Iron Overload in Diabetic Retinopathy: A Cause or a Consequence of Impaired Mechanisms?

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Iron is an essential ion for life, playing a central role in many metabolic processes. The most important property of free iron is its capacity to be reversibly oxidized and reduced, but at the same time this make it highly pro-oxidant molecule. In this regard, iron is able to generate powerful reactive oxygen species (ROS). Therefore, the maintenance of iron homeostasis in the organism is crucial, and high levels of free iron could be harmful.

In the human retina, iron levels increase with age in both men and women. However, women have significantly more...
retinal iron than men at all ages, in spite of having a higher incidence of anemia, which suggests tissue-specific mechanisms of iron regulation [19]. Abnormalities in local iron homeostasis have been implicated in several degenerative diseases, including Parkinson’s, Alzheimer’s, and age-related macular degeneration, where it has been hypothesized that oxidative stress contributes to cell death [20–22]. In addition, iron participates in other ocular diseases such as glaucoma and cataract [23, 24]. However, the potential role of dysregulation of iron metabolism in the pathogenesis of DR remains to be elucidated. Here we present an overview of the intricate network of proteins involved in retinal iron handling, and we discuss evidence which suggests that iron may contribute to retinal degeneration observed in DR.

2. Iron Homeostasis in the Retina

Since iron is highly toxic due to its ability to generate free radicals, homeostatic mechanisms maintain iron levels by regulation of the proteins involved in iron import (transferrin, transferrin receptor, divalent metal transporter-1), storage (ferritin), and export (ceruloplasmin, hephaestin, ferroportin, and hepcidin) [16, 25–34]. The opposing requirements and toxicities of iron are managed by an iron-responsive mechanism of posttranscriptional regulation of key iron-handling proteins [35]. This regulation allows individual cells to regulate iron uptake, sequestration, and export according to their iron status. Iron-regulatory proteins (IRPs) register intracellular iron status and, in cases of intracellular iron deficiency, bind to iron-responsive elements (IREs) on the mRNA of the regulated protein [36–39].

Iron circulates in the blood stream by being incorporated in the heme molecule of hemoglobin and mioglobin, and most nonheme iron is bound to transferrin, a protein capable of binding two molecules of ferric iron. Iron uptake by cells involves the transferrin binding to its receptor (TF-R) and subsequent endocytosis. After acidification of the endosome, transferrin releases its iron and is recycled to the membrane where it is released to the extracellular space. Iron in the endosome is then transported out through ferroportin or by divalent metal transporter-1 (DMT1).

Transferrin is present in the vitreous fluid of rabbits at a higher relative concentration found in the plasma or in the aqueous humor [40]. In fact, transferrin makes up about 25% of the total protein in the rabbit’s vitreous humor [41]. In animal models and the human retina, the main site of transferrin synthesis is the retinal pigment epithelium (RPE) [33, 40, 42, 43].

Transferrin may protect the retina from the potentially toxic effects of unbound iron, because iron bound to transferrin does not cause oxidative stress [44]. Transferrin probably helps to transport iron to photoreceptors through their TF-R [33]. Finally, transferrin may also have neurotrophic effects that are essential for normal retinal functioning [45]. In the rat, retina TF-R has been detected in the RPE, the inner segments of photoreceptors, the outer plexiform layer, inner nuclear layer, inner plexiform layer, and in the ganglion cell layer [33]. Tf-Rs are located on both the basolateral and apical surfaces of RPE cells, suggesting that there is a bidirectional iron stream in the blood-retinal barrier depending on the iron status in the epithelial cells [16, 33].

Once into the cell, the iron is rapidly uptaken by cytosolic ferritin, a protein capable of incorporating 4,500 iron molecules. Ferritin is composed of 24 subunits of two chains: H-ferritin (heavy chain, or “heart ferritin”) which possesses a ferroxidase function which reduces the ferric form to the ferrous one, mainly localized in the heart, and L-ferritin (“light” or “liver” ferritin) which does not have ferroxidase activity [31]. The ability of cells to store and retrieve iron from ferritin is dependent on the ratio of H:L ferritin chains, but the mechanisms that regulate this ratio are not fully understood. H-ferritin is not only responsible for iron oxidation and uptake, but also has other functions such as reducing the cell proliferation rate and apoptosis [46, 47]. Another form of ferritin, mitochondrial ferritin, has been identified. Mitochondrial ferritin is 80% homologous to H-ferritin found in the cytoplasm and stores iron more efficiently than cytoplasmic ferritin [48, 49]. In the murine retina, mitochondrial ferritin has been found in the photoreceptor inner segments and diffusely throughout the inner retina [50].

By reducing the intracellular level of free iron, ferritin is capable of reducing oxidative stress. There are some factors such as ascorbic acid (Vitamin C), alpha-lipoic acid, or UVB irradiation that can affect iron metabolism [51]. Ascorbic acid is present in the retina at a high concentration compared with its presence in other human organs, and it is able to protect the retina against oxidative damage [52–54]. In this regard, we have recently found ~20-fold higher levels of ascorbic acid in the vitreous fluid than in serum. In addition, the vitreous fluid from PDR patients contained a significant lower amount of ascorbic acid in comparison with vitreous samples from nondiabetic subjects [55]. Moreover, it has been demonstrated that ascorbic acid causes large increases in ferritin synthesis and increased loading of iron into ferritin in cultured epithelial cells of the lens [56, 57]. Therefore, the effect of ascorbic acid in iron metabolism contributes to its antioxidant properties, and the reduced levels detected in the eyes of diabetic patients could be involved in the pathogenesis of DR.

The iron which is not used by the cell needs to be returned to the blood stream. Only ferrous iron can pass through the plasma membrane into the blood, and only ferric iron can be incorporated into transferrin [58]. Therefore, the iron is transported out of the cell bound to ferroportin (a cell membrane protein), and it is then oxidized by the ferroxidases ceruloplasmin and hephaestin, thus making it available to be bound to transferrin.

3. Iron-Dependent Regulation of Retinal Functions

The RPE constitutes the outer blood-retinal barrier and regulates the flow of iron between the choroidal vasculature
and the outer retina. Of all the retinal cell types, RPE cells are theoretically the most susceptible to oxidative damage because of their proximity to the choriocapillaries. In fact, in the human retina, the highest levels of iron are found in the choroid, RPE, and photoreceptor segments [50]. However, other cell types such as pericytes, endothelial cells, retinal Muller cells, ganglion cells, and astrocytes can be, affected even earlier than the photoreceptors and RPE cells by oxidative damage [12]. Iron in the eye is important for the phototransduction cascade. Indeed, iron is an essential cofactor for the enzyme guanylate cyclase, which synthesizes cGMP, the second messenger in the phototransduction cascade [59]. In addition, isomerization of all-transretinal within the retinal pigment epithelium (RPE) in the visual cycle requires iron for the activation of RPE65, an enzyme involved in the visual cycle pathway [60].

The citosolic aconitase system, a dual-function protein involved in the metabolic regulation of iron that is found in all mammalian cell types studied, is located in the RPE of the retina [51]. When iron is scarce, c-aconitase functions as an iron regulatory protein (IRP) controlling the translation of numerous proteins. However, when iron is abundant, the IRP triggers aconitase activity and regulates L-glutamate production, a neurotransmitter involved in retinal neurodegeneration [61]. Thus, cultured lens epithelial cells (LECs), retinal pigment epithelial (RPE) cells, and retinal neurons synthesize and secrete L-glutamate, and this process is regulated by iron by way of its effect on c-aconitase [61]. Elevated levels of glutamate in the retina have been found in experimental models of diabetes, as well as in the vitreous fluid of diabetic patients with PDR [62–65]. The heme oxygenase (HO) system acts as an antioxidant. There are 2 main isoforms of HO: HO-1, a “heat shock protein”, which is very sensitive to oxidative stress [66], and HO-2, which is expressed constitutionally in the endothelial, neural, retinal and testicular cells [67]. HO-1 is an inducible enzyme whose activity increases in response to iron as well as heme, light, oxidative stress, and inflammation. It degrades heme to iron, carbon monoxide (CO), and biliverdin. The release of iron upregulates the synthesis of ferritin as a cytoprotective mechanism (see below). CO has important roles in vasodilatation, and biliverdin is subsequently converted to the antioxidant bilirubin. It has been demonstrated that increasing HO-1 promoter activity in RPE cells could trigger a protective response [68]. In the retina, overexpression of HO-1 in photoreceptor cells provided protection from light damage [69]. In murine models, HO-1 and HO-2 were localized in the outer segment of the photoreceptor layer, inner plexiform layer, ganglion cell layer, glial fibres, and capillary endothelium [70].

However, during hemorrhage, the excessive generation of iron and bilirubin, that is neurotoxic, has deleterious consequences.

Severe hypoxia due to capillary occlusion is the main condition for the initiation of neovascularization in PDR. Hypoxia upregulates the expression of angiogenic factors directly or through the hypoxia-inducible factor (HIF-1). HIF-1 activates several genes related to iron metabolism such as HO-1, transferrin, transferrin receptor, and ceruloplasmin [71–73]. In addition, it has been recently demonstrated that ischemic preconditioning of the retina is highly effective in preventing subsequent injury caused by iron-dependent free radical burst after prolonged ischemia. This protection appears to be provided by increased ferritin levels [74].

4. Disruption of Iron Homeostasis and Oxidative Damage in DR

It has been demonstrated that intravitreal levels of iron in PDR are 2.5 times the normal levels [75]. In addition, transferrin concentrations have been found elevated in the vitreous fluid and retinal membranes from patients with proliferative vitreoretinopathy diseases including PDR [76, 77].

There are several mechanisms that could explain iron overload in diabetic eyes (Figure 1). First, it has been demonstrated in vitro that hyperglycaemia causes a complete destruction of heme molecules from hemoglobin and myoglobin, releasing free iron into the interstitial space [78, 79]. Second, intraretinal and vitreal hemorrhages could contribute to iron overload in PDR. Finally, angiotensin II stimulates the local gene expression of proteins related to iron metabolism (TfR, DMT1, ferroportin, and hepcidin) in the rat kidney, thus contributing to the production of high levels of iron transporters and facilitating iron uptake by the cells [80, 81]. In this regard, it is worth mentioning that the major components of the renin-angiotensin system, including angiotensin II, have been identified in glial cells, neurons, and blood vessels from murine retinas [82–84] and are overexpressed in diabetic rats [85, 86]. In human beings, vitreal concentrations of prorenin, renin, and Ang II are elevated in patients with DR [87, 88]. In addition, proteomic analysis of vitreal samples taken from patients with diabetes revealed that angiotensinogen was found in greater concentration in samples taken from those with PDR, compared with those with no DR or non-diabetic control subjects [89].

The consequences of iron overload in the diabetic eye are complicated to evaluate because, as mentioned above, there are multiple forms of iron with different reactivity and several proteins that modulate their levels and actions. However, among the potential mechanisms of iron-induced damage, it seems that oxidative damage is the most important (Figure 1).

Increased intraocular levels of iron cause oxidative damage to photoreceptors with greater damage to cones than rods [90]. In addition, it has been shown that iron chelation protects the RPE cells against cell death induced by oxidative stress [91, 92]. In the retina of human donors with DR high levels of peroxidized lipids in Bruch’s membrane promoted by the local ferric iron involved have been demonstrated [93]. This happens also in the eyes of patients suffering from vitreal bleeding in the course of PDR, after which there is an important reactivation of superoxide generation catalyzed by the locally released free iron [94]. In addition, it should be noted that iron ion catalyses the binding of the AGEs (advanced glycation end products) to the specific receptor, that is, a crucial step in the pathogenesis of the DR [95].
It has been demonstrated that impaired retinal iron homeostasis is associated with defective phagocytosis in both murine models [96] and in ARPE-19 cell cultures [97]. An impairment of phagocytosis has been described in long-term diabetes [98] and, therefore, it is possible that this could also happen to RPE cells. Thus, iron overload could contribute to the phagocytosis defect associated with diabetes. This defect implies a delayed and impaired phagocytosis of both the apoptotic cells and the local detritus, which generates the accumulation of autoantigens and the synthesis of proinflammatory cytokines.

As mentioned above, HO may respond to oxidative stress and upregulation of the HO system (HO-1 and HO-2) in rats with streptozotocin-induced diabetes has been demonstrated [70]. In diabetic rats, increased retinal HO-1 mRNA expression has been shown to be preventable with antioxidant therapy [99], and HO-1 overexpressing neurons have shown reduced levels of apoptosis [100]. However, in human eyes with long-term diabetes, reduced HO-1 mRNA expression in RPE cells has been demonstrated [101], thus suggesting that increased HO activity induced by diabetes is dependent on diabetes duration.

Although the HO system has been generally accepted as having an antioxidant role in several tissues, HO also could exhibit pro-oxidant activity in the vascular endothelial cells. For example, it has been demonstrated that in the endothelial cells, HO increases the expression of nitric oxide (NO), endothelin-1, and VEGF [102–104], all of which are relevant factors in the pathogenesis of PDR [2].

In cellular cultures, free iron stimulates the expression of adhesion molecules and monocyte endothelial adhesion, [105–107] key steps in the development of DR. Finally, in murine models, iron overload is associated with RPE hypertrophy and hyperplasia due to the stimulation of citosolic-aconitase system, which acts as an enzyme initiating the proliferation cascade [108].

In summary, careful control of iron availability is central to the maintenance of normal cell functions. Iron overload seems to be caused by several processes involved in the pathogenesis of DR. However, at the same time iron causes retinal damage mainly by increasing oxidative stress. Further studies addressed to exploring the role of iron in the pathogenesis of DR are necessary not only to improve our knowledge on this issue, but also to design novel therapeutic strategies based on the regulation of iron proteins.

References


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