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## Prions and Prion diseases: insights from the eye

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## Abstract

Prion diseases are invariably fatal neurodegenerative disorders that have gained much publicity due to their transmissible nature. Sporadic Creutzfeldt-Jakob disease (sCJD) is the most common human prion disorder, with an incidence of 1 in a million. Inherited prion disorders are relatively rare, and associated with mutations in the prion protein gene. More than 50 different point mutations, deletions, and insertions have been identified so far. Most are autosomal dominant and fully penetrant. Prion disorders also occur in animals, and are of major concern because of the potential for spreading to humans. The principal pathogenic event underlying all prion disorders is a change in the conformation of prion protein (PrP<sup>C</sup>) from a mainly  $\alpha$ -helical to a  $\beta$ -sheet rich isoform, PrP-scrapie (PrP<sup>Sc</sup>). Accumulation of PrP<sup>Sc</sup> in the brain parenchyma is the major cause of neuronal degeneration. The mechanism by which PrP<sup>Sc</sup> is transmitted, propagates, and causes neurodegenerative changes has been investigated over the years, and several clues have emerged. Efforts are also ongoing for identifying specific and sensitive diagnostic tests for sCJD and animal prion disorders, but success has been limited. The eve is suitable for these evaluations because it shares several anatomical and physiological features with the brain, and can be observed in vivo during disease progression. The retina, considered an extension of the central nervous system, is involved extensively in prion disorders. Accordingly, Optical Coherence Tomography and electroretinogram have shown some promise as pre-mortem diagnostic tests for human and animal prion disorders. However, a complete understanding of the physiology of PrP<sup>C</sup> and pathobiology of PrP<sup>Sc</sup> in the eye is essential for developing specific and sensitive tests. Below, we summarize recent progress in ocular physiology and pathology in prion disorders, and the eye as an anatomically accessible site to diagnose, monitor disease progression, and test therapeutic options.

Author contributions

The authors declare no competing interest.

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NS drafted the manuscript, SC and AA revised the manuscript, EL provided critical feedback. All authors approved the final version of the manuscript.

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Declaration of competing interest

#### Keywords

Prion protein; glaucoma; extracellular matrix; hepcidin; iron; TGFβ2; oxidative stress

#### 1. Introduction

Prions or PrP<sup>Sc</sup> is a conformational isoform of a normal glycosylphosphatidyl inositol (GPI) linked protein expressed on the plasma membrane of all cells, but most abundant on neuronal cells (Prusiner, 1998). The triggers that initiate its conversion from PrP<sup>C</sup> to the PrPSc form, the mechanism by which it spreads, and the underlying cause of neuronal death have been the focus of several studies, and significant progress has been made on all fronts (Scheckel and Aguzzi, 2018; Sigurdson et al., 2019; Singh et al., 2014). For example, it is known that point mutations in the PrP<sup>C</sup> gene increase its susceptibility for conversion to the PrPSc form (Singh et al., 2010). In infectious disorders, exogenous PrPSc initiates the conversion of PrP<sup>C</sup> to PrP<sup>Sc</sup>, which then spreads to distant sites (Aguzzi and Glatzel, 2006; Prusiner, 1998). Factors that contribute to the susceptibility of a particular PrP<sup>C</sup> form for conversion to PrP<sup>Sc</sup> are not fully understood, nor is the mechanism by which PrP<sup>Sc</sup> causes neuronal death. These, and other questions need to be answered because of the spread of prion disease in free ranging deer and elk as Chronic Wasting Disease (CWD), and the danger of potential spread to humans (Hannaoui et al., 2017). Although prion disorders have been known in sheep and goats for several years, the spread to farm and wild animals is relatively new. The emergence of Bovine Spongiform Encephalopathy (BSE) in cattle and its onward spread to humans was a rude reminder that protection offered by the species barrier is only partial, and new strains are likely to emerge (Prusiner, 1997). The eye provides an opportunity to address these questions because of structural similarities to the brain, and the ease of access during disease development.

Natural pathways of transmission of PrP<sup>Sc</sup> include the oral route from the gastro-intestinal tract to the brain, peripheral blood, placenta, mucous membranes, and the cornea (Weissmann et al., 2002). Accidental transmission of BSE to humans occurred by eating contaminated beef, and is under control since regulations on the processing of animal feed were put into place. The cornea has been of some concern because of transmission through corneal transplants from subjects exposed to BSE (Maddox et al., 2008). Currently, CWD is spreading among the deer and elk population from contaminated water reservoirs and common grazing grounds (Saunders et al., 2012), and could spread horizontally and to other species from corneal abrasions since corneal epithelium expresses PrP<sup>C</sup>, and infectious PrP<sup>Sc</sup> has been detected at that site (Ashok et al., 2018).

In addition to expression of PrP<sup>C</sup> (Büeler et al., 1992), a certain degree of sequence homology is essential for transmission (Hagiwara et al., 2013), and is the basis of species barrier. Other prominent factors that contribute to disease pathogenesis include inflammation and brain iron dyshomeostasis (Carroll et al., 2015; Singh et al., 2009b; Singh et al., 2010). Whether these play a primary role, or amplify the pathology by other secondary mechanisms has been difficult to evaluate. It is likely that several factors contribute to the final outcome, making it difficult to discern which comes first. One of the hypotheses is the loss of

physiological function of PrP<sup>C</sup> combined with gain of toxic' function by PrP<sup>Sc</sup> in prion disease pathogenesis.

Several functions have been ascribed to PrP<sup>C</sup>. In the eye, overexpression of PrP<sup>C</sup> protects the photoreceptors from damaging light (Frigg et al., 2006; Williams et al., 2011), and plays a role in iron transport across retinal pigment epithelial cells that form the outer blood-retinal barrier, and non-pigmented epithelial cells of the ciliary epithelium that form the bloodaqueous barrier. PrP<sup>C</sup> is known to function as a ferrireductase partner for the Zirt, Irt-like (ZIP) family of iron transporters on the plasma membrane and the endosome membrane (Ehsani et al., 2012; Singh et al., 2015). It is therefore logical to assume that loss of function of PrP<sup>C</sup> due to conversion to PrP<sup>Sc</sup> will induce a certain degree of iron deficiency, an essential metal necessary for vital enzymatic reactions (Gasperini et al., 2016; Singh et al., 2014). Compensatory upregulation of iron uptake proteins creates iron dyshomeostasis and a potentially toxic microenvironment. PrPC is known to stabilize the extracellular matrix, and loss of this function interferes with neuritogenesis (Alleaume-Butaux et al., 2013; Kleene et al., 2007; Ramljak et al., 2015) in the brain, and induces primary open angle glaucoma in the eye (Ashok 2019). These are a few examples where the eye has helped to confirm physiological functions of PrP<sup>C</sup> in the brain, and uncover pathological implications of dysfunction in the eye.

An important area of investigation concerns the role of inflammation in PrP<sup>Sc</sup>-mediated neurotoxicity (Carroll et al., 2015). Several studies have demonstrated upregulation of various cytokines during prion disease progression. Most studies are in mouse models, and use different strains of scrapie to demonstrate the universality of the phenomenon (Carroll and Chesebro, 2019; Carroll et al., 2015). Although some cytokines are beneficial, these are known to upregulate hepcidin, a peptide hormone secreted mainly by the liver and known to regulate systemic iron by downregulating ferroportin (Fpn), an iron export protein (Drakesmith et al., 2015). In that respect, identification of extra-hepatic, local expression of hepcidin in the brain and the eye adds a new dimension to iron management (Ashok 2020), organs separated from systemic circulation by the blood-brain and blood-retinal barriers respectively.

Progress on these fronts has been slow, partly because of the structural and functional complexity of the brain. The eye provides a suitable alternative because disease-specific changes can be visualized *in vivo* during disease progression, and the mechanism of toxicity by PrP<sup>Sc</sup> can be parsed out in a way that is experimentally not possible in the brain. Thus, using the eye as a 'window' to the brain, several new observations have emerged. Some reinforce previous findings from human and animal brains, while others provide new perspective on the functional role of PrP<sup>C</sup> and consequences of dysfunction. Some of these observations are discussed below.

#### 2. The eye as a 'window' to the brain

Thus far, the eye has been studied mainly as a site where  $PrP^{Sc}$  accumulates and causes disease (Greenlee et al., 2016). It is only recently that the physiological function of proteins implicated in diseases of the brain such as  $PrP^{C}$ , the main cause of sCJD, amyloid precursor

protein (APP) and tau, involved in Alzheimer's disease (AD), and  $\alpha$ -synuclein, the cause of synucleinopathies have been evaluated in the eye. The retina provides a convenient site to examine loss of function and gain of toxic function by these proteins in vivo during disease progression, and has been used successfully for the early diagnosis of some of these disorders. For example, Optical Coherence Tomography (OCT) is used for the early diagnosis of prion disorders, AD, PD, amyotrophic lateral sclerosis, and multiple sclerosis, to mention a few, and the sensitivity and specificity of the test is improving consistently (Adhikary et al., 2010; Doustar et al., 2017; Torrent et al., 2010). Likewise, retinal function measured by electroretinogram (ERG) is emerging as a sensitive test for retinal function in the early stages of these disorders (Ngoo et al., 2019). The retina is more conducive to analysis than the brain, and has yielded important information about the physiological function of PrP<sup>C</sup>, APP, and a-synuclein. Like the brain, the retina is separated from systemic circulation by the outer and inner blood retinal barriers comprised of retinal pigment epithelial (RPE) cells and capillary endothelial cells. RPE cells form a monolayer between choroidal sinuses and the outer retina, and expresses proteins and receptors in a polarized manner to allow unidirectional transport of select substances. Likewise, capillary endothelial cells form a tight monolayer between the peripheral circulation and the inner retina, and allow selective transport of essential substances to the inner neuroretina. The RPE cell layer is followed by photoreceptors, which rely on these cells for the daily turnover of outer segments by autophagy. This layer is followed by the outer and inner plexiform and nuclear layers, which are traversed by Muller glia. The ganglion cells form the innermost layer of the retina, which form the retinal nerve fiber layer (RNFL) that carries signals to the brain through the optic nerve. PrP<sup>C</sup> is expressed in all layers of the retina, including the outer and inner blood-retinal barriers, and blood aqueous barrier formed by non-pigmented ciliary epithelial cells. It is also expressed widely in the anterior segment, including the ciliary body, lens epithelium, corneal endothelium and epithelium, and the trabecular meshwork (Ashok et al., 2018; Asthana et al., 2017; Frederikse et al., 2000). One might argue that PrP<sup>C</sup> is of neuroectodermal origin, and it is not surprising to expect widespread expression in ocular tissues. However, systematic studies on different regions of the eye have revealed important physiological functions of PrPC, and deleterious consequences of loss of this function.

The eye is involved early in prion diseases (Chesebro et al., 2005; Striebel et al., 2019). Several studies have examined the spread of prion disease from the brain to the neuroretina, and total destruction of the latter by end stage disease. Approximately 40% of patients with sCJD develop visual symptoms, providing a window of opportunity for early diagnosis and intervention. PrP<sup>C</sup> is expressed, and is converted to the PrP<sup>Sc</sup> form in the cornea, sclera, extraocular muscle, choroid, optic nerve, and lens (Figure 1) (Kercher et al., 2007; Orru et al., 2018). Soluble PrPC is present in the aqueous humor and vitreous humor (Ashok 2019). Iatrogenic spread of prions through corneal grafts has been reported, though it is relatively rare (Armitage et al., 2009; Head et al., 2003). In transgenic mice expressing CWD prions, PrP<sup>Sc</sup> was detected in the cornea in addition to all layers of the retina (Asthana et al., 2017). The presence of PrP<sup>Sc</sup> in dust particles, grass blades, and water reservoirs have prompted studies on peripheral routes of entry from the conjunctiva with positive results (Gough et al., 2015). Thus, the eye is a source of PrP<sup>Sc</sup> transmission, all be it at a much slower state

(Atkins et al., 2018; Gnanajothy et al., 2013). This is an important question since horizontal spread of animal prions through corneal abrasions is a distinct possibility (Asthana et al., 2017).

The eye offers several other advantages. For example, progression of the disease can be visualized *in vivo* in mouse models infected with prions, and the specific tissue analyzed after harvesting the eye. This allows a spatiotemporal evaluation of disease progression, and correlation between loss of function and gain of toxic function with disease progression. By generating tissue-specific PrP<sup>null</sup> mice, it is possible to evaluate the progression of disease, and the role of PrP<sup>C</sup> at specific sites. Though several functions of PrP<sup>C</sup> have been described, implications of their dysfunction varies based on the tissue or organ in question. Some of these are exemplified by the eye because of its unique function in transmitting light signals. We provide some examples below.

#### 3. The loss of physiological function and gain of toxic function

#### hypothesis.

Of the many functions that PrP<sup>C</sup> participates in, we will focus on those where the eye has helped to resolve outstanding questions or reinforce known facts. Below are some examples.

#### Iron transport

Transport of iron to the eye has been the subject of several studies because of its vital role in the eye, and the inherent toxicity due to its redox-active nature (García-Castiñeiras, 2010; Song and Dunaief, 2013). The eye is particularly susceptible to iron induced toxicity because of constant exposure to light, which reduces iron from its relatively stable ferric form to the redoxactive ferrous form (Song et al., 2014). Transport of iron to the eye is therefore regulated by the outer and inner blood-retinal barriers, and the blood-aqueous barrier by specific proteins. In addition, relatively high concentrations of citrate, glutathione peroxidase, superoxide dismutase, and other reducing agents and enzymes protect the eye from iron-mediated toxicity (Goralska et al., 2009). It is believed that iron enters the eye from the inner blood retinal barrier, and leaves from the outer blood retinal barrier.

 $PrP^{C}$  is a ferrireductase, and facilitates the transport of transferrin (Tf)-bound and non-Tf bound iron across biological membranes. Both forms of iron are in the oxidized, ferric  $(Fe^{3}+)$  form, and need reduction to the ferrous  $(Fe^{2}+)$  form for transport through metal transporters that span the membrane. Tf-bound iron taken up by cells *via* the Tf/transferrin receptor (TfR) pathway is released from Tf at the low pH of endosomes, and reduced to the ferrous form by membrane bound reductases such as  $PrP^{C}$  or Steap3 before transport to the cytosol across divalent metal transporter 1 (DMT1) (Asthana et al., 2017; Ohgami et al., 2006; Singh et al., 2013). Likewise, ferric iron reduced at the plasma membrane by  $PrP^{C}$  or Dcytb is transported through the ZIP family of divalent metal transporters to the cytosol (Ashok and Singh, 2018; Lane et al., 2015; Leung et al., 2008) (Figure 2). Here, it is utilized for metabolic purposes, and excess is transported out through the combined action of Fpn and a ferroxidase such as ceruloplasmin or haphestin. The redundancy in ferrireductase proteins emphasizes the significance of this function to cell viability.  $PrP^{C}$  is expressed on

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capillary endothelial cells that form the inner blood-retinal barrier, RPE cells that form the outer blood-retinal barrier, and non-pigmented epithelial cells that form the blood-aqueous barrier. Absence of PrP<sup>C</sup> in PrP<sup>null</sup> mice creates a phenotype of mild iron deficiency, indicating a role in iron transport (Singh et al., 2009a; Singh et al., 2009b).

In RPE cells, PrP<sup>C</sup> is expressed on the basolateral membrane *in vivo* and *in vitro*. In an RPE cell line, polarization increases intracellular ferritin, indicating uptake of iron. Concomitant cleavage of PrP<sup>C</sup> that releases the N-terminal ferrireductase domain comprising of the octapeptide repeat region suggests regulated uptake of iron (Asthana et al., 2017). It is likely that PrP<sup>C</sup> plays a dual role in iron uptake or export depending on the iron status of the retina. It is interesting that almost all  $PrP^{C}$  in RPE cells is cleaved at the  $\beta$ -site, an event that correlates with polarization and accumulation of ferritin (Ashok et al., 2018; Asthana et al., 2017). It is likely that accumulation of ferritin serves as a signal, and shedding of the ferrireductase domain blocks further uptake of iron. Silencing of PrP<sup>C</sup> in RPE cells lowers, while overexpression upregulates intracellular ferritin, supporting this claim (Asthana et al., 2017). The reason for  $\beta$ -cleavage of PrP<sup>C</sup> at amino acid ~90 instead of the physiological cleavage at amino acids 111/112 or the  $\alpha$ -site is unclear at present (McDonald et al., 2014; Watt et al., 2005). Although both cleavage events would remove the N-terminal ferrireductase domain,  $\beta$ -cleavage, an event associated with oxidative stress, is evident in most ocular tissues (Haigh et al., 2015; Watt et al., 2005). The significance of this observations is unclear at present.

The presence of  $PrP^{C}$  on the basolateral membrane of non-pigmented epithelial (NPE) cells suggests transport of iron from this site as well (Asthana et al., 2017). Local expression of hepcidin, Fpn, and ceruloplasmin on NPE cells support this assumption (Ashok et al., 2020b; Ashok et al., 2018). Moreover, expression of local hepcidin in the ciliary epithelium, corneal endothelium, lens epithelium, and trabecular meshwork further supports local regulation of iron at these sites (Ashok et al., 2020b). This contrasts with the belief that the anterior segment receives iron from the retina by diffusion, and supports the presence of an independent source of iron for the anterior segment (García-Castiñeiras, 2010). It is notable that  $PrP^{C}$  is mainly full-length in the corneal endothelium, and  $\beta$ -cleaved in RPE, NPE, and trabecular meshwork (TM) cells (Ashok et al., 2018). Further investigations are necessary to evaluate whether this has functional implications, or is a cell specific phenomenon.

How does loss of function of PrP<sup>C</sup> and gain of toxic function impact the functioning of the eye? Although iron deficiency or excess alters ocular function in defined ways, the effect of PrP<sup>C</sup> on iron homeostasis has been difficult to discern. Loss of PrP<sup>C</sup> in PrP<sup>null</sup> mice has minimal effect on functioning of the eye, though a mild phenotype of iron deficiency is noted (Asthana et al., 2017). It is likely that other iron uptake proteins compensate for the loss of PrP<sup>C</sup>, minimizing the effect of loss of function of PrP<sup>C</sup>. However, PrP<sup>Sc</sup> results in sequestration of iron in ferritin, which co-aggregates with PrP<sup>Sc</sup> (Singh et al., 2009a). This could result from different scenarios, which are not mutually exclusive. 1) PrP<sup>Sc</sup> is transported to lysosomes for turnover, where it accumulates because it is resistant to lysosomal hydrolases. Ferritin is also transported to lysosomes for releasing stored iron for utilization by the cell. It is likely that the hydrophobic nature of ferritin and PrP<sup>Sc</sup> results in co-aggregation of the two, resulting in PrP<sup>Sc</sup>-ferritin aggregates. Such aggregates have been

observed in neuroblastoma cells infected with  $PrP^{Sc}$ . The ferritin is rich in iron and shows a Perls' positive reaction, leaving no doubt that iron is indeed increased but biologically unavailable. This creates a phenotype of relative iron deficiency despite excess brain iron, and upregulation of iron uptake proteins Tf and TfR (Singh et al., 2009a). 2) Neuro-inflammation accompanies prion infection, and the cytokines released are toxic of their own accord. In addition, IL-6, IL-1 $\beta$ , and the TGF family of cytokines are transcriptional triggers of hepcidin. Downregulation of Fpn by hepcidin is likely to increase intracellular iron and upregulate ferritin, resulting in the accumulation of iron rich ferritin. This is likely to create a feed-forward loop by further activating microglia, and the release of cytokines. In scrape infected retina, activation of Müller glia occurs prior to accumulation of ferritin, supporting this concept (Asthana et al., 2017; Carroll et al., 2015). Further studies are necessary to understand this phenomenon fully, and determine its role in PrP<sup>Sc</sup>-induced toxicity.

One of the cytokines that upregulate hepcidin is transforming growth factor  $\beta 2$  (TGF $\beta 2$ ). This is of special interest in the TM because of the association of TGFB2 with primary open angle glaucoma (POAG) (Hill et al., 2018). The cause of POAG is not completely understood, but TGF<sup>β</sup>2-induced upregulation of fibrillogenic proteins increases the resistance of aqueous outflow from the TM, elevating intraocular pressure (IOP). A persistent increase in IOP induces ganglion cell death, resulting in glaucoma (Montecchi-Palmer et al., 2017). There are two main pathways by which TGFB2 brings about this change; the canonical or Smad dependent pathway, and the non-canonical or Smadindependent pathway (Hata and Chen, 2016; Prendes et al., 2013). The Smad-dependent pathway functions through pospho-Smad2/3, which along with co-Smad 4, is translocated to the nucleus and activates various fibrillogenic proteins. These change the characteristics of the extracellular matrix and increase the IOP. One of the genes upregulated by phospho-Smad4 is hepcidin. Since the TM expresses hepcidin locally (Ashok et al., 2020b; Ashok et al., 2019), this raises the intriguing possibility of upregulation of hepcidin by TGF $\beta$ 2, which downregulates Fpn and causes iron accumulation and upregulation of ferritin. The oxidative stress induced by sequestration of iron in ferritin upregulates TGF $\beta$ 2, creating a positive feed-forward loop. Heparin, a hepcidin antagonist, and N-acetyl carnosine, an antioxidant, are able to disrupt this loop, bringing iron-mediated oxidative stress to the fore-front of glaucomatous change (Poli et al., 2017) (Figure 3).

## PrP<sup>C</sup> and extracellular matrix

PrP<sup>C</sup> plays a prominent role in endothelial to mesenchymal transition (Endo-MT), a change associated with POAG. In contrast to upregulation of hepcidin that occurs through the canonical pathway, Endo-MT is mediated by the non-canonical pathway, and is associated with neurotoxicity by interfering with neuronal polarity and neuritogenesis (Ghodrati et al., 2018; Hajj et al., 2007; Hartmann et al., 2013). PrP<sup>C</sup> interacts with the extracellular matrix through β1 integrin, and downregulation of PrP<sup>C</sup> or dysfunction due to change in conformation to the PrP<sup>Sc</sup> form induces increased synthesis of fibrillogenic proteins including fibrinogen, α-smooth muscle actin, collagen 1A, vimentin, and laminin, indicating transition to a mesenchyme-like phenotype (Ashok et al., 2019; Marbiah et al., 2014; Mehrabian et al., 2014).This changes cell-cell interaction to cell-substrate interactions, interfering with neuronal polarity and neuritogenesis in the neurons, and pliability of the

extracellular matrix surrounding the TM. This change interferes with fine-tuning of the Ras homolog gene family member A (RhoA)-associated coiled-coil containing kinase (ROCK) signaling pathway, resulting in over-activation of ROCK. This induces a change in the pliability of the TM such that it does not react to stretch, resulting in elevation of the IOP (Arantes et al., 2009; Loubet et al., 2012; Zhong et al., 2018) (Figure 4). Inhibitors of the RhoA-Rock pathway are being suggested as a therapeutic option for both prion disorders and POAG, supporting a key role of PrP<sup>C</sup> in regulating cytoskeletal homeostasis (Alleaume-Butaux et al., 2013; Alleaume-Butaux et al., 2015; Kim et al., 2017).

## Conclusions

The eye offers several advantages over the brain for evaluating and understanding the pathobiology of prion disorders. The physiological functions of PrP<sup>C</sup> are reproducible in the eye, and so is their corruption by PrPSc. Iron transport across the outer and inner bloodretinal and blood aqueous barriers is easier to evaluate in the eye relative to the brain because of its less complicated structure, and individual proteins participating in this process can be identified with ease. The anterior segment provides a unique opportunity to study the exchange of iron through monolayers of cells, and determine the directionality of transport. Expression of iron transport proteins and hepcidin, and polarized expression of Fpn, and ceruloplasmin on monolayers is identifiable only in the eye. Upregulation of hepcidin by TGFβ2, and reciprocal upregulation of TGFβ2 by hepcidin in the TM has been possible in the eye, and is likely to have broader implications in the brain. The role of PrP<sup>C</sup> as an extracellular matrix stabilizing protein and its corruption by PrPSc has been described in the brain, but is easier to understand in the eye where it causes glaucomatous alterations by similar mechanisms. PrP<sup>C</sup> is likely to perform other functions in the eye, including the cornea, lens epithelium, iris, and other structures. A detailed analysis of downregulation of PrP<sup>C</sup> and change in conformation to PrP<sup>Sc</sup> is likely to reveal other important functions of PrP<sup>C</sup>, and the mechanism by which their corruption causes disease. Additional research is necessary to understand how PrPSc causes disease-specific changes in the retina and perhaps other regions of the eye, which have remained unexplored due to the rapid course of this disease. A clear understanding of these processes will help in the development of a specific and sensitive diagnostic test for prion disorders, and therapeutic options in the future.

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#### Glossary

PrP <sup>C</sup>	Prion protein
Fpn	Ferroportin
sCJD	Sporadic Creutzfeldt-Jakob disease
PrP <sup>Sc</sup>	Prion protein scrapie

OCT	Optical Coherence Tomography
ERG	Electroretinogram
CWD	Chronic Wasting Disease
BSE	Bovine Spongiform Encephalopathy
ZIP	Zirt, Irt-like protein
APP	Amyloid precursor protein
RPE	Retinal pigment epithelium
RNFL	Retinal nerve fiber layer
NPE	Non-pigmented epithelium
TM	Trabecular meshwork
TfR	Transferrin receptor
DMT1	Divalent metal transporter 1
TGFβ2	Transforming growth factor beta 2
POAG	Primary open angle glaucoma
IOP	Intraocular pressure
Endo-MT	Endothelial to mesenchymal transition
RhoA	Ras homolog gene family member A
ROCK	Rho associated coiled-coil containing kinase
ECM	Extracellular matrix

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## Highlights

- **1.** Prion protein (PrP<sup>C</sup>) is expressed in the anterior & posterior segment of the eye.
- 2. Absence of PrP<sup>C</sup> promotes endothelial to mesenchymal transition in TM cells.
- **3.** PrP<sup>C</sup> helps to maintain iron homeostasis in the anterior segment.
- 4. Anterior segment maintains iron homeostasis independent of the retina.
- **5.** TGFβ2 and hepcidin form a positive feed-forward loop fueled by ironcatalyzed ROS.



## Figure 1. Expression of PrP<sup>C</sup> in the eye.

Immunoreaction of human eye shows a positive reaction for  $PrP^{C}$  in the cornea epithelium and endothelium, TM, CB, iris, lens epithelium, retinal ganglion cell layer, and RPE cells (Ashok et al., 2018; Asthana et al., 2017).  $PrP^{C}$  is expressed on the basolateral surface of NPE and RPE cells. Soluble  $PrP^{C}$  is present in the aqueous humor and vitreous humor. Red dotted line represents a positive reaction for  $PrP^{C}$ .



## Figure 2. PrP<sup>C</sup>-mediated iron homeostasis.

 $PrP^{C}$  facilitates uptake of Tf-Fe<sup>3+</sup> and non-Tf-bound iron (Fe<sup>3+</sup>) by its ferrireductase activity. Tf iron released in the endosome is reduced to Fe<sup>2+</sup> for transport through DMT1. Non-Tf bound iron is reduced at the plasma membrane for transport through ZIP14 and other divalent metal transporters. Loss of  $PrP^{C}$  function reduces iron uptake, and creates iron deficiency Singh 2014).



#### Fig 3. TGFβ2-Hepcidin feed-forward loop.

TGF $\beta$ 2 upregulates transcription of hepcidin by the canonical, Smad-dependent pathway. Autocrine and paracrine activity of hepcidin downregulates Fpn, resulting in intracellular accumulation of iron, compensatory upregulation of ferritin, and increase in ROS, which upregulates TGF $\beta$ 2. Heparin, a hepcidin antagonist, and antioxidants inhibit upregulation of TGF $\beta$ 2 by decreasing ROS, disrupting the cycle (Ashok et al., 2020a).



## Fig 4. PrP<sup>C</sup> stabilizes the ECM.

PrP<sup>C</sup> maintains cell-ECM interactions through β1-integrin and other proteins. Downregulation of PrP<sup>C</sup> causes aggregation of β1-integrin, activating the RhoA-ROCK pathway, and upregulation of fibronectin, collagen 1A, α-SMA, vimentin, and laminin that deposit in the ECM. It also modulates LIM kinases, which play a major role in maintaining cytoskeletal dynamics *via* phosphorylation of the cofilin family of proteins (Alleaume-Butaux et al., 2015; Ashok et al., 2019; Kim et al., 2020).