/dl decreased net blood consumption and was associated with improved control of iron overload in Italian patients.²⁵ This optimal balance may not be universal and may depend on the severity of thalassemia genotype. In the pre-chelation era, LICs of 40mg/g dw were typically seen by 10 years of age.²⁶ Failure to control these levels risks extra-hepatic spread of iron (see below).

Sickle cell disease

The age of commencing blood transfusion, transfusional ILR, and the nature of the transfusion regimen itself, all affect the rate and extent of iron overload

in SCD and often differ considerably from TM. Net iron accumulation from transfusion in SCD is slower than TM, firstly because of differences in transfusion practice between these conditions and secondly because SCD patients tend to be in negative iron balance in the absence of transfusion. In SCD there is considerable intravascular hemolysis leading to iron loss via urine²⁷⁻²⁹ (as in PNH) and possibly bile³⁰. Urinary iron loss in SCD may reach as much as 15 mg/d²⁸ (approx. 0.2mg/kg/d i.e. comparable to average SCD transfusional ILR). Furthermore the marrow is less expanded in SCD than in TM or NTDT, leading to less hepcidin suppression and less tendency for increased iron absorption (Porter 2014, in preparation). Under conditions of hypertransfusion, where synthesis of HbS is suppressed, or under vigorous chronic automated exchange procedures, were the %HbS is maintained at low levels, intravascular hemolysis will also be suppressed and thus the tendency to lose iron through this mechanism will be diminished.

Historically, blood transfusions were typically sporadic and given by simple transfusion or by some form of partial exchange procedure in response to acute episodes, which over a lifetime would lead to significant iron overload. Transfusion has been increasingly given to prevent primary and secondary stroke.³¹ This approach, together with a wider use of transfusion to prevent or treat other complications, such as chest syndrome, or in preparation for major surgery, puts an increasing proportion of patients at risk of TIO. In a large multicenter international study, where most patients received simple (60%) or exchange transfusions (20%), the mean ILR was 0.22mg/kg/day³², notably lower than in TM. Manual exchange procedures, where about 1/3 of the blood volume is exchanged, lead to ILR of about 40% of simple transfusions, as estimated from ferritin increments.³³ With automated erythrocytapheresis, ILR was only 0.053mg/kg/day with a target pre-transfusion HbS<50%;³⁴ this compared with 0.39mg/kg/day for simple transfusion with a target HbS<30% and 0.29mg/kg/day with a target HbS<50%.

Other conditions

In other forms of TIO, the rates of ILR again vary considerably; for example a mean of 0.4mg/kg/day was found in transfusion-dependent DBA patients with 0.28mg/kg/day in MDS patients in the same study³⁵. Patients who receive repeated myelo-ablative chemotherapy cycles for leukemias or lymphomas can accumulate over 100 units of transfused blood or 20g of excess body iron that will eventually require removal if long-term iron toxicity is to be avoided.

MECHANISMS OF IRON TOXICITY IN TRANSFUSIONAL OVERLOAD

The pathophysiological consequences of TIO are summarized in **Figure 1**. These are broadly observed in tissues in which storage iron accumulates at the highest concentrations. Ferritin within cells is degraded in lysosomes or proteasomes, the iron is released into LIP and this iron is reincorporated into new ferritin synthesis or made available for synthesis of essential iron containing proteins.²⁰ Once the LIP reaches a critical concentration,³⁶ the iron

can redox cycle between ferric Fe(III) and ferrous Fe(II) forms through the donation or acceptance of an electron and enhance the generation of reactive oxygen species (ROS), with a cascade of consequences (**Figure 1**). Both the concentration of LIP, as well as the capacity of cells to accommodate increased levels of iron are likely to vary between cell type and the exact nature and redox state of LIP remain unresolved. For example in the human K562 cell line LIP concentrations of 0.24-0.4µM have been estimated using the fluorochrome calcein as a probe³⁷. However using EPR spectroscopy which detects Fe(III) and requires no manipulation of cells, an intracellular EPR-detectable high-spin ferric iron signal was found at approximately 3.2µM^{38,39}. More recently, increased levels of LIP have been linked to increased ROS production and potentially oncogenic effects.⁴⁰ Ferritin acts as a sink for LIP by decreasing its magnitude and its potential toxicity. For example, murine erythroleukemia cells overexpressing H-ferritin displayed lower levels of LIP and ROS⁴¹.

Not all ROS are necessarily toxic to cells however. Large quantities of superoxide are produced naturally by respiration (about 30g/day of superoxide) but their toxic potential is controlled by their conversion to water by superoxide dismutase and glutathione peroxidase. It is the favorable redox potential of the Fe(II)/Fe(III) couple (between +0.35 and -0.5V) that allows it to redox-cycle and thus to catalyze the interaction of superoxide with hydrogen peroxide (H₂O₂) through the Haber-Weiss reaction, generating highly reactive hydroxyl free radicals (OH).⁴² The hydroxyl radical has a great affinity for electrons, will oxidize all substances within its immediate vicinity (diffusion radius of 2.3nm),⁴² and has been shown to promote lipid peroxidation,^{43,44} with damage to organelles such as lysosomes⁴⁵ and mitochondria.^{46,47} The interaction of the hydroxyl radical with lipid proceeds though the initial abstraction of a hydrogen atom (to yield a water molecule), molecular rearrangement of the lipid with peroxidation, and the formation of a peroxyl radical which is able to propagate further lipid peroxidation in a chain reaction. The end result is decomposition of lipid molecules with concomitant effects on the integrity of organelles. While organelle damage may lead directly to apoptotic cell death, 48,49 this may also encourage fibrogenesis as iron-induced aldehyde lipid peroxidation products such as MDA⁵⁰ and 4-HNE⁵¹ promote collagen gene expression. Fibrogenesis is also associated with autocrine production of TGF β -1 in stellate cells⁵² (**Figure 1**). ROS also damage DNA, risking genomic instability, mutagenesis and cell death or neoplasia. ROS also directly activate caspases, thereby accelerating apoptosis,53 but, paradoxically, may also have anti-apoptotic effects by activating NF-kB,54 which may contribute to MDS transformation and to ironmediated neoplasia, such as hepatoma.

An important, often neglected, mechanism of toxicity from iron overload is that of the increased risk of infection, which is the second commonest cause of death in TM. Several mechanisms come into play, the most important being transferrin saturation. This protein, in addition to its pivotal role in supplying iron to the erythron and other tissues, naturally exists where only an average of 1/3 of its two iron binding sites are occupied with Fe(III). A key role of Tf is to deprive bacteria of the iron that these microorganisms require to grow. While some bacteria have adapted to utilize transferrin iron, most have not, and so there is a paradigm shift in the availability of iron to microorganisms once transferrin becomes saturated. Other mechanisms, such as effects on neutrophil function have been postulated to be affected by TIO.⁵⁵ Recent work has shown that following blood transfusion, NTBI is liberated from the rapid catabolism of a proportion of non-viable red cells⁵⁶. In principle, this and other forms of NTBI in plasma will be more available to microorganisms than transferrin iron.

DISTRIBUTION AND CONSEQUENCES OF TIO

Iron distribution and consequences in Thalassemia Major

The impact of chronic blood transfusion on body iron distribution is most completely described in TM, where transfusion typically begins in the first year of life. Transfused iron initially accumulates as storage iron in spleen, liver and BM macrophages and later in hepatocytes, with 80% of storage iron in the liver. The storage capacity of the macrophage system following blood transfusion has not been recently studied, but historical sources estimate it at about 10g. Histological descriptions using Perl's stain show that with increasing TIO, increasing proportions are seen in hepatocytes once the macrophage system is saturated. Interestingly, particularly with optimal chelation therapy, TM patients today have low iron concentrations in hepatocytes while macrophage iron remains present⁵⁷. This contrasts with NTDT where iron accumulates through the portal system and concentrates in peri-portal hepatocytes with macrophage sparing. This distribution in NTDT is thought to be influenced further by low hepcidin levels, and therefore high macrophage ferroportin, due to high levels of IE typical in thalassemia.^{57,58}

As TIO evolves, particularly with sub-optimal chelation therapy, a variable proportion of iron 'escapes' from the liver into the endocrine tissues and heart. This gives rise to the classic pathology, morbidity and mortality historically associated with TIO. An understanding of the effects of blood transfusion on body iron distribution is best appreciated from *post-mortem* obtained during pre-chelation era.⁵⁹ data because iron chelation fundamentally alters body iron distribution, being relatively tropic for hepatocellular iron compared with extra-hepatic iron. Data obtained under these circumstances showed that in patients dying from complications of TIO, iron was unevenly distributed in the body, with high concentrations present in liver, heart and endocrine tissues, very low in striated muscle and none in the brain.⁶⁰ Remarkably, these patients typically died of heart failure in the second and third decades of life, although the myocardial iron concentration (MIC) was a fraction of that in the liver. This observation has recently been supported by MRI evidence;⁶¹ examination of myocardial tissue both biochemically and by MRI at *post mortem* in patients dying from iron-induced cardiomyopathy showed an average MIC of only 5.98±2.42mg/g dw. Evidently, the heart is less adapted to accommodating high concentrations of storage iron than the liver, even though the storage iron is not directly toxic to cells (see below).

Prior to the introduction of cardiac MRI monitoring and newer chelation regimens, the frequency of heart failure, diabetes, hypothyroidism and falling⁶². hypoparathyroidism were all Hypogonadism (typically hypogonadotropic) is still an early and common feature of iron overload in TM, presenting with primary or secondary amenorrhea in females or poor growth and delayed puberty.⁶² Since the introduction of cardiac MRI imaging and the intensification of chelation therapy in selected patients with increased MIC (mT2*<20ms), the incidence of heart failure has fallen further. Indeed in a recent cohort analysis of patients followed for a decade by cardiac MRI and receiving individually tailored chelation, heart failure was no longer the leading cause of death and the proportion of patients with myocardial iron (mT2*<20ms) fell from 60% to 23%⁶³. Cirrhosis, which develops one or two decades after heart failure, is becoming more common as patients live longer, being present in about 50% of patients at *post mortem* and is particularly common in patients with chronic hepatitis. Similarly, hepatocellular carcinoma⁶⁴ is becoming more common.

The relationship between the accumulation of liver iron and the risk of extrahepatic spread has been the source of intense debate. Early work suggested a close relationship in TM between the control of LIC with deferoxamine and long-term outcome from cardiomyopathy⁶⁵. *Post mortem* data in other diseases in the absence of chelation also suggested a relationship between transfusional iron load rate (ILR), LIC and MIC.66,67 When cardiac T2* became available in patients receiving a variety of chelation regimens, only a weak correlation between LIC and mT2* was seen, and it was argued therefore that control of LIC was not important to control of MIC in TM and therefore to limiting potentially fatal cardiomyopathy.⁶⁸ The interpretation of the UCLH group was that this lack of correlation was mainly due to the high proportion of patients in this study having been on intensive chelation therapy with DFO, which was subsequently shown to decrease LIC at a faster rate than myocardial iron.68,69 This would therefore mask a potentially important relationship between these variables. Noetzli and colleagues⁷⁰ have somewhat clarified this issue by demonstrating the importance of longitudinal rather than cross-sectional analysis of the relationship. They showed that failure to control LIC over several years with chelation increased the risk of myocardial iron deposition. Conversely, LIC reduction had a delayed effect on decreasing the MIC. Longitudinal UK studies of the LVEF relationship to mT2* show that the risk of heart failure increases for mT2*<10ms⁶⁸, which approximates to MIC>2.7mg/g dw⁶¹. It can be concluded that LIC control in TM is important both to limiting liver damage and to controlling MIC, thus markedly reducing the risk of iron-mediated cardiomyopathy with heart failure.

Iron distribution and consequences in sickle cell disease

In patients receiving sufficient repeated transfusions to cause TIO, clinical consequences begin later than in TM, and thus effects on growth and sexual development are relatively uncommon. With transfusion, iron from erythrocyte catabolism initially accumulates in macrophages (Kupfer cells, sinusoidal compartment), but later, when the LIC exceeds 7mg/g dw, in hepatocytes⁷¹ (based on the Angelucci formula⁷², being equivalent to about 5g of transfused

iron in a 70kg adult, or about 25 units of transfused blood). Hepatocellular iron stores in SCD only approached those of the sinusoidal compartment when total liver iron levels were high (>15 mg/g dw or about 50 units of transfused red cells)⁷¹. Accumulation of liver iron without adequate chelation therapy risks fibrosis and cirrhosis,^{73,74,75}. Fibrosis has been reported as early as 2 years after initiation of transfusion and in about 1/3 of patients with LIC>9mg/g dw and in direct proportion to the LIC⁷⁵, correlating with LIC⁷⁴ in the absence of hepatitis C infection^{75,76}. The true frequency of cirrhosis in multi-transfused adult SCD patients is not clear. *Post mortem* studies found cirrhosis in 11% of all patients and in nearly half of patients who died with severe liver siderosis⁷⁷.

The extra-hepatic consequences of iron overload, particularly endocrine and cardiac effects, appear to be later or more delayed in SCD than TM. Myocardial iron deposition judged by T2* is rare^{78,79} and after more transfusion episodes in SCD than TM⁸⁰. However, post-mortem studies show iron deposition in the heart in heavily transfused patients⁷⁷. In SCD and TM patients matched for LIC, the incidence of heart disease, gonadal failure and endocrine disturbances including growth delay, appear to be less for age<20 years in SCD⁸¹. Despite these differences from TM, SCD patients are unlikely to be completely protected from the extra-hepatic effects of TIO and indeed cases of myocardial iron are reported by MRI⁸². Recent preliminary studies from the Multi Center Study of Iron Overload (MCSIO) group found evidence of increased pituitary iron in SCD with highest LIC⁸³. In one study patients with the lowest bone mass also had the highest serum iron values, although SF was within normal limits in these patients⁸⁴. Furthermore there was an inverse correlation between the estimated pituitary iron and the pituitary volume, and hence its endocrine reserve. MRI may in principle identify early pituitary iron deposition in SCD before clinical manifestations are apparent.⁸³ Increased iron has also been identified by MRI in the kidney.^{85,86} This signal is highest in non-transfused patients with high levels of LDH, lacks correlation with LIC and is higher than in TM patients, suggesting it originates from iron taken up by kidney from hemoglobin freed during intravascular hemolysis rather than iron delivered by transferrin or NTBI as a consequence of TIO. This also suggests that kidney R2* may be a biomarker for chronic hemolysis-mediated vascular complications in SCD. The extent to which this mechanism is implicated in renal damage in SCD is not clear.

Iron distribution in other forms of TIO.

TIO is seen in an increasing number of underlying conditions (**Table 1**). Iron accumulation in MDS may start even before patients become transfusion-dependent because of IE, which although variably counterbalanced by increased levels of cytokines up-regulated during infections, e.g. IL-6, may still inhibit hepcidin production⁸⁷ with subsequent increased iron absorption from the gut, as described earlier. Once transfusion begins, as with other forms of TIO, iron initially accumulates in RES and then in the liver. A key question with respect to the pathophysiology of iron overload is how rapidly iron spreads extra-hepatically and how rapidly this is likely to be problematic and hence require chelation treatment. In the pre-chelation era, patients

examined at *post mortem* had an increased risk of myocardial iron with increasing numbers of blood transfusions⁶⁶. Early studies with cardiac MRI showed and increasing risk of myocardial iron as the number of transfusions exceeded 50 units^{88,89}. Analysis of the relationship between transfusion and myocardial iron is complicated since patients have received chelation therapy in some but not all reports. Overall, in non-chelated MDS patients, iron spread to myocardium occurs after approximately 70–100 units of blood (containing 14–20g iron).^{66,88}

The prognostic impact of the resulting moderate degree of iron overload⁸⁷ for overall survival is not clear because it is difficult to clearly separate the effects of TIO from other co-morbidities associated with BM failure without prospective controlled data. The prognostic importance of transfusion dependency for overall survival in patients with Low- and Intermediate-1-risk MDS has been examined in a retrospective analysis of European MDS and AML registry data,⁹⁰ which showed that patients with >20 units transfused had a higher mortality rate (30%) within 2 years of diagnosis than transfusionindependent patients (5%). Among 705 patients followed for 2 years or until death, cardiac comorbidities were seen in 79% of chronically transfused patients versus only 54% of non-transfused MDS patients and 42% of a Medicare control population⁹¹ Although some retrospective data suggest a prognostic advantage to chelation therapy in MDS,⁹²⁻⁹⁵ this has not yet been reported in prospective studies. It is clear that response with iron balance to chelation is similar to other forms of TIO. Improved hematopoiesis has been observed in 20% of patients receiving deferasirox for 1 year.⁹⁶ Responders showed a greater decrease in serum ferritin levels, suggesting that removal of iron from the BM plays a role in hematologic improvement.

There are a large number of disparate anemias that require chronic blood transfusion (Table 1). Patient numbers reported are insufficient to draw clear conclusions about whether the risks of TIO differ from those of TM. DBA is perhaps the most common of the rare anemias, but this is a heterogeneous group of conditions with a presumed shared pathology of ribosomal protein dysfunction. Recent work by the MCSIO group suggests transfused DBA patients are particularly susceptible to the extra-hepatic consequences of iron overload (Porter 2014, in preparation). The mechanisms and implications are discussed below.

MECHANISMS UNDERLYING DISTRIBUTION OF TRANSFUSED IRON

Because the distribution of storage iron appears to be key to the pathophysiology of TIO, some of the putative mechanisms determining this distribution will be considered further.

Transferrin iron is delivered to tissues expressing TfR1 in a controlled way through receptor-mediated endocytosis. The expression of TfR and hence iron acquisition from transferrin depends on iron homeostatic mechanisms (see above). The erythron and hepatocytes are rich in these receptors as are cells undergoing proliferation where its expression is cell cycle dependent. As iron overload develops, transferrin becomes increasingly saturated, with the eventual appearance of NTBI, typically when TfSat>75%.^{97,98} The uptake of NTBI is less regulated than uptake from Tf and the distribution is also substantially different.^{99,100} This may account for the pattern of iron deposition, and hence its toxicity in advanced TIO. In experimental models, NTBI is rapidly taken up by hepatocytes¹⁰¹ and myocytes.¹⁰² In cultured heart cells, NTBI species are taken up at 200 times the rate of transferrin iron and generate free radicals, lipid peroxidation, organelle dysfunction and abnormal rhythmicity.^{102,46} L-type calcium channels⁹⁹, and zinc transporters¹⁰³ have been implicated in NTBI uptake which appears to be restricted to tissues known to accumulate iron. Plasma NTBI (or its sub-fractions) can also promote lipid peroxidation through the generation of free radicals¹⁰⁴ and associate with depletion of plasma anti-oxidants.¹⁰⁵

The presumed relationship between NTBI levels and extra-hepatic iron distribution has not been convincingly demonstrated clinically. A weak correlation between NTBI and mT2* was reported by Piga in TM patients¹⁰⁶. However a careful analysis of NTBI levels in TM patients at UCLH has failed to find such an association (unpublished data), suggesting that the NTBI species heterogeneity may in principle affect the pattern of tissue iron uptake. NTBI species consist of iron citrate monomers, oligomers and polymers, as well as protein bound forms and these may differ in their rates of uptake into tissues^{167,108}. The various assays used to quantitate NTBI may measure different species that have variable importance to tissue iron uptake. It is not yet clear which assay is most appropriate to predicting extra-hepatic iron distribution. The classic NTBI assay captures both directly chelatable iron, as well as a fraction of iron species that are only slowly chelatable.^{108,109} Furthermore, the assay detects iron chelates of deferiprone¹¹⁰ or deferasirox.¹¹¹ The LPI assay measures a fraction of NTBI that is redox-active and that is removed by the presence of iron chelators in plasma. The effect of iron chelate complexes is not clear but likely to affect this assay less than the NTBI assay. New assays for NTBI species are under development with the intention of obtaining an assay that has clear utility for predicting the risks of extra-hepatic iron distribution.

A further reason why NTBI levels are difficult to link precisely to extra-hepatic iron distribution is that NTBI is generated by factors other than iron overload. Two identified factors that generate NTBI are low levels of erythropoiesis and/or IE. Suppression of erythropoiesis, e.g. following myeloablative chemotherapy, leads to decreased clearance of transferrin iron and the rapid appearance of NTBI,¹¹³ which is quickly reversed following regeneration of erythropoiesis. In DBA, where recent work shows an absence of soluble transferrin receptors and hence of clearance of Tf iron by the erythron, NTBI iron levels are particularly high and this condition is associated with a high propensity to myocardial iron accumulation (Porter 2014, in preparation). Conversely, a high utilization of transferrin iron such as in the highly expanded erythron in NTDT would de-saturate small concentrations of transferrin, which in turn could inhibit NTBI iron uptake into target tissue. This effect would also be active in TM patients where the transfusion regimen

does not completely suppress erythropoiesis. This hypothesis is currently under investigation (Garbowski 2014, in preparation).

Patients with SCD have low levels of NTBI^{112,113} and LPI^{114,115} compared with TM patients with similar levels of iron overload. It is an attractive hypothesis to link the low propensity for extra-hepatic iron distribution in SCD to this observation. Of note LPI is also low in SCD relative to other forms of TIO at similar levels of iron loading. However patients with NTDT have high levels of NTBI^{116,117} and LPI¹¹⁸ but a very low risk of myocardial iron,¹¹⁹ suggesting that absolute NTBI levels are not the only consideration. When a comparison of LPI in rare anemias resulting from hemolysis or from decreased red cell production was made,¹²⁰ both were associated with raised LPI, which was removed following chelation therapy. Analysis of the same patients showed a significant correlation of LPI levels with both the degree of iron overload (ferritin) and transfusional ILR (**Figure 2**). This would support the notion that with increasing transfusional ILR, LPI and possibly the potential to extrahepatic iron loading increases. A systematic study of the relationship of transfusional loading rate, NTBI speciation and extra-hepatic complications of iron overload is warranted.

KEY POINTS

- The pathophysiological consequences of TIO are best understood in TM and broadly reflect the distribution of excess storage iron to heart, endocrine tissues and liver.
- The pattern of excess iron distribution reflects the pattern of NTBI uptake to these tissues.
- Storage iron does not directly damage cells but its intracellular turnover contributes to labile intracellular iron pools that generate harmful free radicals
- TIO also increases the risk of infection, due to increased availability of labile iron to microorganisms
- In other conditions such as SCD, DBA and MDS the propensity to the extra-hepatic iron distribution and its consequences vary compared with TM
- The mechanisms underlining this variability may reflect differences in the transfusional iron loading rates, age of commencing transfusion as well as differences between transferrin iron utilization and NTBI generation.

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FIGURES

Figure 1



Adapted from Porter JB. Hematol Oncol Clin North Am. 2005;19(suppl 1):7-12

Figure 1. Pathological mechanisms and consequences of iron overload. In iron overload resulting from repeated blood transfusions or long-term increased iron absorption, iron that is not liganded to naturally occurring molecules such as transferrin or ferritin or to therapeutic iron chelators, generates a variety of reactive oxygen species (ROS), most notably hydroxyl radicals. This occurs in cells where labile plasma iron is accumulated (especially liver, endocrine tissues and myocardium) thereby increasing levels of both storage and labile cellular iron. ROS increase lipid peroxidation and organelle damage, leading to cell death and fibrogenesis mediated by transforming growth factor, TGF β 1. (Porter JB. Hematol Oncol Clin North Am. 2005;19(suppl 1):7-12.) ROS also damage DNA, risking genomic instability, mutagenesis and cell death or neoplasia. ROS directly activates caspases thereby accelerating apoptotic death (Zuo Y, et al. Cell Res. 2009;19:449-57). Paradoxically, ROS may also have anti-apoptotic effects by activating NF-kB (dashed lines) (Aggarwal BB. Cancer Cell. 2004;6:203-8.) which may contribute to MDS transformation and to iron mediated neoplasia such as hepatoma.





(A) The relationship between baseline pre-dose labile plasma iron (LPI) and transfusion rate in the year prior to study entry. There is a

significant correlation (R = 0.58, P = 0.0005, n = 32) between transfusion rate in the year prior to study entry and baseline pre-dose LPI in all

patients. Hemolytic anemias are shown in circles and production anemias in diamonds. The grey area denotes the healthy reference range.

(B) The relationship between baseline pre-dose LPI and baseline serum ferritin. There is a significant relationship (R = 0.47, P = 0.004) between

baseline ferritin and baseline pre-dose LPI for all patients. Hemolytic anemias are shown in circles and production anemias in diamonds. The

grey area denotes the healthy reference range.

From Porter JB, Lin KH, Beris P, et al. Response of iron overload to deferasirox in rare transfusion-dependent anaemias: equivalent effects on serum ferritin and labile plasma iron for haemolytic or production anaemias. *Eur J Haematol.* 2011;87(4):338-348. Copyright © 2011 John Wiley & Sons A/S

TABLES

Condition	Underlying mechanism for iron overload	Typical distribution and mechanism	Consequences
Inherited anemias			
Thalassemia Major (TM)	Blood transfusions for anemia (+++) Increased iron absorption (+)	High NTBI (++) Liver, heart, endocrine	Cardiomyopathy Endocrinopathy Infection Cirrhosis, hepatoma
Sickle Cell Disease	Intermittent transfusion (++) Intravascular hemolysis with iron loss (+)	Liver predominantly High erythron iron utilization Low NTBI (+/-)	Cirrhosis (Extra-hepatic Iron)
Congenital Dyserythropoietic	Ineffective erythropoiesis Variable transfusion	NTBI ¹²¹ (+)	Not systematically described
Anemia	 70% of patients w CDA-i, 5% with CDA-II, 50% with other CDA variants 		
Sideroblastic Anemia Pyruvate Kinase Deficiency (severe)	Transfusion Ineffective erythropoiesis Transfusion dependence Ineffective erythropoiesis Extravascular hemolysis	Hepatic, myocardial High NTBI Like NTD unless transfused	Hepatic Myocardial ()() Like NTDT unless transfused
Diamond Blackfan Anemia (DBA)	Transfusion dependence (+++)	Hepatic and extra-hepatic Low erythron iron utilization High NTBI (+++)	As with TM
Pure Red Cell Aplasia	As above	As above	As above
Acquired anemias			
Aplastic Anemia	Blood transfusions for anemia	Liver, heart, endocrine	Abnormal LFTs Heart ¹²² and liver failure ¹²³ LIC reduced by chelatio
Fanconi anemia	Regular transfusion	Not well described	As above
Myelodysplasias ¹²⁴ (MDS)	Variable Blood transfusion late onset Ineffective erythropoiesis(++)	Hepatic Extra-hepatic >60-200 units	Hepatic ¹²⁵ Cardiomyopathy ¹²⁵
Myelofibrosis	High Blood transfusion late onset Massive extra-vascular hemolysis	high LPI ??	Infection not described
Multiple myeloablative chemotherapies	Intermittent transfusion	myocardial iron >100 units	cardiomyopathy

Table 1. Conditions associated with transfusional Iron overload

Table 2. Abbreviations

4-HNE	4-hydroxy-nonenal	LVEF	Left Ventricular Ejection Fraction	
AML	Acute Myeloid Leukemia	MAP	Mitogen-activated Protein Kinase	
вм	Bone Marrow	MDA	Malondialdehyde	
BMP	Bone Morphogenic Protein	MDS	Myelodysplastic Syndrome	
CDA	Congenital Dyserythropoietic Anemia	МІС	Myocardial Iron Content	
DBA	Diamond-Blackfan Anemia	MRI	Magnetic Resonance Imaging	
DcytB	Duodenal cytochrome B	MCSIO	Multi Center Study of Iron Overload	
DFO	Deferoxamine	NF-кВ	Nuclear Factor Kappa-light-chain-enhancer of activated B cells	
DMT1	Divalent Metal Transporter-1	NTBI	Non-Transferrin-Bound Iron	
dw	dry weight	NTDT	Non-Transfusion-Dependent Thalassemia	
EPR	Electron Paramagnetic Resonance	PNH	Paroxysmal Nocturnal Hemoglobinuria	
ERK	Extracellular Signal-regulated Kinases	RES	Reticulo-Endothelial System	
GDF15	Growth Differentiation Factor-15	ROS	Reactive Oxygen Species	
Hbd	Hemoglobin-deficit (mouse strain) caused by Sec15I1 gene defect in the vesicular trafficking pathway	SCD	Sickle Cell Disease	
HbS	Sickle Hemoglobin	SF	Serum Ferritin	
HFE	Hemochromatosis gene (High Iron Fe)	SMAD	Small Mothers Against Decapentaplegic transcription factors pathway	
IE	Ineffective Erythropoiesis	STAT	Signal Transducer and Activator of Transcription	
IL-6	Interleukin 6	Tf	Transferrin	
ILR	Iron Load Rate	TfR1	Transferrin Receptor-1	
IRE	Iron Responsive Element	TGFβ-1	Transforming Growth Factor β -1	
IRP	Iron Regulatory Protein (iron responsive element binding protein)	τιο	Transfusional Iron Overload	
LDH	Lactate Dehydrogenase	тм	Thalassemia Major	
LFTs	Liver Function Tests	UCLH	University College London Hospitals	
LIC	Liver Iron Content	UTR	Untranslated Region	

LIP Labile Intracellular Iron Pool