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MODULATION OF INFLAMMATORY PATHWAY BY INFLAMMATORY INHIBITOR

JSH-23 MITIGATES DIABETIC RETINOPATHY

By

Rubens Petit Homme B.S., Florida Southern College, 2016 M.S., University of Louisville, 2019

A Dissertation Submitted to the University of Louisville School of Medicine In Partial Fulfillment of the Requirements for the Degree of

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Department of Physiology University of Louisville Louisville, Kentucky

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DEDICATION

To my brothers Robens Manno & Ferry Petit Homme, Juan Luis Garcia, my sisters Brindy, Luchandley, Guerlie, and Rose My Parents, Mrs. Elicienne and Mr. Bienera Petit Homme, My best men & women, Tervens, Clay, Senatus, Anne, & Frederick The moms and dads that showed me my left foot from my right, aunt Clarine, Guerda, Ginette, Roseline, Mrs. Honors, Ms. Evon, Grandma Dada, Dr. Gauthier, Dr. Lee, Dr. Eubank, Dr. Crowe, Mr. Ferdinand, Coach Locke, and of course Dr. Adrienne Bratcher

To my city, La Pointe Des Palmistes, My guardian angels, Terancia, Aunt Paulna, & Mrs. Serafin, My Church, First Baptist Church of Nazarene in Lakeland, My alma matter, the Red Devils and the Moccasins To those that want but never was as privileged as I am "Habacuc Milord."

To my family that has walked, continue to walk, and will always walk with me through all the paths of my life.

It's a great remorse that all your names cannot fit within one page. All the cells and water in my body would like to say, "Thank you for sharing the love with me all the days of my life!"

My people continue to share the love, and I'm the living proof that your grace has no limit to the success it can bring to someone.

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ABSTRACT

MODULATION OF INFLAMMATORY PATHWAY BY INFLAMMATORY INHIBITOR (JSH-23) MITIGATES DIABETIC RETINOPATHY

Rubens Petit Homme

August 11, 2020

Diabetic retinopathy (DR) is the most common microvascular complication of diabetes mellitus (DM). Its pathology is consisted of the breakdown of the blood-retinal barrier (BRB), particularly the inner blood-retinal barrier (iBRB). The exact mechanism that leads to the iBRB impairment and the development of DR has not been elucidated yet. However, chronic inflammation that is hallmark of the type I DM has recently become a key focus in DR study. Hyperglycemia causes dysregulation of nuclear factor-kappa B (NF-κB); and its constant activation increases the expression of inducible nitric oxide synthase (iNOS), cyclooxygenase-2 (COX-2), and intercellular adhesion molecule-1 (ICAM-1) in the retina and vitreous of DR patients and animal models. This increase in inflammatory mediators leads to the degradation of junctional molecules, occludin, connexin-43, death of retinal cells via pyroptosis, and thus disruption of the iBRB. Derailment of the iBRB induces fluid accumulation into the retina of DR patients, which eventually progresses into vision impairment. The purpose of this study

is to investigate whether chronic inflammation in DM induces disruption of iBRB; and whether inhibition of the expression and activity of NF-κB by Methyl-N¹-(3-phenyl propyl) benzene-1, 2-diamine (JSH-23) would mitigate this iBRB disruption.

Methods: Type 1 diabetic mouse model (Akita) on a C57BL/6J background along with non-diabetic littermate were used in the experiments. All mice were 22-24 weeks of age at the start of the experiments. The experimental and the control animals were treated for four weeks on alternate days with JSH-23 @5mg/Kg body weight. The intraocular pressure (IOP), blood pressure, blood glucose, and body weights of all animals were measured before and after treatment. Fluorescence angiography (FA), optical coherence tomography (OCT), and dark-adapted electroretinogram were performed before and after the treatment. Also, intraperitoneal glucose tolerance test (GTT) were performed in the animals before and after the treatment. Then, the retinas and plasma were collected. Western blotting and real-time polymerase chain reaction were performed to analyze protein expression and the mRNA levels of NF-κB p65, iNOS, COX-2, ICAM-1, occludin, connexin-43, NLRP3, caspase-1, and IL-1β. Additionally, plasma insulin levels were also analyzed via ELISA kit.

Results: The body weight of Akita mice was significantly lower than the C57BL/6J mice. The treatment with JSH-23 did not have any effect on body weight. The blood glucose of Akita mice was significantly higher than the blood glucose of non-diabetic littermate control mice. The treatment substantially decreased blood glucose in Akita mice. Also, the plasma insulin level for the Akita mice was significantly lower

compared to non-diabetic littermate control mice. The intraperitoneal

GTT showed that the Akita mice were unable to lower their blood glucose levels. The total and phosphorylated protein expression levels of NF- κ B p65 were significantly decreased after treatment with JSH-23 in Akita mice, and the gene expression of NF- κ B p65 also reduced. The protein and gene expression levels of iNOS, COX-2, and ICAM-1 were reduced after treatment in Akita mice, and the treatment significantly lowered the expression of pyroptosis marker NLRP3. The gene expression of other proteins in the pyroptosis pathway were also reduced, including caspase-1 and IL-1 β . On the other hand, the protein and gene expression levels of junctional molecules occludin, connexin-43 were significantly increased in Akita mice after treatment with JSH-23.

The blood pressure of the animals was normal and the treatment did not have any effect on the blood pressure. FA depicted an increase in retinal capillary permeability in Akita mice, and the treatment significantly reduced that permeability. The OCT showed retinal layers thickening in the retina of Akita mice, and treatment increased the retinal layers thickness. Electroretinogram (ERG) showed an increase in the amplitude of the A-wave and the B-wave in Akita mice.

Conclusions: The results showed that inhibitor (JSH-23) improved visual function by decreasing the retinal vascular permeability, IOP, and the retinal inflammation in Akita mice.

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CHAPTER I

GENERAL INTRODUCTION

<u>Retina</u>

Phototransduction carried out by retinal neuronal cells, supported by retinal glial cells (RGC), is the primary function of the retina [1-4]. Phototransduction is divided into three parts: a receptor (rhodopsin), G-protein (transducin), and an effector (cGMP phosphodiesterase or PDE/PDE6). It starts with 11-cis retinal absorption of a photon that leads to a conformational change in 11-cis retinal to 11-trans retinal. Rhodopsin is activated (metarhodopsin II), which then enables transducin [3, 5].

Transducin activate cGMP phosphodiesterase (PDE). PDE, hydrolyze cGMP, lead to the closing of a cGMP-gated ion channel in the membrane. PDE decreases cGMP concentration; as a result, the cGMP-dependent gated channels in the plasma membrane closed. Uptake of cation reduces in the photoreceptor outer segment (POS); the membrane hyperpolarizes. That decreases the secretion of glutamate. Termination of phototransduction involves inactivation of rhodopsin, transducin, PDE, and increasing concentration of cGMP. Rhodopsin kinase (GRK1) phosphorylated rhodopsin; protein arrestin (Arr1) complete the inactivation by binding to the receptor [6, 7]. G-PDE inactivated by RGS9-1- G-protein signaling protein [8]; Guanylyl cyclase restored the concentration of cGMP [9, 10].

The vision cycle is divided into dark and light periods. The dark period leads to a

constant flow of Na⁺ into the photoreceptor, thus depolarizing the plasma membrane. Depolarization triggers the release of glutamate however during the light cycle, the opposite happens, the flow of Na⁺ stops, and thus the plasma membrane hyperpolarizes and the glutamate secretion stops [11-13].

Rods photoreceptors are sensitive to blue-green light (peak ~ 500 nm wavelengths). In fact, they are very sensitive and at the same time highly efficient in capturing a single photon [14]. Visual pigments in cones are the opsin molecules, and cones sensitivity depends on the molecular structure of opsin for the long wavelengths (red light), short-wavelength (blue light), and medium wavelength (green light).

Anatomy and Physiology of the Retina

The retina is a semi-circle with a diameter between 30 and 40 mm, 0.5 mm thickness, lining the posterior part of the eye. All the major blood vessels of the retina converge towards the center to form the optic nerve. The foveal (the center of the macula) contains no blood vessels and only cone photoreceptors are located to the left of the optic nerve. The retina is divided into two areas: the central (macula) and the peripheral retina. The central retina is the circular area with a 6 mm diameter from the fovea, while the peripheral retina is everything outside of the central retina [15].

The retina is further divided into ten different layers of cells and cell bodies such as outer nuclear layer (ONL), inner nuclear layer (INL), inner plexiform layer (IPL), outer plexiform layer (OPL), ganglion cell layer (GCL), outer limiting membrane (OLM), nerve fiber layers (NFL), inner limiting membrane (ILM), and the retinal pigment epithelium (RPE). The cell layers contain ganglion, photoreceptors, and epithelial cells. The ganglion cells (GCs) are closest to the retina and serves as the output of photoreceptors and input to the optic nerve. The photoreceptors capture and transduce light, while the epithelial performs many functions to maintain retinal homeostasis.

The cell bodies of first-order neurons (rods and cones photoreceptors) reside in the ONL. The INL houses the cell bodies of second-order neurons (bipolar, horizontal, and amacrine cells). First-order and second-order neurons synapse in the OPL. GCs and second-order neurons synapse in the IPL. The OLM contains adherens junctions between photoreceptors and Müller cells, while the ILM houses the Müller cells. The OLM forms a barrier between the outer retina and the subretinal space, while the ILM forms a barrier between the inner retina and the vitreous humor. All retinal cell layers are perfused mainly by the central retinal artery (CRA).

The retina has two main blood supplies: choroid (perfused photoreceptors) responsible for about ³/₄ of retinal blood supply, and the CRA (perfused inner retina), responsible for the last ¹/₄ of the retinal blood supply [16]. The retinal layers are perfused via retinal capillaries. The retinal capillaries form a network throughout the retina, radial peripapillary capillaries (RPCs), inner and outer layer capillaries. The RPCs is located within the NFL and is alimented about 5 mm from the optic disc, inner capillaries alimented the GCL, while the outer capillary aliments the IPL, OPL, and the INL [16, 17].

Outer Plexiform Layer (OPL). Second-order neurons synapses with photoreceptors in the OPL to propagate the signals. The OPL contains mainly two types of cells: bipolar and horizontal cells. The retina contains 11 different types of bipolar cells, all except one synapse with cones photoreceptors [18-21]. Bipolar cell bodies reside in the INL, with dendrites located in the OPL. Some photoreceptors synapse with one bipolar cell while other with many bipolar cells. There are also some bipolar cells that are photoreceptors specific, e.g., S-cone. Bipolar cells designated as either ON or OFF [19]. Bipolar cells

responded differently to glutamate. Based on the receptor, it could lead to depolarization or hyperpolarization of the plasma membrane. Depolarization indicates an ON-bipolar cell, while hyperpolarization indicates an OFF-bipolar cell [22]. The ON-bipolar cell has mGluR6 glutamate receptors [23, 24] while OFF-bipolar cell has iGluR AMPA-kainate glutamate receptors [25]. The retina has only two distinct types of horizontal cells, which are layered horizontally in the OPL [21]. In the presence of glutamate, horizontal cells hyperpolarize. The IPL further carries the signal to amacrine and GCs.

Inner Plexiform Layer (IPL). Bipolar cells synapse with amacrine and GCs within the IPL. There are various types of amacrine cells [19, 26]. They used serotonin, acetylcholine, and other neurotransmitters for further transducing phototransduction towards GCs [27-29]. They are various types of GCs as well [30, 31]. GCs are divided into alpha, beta, gamma, and G4 to G23 [31]. Beta GCs are the most prevalent, approximately half of all the GCs [32, 33]. The IPL separates the ON-bipolar and OFF-bipolar cells by dividing the GCs into two compartments below the amacrine cells. The ON-bipolar cells interact only with a compartment while the OFF-bipolar cells interact with the other compartment [34-39]. The optic nerve synapses with the GCs and sends the information to the brain.

Visual Responses of Ganglion Cells. Similar to bipolar cells, GCs are divided into ON, OFF, and ON-OFF centers cells based on their response to light. ON-GCs fires a burst of action potential (AP) upon stimulation then maintain the firing at a short interval. ON-OFF-GCs fires AP when activated and then at the end of the stimulus, while OFF-GCs centers fire AP at the end of the stimulus [40, 41]. OFF-GCs cells synapse with axon terminal of OFF-bipolar cells in the proximity of the amacrine cell bodies, ON-GCs cells

synapses with ON-bipolar cells, while ON-OFF cells do both [36, 42]. As stated earlier, GCs receive the final output of the signals, and they then send that output through the brain via the optic nerve for interpretation. Another structure critical to retinal function is the RPE.

Retinal Pigment Epithelium. The RPE located between the choroid and the POS. RPE apical side faces the POS, while its basolateral side faces the Bruch's Membrane. It has pigments granules in a single sheet of cell shape that resembles a hexagon that links together via tight junctions. The RPE contains organelles capable of digesting POS into phagosomes. The RPE functions to absorb light, buffer ion, phagocytoses, epithelial transport, secretion, and immune modulation.

Phototransduction results in a high amount of photo-oxidative energy within the retina [63, 110]. The RPE located in the middle of the outer retinal blood flow (choroid) [117]. The choroid perfuses substantially higher than the kidney [111, 112], while the retina extraction of oxygen is significantly low, thus the choroidal venous blood is 90% saturated with oxygen. RPE also phagocytoses POS. These factors increase the free radicals levels in the RPE [63, 110]. RPE uses pigmented granules to defend itself from the oxidative stress (OS) [113] that move to the apical side during light absorption [114, 115]. Aside from the pigmented granules, RPE uses antioxidants, and it has a high capacity to repaired macromolecules such as lipids, DNA, and proteins that damaged due to an increase of OS in the retina [113, 116].

The RPE separates the systemic and the subretinal flow, facilitates the transport of nutrients to the retina, balances the ion in the retinal space, and removes metabolic wastes from the retina. It also transports key phototransduction metabolites to the retina such as glucose, retinal (retinaldehyde; a form of vitamin A), and ω -3 fatty acids. Apical and basolateral glucose transporter 1 (inducible to meet metabolic needs) and glucose transporter 3 facilitate glucose transport into the retina [118]. Retinal transportation initiates with retinal uptake and then induces in the visual cycle— absorption of 22:6 ω 3 fatty are acids used to transport ω -3 fatty acid. The 22:6 ω 3 fatty are acids used to transport ω -3 fatty acid. The 22:6 ω 3 fatty are acids used to transport ω -3 fatty acid. The 22:6 ω 3 fatty acids are also used to synthesize glycerophospholipids for incorporation into the membrane of photoreceptors [119].

The retina contains the highest cell density of any tissue in the human body and its cells have a substantial metabolic turn-over, which increases the production of water and thus its accumulation into the retina. Also, intraocular pressure (IOP) orient movement of water towards the retina, further contributing to its accumulation into the retina. There is also a substantial production of lactic acid in the retina (19 mM) due to retinal metabolic activity [121]. RPE remove water via active transport through aquaporin-1 located both in the apical and basolateral membrane of the RPE [122, 123]. The energy for that transport is obtained via the ubiquitous Na^+/K^+ pump located on the apical front of the RPE which is located on the basolateral membrane. Lactic acid is eliminated via mono-carboxylate transporter-1 (MCT1) (apical) and mono-carboxylate transporter-3 (MCT3) (basolateral). The pH regulation is mandatory for facilitating the transport. MCT1 and MCT3 use H⁺ for the transport and is regulated via apical Na⁺/H⁺exchanger and Na⁺/HCO³⁻ (1:2) co-transporter and basolateral Cl/HCO³⁻ exchanger [124]. Through the transport of ions across its epithelial junctions, the RPE regulates the balance of ions in the subretinal space. Additionally, to maintain homeostasis, RPE needs to compensate for the fast change in ions in the subretinal space [125].

K⁺ ions are the main ion concentration that must be in check. During the dark cycle, photoreceptors channels on the POS uptake Na⁺. The influx is balanced via the apical membrane of the membrane of the RPE by releasing K⁺ ions into the subretinal

space. During the light cycle, the cations channels are closed; thus, there is a decrease in the outflow of K^+ from the inner segment. Also, the Na⁺/K⁺-ATPase pump of the POS takes up K⁺. All these events combine to decrease the K⁺ concentration in the subretinal space. RPE K⁺ channels such as Kir7.1 in the apical membrane release K⁺ into the subretinal space to maintain [K⁺] concentration [125, 126].

The RPE is responsible for the conversion of 11-trans to 11-cis retinal to maintain proper visual conduction [129]. First, the trans-retinal is transported through the POS from the intradiscal space via an ATP-driven flippase. Trans-retinal is converted to N-retinylidinephosphatidylethanolamine by reacting with phosphatidylethanolamine. Then retinol-dehydrogenase reduces trans-retinal to trans-retinol in the membrane. Transretinol move to the subretinal space and the interphotoreceptor retinoid-binding protein (IRBP) carries the trans-retinol to the RPE. In the RPE, cellular retinol-binding protein transfers the trans-retinol to a protein complex. This complex contains three enzymes that mediate the conversion of trans-retinol to cis-retinal. Lecithin retinol transferase (LRAT) adds an acyl to the retinol. RPE65 (RPE protein with 63 kDa) catalyzes the reisomerization of acyl-retinol. And the 11-cis-retinol dehydrogenase oxidizes retinol to 11-cis retinal.

Photo-transduction destroys the POS, which is then replenished by the RPE via RPE shredding of the damaged part and phagocytosis [130]. This process is regulated by the circadian rhythm. It is a daily process that occurs early in the day and activated by light [131]. The phagocytosis mechanism is regulated via three receptors: $\alpha V\beta 5$ integrin, CD36, and receptor-tyrosine kinase c-mer (MerTK) [132]. The $\alpha V\beta 5$ integrin mediates the binding of POS to the RPE, MerTK is phosphorylated via focal adhesion kinase [133]. That leads to the production of inositol-1, 4, 5-triphosphate leading to an increase

in Ca^{2+} concentration inside the cells, and CD36, thus activating the phagocytosis. The blood-ocular barrier is another factor that plays a crucial role in the homeostasis of the retina.

Figure 1. Schematic of Retinal Layers. The retina function to transmit light from the outside world to the brain via the photoreceptors and second-order neurons. The retina contains many different layers of cells and synapses, each played a distinct role in phototransduction. From the outer layer to the inner layers, OLM, RPE, ONL, OPL, INL, IPL, GCL, NFL, and the ILM.



Blood Ocular Barrier

The blood-ocular barrier (BOB) system is divided into two parts: blood-aqueous and blood-retinal barriers. BOB restricts the passage of ions and other blood-borne molecules from penetrating the retina through the aqueous humor and the vitreous [43]. Breakdown of the BOB allows molecules and other drugs to diffuse to the retina and induce inflammation. Another factor that contributes to retinal homeostasis is the downregulating immune environment (DIE) in the eye [44]. Degradation of the bloodretinal barrier (BRB) and loss of the DIE are hallmarks of retinal diseases [45]. The BOB system, along with the DIE, is critical in downregulating excessive inflammatory response and in maintaining the retinal homeostasis.

The BOB contributes to the eye being a privilege tissue; thus helping the eye abstains from variations in systemic circulations. The BOB maintains an ideal environment for the avascular tissues in the eye and regulates the diffusion of chemicals into the retina. Because once the molecules pass the ocular barriers, there is nothing that can prevent their dissemination to the retina [46]. The blood-aqueous barrier is formed by the ciliary body non-pigmented epithelium, Schelemm's canal endothelium, and the endothelium of the Iridal capillaries [47]. Whereas, the BRB consists of tight junctions that restricts the movement of molecules in and out of the retina.

Blood Retinal Barrier

Blood Retinal Barrier. BRB consists of the inner and outer BRB. The inner BRB (iBRB) formed by tight junctions (e.g., Occludin) between the adjacent endothelial cells (ECs), and this endothelial layer rests on a basal lamina that is covered by pericytes and glial cells (astrocytes, and Müller cells) [48, 49]; whereas the outer BRB is maintained by tight junctions between adjacent RPE cells [48, 50]. The BRB functions to control the

movement of molecules and fluid into the retina also prevents leakage of macromolecules and harmful agents into the retina.

Endothelial Cells. The endothelial cells (ECs) provide the necessary nutrients and oxygen the retina needs to maintain its physiological functions while limiting other blood circulating agents from penetrating the retina. ECs mitigate transcellular and paracellular transport into the retina. Transcellular transport occurs through diffusion across the lipid bilayer for a host of small lipophilic molecules. The other mechanism of transportation is through receptor-mediated or constitutive transcytosis. Caveolin-1 contains receptors for many substances, including interleukin 1, vascular endothelial growth factor (VEGF), albumin, advanced glycation end-products (AGES), and protein kinase C (PKC). Vesicular transport under normal conditions is minimal [51]. Tight junctions restricted the paracellular traffic.

Pericytes. In the formation of retinal vasculature, ECs express a very high level of platelet-derived growth factor B (PDGF-B). It leads to the recruitment of pericyte cells, which then wrap ECs in a 1:1 to ratio. They also form cell-cell contact via Cx43, fibronectin adhesion plaques, and N-Cadherin. Pericytes also secrete angiopoietin-1 which acts via the Tie2 receptor on ECs to promote differentiation, remodeling, and maturation. Pericytes also secrete Angiopoietin-2 (Ang-2) which moves together with VEGF to promote angiogenesis. Overexpression of Ang-2 causes vascular regression, impairs barrier function in part by inducing degradation of TJ (Cadherin). Pericytes also secrete transforming growth factor-beta (TGF- β) which inhibits migration and proliferation of ECs [51].

12

Junctional Molecules

ECs form a restricted lining of the vascular wall to regulate the passage of substances circulating in the blood and the interstitial space whereas, paracellular limit transport via adhesive cell-cell junctional proteins that include gap junction (GP) and tight-junction (TJ) [52].

Gap Junctions. GJ allowed cell-to-cell communication via hemichannels that are consisted of six connexins molecules. Each cell contributes a hemichannel to the connection, which opens the plasma membrane of each cell to the other. GJ allowed the transfer of small molecules (< 1 kDa) such as ions, glucose, amino acids, and neurotransmitters. Most cell types in the retina has GP, including photoreceptors [53], endothelial, and pericytes [54]. Connexin-43 (Cx43) is the most prevalent connexin molecule in the retina [55].

Cx43. Connexin generally localizes in the plasma membrane where it functions in intercellular communications via GJ. Cx43 has various phosphorylation sites in its carboxy-terminal for protein kinases, e.g., PKC. Cx43 is synthesized as a 40 to 42 kDa protein which is then repeatedly phosphorylated through serine residues, and is assembled in the trans-Golgi network, and then transported and inserted through the plasma membrane. The phosphorylation usually functions in intercellular communications via various molecular mechanisms related to assembly, trafficking, and closing of the GJ channel [56, 57]. Cx43 hemichannels open during low [Ca²⁺] extracellularly, while PKC can block its opening.

Tight Junctions: TJ are formed in proximity between two adjacent cells. TJ restrict

transports of blood circulating molecules paracellularly. TJ also regulates lipid flip flop across the lipid bilayer. They are consisted of more than 40 proteins that are generally divided into transmembrane proteins and cytoplasmic scaffolding proteins [51].

Occludin: Occludin is a part of the transmembrane class of TJ proteins and is a 60 kDa tetraspan transmembrane component protein with two loops facing the extracellular space while both the amino and carboxy-termini in the cytoplasm. The amino terminus regulates occludin degradation via interaction with E3-ubiquitin ligase; while the carboxy terminus binds to one of the three Zona Occludentes proteins. Occludin functions more in regulating barrier properties rather than structural properties between ECs. Overexpression of occludin leads to decrease flux into the interstitial space. Downregulation of occludin increases paracellular permeability. Occludin knockout mice depicted that occludin is not required for barrier formation, but it does play a key role in TJ trafficking and stabilization [51]. Occludin and claudins are the two protein/protein family that link the ECs together [58].

Glia cells and the BRB

There are three different types of glial cells that populated the retina: Müller cells, astrocytes, and microglia. The macroglial cells (Müller and astrocytes) play significant roles by maintaining the function(s) of the iBRB.

Müller Cells. Müller cells are the primary macroglial cells in the retina and constitute 90% of the glial cells in the retina. Müller cells act as a linking cells for metabolites

exchange between the neuronal cells and the blood vessels [59]. They function in neurotransmitters recycling, K⁺ uptake, regulation of blood flow, metabolism control, nutrients supply to retinal neurons, anti-oxidative support [60], glycogen storage [61], sheath axons [59], and in maintaining the BRB. They can also interact closely with both retinal neurons and blood vessels due to their long shape [62]. Docosahexaenoic (DHA) protects photoreceptors disk membranes from oxidative damage. DHA also induces increased expressions of anti-apoptotic factors via the Bcl-2 family. Müller cells function in uptaking and delivering DHA to photoreceptors [59]. The depletion of Müller cells leads to the breakdown of the BRB [61]. These cells secreted anti-angiogenic factors to maintain pro and anti-angiogenic environment and thus BRB health.

Astrocytes. Astrocytes are non-prominent in avascularized area, generally located within the NFL. They are wrapped around blood vessels. Astrocytes primarily communicates via GJ [59]. Astrocytes are the primary producer of VEGF and thus play a key role in neovascularization. Other physiological functions of astrocytes include mechanical support for axons, neurotrophic support, and maintenance of the BRB. Astrocytes contained aquaporin4 channels within their plasma membrane which help regulate the fluid in the extracellular space, buffering potassium, and clear the metabolic wastes [59].

Figure 2. A Brief Schematic of the iBRB. iBRB consists of TJ between adjacent ECs that are covered by pericytes. The capillary cells are wrapped by glial cells (astrocytes, and Müller cells). Although pericytes and glial cells are critical to the iBRB function, the transcellular and paracellular transport into the retina is regulated by the ECs and TJ together.

FIGURE 2



Downregulatory Immune Environment (DIE). The immune-privileged status of the eye is maintained by the BOB and DIE. Transportation of antigens and tissue drainage occurred via the blood route instead of the lymphatic pathway [63]. The inflammatory response in the eyes is particular to the eye. To protect the physiological functions, the eye regulates immune response strictly and manages its inflammation. The retina and choroid contain many immune cells that include choroidal macrophages, microglia, dendritic cells, and the mast cells population [64]. These cells introduce antigen to T cells and produce cytokines to non-specific stimuli such as bacterial liposaccharides (LPS). The presence of antigen-non-specific inflammation can be disastrous to host tissue. The BRB protects the retina from the aqueous chamber, and the vitreous body via immunosuppressive functions. Many factors played a crucial roles in maintaining the DIE state in the eye, including Müller cells, RPE, and the microglia. The normal retina has low major histocompatibility complex (MHC) molecules and there is an absence of the bone marrow-derived cells that function with tissues as Antigen Presenting Cells (APCs).

Anterior Chamber Associated Immune Deviation (ACAID). ACAID is the most studied immune privilege mechanism in the eye. It is unique to the anterior chamber, but there is evidence that the same mechanism exists in the vitreous and the subretinal space as well [65]. It contains TGF- β_2 which is capable of reprogramming the APCs to produce less IL-2, less amount of CD-40 in the microenvironment of the eye, and also help produce more TGF- α which suppresses T cell activation [46].

BRB and DR. DR is commonly defined as degeneration and occlusion of the microvessels, which caused fluid accumulation and abnormal neovascularization in the

retina [66]. Loss of pericytes in retinal capillaries is one of the factors that leads to iBRB breakdown [67]. Hyperglycemia (HgA) causes retinal hypoxia and vasodilation increases retinal blood flow in response to the hypoxia temporarily. Exhaustion of autoregulation causes apoptosis of retinal capillary pericytes. The death of pericytes disrupts pericytesendothelial complex which then leads to an impairment in the endothelial TJ function. These events together caused the breakdown of the iBRB.

Retinal ECs in the iBRB are potential targets of the inflammatory markers such as eicosanoids, cytokines, and the growth factors. Chronic HgA upregulates inflammatory mediators which in turn induces leukocytes adhesion to ECs which further injures the iBRB [68-70]. Breakdown of the iBRB as a result of cytokines release generate an inflammatory response which further increases the release of the inflammatory mediators, thus additionally exacerbating the situation. Leukocytes adhesion also induced retinal hypoxia, the retinal hypoxia also leads to abnormal blood vessel growth concurrently [71, 72].

Diabetic Retinopathy

The earliest sign of DR is a dysfunctional iBRB. DR is characterized by loss of pericytes [73, 74], basement membrane thickening, microaneurysms [75], acellular capillary, and iBRB breakdown. The factors that contribute to capillaries degeneration are occlusion of the capillary's lumen by white blood cells or platelets, and death of capillaries cells due to hyperglycemia. There are many biochemical mechanisms proposed for the pathogenesis of DR, although the etiology is yet to be precise. Chronic HgA is the main factor contributing to the development of DR, and its complications. The earliest retinal changes due to HgA are biochemical, cellular, or hemodynamic in nature. Metabolic deregulation affects

the neurons and the support cells of the retina. However, it is the microvessels that in fact are affected the most. They [capillaries] dominate clinical events of the disease and are also the focal point for the development of macular edema and neovascularization. In the early stages of DR, functional cells of the retina exhibit changes such as glial cell reactivity, alterations in glutamate metabolism, and neuronal cell death [76-78]. Until the development of retinopathy, vision is normal in diabetics patients if they don't have any other concurrent eye problems. DR is divided into two stages: Non-proliferative DR (NPDR) and Proliferative DR (PDR). After five years of living with diabetes, NPDR symptoms start to appear. NPDR is intraretinal microvascular derangement that could progress to diabetic macular edema (DME) (accumulation of fluid in the macula) or PDR (extraretinal neovascularization). Progressive retinal ischemia and the rise in local growth factors are the primary mechanisms outlined in the development of PDR. PDR is new growth of blood vessels in the retina or optic disc that extends along the retinal surface or into the vitreous cavity [79, 80]. Pathogenesis of PDR involved hypoxia that results in low perfusion of retinal tissue, thus prompting neovascularization (directly related to VEGF). Initially, the new blood vessels are barely visible, then they gradually increase in size, and could become as large as a retinal vein. New blood vessels are usually asymptomatic [81, 82]. The new growth of blood vessels is gradually accompanied by fibrous proliferation, and it is this contraction of the fibrous tissue that can can lead to retinal detachment and vitreous hemorrhage. These two traits more often than not lead to vision loss in PDR.

The cornerstone of NPDR treatment is glycemic control while managing PDR has the following aims: decrease the growth of blood vessels, contraction of vitreous, and fibrovascular proliferation. Photocoagulation is the gold standard for PDR treatment. However, photocoagulation comes with complications since high density/intensity laser may cause loss of night and colored vision [83] and reduced contrast sensitivity [84]. Other treatments for PDR include VEGF Inhibitors and Vitrectomy [85].

Many biochemical mechanisms have been put forward to explain the vascular pathology of DR. Among the proposed pathways are aldose reductase (AR), AGEs formation, reactive oxygen intermediates elevation, PKC, and the inflammation.

DR-Biomolecular Mechanisms

AR Pathway. Retinal capillary cells (endothelial, and pericytes), and glial cells expressed AR and thus are subjected to the polyol pathway [86]. The polyol pathway or the AR pathway is induced during high blood glucose levels [87]. First, AR reduces glucose into sorbitol (nicotinamide adenine dinucleotide phosphate hydrogen; NADPH serves as a cofactor). Next, sorbitol dehydrogenase (SD) reduces into fructose (nicotinamide adenine dinucleotide (NAD⁺) serves as a cofactor). Since sorbitol cannot cross across the cell membrane it accumulates intracellularly inducing an osmolality effect [87]. Next, as the substrates use NADPH and NAD⁺ in this pathway, they are quickly depleted thus decreasing the body's capacity in responding to the OS. For example, the use of NADPH hinders its bioavailability for glutathione reductase [88]. NADH/NAD+ ratio is increased due to the use of NAD⁺ by SD. Further, NADH oxidase could also use NADH as a substrate to increase the generation of ROS [89]. Additionally, fructose can further metabolize into fructose-3-phosphate, and 3-deoxyglucosone, both of which can become the glycosylating agents [90, 91]. This pathway generates osmolality effects, glycating agents, and OS. Studies have shown that inhibition of AR in diabetic animal models prevents retinal injuries [92]. Although clinical trials with AR inhibitor have not been significantly fruitful [93].

AGEs. Amine group modification of a protein by reducing sugars such as glyoxal, 3deoxyglucosone, and methylglyoxal, generate AGEs [94]. Retinal levels of AGEs increase in diabetic's rat model and humans [95]. The presence of AGEs on the basement membrane interferes with pericytes and ECs communication. AGEs bind to their receptors, activate NF- κ B, and generate ROS.

PKC: PKC is a serine/threonine-protein kinase that mediates many cellular signal transduction pathways [96]. The various isoforms of PKC are activated via diacylglycerol (DAG), calcium, and other substrates. HgA increases the expression level of DAG [97]. Studies reported that DAG levels have increase in the retina of the diabetics [98]. Activation of PKC increases albumin permeability across ECs [99, 100] and hinders vasodilation via depleting nitric oxide (NO) levels. PKC also caused the loss of pericytes [101].

OS: The retina is perfused 300 times more than both the heart and the cerebral cortex, thus making it more susceptible to OS; phototransduction, and shedding of POS by the RPE also contribute. ROS, free radicals in the form of hydroxyl radicals, and superoxide anion are very reactive, mainly due to the unpaired electrons in their valence shell [102]. Many factors lead to the generation of ROS including the mitochondria electron transport chain [103]. Under normal conditions, the mitochondria contribute the most to ROS levels. The process of aging, and inflammation can disrupt homeostasis by generating more ROS than the body can remove thus inducing the process called OS [104, 105]. The body uses antioxidant (vitamin C and E, superoxide dismutase; SOD, and catalase) to neutralize ROS [102]. Once the body anti-oxidant system is overloard that create OS.

HgA activates PKC, polyol flux, and AGEs that generate ROS [106].
Increased ROS injure ECs, activate NF- κ B [107-110], and recruit neutrophils [111]. There is evidence that OS is the main factor that triggers the chronic inflammation response in the retina [112]. Inflammation increases expression of inflammatory mediators, promotes leukocyte adhesion to retinal capillary walls, and occludes retinal capillaries causing hypoxia. Eventually, breakdown of the iBRB happens.

Inflammation

Basics Concept in Immunology. Immune privilege of the eye is maintained by the BOB, DIE, and the lack of direct lymphatic drainage from the eye [113]. The immune system consists of two parts: innate and adaptive. The innate immune pathway is immediate and nonspecific. Pattern recognition receptors (PRR) recognizes the conserved microbial patterns and triggered a nonspecific inflammatory reaction; for instance, toll-like receptors (TLR) recognize LPS of the gram-negative bacterial outer membrane [114] which activate immune cells, monocytes, macrophages, natural killer (NK) cells, polymorphonuclear cells, eosinophils, and basophils. The adaptive immune response, however, is more complicated and elaborated below.

Adaptive Immunity. The key components include B and T lymphocytes, which contain receptors (immunoglobulins; Igs) for recognizing antigenic epitopes [115]. Igs are expressed by B cells and T cells in the absence of exogenous stimulation. Igs contains unique configurations that recognize specific antigens. Each group of lymphocytes depicts the same cell surface receptors to identify multiple epitopes, and to accomplish this billions of lymphocytes are generated.

Figure 3. Mechanism of Diabetic Retinopathy Pathology. Hyperglycemia during diabetes activates many pathways such as PKC, AGEs, and polyol. These pathways induce oxidative stress, and inflammation in the retina. Inflammation contributes to retinal vascular occlusion, and dysfunction of retinal capillary cells that lead to the development of DR. DR starts as NPDR, then gradually progress to PDR or DME or both.





After maturation in the thymus (T cells) or bone marrow (B cells), lymphocytes remain in the dormant or G0 phase of the cell cycle until an antigen binds to its receptor (called naïve). Molecule binding to a B or T cells triggers an immune response to destroy the bound molecule or the antigens. If the molecule is pathogen-derived then it leads to the destruction of the pathogen. If it is a cell-derived then it leads to autoimmunity, and therefore occurrence of the tissue injury.

Activation of a naïve lymphocyte leads to the proliferation of that lymphocyte which recognizes a specific epitope. T cells (CD4⁺ and CD8⁺) become cytotoxic once activated or differentiate into effector cells. CD8⁺ T cells become cytotoxic once activated. CD4⁺ T cells via interleukins (IL) differentiate into one of three effector T cells (Th1, Th2, and Th17). IL-4 leads to Th2, IL-12 leads to Th1 [116], whereas IL-1 and IL-23 lead to Th17 differentiation [117]. Th1 and Th2 cells become memory cells that trigger the immune response if the antigens reappear. Th1 secretes cytokines that target macrophages, and other cells involved in the immune response. Th2 stimulates B cells proliferation to produce antibodies to kill the invader. Th17 cells produce IL-17 [118], and IL-23 to fight fungal and bacterial infections.

The adaptive immune response is divided into three parts: antigens recognition, transportation, and presentation to lymphocytes. Antigens are recognized, processed, and presented (through MHC) to lymphocytes by dendritic cells, and macrophages. There are differences in the process of antigen presentation. B cells can engage with naïve antigen directly but T cell receptor (TCR) can only recognize peptide fragments presented with MHC molecules (Class I or II). MHC Class I molecules (expressed in most cells) present peptides derived from protein degradation in the cytoplasm (intracellular antigens). MHC Class II molecules (expressed in APCs and lymphocytes) present peptides derived from phagocytic vesicles (extracellular antigens). APCs provide co-stimulation needed for lymphocytes activation [119]. APCs upregulates CD80, CD86, ICAM-1, and CD40 molecules to function as ligands for lymphocytes receptors. APCs also secrete soluble cytokines such as IL-12, IL-6, IL-10, and IL-1 α to stimulate lymphocytes further.

CD4⁺ T-cell activation leads to CD4⁺ proliferation and cytokines secretion such as IL-2, IL-3, granulocyte-macrophage colony-stimulating factor, interferon-gamma (IFN- γ), and IL-4 which serves as growth and stimulation factors for lymphocytes and APCs. IL-2, IFN- γ , and IL-4 promote differentiation of CD8⁺ T lymphocytes to mature cytotoxic T lymphocytes. Cytokine production by CD4⁺ T cells also promotes activation and differentiation of B lymphocytes. IL-2, and IFN- γ stimulate B cell to produce immunoglobulin g (IgG) while IL-4, 5, 6, and 10 stimulate the production of IgG, IgE, and IgA antibodies.

Final products of processing in the lymphoid tissues are activated effector CD4⁺, CD8⁺, and B cells. These cells travel through the blood to execute the response. Activated B cells secrete antigen-specific antibodies to kill the target directly. Activated CD4⁺ T cells secrete cytokines, and IFN- γ , which in turn recruit the cells of the innate immune system to the site of the antigen. Activated CD8⁺ lyse target cells and produce proinflammatory cytokines, especially IFN- γ . Cytokines involved in the response perpetuate a positive feedback loop at the target sites. Blood vessels become leaky, vascular endothelial cells display ligands that bind to immune cells receptors, and chemokines secreted by local inflammatory cells to attract more immune effector cells and the cascade continues [119].

Inflammation in DR

DR is regarded as an inflammatory disease due to the presence of inflammatory

mediators that are found in the vitreous and the retina of diabetic animals models and the patients. The increase in expression of these markers is associated with a dysregulation in nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) [120].

NFKB. NF- κ B is a family of five transcription factors that has structurally related subunits: NF- κ B 1 (p50), NF- κ B 2 (p52), RelA (p65), RelB, and c-Rel. Once activated, they form either hetero or homodimers and get translocated into the nucleus. In the nucleus, they bind to κ B DNA binding sites and increase the expression of NF- κ B targeted genes [121].

NF-κB regulates gene transcription that plays key roles in inflammation, immune cell development, cell cycle, proliferation, and cell death [122, 123]. NF-κB is pertinent for both innate and adaptive immune cell functions. It regulates survival, activation, and differentiation of both innate and adaptive immune systems cells. NF-κB activation induces expressions of pro-inflammatory genes, cytokines, chemokines, immunoreceptors, apoptosis, proliferation, and inflammasome [120].

Stimuli that activate NF- κ B include TLRs, antigen receptors, OS markers, genotoxic stress, cytokine, and growth factors. There are two types of NF- κ B activation: specific and non-specific. Both are influential in regulating inflammation and immune response [124]. The NF- κ B subunits reside in the cytoplasm where they remain inactive via inhibitory proteins of the I κ B (I κ B α) family [125]. Activation of NF- κ B requires I κ B kinase (e.g., IKK α) to phosphorylate the I κ B α which then is recognized by the E3 ligase to degrade the inhibitor. This process induces the translocation of NF- κ B heterodimers (e.g., p50/ReIA) into the nucleus [126-128], where they bind to DNA sequences known as κ B sites [121].

Figure 4: Immune System Mechanism. The immune system consists of two parts: innate and adaptive. The innate system is the primitive and nonspecific one and uses dendritic cells, monocytes, macrophages, and other immune cells to destroy the invading pathogen(s). On the other hand, the adaptative system uses the bone marrow-derived (B cells) and thymus-derived Treg cells, innate immune cells, antibodies, and immune ligands (e.g., cytokines) to lyse antigens.



In contrast, the non-specific pathway responds to a specific set of stimuli, such as receptor-activated nuclear factor kappa B (RANK) [129, 130]. The subunits activated in this pathway are RelB and p52 and require IKK α and cleavage of p100 to p52. First, processing of NF- κ B 2 (p100) occurs [129] followed by translocation of p52/RelB into the nucleus [129, 131]. The specific pathway is involved in all aspects of the immune system, and the non-specific is complementary to the adaptive immune system regulation [130]. NF- κ B is regulated primarily via a negative feedback loop wherein activation of NF- κ B induces production of I κ B α [132]. Activation of NF- κ B mainly heterodimerizes into p50-p65 thus transcribing the pro-inflammatory proteins (iNOS, ICAM-1) [133].

Inflammation serves as a protective mechanism against tissue damages and infections under physiological conditions, however it may be detrimental in the long term. Deregulation of inflammatory responses causes damages to host tissues that could lead to the development of short or long-term inflammatory diseases. Chronic inflammatory conditions such as diabetes involve deregulation of NF-κB activation. Dysregulation of NF-κB induces an excessive increase in inflammatory proteins. Recent studies have shown an increase in inflammatory mediators such as COX-2, and iNOS in the retina and vitreous of DR animal models and the patients, as stated earlier. These molecules are consistent with the innate immune pathway. Also, other studies have also shown that retinal inflammation is also associated with the activation of NF-κB [134].

iNOS. Inducible Nitric Oxide Synthase (iNOS) meditates NO production from Larginine. Regular production of NO is generated via two NO synthases (neuronal and endothelial). While iNOS generation of NO involves solely in inflammation. An elevated level of NO is associated with many inflammatory diseases, e.g., inflammatory bowel disease [135-137]. iNOS has high expression in activated immune cells and low to no expression in most resting cells [135]. Activated NF- κ B induces transcription of iNOS [120]. Studies in the literature shows an increased iNOS levels in the retina of diabetic animals, which associated with apoptosis of retinal cells, leukostasis, and iBRB breakdown [138]. iNOS elevation is very well present in the retina of human patients [139-142]. iNOS has also been shown to mediate the production of oxygen radicals.

iNOS generates superoxide anions in macrophages [143]. NO could react with superoxide anion to form peroxynitrite, and injure macromolecules inside the cells. Increase production of NO could also result in nitration and nitrosylation of the retinal proteins, which is also toxic. NO elevation generated by iNOS causes the loss of pericytes, and retinal thinning (GCL, IPL, and OPL). Studies of retinal cells exposed to high glucose (HG) demonstrated that NO (particularly iNOS generated) regulates prostaglandin E₂ (inflammatory marker generated by COX-2) expressions. iNOS is also associated with GCs loss in Akita mice after 22 weeks of diabetes [144]. Inhibitor of iNOS (aminoguanidine) has shown to decrease vascular injuries associated with DR in rodents [145-147]. Studies of retinal cells exposed to HG demonstrated that NO (particularly iNOS generated) increases the expression of cyclooxygenases-2 (COX-2) [144].

COX-2. Eicosanoids are signaling molecules that are derived from arachidonic acids such as prostaglandin (PG), thromboxane, and leukotrienes. Cyclooxygenase enzymes mediate the first two steps of eicosanoids synthesis. Many isoforms have been identified, such as COX-1, and COX-2. COX-1 is expressed in most tissues and it functions to maintain PG production. PG is known to regulate renal blood flow, immune responses, and inflammation [148-152]. In the eye, it is involved in ocular inflammation [153], IOP

[154], corneal neovascularization [155], corneal edema [156], retinal blood flow [157], iBRB, and BOB impairment [158].

COX-1 maintains physiological productions of eicosanoids, whereas COX-2 eicosanoid production occurs in the presence of inflammation. COX-2 is generally absent from most tissues, but its expression is increased in the presence of bacterial LPS and proinflammatory cytokines; thus, COX-2 has high expression in immune cells (monocytes, macrophages, and ECs) [159]. NF-κB activation upregulates COX-2 gene expression in retinal cells [160]; and is associated with iBRB breakdown [161]. Interestingly, COX-2 expression is elevated in the retina of Akita mice [162]. Another inflammatory mediator that is positively regulated by NF-κB and has high expressions in the retina of DR is the intercellular adhesion molecule-1 (ICAM-1).

ICAM-1. ICAM-1 is a membrane-bound glycoprotein and serve an important function in immune responses, e.g., leukocyte trafficking [163]. ICAM-1 is consisted of five Ig domains, a transmembrane domain, and a linker in the cytoplasm that interacts with actin molecules [164]. It is expressed in ECs, leukocytes, platelets, epithelial cells, and glial cells. Its expression is low when inflammation is not present. It performs functions in immune cells such as adhesion, migration, and aggregation [165]. Leukocytes adhesions (leukostasis) to retinal endothelium is associated with iBRB breakdown [166].

Leukostasis is elevated in the retina during hyperglycemic conditions [166]. Leukostasis in retinal vessels is mediated by elevation of ICAM-1. Activation of NF-κB increases ICAM-1 expression [167]. Leukostasis elevation is linked with an increase in vascular permeability [168]. Inhibition of ICAM-1 reduces leakage and leukocytes adhesion in the retina of diabetic rats [169]. Leukostasis in retinal vessels is mediated by elevation in ICAM-1 and CD18 leucocytes integrin [170]. Streptozotocin (STZ) induced diabetes in mice that are deficient in ICAM-1 shown less leukostasis in retinal vessels [171]. STZ induced WT C57BL/6 J mice exhibited increased expression of ICAM-1 in their retina [169]. Activated leucocytes injected into the retina of healthy mice show increased leukostasis via ICAM-1 [172] and inhibition of ICAM-1 decreases leukostasis in the retina of diabetic rats [173].

Pyroptosis

Pyroptosis is a cell death pathway due to inflammation. It involves the assembly of an inflammasome, activation of caspase 1 (Casp-1), IL-1β, and IL-18. Pathogenassociated molecular patterns (PAMPs) and danger-associated molecular patterns (DAMPs) induce inflammasome assembly [174]. Inflammasome contains a nod-like receptors protein (NLRP) family receptors (e.g., NLRP3), adaptor protein ASC (apoptosis-associated spec-like protein containing CARD) and pro-caspase 1. Inflammasome plays a vital role in the innate immune system against infections, but its deregulation could lead to the development of autoimmune and inflammatory diseases also [175]. NF- κ B regulate inflammasome assembly [176, 177]. NF- κ B is a known activator of nod-like receptor pyrin domain 3 (NLRP3) [176, 177]. NF- κ B induces pyroptosis through activation of NLRP3 [178].

NLRP3. Nod-like receptors (NLRs) are PRR and generally located in the cytoplasm, thus making it possible to detect pathogenic invaders inside the cell [179]. Most NLRs are

Figure 5. Nuclear Factor Kappa B Activation. NF- κ B homo/heterodimerizes and translocate into the nucleus once activated. They bind to κ B DNA binding sites and induce the transcriptions of inflammatory mediators such as iNOS, COX-2, and ICAM-1.



active during the innate immune responses, and are activated via PAMPs, DAMPs, cell stress, or ROS. NLRs family have 14 distinct members, named NLRP1 to NLRP14 [180]. NLRP3 can also be activated by extracellular ATP, and UVB radiation [180]. NLRs form complexes known as inflammasomes, which cleave casp-1, and casp-1 cleaves interleukins-1 beta (IL-1β), and interleukins 18 (IL-18) [181]. NLRP3 is the most studied NLRs and various stimuli could trigger its activation. It is involved in multiple pathologies, e.g., type II diabetes, and obesity [182]. NLRP3 structure has three distinct domains: pyrin domain (PYD), leucine-rich repeat (LRR) domain, and a nucleotidebinding domain (NBD) [183]. NLRP3 inflammasome is consisted of NLRP3, ASC, and Casp-1. Stimuli activate NLRP3 via interaction with the LRR, then the PYD and the NBD are free to interact with other entities. Through the NBD domain, NLRP3 oligomerizes and then recruit ASC to form the NLRP3-ASC complex [184]. ASC then recruits casp-1, casp-1 auto processed, and cleaved pro-IL-1 β to IL-1 β and IL-18 [185, 186]. The NLRP3 activation mechanism is yet to be defined though. A common mechanism has been suggested that the extracellular ATP stimulates purinergic P2X7 ATP-gated ion channel leading to the efflux of K⁺ ions. Pannexin-1 membrane pore protein is recruited into the target cell and that can lead to the activation of the assembly of the NLRP3 inflammasome [180].

HgA can upregulate NLRP3 expression in cultured Müller cells [180]. The role NLRP3 and inflammasomes play in DR is still under intense investigation. Hui Chen et al. reported a high level of NLRP3 expression in the blood from diabetic patients. They speculated that its activation could be due to elevation in ROS that could presumably be a byproduct of other molecular mechanisms, e.g., polyol pathway [187].

Casp-1. Casp-1 is part of the caspase family of cysteine proteases that involve either (a) an increase activation of proinflammatory cytokines or (b) cellular death by apoptosis. However, casp-1 can participate in both cytokines activation and apoptotic cell death. Casp-1 resides in the cytoplasm until activation by NLRP3, and then it heterodimerizes into two subunits (10 & 20 kDa) [180]. Casp-1 activation induces expression of both IL-1 β and IL-18 [188]. HgA has been shown to activate casp-1 in the retina of diabetic mice, patients, and Müller cells [188]. Inhibition of casp-1 by minocycline prevents the degeneration of retinal capillaries cells induced by the development of diabetes [188]. Casp-1 activation induces expression of both IL-1 β and IL-18 [188].

IL-1 β . IL-1 β is part of the IL-1 family which is consisted of 11 different ligands and receptors. These ligands function to induce inflammation or anti-inflammation independently in the blood or tissues. IL-1 β is one of the first ILs discovered. IL-1 β is synthesized as pro- IL-1 β , which remains inactive until it is cleaved via casp-1 to produce mature IL-1 [180]. IL-1 β binds to its receptor (IL-1 receptor type 1; IL-1R). IL-1 β action is regulated via IL-1R antagonist, which competes for the same receptor. IL-1 β functions in the extracellular milieu where it resides within the cytosol and is secreted upon stimulation or cell death. Many different retinal cells produced IL-1 β , including microglia, and the astrocytes [189]. IL-1 β played a crucial role in innate immune response, helps fighting pathogens, and tissue repair. IL-1 β induces the activation of microglia and other retinal glial cells along with proinflammatory mediators. Thus, prolonged exposure or overexpression of IL-1 β can lead to tissue damage. Elevation of IL-1 β expression was reported in the retina of diabetic rodents experimental models as early as eight weeks after the onset of HgA. IL-1 β has also been reported in the

vitreous of patients with DR. IL-1 β administration in the vitreous of nondiabetic rats led to apoptosis of retinal capillary cells via activation of NF- κ B [189, 190]. Susana Vallejo et. al reported that anakinra (inhibitor of IL-1R) prevented impairment of ECs in STZ induced diabetic rats via NF- κ B activation and subsequent increase in iNOS production [191]. Literature also reported that constant exposure to IL-1 β led to ECs impairments and degeneration of blood vessels [188]. Other studies reported that HG increased the expression of IL-1 β in Müller cells [192]. In the retina of diabetic mice, casp-1 and IL-1 β have been detected as well [193]. These markers have also been shown to be implicated in the death of microglia and Müller cells [194]. Furthermore, IL-1 β increases the expression of COX-2 and adhesion molecules in ECs [180].

iBRB and DR

Recent studies reported existence of reduced gap junction intercellular communication (GJIC) thus highlighting the iBRB dysfunction. Cx43 expression in numerous retinal cells is downregulated during HG environment [195]. Studies have shown that HG downregulates Cx43 expression levels in the mitochondria, potentially promoting endothelial and pericytes cell death [196, 197]. HgA showed to impair Cx43 expression, thus decreasing the activity of GJ, and the decrease in expression of the TJ molecules in ECs, and a subsequent cell death phenotype [55].

Studies in astrocytes depicted that activation of NF-κB by TLR-3 protein leads to a decrease in the expression of Cx43, thus reducing GJIC [198]. Kunihiko Nagasawa et al. reported that Cx43 is associated with TJ proteins in the blood-brain barrier (BBB) [199]. Studies in rat retinal ECs depicted that HG decreased Cx43 expression, and this decrease in Cx43 expression downregulated TJ protein, particularly occludin, thus impairing cell permeability but the mechanism(s) is yet to be discovered [200].

HgA decreases expression and redistribution of occludin [58, 201]. Ermelindo Leal et al. found that increased expression of NO generated by iNOS reduced the expression level of occludin in STZ induced diabetic rats, therefore leading to the breakdown of the iBRB [201]. NF- κ B activation induced by TNF- α can internalize occludin which may disrupts the TJ barrier between epithelial cells in inflammatory bowel disease [202].

Loss of pericytes occurs during HgA in part due to a reduction in PDGF-B levels. Additionally, a concomitant loss of both the pericytes and ECs could result in acellular capillaries (no ECs or pericytes). This particular aspect increases the likelihood of microaneurysm formation and bleeding in the retina which would indicate iBRB breakdown. In the presence of IL-1 β , activated pericytes secrete various cytokines and chemokines, including IL-1 β . It secretes these factors to attract immune cells to the sites of inflammation. Pericytes increase the expression of ICAM-1, other adhesion molecules, and at the same time downregulate the TJ molecules [203] thus facilitating immune cell passage through the endothelium.

HG leads to the death of Müller cells via activation of casp-1 and IL-1 β [192]. HgA increased the expression of IL-1 β in Müller cells [61] and its secretion. IL-1 β then bound to IL-R1 on ECs and increased the expression of ICAM-1 [60]. Studies have shown that HgA impaired Müller cells uptake of glutamate, which could lead to glutamate toxicity in diabetic rats. HgA also reduced potassium uptake, further exacerbating the glutamate toxicity by increasing the neuronal excitation [61]. Diabetes impairs Müller cells function via decreasing expression of glutamine synthase, and the activity of the glutamate transporters [204]. Selective ablation of Müller cells results in BRB in Rlbp-LacZ mice which also impairs the A-wave and B-wave responses of the mice [204]. HgA significantly impaired communication between astrocytes by reducing Cx43 expressions and other GJ as early as four weeks post STZ treatment in rats. These animals also depicted the death of astrocytes after six weeks of HgA [205].

Inflammation and Intraocular Pressure

The aqueous humor is secreted by the ciliary body (CB) epithelium and it is the key factor that determines the dynamics of intraocular pressure (IOP). It is the plasma components of the CB that help secrete the aqueous humor into the posterior chamber (PC) which then circulate into the anterior chamber (AC). From there the aqueous humor drains into the trabecular meshwork and Schlemm's canal. The cells in the CB synthesized the aqueous humor in three steps. First, when the blood flows through the CB the plasma in form an ultrafiltrate is made due to the pressure gradient between blood flow and the interstitial of the ciliary body. Last, the epithelium of the CB transports the aqueous humor from its basal end to its apical surface actively (and selectively) and then pumping it into the PC. Systemic blood pressure has very little to do with aqueous humor synthesis, although the first step has to do with blood flow, only about 4% of plasma is being filtered [206].

Circulation of the aqueous humor generates the IOP within the AC. IOP plays a crucial role in the pathophysiology of glaucoma. An increase in IOP is a key risk factor for glaucoma, although some patients with normal IOP may have glaucoma, while others with high IOP may not have glaucoma. One factor that is associated with an increase in IOP is increased resistance in outflow [206].

HgA and IOP. Multiple demographics studies concluded that there is an increase

in IOP in diabetic patients than nondiabetic patients, and that an increase in IOP is independent of open-angle glaucoma. The mechanism has not yet been determined though. It is speculated that HgA causes an osmotic gradient that shifts fluid between plasma and intraocular fluids thus elevating the IOP [207]. A Japanese study found a significant correlation with IOP and HBA_{1c} levels in patients with DR [207]. Similarly, a Chinese population study found an association between DM with high IOP (>21 mmHg) [208]. Demographic study in Singapore also found an association between elevated IOP and DM. Even after accounting for the difference in central corneal thickness, which is higher in diabetics individuals, the increase in IOP was independent of the development of glaucoma.

JSH-23 Mechanism

4-methyl-N1-(3-phenyl propyl)-1, 2-benzenediamine (JSH-23) is a selective inhibitor of NF-κB-p65/p50. It stops nuclear translocation of NF-κB-p65/p50 and thus helps stop the transcription of its target genes [209]. JSH-23 has an IC₅₀ of 7.1 μ M for NF-κB nuclear translocation and transcriptional activity on macrophages RAW 264.7 stimulated by LPS. Kumar et al. showed that oral administration JSH-23 inhibits NF-κB in STZ induced male Sprague Dawley rats, and was able to decrease the expression of COX-2, iNOS in the sciatic nerve and improve nerve function [210]. When JSH-23 was used in a chronic mild stress model to inhibit the effect of NF-κB, it was able to decrease inflammatory mediators and increase anti-oxidant defense in the hippocampus [211]. After sepsis, LPS induced cardiac injury through differentiation of M1 macrophages. JSH-23 inhibits LPS induced cardiac injury in a sepsis animal model [212]. Transition metal ions activate NF-κB and NLRP3 in bone marrow-derived macrophages, thus pyroptosis and again JSH-23 was shown to inhibit NLRP3 activation via inhibiting phosphorylation of NF- κ B in macrophages after transition metal activation [213]. Cisplatin activates NF- κ B and induces acute kidney injury by necrosis and apoptosis of proximal tubular epithelial cells. JSH-23 reduced many of the response genes of the NFκB activation, including casp-1 in C57BL6J mice treated with cisplatin [214]. To our knowledge, JSH-23 has not yet been used to decrease NF-κB activation in DR or Akita Mice. Therefore, to test our hypothesis, we used the *Ins2*^{Akita} mouse (Akita) that has been well-characterized as an excellent model system to study T1D. Akita harbors a missense mutation in the *Insulin 2* gene, which leads to a conformational change in insulin protein, causing it to accumulate in pancreatic β-cells. Accumulation causes loss of cellular function, decreased cell density, and, finally, cell-death leading to HgA because of insulin deficiency [144, 215-217]. These mice become hyperglycemic after four weeks of age. HgA-induced retinal pathology ensues after 12 weeks, starting with appearance of acellular capillaries, and infiltration of the leukocytes into vascular walls [144, 218, 219], increased in vascular permeability, degeneration of astrocytes and microglia, thinning of the IPL, INL, and loss of RGCs [219]. The Akita model has many vital advantages. It has an autosomal dominant mutation with stable hypoinsulinemia, and lacks systemic immunologic alterations. It is fertile, and its background does not contain the rd (pde6) variations; such modification are responsible for photoreceptors degeneration [144].

Summary. HgA in Type I Diabetes (T1D) induces retinal inflammation via NF- κ B activation. Retinal capillary cells such as endothelial, pericytes, and glial cells are critical parts of the iBRB but remain potential targets of inflammatory mediators. Inflammatory mediators also targeted junctional molecules thus impairing the cell-cell

communication and the paracellular transport. NF- κ B increased the pro-inflammatory genes, which induced iBRB breakdown. Thus, we proposed that the inhibition of NF- κ B p65 subunit with JSH-23 to prevent retinal microvascular impairment in the retina of Akita mice.

Figure 6. Molecular Structure of JSH-23. JSH-23 inhibits the nuclear translocation of

NF-κB p65 subunit and its transcriptional activity.



Figure 7. Summary. iBRB is consisted of endothelial, pericytes, and glial cells. The ECs and their TJ regulate transcellular and paracellular transport into the retina. HgA activates NF- κ B, which then increases the expression of inflammatory mediators thus causing leukostasis, vascular occlusion, and the iBRB breakdown.

FIGURE 7



Figure 8. Summary of Background. HgA (in T1D Akita mouse model) dysregulates NF- κ B. NF- κ B overexpression increases the production of pro-inflammatory molecules from its target genes that lead to degradation of the junctional molecules in the affected retinae. As a result, the iBRB is impaired, which eventually results in the vision loss or even blindness in the susceptible diabetic patients.

FIGURE 8



CHAPTER II

HYPOTHESIS AND SPECIFIC AIMS

Key Objective

The goal of this study is to determine the effects of chronic inflammation in impairing retinal capillary vessel in Akita (T1D model) and to investigate whether JSH-23 can inhibit these injuries.

Hypothesis

During Diabetes (T1D), vision loss is due to retinal vascular leakage by an increase in inflammatory mediators (NF- κ B p65, COX-2, INOS, and ICAM-1) which lead to an increase in retinal vascular permeability, intraocular pressure, and vision loss. Therefore, treatment with inflammatory inhibitor (JSH-23) will mitigate this vision loss.

Specific Aims

Specific Aim 1: To determine whether there is an increase in inflammatory mediators (NF- κ B p65, COX-2, INOS, ICAM-1) in Retina of T1D mice and

whether inflammatory inhibitor, JSH-23, mitigate this increase in inflammatory mediators. (Chapter 3)

Specific Aim 2: To determine whether T1D increases retinal vascular permeability by decreasing expression of junctional molecules (Occludin, Connexin 43) and whether treatment with JSH-23 mitigate this increase in retinal vascular permeability. (Chapter 3)

Specific Aim 3: To determine whether T1D leads to vision loss in part by increase retinal vascular permeability and ocular pressure and whether treatment with JSH-23 mitigate this vision loss and retinal vascular permeability. (Chapter 4)

CHAPTER III

JSH-23 DOWNREGULATES NF-κB AND INCREASE EXPRESION OF JUNCTIONAL MOLECULES IN THE RETINA OF AKITA MICE

Introduction

T1D is a chronic illness that is diagnosed during childhood. It occurs in ~1:400-600 American children [220]. A recent disturbing trend is a 15-20% increase in new diagnoses of T1D under age 5 [221, 222]. The reason for this alarming increase in youths is debatable. Clinically, diabetes management has proven challenging for a variety of reasons, including physiological factors such as increased insulin sensitivity, especially in children [223]. T1D microvascular complications included DR which is divided into NPDR and PDR. DME can develop in either of the two stages [224]. Other serious complications include retinal detachment, and blindness.

DR is a leading cause of blindness in the working age adults. In its advanced form, DR can lead to blindness due to the development of DME, and cataract. Current clinical interventions vary depending on the extent of disease progression. Tight bloodglucose control and dietary measures may help slow down the disease progression in some people however; glucose control does not stop disease progression to DME or PDR. Patients may require laser surgery to seal the leaking blood vessels in their eyes or to discourage other vessels from leaking further. VEGF inhibitor injection into the eye may also be needed to decrease unabetted underlying inflammation or to stop the formation of abnormal vessels. Leakage from retinal vessels leads to a fresh wave of inflammatory swelling that invariably results in a cloudy or blurred vision. Even worse, a surgical procedure may be warranted to repair a detached retina. If left untreated, DR causes permanent blindness [225, 226].

Current interventions do not adequately address the "retino-vascular" signature pathology that underlies the initiation, progression, and maintenance of the DR phenotype. Furthermore, they fail to restore loss vision in the majority of patients. The etiology of DR is not fully understood although many pathways have been proposed for the development of DR pathology such as polyol flux, hexosamine, AGEs, and OS. DR is regarded as an inflammatory disease due to the presence of various inflammatory mediators found in the vitreous and the retinas of diabetic animals and humans [227]. An increase in the expression of inflammatory mediators is directly associated with NF- κ B activation [120]. NF- κ B is a dynamic transcription factor that orchestrates complex biological processes, including the inflammatory response itself [228]. Owing to its upstream role in cell-signaling, NF- κ B is considered a central component in the inflammatory response, directing the elaboration of cytokines, and responses to stress, free radicals, heavy metals, ultraviolet irradiation, oxidized LDL, and bacterial, and viral antigens.

NF-κB is the first responder to harmful stimuli for cells/tissues. It shuttles between cytoplasm and nucleus to initiate and then help execute a robust genetranscription program. It consists of a family of structurally related transcription factors viz. NF-κB1(p50), and RelA (p65) that can form hetero- and homodimers and translocate to the nucleus upon activation [121, 209, 229, 230]. There is a presence of a chronic inflammation in diabetics individuals, which could be mediated by dysregulations in NF- κ B signaling [120]. It has been shown that NF- κ B increased the pro-inflammatory genes in the retina of both diabetic animal models and patients [231]. It also induced the activation of pyroptosis [194]. It seems that the sustained inflammation could lead to degradation of the iBRB. The BRB consists of the iBRB and oBRB which regulates the movement of ions, and solutes into the retina. The iBRB maintains harmony between TJ molecules, and retinal ECs and any kind of impairments can lead to fluid accumulation in the retina [232]. In diabetes, there is a decrease in expression in junctional molecules in the presence of sustained inflammation [233].

As DR is a significant complication of an underlying autoimmune disease, we tested our hypothesis that systemic inhibition of NK-κB activity in a T1D model could help mitigate inflammation-induced microvascular changes in the retina that we believe are responsible for vision problems. And we know that the inflammatory mediators are elevated in the retina of DR; and are associated with iBRB breakdown [234]. Therefore, we hypothesized that pre-empting inflammation by inhibiting NK-κB activity with JSH-23 would certainly help mitigate the microvascular alterations in the diabetic retina. Moreover, JSH-23 has been demonstrated to inhibit the nuclear translocation of NF-κB-p65 and thus inhibit its transcriptional activity [209]. The objective of this study is to investigate the effect of how suppressing inflammation could potentially prevent the development of DR in the Akita mouse model.

Methods

Animal Genotyping and Maintenance of the Animals. Male C57BL/6J (WT) and Ins2^{Akita+/-} mice (Akita) were purchased from the Jackson Laboratory (Bar Harbor, ME). All animals were ~24-26 weeks old and were maintained in a 12:12 hour light-dark cycle

with a regular mouse diet at the University of Louisville School of Medicine animal facility. Mice were cared according to ARVO guidelines, as approved by the Institutional Animal Care and Use Committee (IACUC) of the University of Louisville, Louisville, KY. The experimental protocols were performed following the National Institute of Health guidelines (NIH Pub. No. 86–23, revised 1985). Genotyping of the mice was carried out by collecting the tail biopsy. DNA was isolated using the DNeasy Blood and tissue kit (Qiagen, Germantown, MD). The cross-breeding yielded Ins2^{Akita} with high and low glucose levels, as measured by blood glucose meter Ultra Touch 2 (Life Scan, USA). The high glucose level category consists of mice with blood glucose levels that were greater than or equal to 300 mg/dL, and the low glucose levels were animals exhibiting less than or equal to 250 mg/dL. Only mice with high glucose levels used in the experiments.

Mice were divided into four different groups: (a) WT, (b) WT + JSH-23, (c) Ins2 ^{Akita}, and (d) Ins2^{Akita} + JSH-23. JSH-23 purchased from Sigma (Sigma Aldrich, St. Louis, MO; CAS# 749886-87-1). A stock solution of JSH-23 was prepared by diluting 25 mg in 1000 microliter (μ L) of 200 proofs ethanol (Decon Labs, King of Prussia, PA; CAS# 64-17-5) and stored at -70 degrees Celsius. A 100 μ L of the stock solution diluted with 4900 μ L of 1X sterile phosphate-buffered saline (PBS). The control solution prepared with 4900 mL of 1X PBS and 100 mL of 200 proofs ethanol. Akita and wild type (WT) C57BL/6J mice strains were treated with or without a specific and selective inhibitor for the translocation of the NF- κ B (i.e. with JSH-23) intraperitoneally (IP) @ 5mg/Kg body weight on alternate days for a total of 4 weeks. Blood pressure (CODA Instrument), blood glucose (Ultra Touch 2), body weights (OHAUS, CS-2000 Balance), and intraocular pressure (iCare, Tonolab) were measured before and after the injections of the

NF- κ B inhibitor (*Fig. 9*).

Measurement of body weight, blood glucose, blood insulin, and glucose tolerance test. Body weights of all the animals recorded using an electronic weighing balance (OHAUS, CS-2000). *ELISA*. Mouse INSULIN ELISA kit (EMINS) from Thermo Fischer Scientific (Waltham, MA) was used to analyze plasma insulin levels. The test performed according to manufacture instructions.

Intraperitoneal Glucose Tolerance Test. The glucose tolerance test used to measure the clearance of an IP injected glucose load from the body and to detect disturbances in glucose metabolism that linked to diabetes or metabolic syndrome. Glucose (200 mg/ml) was injected IP (@ 2g per kg body weight after 6 hours of fasting. Blood glucose was measured (tail vein) at 0, 30, 60, 90, and 120 minutes after the injection using Ultra Touch 2 [235].

Reagents and Antibodies. Antibodies and chemicals were purchased from trusted sources such as Sigma-Aldrich, Abcam, Cell Signaling Technology, or Santa Cruz Biotechnology. Details for the antibodies are; iNOS (ab15323) and NLRP3 (ab214185) purchased from Abcam (Cambridge, MA, USA). COX-2 (4842S), Total NF-κB p65 (8242S), and p-NF-κB p65 (3033S) purchased from Cell Signaling Technology (Danvers, MA, USA). ICAM-1 (sc-107) purchased from Santa Cruz Biotechnology (Dallas, TX). The secondary antibodies also purchased from Santa Cruz Biotechnology, namely rabbit anti-mouse (sc-358914), mouse anti-rabbit (sc-2357), and mouse anti-goat (sc-2354). GAPDH (MAB374) was purchased from EMD Millipore (Burlington, MA). All reagents/antibodies were used for Western Blotting following the manufacturers'

recommended protocols.
Figure 9. Treatment plan. The blood glucose, blood pressure, IOP, and body weight of the animals were measured before and after treatment with JSH-23. The treatment was carried on alternate days for four weeks.



Western Blotting. Protein expression profiles were performed via Western blots analyses. Retinas were separated from enucleated globes from the euthanized mice, and the retinal protein lysates were used immediately or snap-frozen in liquid nitrogen and stored at – 80°C until further use. Protein were extracted via homogenizing in RIPA buffer (Boston BioProducts, Worcester, MA). These cocktails were added to the RIPA buffer; a cocktail of 1 mmol/L phenylmethylsulfonyl fluoride (Sigma, Saint Louis, MO), 1% protease inhibitors cocktail (Sigma). Then the samples were sonicated with the Sonifier 450 (Branson Ultrasonics, Danbury, CT). The homogenates were centrifuged at 15,000 rpm for 15 min at 4° C, and the resultant supernatants were collected and stored at -80° C. Bradford assay was used to estimate the total protein contents. Per sample, 40 μ g of total protein amount resolved on a 10% SDS-PAGE and then transferred to polyvinylidene difluoride (PVDF) membrane. The membranes were incubated with primary antibodies overnight and then with the secondary antibodies were added the next day for 3 hours before visualization with ECL Luminata Forte (Millipore, Temecula, CA) via a Bio-Rad ChemiDoc system (BioRad Laboratories, Des Plains, IL, USA). Band intensities were normalized to GAPDH for all the target proteins analyzed, and their respective quantifications were performed through Image Lab[™] Software (Bio-Rad, Hercules, CA).

Quantitative PCR. RNA was isolated with TRIzol® reagent (Life Technologies, Carlsbad, CA, USA) based on the instruction of the manufacturer. Nanodrop-1000 (Thermo Scientific, Waltham, Ma) was used to analyze RNA quantification and purity. Im Prom-II TM Reverse Transcription System (A300, Promega, Madison, WI) was used to reverse-transcribe the total RNA into cDNA per instructions of the manufacturer. Quantitative PCR (qPCR) was performed for different transcripts (ReIA, iNOS, COX-2, ICAM-1, Cx43, Occludin, NLRP3, Casp-1, IL-1 β). The final reaction included 10 µl of nuclease-free water, 8 µl of Bullseye EvaGreen qPCR Mastermix (BEQPCR-S, MIDSCI, Valley Park, MO), 50 picomoles of forward and reverse primers, and 1µl of cDNA (*Table 1*). The data were normalized with housekeeping gene GAPDH.

Statistical Analyses. All values are reported as mean \pm SEM. One-way ANOVA was used to analyze the difference between the groups, including a Tukey's post hoc analysis for groups' comparisons. The tests were performed with p<0.05 and a total number of mice (n) = 5-20 were subjected toanalysis depending upon particular experiments from each group. GraphPad Prism (Ver 8.1, GraphPad Software) used during analyses.

Results

Confirmation of Ins2^{Akita} *Mutation in the Expanded Mouse Colony and Effect of JSH-23 on Glucose Metabolism.* Mice tail DNA samples were analyzed using PCR employing specific primers to determine the mice genotype. Genotyping revealed the mutated version of the insulin gene exhibiting two discrete bands: 280 and 140 bp representing the heterozygous status while wild-type (WT) mice showed one 140 bp band (*Fig. 10A*). To generate mice for the study, Akita female with low blood glucose (LG) were cross-breed with male Akita mice with high glucose (HG). Cross-breeding yielded Akita progeny with HG and LG levels as measured by blood glucose (BG) meter Ultra Touch 2 (Life Scan, USA). Only Akita mice with a BG level <300 mg/dL were used in this study. Akita mice have a mean BG level of 596.47 mg/dL, significantly higher than C57BL/6J BG of 173.47 mg/dL (p<0.0001, n=15 per group). JSH-23 significantly decreased

 Table 1. Nucleotide sequence. Forward and reverse primers gene sequences for each gene analyzed.

TABLE 1

Gene	Forward Sequence	Reverse Sequence
Rel A[236]	5'- GCCCAGACCGCAGTATCC	5'- GTCCCGCACTGTCACCTG
iNOS [237]	5'- CAGCTGGGCTGTACAAACCTT	5'- CATTGGAAGTGAAGCGGTTCG
COX-2 [238]	5'- CAGAACCGCATTGCCTCTG	5'- TTGTAACTTCTGGTCCTCATGTCGA
ICAM-1[239]	5'- TTCACACTGAATGCCAGCTC	5'- GTCTGCTGAGACCCCTCTTG
Cx43 [240]	5'- CCAAGGAGTTCCACCACTTTG	5'- CCATGTCTGGGCACCTCTCT
Occludin[241]	5'- AGACCCAAGAGCAGCCAAAG	5'- GGAAGCGATGAAGCAGAAGG
NLRP3 [242]	5'- CTTCTAGCTTCTGCCGTGGTCTCT	5'- CGAAGCAGCATTGATGGGACA
Casp-1[242]	5'-GTACACGTCTTGCCCTCATTATCTG	5'- TTTCACCTCTTTCACCATCTCCAG
IL-1β [242]	5'-CAACCAACAAGTGATATTCTCCATG	5'- GATCCACACTCTCCAGCTGCA

Figure 10. Genotyping and Physiological Parameters of Akita, and C57BL/6J Mice.

A. Genotyping of Akita, and C57BL/6J mice. **B.** Body weight of the Akita, and the C57BL/6J, Akita mice body weights were significantly lower than the body weights of the C57BL/6J mice (p<0.0001, n=19 per group). **C.** The blood glucose level of Akita, and C57BL/6J mice the blood glucose levels of Akita mice were significantly higher than C57BL/6J mice (p<0.0001, n=15 per group). JSH-23 significantly decreased blood glucose levels in Akita mice (p<0.0001). **D.** Plasma insulin levels of Akita mice were substantially lower (p<0.0001, n=19 per group). **E.** Intraperitoneal glucose tolerance test depicted the inability of the Akita mice to lower their blood glucose levels (p<0.009, n=5 per group).



Figure 11. Expression of NF-κB p65 in the retinae of Akita, and C57BL/6J mice after treatment with JSH-23. A. Immunoblotting image of total and phosphorylated NF-κB p65. **B.** The total expression level of NF-κB p65 was significantly elevated in Akita mice (p<0.05, n=7 per group). JSH-23 significantly decreased the total expression of NF-κB p65 in Akita mice (p<0.005). **C.** The phosphorylated expression level of NFκB p65 significantly elevated in Akita mice (p<0.05, n=7 per group). JSH-23 could significantly decreased the phosphorylated expression level of NF-κB p65 (p<0.05). **D.** The gene expression of NF-κB p65 was significantly elevated in Akita mice (p<0.0001, n=7). JSH-23 significantly decreased the gene expression of NF-κB p65 in Akita mice (p<0.0002).



Figure 12. Expression levels of ICAM-1, iNOS, and COX-2 in the retinae of Akita, and C57BL/6J mice after treatment with JSH-23. A. Immunoblotting image of ICAM-1, iNOS, and COX-2. B. The expression level of ICAM-1 was significantly elevated in Akita mice (p<0.0218, n=6 per group). JSH-23 significantly decreased the expression of ICAM-1 in Akita mice (p<0.0028). C. The expression level of iNOS was significantly elevated in Akita mice (p<0.0001, n=6 per group). JSH-23 significantly decreased the expression of iNOS in Akita mice (p<0.0001). D. The expression level of COX-2 was significantly elevated in Akita mice (p<0.0001, n=6 per group). JSH-23 significantly decreased the expression of COX-2 in Akita mice (p<0.0001).









A.



p<0.0001

n=6



Figure 13. Gene expression level of ICAM-1, iNOS, and COX-2 in the retinae of Akita mice, and C57BL/6J treated with JSH-23. A. ICAM-1 gene expression was elevated in Akita mice (p<0.0003, n=5 per group). JSH-23 decreased the gene expression in Akita mice (p<0.0011). The iNOS gene expression was elevated in Akita mice (p<0.0012, n=5 per group). JSH-23 decreased iNOS gene expression levels in Akita mice (p<0.0031). COX-2 gene expression was increased in Akita mice (p<0.0004, n=5 per group). JSH-23 reduced the expression of COX-2 in Akita mice (p<0.0001).



C. COX-2 Gene Expression



Figure 14. Pyroptosis Marker in the retinae of Akita, and C57BL/6J mice treated with JSH-23. A. Immunoblotting imaging of NLRP3. B. NLRP3 expression level was elevated in Akita mice (p<0.018, n=6 per group). JSH-23 decreased NLRP3 expression in Akita mice (p<0.0003). C. NLRP3 gene expression was increased in Akita mice (p<0.0025, n=6 per group). JSH-23 decreased the gene expression level of NLRP3 in Akita mice (p<0.0003). D. Casp-1 gene expression increased in Akita mice (p<0.0001, n=6 per group). JSH-23 decreased the gene expression of casp-1 (p<0.0001). E. IL-1 β gene expression was elevated in Akita mice (p<0.0001, n=6 per group). JSH-23 decreased the gene expression level of IL-1 β (p<0.0001).



Figure 15. Expression of Junctional Molecules in the retinae of Akita, and C57BL/6J mice treated with JSH-23. A. Immunoblotting image of Cx43, and occludin. B. Cx43 expression was decreased in Akita mice (p<0.0165, n=5 per group). JSH-23 increased the expression of Cx43 (p<0.0006). C. Occludin expression was decreased in Akita mice (p<0.0068, n=5 per group). JSH-23 decreased the expression of occludin (p<0.0071). D. Cx43 gene expression was decreased in Akita mice (p<0.003, n=5 per group). JSH-23 increased the gene expression in Akita (p<0.0237). E. Occludin gene expression was decreased in Akita mice (p<0.0073, n=5 per group). JSH-23 increased the gene expression of occludin (p<0.006).

n=5



B.





Occludin Expression





WT USH231

N

D.





E.

0.0

Relative to GAPDH

Occludin Gene Expression

ANIA LISH-231

AXIES



the mean (456.2 mg/dL, p<0.0001) blood glucose level in Akita mice (*Fig. 10C*). Akita mice have a mean body weight of 26.60g, substantially lower than C57BL/6J mice with a mean of 30.82g (p<0.0001). JSH-23 did not have any effect on the mean body weights of the Akita mice (*Fig. 10B*). Akita displayed significantly lower plasma insulin levels (p<0.0001) (*Fig. 10D*). As expected, results from IP administered glucose tolerance test (ipGTT) showed high basal glucose levels in Akita in comparison to C57BL/6J mice. Furthermore, ipGGT also revealed that the Akita strain was unable to lower its glucose level (p<0.0090) in their blood (*Fig. 10E*).

JSH-23 Diminishes Key Inflammatory Molecules in the Diabetic Retinae. Western blotting and qPCR analyses were used to examine the effects of systemic inhibition of NF- κ B in mice retinae. The total (p< 0.0353) and phosphorylated amounts (p<0.0463) of expression levels of NF- κ B p65 were elevated in the retinae of Akita mice. NF- κ B p65 subunit gene expression (p<0.0001) was also elevated in the retina of Akita mice. JSH-23 decreased the total (p<0.0053) and phosphorylated (p<0.0254) and gene expression (p<0.0002) levels of NF- κ B p65 (*Fig. 11*).

ICAM-1 expression level (p<0.0218) and the gene expression (p<0.0003) were elevated in the retinae of Akita mice. JSH-23 decreased the expression level (p<0.0028) and gene expression (p<0.0001) of ICAM-1 in the retinae of Akita mice. The iNOS expression level (p<0.0001) and gene expression (p<0.0012) were increased in the retinae of Akita mice. JSH-23 decreased the expression (p<0.0001) and gene expression (p<0.0031) levels of iNOS in the retinae of Akita mice. COX-2 expression (p<0.0001) and gene expression levels (p<0.0004) were increased in the retinae of Akita mice. JSH-23 decreased expression level (p<0.0001) and gene expression levels (p<0.0001) of COX-2 in the retinae of Akita mice (*Fig. 12 & 13*).

JSH-23 Decreases Expression of Pyroptosis Markers in the Diabetic Retinae. NLRP3 expression (p<0.0180) and gene expression levels (p<0.0025) were increased in the retinae of Akita mice. JSH-23 decreased expression (p<0.0003) and gene expression levels (p<0.0003) of NLRP3 in the retinae of Akita mice. The gene expressions levels of casp-1 (p<0.0001) and IL-1 β (p<0.0001) were increased in the retinae of Akita mice. JSH-23 decreased gene expression levels of casp-1 (p<0.0001) and IL-1 β (p<0.0001) in the retinae of Akita mice (*Fig. 14*).

JSH-23 Augments Integrity of the Tight and Gap Junctions Proteins in the Diabetic Retinae. Cx43 expression (p<0.0165) and gene expression levels (p<0.0003) were elevated in the retinae of Akita mice. JSH-23 increased expression (p<0.0006) and gene expression levels (p<0.0237) in the retinae of Akita mice. Occludin expression (p<0.0068) and gene expression levels (p<0.0073) in the retinae of Akita mice were also modulated as can be seen that JSH-23 increased the expression (p<0.0071) and gene expression levels (p<0.0060) of occludin in the retinae of Akita mice (*Fig. 15*).

Discussion

There are several mouse models of T1D induced DR that vary in etiology, severity, mode of inheritance, progression, and overall disease pathology. Akita strain of mouse becomes diabetic due to a mutation in the insulin 2 (Ins2) gene that results in improper folding of the insulin protein, thus causing a toxic injury to β -cells of the

pancreas. This structural alteration in the insulin molecule impairs insulin secretion, OS, cell death, and an impairs glucose-stimulated insulin release. This also produces toxic byproducts, leading to a severe inflammatory changes in the diabetic retinae causing the retinovascular permeability, and leukostasis [144, 216-218, 243]. NF- κ B is the first responder to harmful stimuli for the cells. Known inducers of NF- κ B are ROS, IL-1 β , bacterial LPS, and ionizing radiation mediated DNA damage [244-248]. NF- κ B is a family of transcription factors, RelA (p65), that hetero/homodimerize, translocate to the nucleus to augment NF- κ B directed genes' expression, particularly the inflammatory mediators [121].

Inflammatory mediators such as iNOS, COX-2, and ICAM-1 are elevated in the retinae of DR, associated with iBRB breakdown, thus increasing the retinal vascular permeability. Increased leukostasis mediated by ICAM-1 is associated with an increase in the vascular permeability [234]. HgA induces leukostasis in retinal vessels in rodents that leads to endothelial apoptosis [249-251]. Leukostasis impairs vision via two proposed mechanisms: (a) iBRB breakdown due to cellular apoptosis in the endothelium and (b) endothelial apoptosis that leads to avascular capillaries; thus, nonperfusion or hypoperfusion which in turn results into neovascularization [251]. NF-κB plays an essential role in inflammation and leukostasis via ICAM-1.

Inflammation of the retinal vasculature was significantly diminished as demonstrated by expression analyses of the critical inflammatory mediators such as ICAM-1, COX-2, and iNOS in the retinae of Akita mice. Also, the total NF- κ B p65, and pNF- κ B p65 activities were significantly reduced, indicating a clear readout of the specific inhibition of the critical transcription factor that is responsible for mounting immune response in the diabetic retinae. In diabetes, leukocyte adhesion to retinal endothelium is the primary mechanism of leukostasis. This adhesion is mediated by the upregulation of ICAM-1 due to NF- κ B activation, which increases ICAM-1 expression. Increased leukostasis is associated with enhanced vascular permeability [166-168]. Furthermore, inhibition of ICAM-1 has shown to reduce the leakage and leukocytes adhesion along with a decrease in leukocytes adherence in diabetic rats [169]. Prominent inflammatory mediators like iNOS, COX-2, and ICAM-1 were elevated in the diabetic retinas, as shown in this study. These molecules are associated with iBRB breakdown leading to an increase in the retinal vascular permeability during diabetes because of the chronic inflammation that potentially contributes to iBRB breakdown. It was reported that inhibition of NF- κ B activation by calcium dobesilate prevented degradation of occludin in part by decreasing the expression of ICAM-1 [58].

Similarly, the HgA induced downregulation of Cx43 severely affected the GJIC in the blood vessels and that has again been shown to contribute to iBRB integrity during DR pathogenesis [252]. Low expression of Cx43 was concurrently associated with other companion junctional proteins such as occludin and ZO-1 that were also found to be downregulated [195]. Therefore, we analyzed retinal cell lysates for occludin, and Cx43, employing Western blotting and qPCR assays. Their expression levels in JSH-23-treated Akita mice were increased significantly. JSH-23 was shown to be a protective agent by attenuating the loss of Cx43 during the hypoxic environments [253]. Furthermore, others have shown that specific inhibition of NF- κ B by JSH-23 attenuated the pathological sprouting of the lymphatic vessels [254]. NF- κ B activation also activated the assembly of the NLRP3 inflammasome. NLRP3 has been shown to be active during inflammation, and activation of inflammasome in the retinae [242, 255, 256]. Pyroptosis, and inflammatory cell death both activate via the assembly of the NLRP3 inflammasome that further enables casp-1, which then cleaves the IL-1 β . Retinal vascular cell death during HgA causes the breakdown of iBRB. HgA induced activation of casp-1, and IL-1 β has been reported in causing Müller cell death [193]. This study showed an increase in NLRP3 expression level and the gene expression of casp-1, and IL-1 β in the retinae of Akita mice. JSH-23 significantly decreased NLRP3 expression levels and the gene expression of both casp-1 and IL-1 β in the retinae of Akita mice.

The incidence of T1D is rising, and glycemic variability is common among young children these days. This has long-term implications for health and well-being for the diabetics. Research has identified several potential options to cure T1D, but none of them have shown any concrete outcome till today. Thus, targeted interventions are needed to reduce the risk of long-term T1D-related complications, and improve the immediate quality of life in patients. The NF- κ B molecule is a critical player in the inflammatory process; thus, strategies inhibiting NF- κ B has potential therapeutic applications in the fight against diabetes. The result of this study showed that the sustained long-term inhibition of NF- κ B activity could significantly diminish the degradation of cell-junctional molecules in the retinae of Akita mice.

Conclusion: Inflammatory mediators such as ICAM-1, INOS, and COX-2 expression were elevated in the retinae of Akita mice. That upregulation of inflammatory mediators was associated with the activation of NF- κ B p65. Elevated levels of inflammatory mediators are linked with the degradation of junctional molecules, thus impairing the iBRB. JSH-23 decreased NF- κ B p65 expression levels in the retinae of Akita mice along with inflammatory mediators (ICAM1, INOS, and COX-2) expression in the retinae of Akita mice. JSH-23 treatment reduced NLRP3 and other pyroptosis mediators (e.g. casp-1, and IL-1 β). Inhibition of NF- κ B p65 also upregulated the expression of Cx43, and occludin in the retinae of Akita mice. Inhibition of NF- κ B p65 also decreases blood glucose level. That decrease could be due to an increase in insulin sensitivity in the Akita mice.

CHAPTER IV

EFFECT OF JSH-23 ON VISION IN AKITA MICE

Introduction

T1D predisposes patients to long-term complications like kidney failure, neuropathy, and blindness. Lately, the incidence of T1D is rising, and glycemic variability is becoming more common in the younger population. Although T1D is considered an autoimmune disease, it is not clear what exactly causes the immune system to attack the beta cells in the pancreas. Up to 90 percent of T1D patients can develop retinopathy post-diagnosis; as a result, it is becoming a severe medical problem globally. DR is the leading cause of blindness in the working-age adults [250]. Unfortunately, there is no cure for DR.

HgA causes endothelial damage and capillary nonperfusion, ultimately leading to retinal hypoxia. This prompts the growth of abnormal blood vessels in the eye. Early detection and timely intervention can reduce the risk, but vision loss is irreversible. Research in the past has identified several options to treat T1D but not all benefit the majority of patients. Thus, more study needed to address the vascular pathology that occurs in the retinae.

HgA is known to cause sustained ROS generation, ATP depletion, ER stress, and, finally, inflammation as a complication of DR pathogenesis. This eventually leads to microvascular damage, basement membrane thickening, iBRB breakdown, capillary cell death, acellular capillary, neovascularization, and sometimes the detachment of the retina itself.

Inflammatory mediators such as iNOS, COX-2, and ICAM-1 are elevated in the retinae of DR, and are associated with iBRB breakdown, thus increasing the retinal vascular permeability. Increased leukostasis is related to an increase in vascular permeability too [234]. HgA induces leukocytes adhesion in rodents retinal vessels that leads to endothelial apoptosis [249-251]. NF- κ B plays an essential role in inflammation, and leukostasis via ICAM-1.

iBRB breakdown is strongly associated with the release of proinflammatory factors such as iNOS, and elevated levels of inflammatory markers that play a definitive role in raising IOP. Additionally, IOP elevation linked with the upregulation of the hypoxia-inducible factor-1 alpha (HIP-1 α), interleukins, and VEGF [257]. Inflammation contributes to BRB breakdown and thus increases the intraocular pressure (IOP).

In this study, we decided to inhibit NF- κ B, the first responder during the inflammation, employing a T1D mouse model and demonstrate that specific targeting of the NF- κ B activity could help prevent chronic inflammation that is responsible for the degenerative changes in the retinae. Our findings highlight the significance of targeted intervention in reducing the long-term diabetic complications and thus improving the underlying constant inflammatory milieu that is propagated by chronic metabolic dysfunction in the diabetic patients.

Methods

Systemic Blood Pressure (BP) and Intraocular Pressure. The systemic blood pressure

was measured with CODA noninvasive blood pressure instrument (Kent Scientific, Torrington, CT). The IOP was recorded by a Tonometer (iCare, Tonolab). Both pressure were measured before and after the JSH-23 treatment in mice [258].

Fluorescein Angiography. Mice injected by IP with 100 µl of AK-FLUOR[®] (NDC 17478-253-10, Akorn, Inc, Lake Forest, IL) after induction of anesthesia with 300 cc of 2x tribromoethanol (TBE). The fundus was visualized by Micron IV microscope (Phoenix Technology Group, Pleasonton, CA). Images were analyzed via ImageJ for the presence of fluorescence outside of the retinal capillaries [259].

In vivo Electroretinography (ERG), and Optical Coherence Tomography (OCT). Electroretinograms (ERGs) were performed on C57BL/6J and Akita age-matched mice. Briefly, mice were dark-adapted overnight, and anesthetized by IP using TBE. After pupils were dilated, full-field scotopic ERGs were recorded using the Tucker–Davis System Workstation (Tucker–Davis Technologies, Alachua, FL) for the real-time visualization of the averaged a- and b-waves. The BioSigRZ system was used for the quantification, and further analysis. OCT was performed to measure the anatomical differences in the retinal layers before and after treatment with JSH-23, employing the Image-Guided OCT2 equipment (Phoenix Technology Group, Pleasonton, CA)

Light-dark Box Chamber Test. The animals were subjected individually one by one for the light-dark box test into the light chamber with the light close. Then, PACS Shuttle Box Version 3.40 was set up with Experiment Mode as "Passive Avoidance", UCS Grid Intensity (Amperes): 0, Exploration duration: 30

seconds, Maximal Trial Duration: 5 minutes, CS Light Intensity: 10 volts, UCS Grid Duration: 2 seconds. After 30 seconds of exploration, the light flickered on. Then the time it took the mice to transfer from the light chamber to the dark was measured [260].

Statistical Analyses. All the values are reported as mean \pm SEM. One-way ANOVA was used to analyze the difference between the groups, including a Tukey's post hoc analysis for groups' comparisons. The tests were performed with p<0.05 and a total number of mice (*n*) = 5–20 were subjected to experimentations from each group. GraphPad Prism (Ver 8, GraphPad Software) was used during the subsequent analyses.

Results

Monitoring of Systemic Blood Pressure and Intraocular Pressure. We next evaluated systemic and IOP before and after JSH-23 treatment. It is known that JSH-23 increases blood pressure in hypertriglyceridemic (HTG) rats [261], therefore; we sought to determine whether JSH-23 did any alteration to the mice systemic blood pressure. To find that out, we subjected treated and untreated mice to a noninvasive tail-cuff method for recording blood pressure. None of the mice acquired showed any statistically significant change in their blood pressures. For example, the mean blood pressure in Akita mice was 117.15 mmHg, and for C57BL/6J strain it was 90.17 mmHg. JSH-23 did not have any effect on the treated Akita mice with a mean of 102.7 mmHg (*Fig. 16A*). The mean IOP of the Akita mice was 25.33 mmHg vs. 14.55 mmHg for C57BL/6J (p<0.0001, n=12 per group). However, JSH-23 did decrease the IOP in Akita mice, 16.02 mmHg (p<0.0001) (*Fig 16B*).

Figure 16. Systemic Blood Pressure and Intraocular Pressure (IOP) in Akita and

C57BL/6J mice strains. **A.** JSH-23 does not affect the systemic blood pressure (n= 8 per group). **B.** IOP was elevated in Akita mice (p<0.0001, n=12 per group) and JSH-23 treatment decreased IOP in the Akita mice (p<0.0001).



Blood Pressure Measurement for C57BL6/J and Akita Mice Treated with JSH-23

B.

A.







Figure 17. Fundus imaging of Akita, and C57BL/6J treated with JSH-23. The retinae

of Akita mice depicted derailment in their vasculature network and the treatment with

JSH-23 could repair these vasculature derailments in the affected retinae.



Akita

Akita + JSH-23

Figure 18. Fluorescence Angiography in Akita, and C57BL/6J strains treated with

JSH-23. The retinae vasculature of Akita mice showed an increased vascular permeability (p<0.0001, n=6 per group). And treatment with JSH-23 decreased the vascular permeability in Akita mice strain (p<0.0001).

FIGURE 18



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Figure 19: Optical Coherence Tomography in Akita, and C57BL/6J treated with

JSH-23. The retinal layer's thickness decreased in Akita mice (p<0.0001, n=5 per group) and the treatment with JSH-23 prevented the retinal thickening (p<0.0001).

FIGURE 19




Figure 20: Electroretinogram of Akita, and C57BL/6J mice strains treated with JSH-23. A. Electroretinograms of C57BL/6J, and Akita mice treated and non-treated with JSH23. **B.** Electroretinograms superimposed on top of each other on excel spreadsheet **C.** Amplitude of A-wave in the retinae of Akita was found to be decreased (p<0.0001, n=6 per group) and the treatment with JSH-23 prevented this decrease in the A-wave amplitude in Akita (p<0.0001). **D.** The amplitude of B-wave in Akita mice was also decreased (p<0.0001, n=6 per group) and treatment with JSH-23 could prevent this decrease in B-wave amplitude in the Akita mice (p<0.0001).



B.











Figure 21: The light-dark chamber movement analysis of Akita and C57BL/6J mice treated

with JSH-23. A. Assessment of vision-guided behavior employing a light-dark chamber, it took Akita mice significantly longer to move from the dark to the light chamber (p<0.0001, n=8 per group) relative to C57BL/J6. Akita treated with NF-kB inhibitor demonstrated significantly decrease time taken by them to move from light chamber to the dark chamber (p<0.0477, n=10 per group).



p<0.0001

FIGURE 21

Assessment of Retinal Fundus. The retinal fundus images of Akita mice depicted the degeneration of the retinal blood vessels along with yellow spots that appeared to be cellular debris like structure and the treatment with JSH-23 seemed to prevent the degeneration of these inflamed retinal blood vessels in Akita mice post treatment (*Fig. 17*).

Assessment of Retinal Vasculature by Fluorescein Angiography (FA) and Fundoscopy. Retinal complications such as vascular damage is consistent with T1D. We know that HgA-led inflammation causes retinal vascular leakage, and microaneurysm formation. We studied the anti-inflammatory effects of JSH-23 on the retinal vasculature employing *in vivo* FA [218]. The results showed diminished retinal permeability in the JSH-23 treated mice (p<0.0001, n=6 per group) (*Fig. 18*). Interestingly, a decrease in perfusion time (time is taken by fluorescein isothiocyanate; FITC dye to perfuse the retinal vasculature completely) in Akita mice of about 2 minutes was observed after an IP injection of the dye. There was also an increase in microaneurysm formation in the Akita mice as expected (*Fig. 18*).

In vivo Electroretinography and Optical Coherence Tomography (OCT). Functional changes in the retinae was evaluated by full-field scotopic electroretinograms (ERGs) on C57BL/6J, and Akita mice employing the BioSigRZ system. Akita mice demonstrated widened and truncated A and B waves in contrast to the sharp A, and B waves in C57BL/6J mice strain. ERGs of treated, and non-treated of Akita and C57BL/6J showed in *Fig 20A* and *Fig.20B*. Also, the A-wave was significantly reduced in Akita mice compared to WT (p<0.0001 n=6 per group). The average amplitude for Akita mice was 32.88 μ V vs.

66.18 μ V of thw C57BL/6J mice. JSH-23 did increase the magnitude of the A-wave in Akita mice, as shown by mean value of 61.75 mV (p<0.0001) (*Fig. 20C*). On the other hand, the B-wave amplitude showed a significant difference between Akita, and C57BL/6J strains e.g. 103.78 μ V vs. 207.53 μ V (p<0.0001, n=6 per group). JSH-23 treatment increased the amplitude, and mean of 150.2 mmHg (p<0.0001) (*Fig. 20D*). Retinal structural integrity was evaluated employing real-time, image-guided OCT in Akita strain that displayed hyporeflective retinal lucency (blue arrows) along with abnormal thickening of the membrana limitans interna, inner nervous stratum opticum, and GCL. The retinal thickness layer was reduced in the Akita strain with a mean of 128.58 μ m, and 254.25 μ M C57BL/6J mice (p<0.0001, n=5 per group). JSH-23 prevented degradation of the retinal thickness layer in Akita mice, as reflected by the mean value of 254.25 (p<0.0001) (*Fig. 19C*).

Evaluation of Vision Guided Behavior in JSH-23 Treated Mice

Finally, the mice were evaluated for their vision-guided behavior employing a light-dark chamber analysis. It took Akita mice significantly longer to move from the dark to the light chamber (p<0.0001, n=8 per group) relative to C57BL/J6. The results revealed that Akita mice treated with JSH-23 demonstrated significantly decreased time taken by them to move from the light chamber to the dark chamber post treatment (p<0.0477, n=8 per group).

Discussion

We hypothesized that activation of NF-KB, and its nuclear translocation could be

separated from other biological processes, thus giving us an opportunity to test the potential strategy of targeting specifically NF-kB activity for a longer duration on a sustained basis, and study its beneficial effect(s) in a well-characterized T1D mouse model. The model exhibits degenerative changes in the retinal architecture, including its blood vessels leading to the development of symptoms mimicking DR pathology [262]. By this account, treatment with inflammatory inhibitor such as JSH-23 should in principle, prevent the development of DR phenotype. To test this hypothesis, C57BL/6J and Ins^{2Akita} (Akita) strain of mice were treated with JSH-23 or without for a total period of four weeks continuously. Several disease indicators, such as IOP, blood pressure, visual function, were evaluated systematically one by one. For assessing the structural, and functional aspects of the retinae methods such as FA, OCT, and ERG with or without JSH-23 treatment were employed. Our findings revealed that JSH-23 significantly decreased the retinal vasculopathy in the diabetic retinae, and retinal structure was successfully protected as evidenced by OCT, and a concurrent decrease in the vascular leakage as shown by fundus photography and FA analyses. Further, the data indicated that visual function were affected in the Akita mouse and is consistent with previously described results [218]. Similar changes in the mouse models of hypertension, and retinal neovascularization depicted severe capillary non-perfusion, vessel constriction, beading, neovascularization, fibrosis, and edema that have been reported in the literature [263]. These changes are associated with inflammation and activation of NF- κ B. Our findings indicate that mitigation of DR severity and its pathology are possible via timely inhibition of inflammation on a sustained basis using a selective inhibitor of inflammation.

Chronic inflammation during diabetes is associated with iBRB breakdown that leads to an increased retinal vascular permeability, and thus elevate IOP, thereby decreasing vascular permeability. In our study the IOP also improved, as did visual function which were revealed by the ERG analyses when inflammation was successfully arrested. Notably, the inflammatory response to injury has also been shown to be effectively attenuated by the JSH-23 treatment in another model system, as well [264]. In a nutshell, this study advances our understanding of the role of inflammation in DR, and sheds important light on one of the approaches to further reduce the inflammatory insults in a privileged organ-like the eye. However, additional studies are needed on how best to deliver the JSH-23 treated mice, again confirming that inhibition of inflammation via NF- κ B inhibition could help protect the retinal vasculature in the diseased eyes. Therefore, the results from the present work indicate that mitigation of DR severity, in fact, is possible via inhibition of inflammation using a selective inhibitor.

Conclusions. Treatment with JSH-23 successfully decreased IOP in Akita mouse strain without change in systemic BP noticed. One of the significant findings that came out of this work is that the inhibition of inflammation perhaps helped improve the vision in Akita mice thus indicating that controlling inflammation could potentially decrease retinal inflammatory insults, and therefore improve the structural, as well as, functional attributes of the diabetic retinae in a chronic disease such as diabetes.

CHAPTER V

CONCLUDING DISCUSSION

T1D or insulin-dependent diabetes is a chronic illness that is usually diagnosed during childhood [220-222]. Clinically, diabetes management has proven challenging for a variety of reasons including physiological factors such as increased insulin sensitivity, especially in children [223]. T1D leads to the degeneration of microvascular network. DR costs the US government 1/2 a billion dollar in medical cost every year [265]. According to the National Eye Institutes (NEI), in 2010 there were approximately 7.7 million individuals living with DR, and that number is projected to double by 2050. Although more Caucasians are affected by the disease, it is more common in African-Americans and Hispanics communities [266]. DR affects both type I and type II DM patients, however DR is twice more common in T1D [225].

DR is the leading microvascular damage in Diabetes. It is defined by leakage of the blood and other body fluids into the retinae from nearby microvessels followed by abnormal angiogenesis, increase in permeability, micro aneurysms, loss of pericytes, and thickening of the basement membrane [226]. DR starts as NPDR, gradually progresses to DME and or PDR [224]. NPDR is characterized by the presence of hemorrhage, microaneurysms, abnormality in the retinal microvessels, and the cotton wool spots (fluffy white patches in the retina). On the other hand, PDR is characterized as an abnormal growth of blood vessels in the retina [267]. Untreated DR could develop into DME (fluid accumulation into the macula) or PDR (meovascularization). DME not PDR is more prevalent in type II diabetes [268]. Other complications of DR include retinal detachment. The development of vessels' growth into the area that drains fluid from the eye raises IOP to the point that can damage the optic nerve. If left untreated, PDR causes severe vision loss or even blindness.

Treatment of DR depends on the stage of the disease itself since the goal remains to slow or stop further disease progression. Many times, laser (photocoagulation) is used to create burns in the retina to seal the leaks in order to avoid further leakage. Injections of drugs (anti-vascular endothelial growth factor) in the eye are aimed at stopping the vascular leakage and new vessel formation. These approaches, along with tight bloodglucose control and dietary measures, may help slow down disease progression in some people [225, 226]. Current interventions do not adequately address the "retino-vascular" signature pathology that underlies the initiation, progression, and maintenance of the DR phenotype. Furthermore, they fail to restore vision in the majority of patients. The etiology of DR is not fully understood despite an abundance of literature describing the impact of the polyol flux, hexosamine pathways, AGEs, and OS. However, recently, DR is regarded as a stand-alone inflammatory disease due to the presence of various inflammatory mediators that have been repeatedly found in the vitreous and retinas of both diabetic animals and humans [227]. Inflammation is a response of the immune system after an injury had occurred or to a foreign entity. The resolution of inflammation involves cytokines, which upregulate pro-inflammatory proteins [269]. There is a consistent presence of a chronic inflammatory environment in the diabetics individuals, which could be mediated by dysregulation of the master regulator i.e. NF-kB.

Interestingly, a substantial increase in the expression of inflammatory markers/molecules is directly associated with NF-κB [120], a dynamic transcription factor that orchestrates complex biological processes, including the inflammatory response itself [228]. Owing to its upstream role in cell-signaling, NF- κ B is considered as a central component in the inflammatory response [121, 209, 229, 230]. Dysregulations of NF- κ B induces overexpression of inflammatory mediators (iNOS, ICAM-1, COX-2) in the retinas of diabetic animal models and patients [231]. It induces activation of pyroptosis process (NLRP3, IL-1 β) [194]. This sustained inflammation could easily lead to degradation of the iBRB in many susceptible individuals leading to a substantial decrease in the expression of junctional molecules such as Cx43, and occludin [233]. Furthermore, iBRB impairments also contributes to fluid accumulation in the retina [232].

There is an increase in inflammatory mediators (NF- κ B p65, COX-2, INOS, and ICAM-1) in the retina of T1D mice. These mediators are associated with decreasing the expression of junctional molecules (occludin, and Cx43) leading to iBRB impairment, increased retinal vascular permeability, and increase in the ocular pressure. The objective of this study was to investigate whether treatment with JSH-23 mitigates this vision loss and retinal vascular permeability, thus potentially preventing the development of DR. We therefore hypothesized that pre-empting inflammation by systemically inhibiting NK- κ B activity with, JSH-23 could mitigate microvascular alterations in the diabetic retinae.

Findings from this study suggest that decreasing the inflammation alone could protect the diabetic retinae and its vascular network highlighting the fact that regulating NK- κ B may have broad implications in treating chronic inflammatory diseases, including diabetes and its devastating complications. This study showed a decrease in the blood glucose of Akita mice after treatment with JSH-23. This decreased in blood glucose level could be due to increase in insulin sensitivity, byproduct of suppressing inflammation. JSH-23 decreased the inflammatory mediators (NF- κ B p65, ICAM-1, iNOS, COX-2) in the retinae of Akita mice. It also reduced pyroptosis markers (NLRP3, Casp-1, and IL- 1β) in the retinae of Akita mice. More importantly, JSH-23 also reduced the degradation of junctional molecules in the retina of Akita mice (occludin, and Cx43). Not only it also decreased retinal vascular permeability, IOP, in Akita mice but also increased A-wave and B-wave in Akita mice.

In this study we conclude that during T1D, vision loss is due to retinal vascular leakage by an increase in inflammatory mediators (NF- κ B p65, COX-2, INOS, and ICAM-1), which leads to an increase in retinal vascular permeability, IOP, and the vision-loss. Treatment with an inflammatory inhibitor (JSH-23) most likely helped improve vision in Akita as scored via the vision-guided behavioral test.

Figure 22. Summary. Inhibitor of inflammation (employing the JSH-23) improved visual function by decreasing the retinal vascular permeability, intraocular pressure, and retinal inflammation in the T1D Akita mouse model successfully.



Future Directions

The resident immune cells of the retina are called microglia. They help stabilized or maintain the homeostasis of the retinal environment. They reside in multiple cell layers include the IPL, OPL, and the GCL [59]. Microglia exists physiologically into two states: activated or surveying. It contains receptors for many different neuromodulators or neurotransmitters such as glutamate, dopamine, bradykinin, and norepinephrine. Microglia also secretes many substances in responses to injury, inflammation, or constitutively; NO, ROS, cytokines, and chemokines [59].

Microglial activation is due to extracellular signals, including cell death, neurodegeneration, nucleic acids, which are recognized via TLRs, which then activate the pro-inflammatory pathway of NF- κ B, followed by upregulation of inflammatory mediators [270]. Microglia are enabled in DR, although the mechanism is still at large. Some studies in diabetic rats showed that HgA does not change the density of microglia, but more of them are activated. In human DR patients, the number of microglia significantly increases, and they migrate towards injured areas such as micro-aneurysms. Thus, it is still yet to be clear whether microglia in DR in part are protective [270]. Upon activation, microglia function in presentation of antigen, producing inflammatory markers, and phagocytoses. Active microglia could also become self-destructive via inducing the death of retinal cells [271].

HgA activates microglia via increased ROS, which then enables NF- κ B. NF- κ B increases the expression of its pro-inflammatory genes, including IL-1 β , and ICAM-1 [270]. Hui-yang Zeng et al. reported microglia were activated around the retinal vessels in the retina of diabetic patients. Activated microglia released IL-1 β and other

proinflammatory mediators, which further impaired the retinal vasculature [272], mainly via activation of NF- κ B [273]. In STZ induced rats, Xiaofei Chen et al. found to be a redistribution of microglia cells in the retinal layers, with the GCL and the NFL depicting an increase in the microglia populations. In contrast, the IPL has the opposite effect, which induced retinal thickening [274]. Anti-inflammatory agents, minocycline, was shown to prevent retinal neuronal cell death in STZ-induced rats by inhibiting the activation and proliferation of microglial cells [275].

Literature reported on JSH-23 effect focused on immune cells (Macrophages RAW 264.7). JSH-23 effect on microglial activation in the retinae would be critical to study the impact on JSH-23 on retinal inflammation. Interestingly, there is no study on the effect of JSH-23 directly on retinal ECs or pericytes.

They are other factors that this study did not take into account that could also affect the increase in IOP including the thickness of the cornea and drainage of aqueous humor.

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APPENDIX

Parts of the Chapter II, and Chapter III have been submitted as journal articles and are under review currently.

CURRICULUM VITA

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Professional Organizations:

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Community Involvement:

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Abstracts:

- 1. **Rubens P Homme,** Naira Metreveli, Sudhakar Veeranki, and Suresh Tyagi, Exercise ameliorates the development of diabetic retinopathy, FASEB, Volume 31, Apr 2017
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- <u>Rubens PH</u>, Mahavir S, Avisek M, Akash KG, Kavya N, Harpal SS, Neetu T, David L, and Suresh CT, Remodeling of Retinal Architecture in Diabetic Retinopathy: Disruption of Ocular Physiology and Visual Functions by Inflammatory Gene Products and Pyroptosis, Frontiers in Physiology, Aug 2018
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- Mahavir Singh, Akash KG, <u>Rubens PH</u>, Avisek M, Laha A., Harpal SS, and Suresh CT, Circular RNAs profiling in the cystathionine-β-synthase mutant mouse reveals novel gene targets for hyperhomocysteinemia induced ocular disorders, Experimental Eye Research, May 2018
- Akash KG, <u>Rubens PH</u>, Avisek M, Suresh CT, and Mahavir S, Effect of MMP-9 gene knockout on retinal vascular function, Physiology Genomics, December 2019
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Poster:

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- 2. **Rubens Petit Homme**, Akash George, Mahavir Singh, and Suresh Tyagi, *Modulation of Inflammatory Pathway mitigates Diabetic retinopathy, Kentucky Physiological Society*, April 2018, Louisville
- 3. **Rubens Petit Homme,** Mahavir Singh, Avisek Majumder, Akash George, and Suresh Tyagi, *Inhibition of Inflammation and Pyroptosis mitigate Diabetic Retinopathy via Suppressing Receptor Activated NF-kB Ligand (RANK-l)*, Experimental Biology, April 2018, San Diego
- 4. **Rubens Petit Homme,** Laha Anewesha, Sashi Papu John, Avisek Majumder, Akash George, Mahavir Singh, and Suresh Tyagi, *JSH-*23 Mitigates Diabetic Retinopathy via Reducing Oxidative Stress, Kentucky Physiology Society, March 2019, Northern Kentucky

 Rubens Petit Homme, Laha Anewesha, Sashi Papu John, Avisek Majumder, Akash George, Mahavir Singh, and Suresh Tyagi, Oxidative Stress Mechanisms in Diabetic Retinopathy, Experimental Biology, April 2019, Orlando

Presentation:

- 1. February 2019. Spencer Symposium, Florida Southern College Department of Chemistry and Physics Lakeland FL. *Stopping Inflammation Could Prevent Diabetic Retinopathy*.
- 2. March 2019. Physiology Seminar Series, University of Louisville School of Medicine Department of Physiology, Louisville KY. *Modulation of Inflammatory Pathway by Inflammation Inhibitor* (JSH-23) mitigates Diabetic Retinopathy