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ROLE OF TRIB3 IN PROGRESSION AND PATHOGENESIS OF DIABETIC RETINOPATHY

by

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A DISSERTATION

Submitted to the graduate faculty of The University of Alabama at Birmingham, in partial fulfillment of the requirements for the degree of Doctor of Philosophy

BIRMINGHAM, ALABAMA

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ROLE OF TRIB3 IN PROGRESSION AND PATHOGENESIS OF DIABETIC RETINOPATHY PRIYAMVADA MILIND PITALE VISION SCIENCE GRADUATE PROGRAM

ABSTRACT

Diabetic retinopathy (DR) is reported to be one of the leading causes of blindness in the United States by Center for Disease Prevention and Control. As the estimated prevalence of the disease will likely triple by 2020, the research in this area should focus on identifying novel targets and therapeutic approaches. One of the therapeutic approaches for DR is the reprograming of retinal metabolism to delay the progression of the disease. The unfolded protein response (UPR) is recognized as a cellular pathway activated in diabetic retina which interacts with key transcription factors to physiologically regulate glucose, lipid homeostasis, and angiogenic signaling. During development and progression of DR, persistent activation of UPR occurs. Tribbles homologous 3 (TRIB3) is one of the UPR mediators activated in response to stress. My current project is focused on exploring TRIB3 protein as a novel therapeutic target for DR treatment. We hypothesize that TRIB3 is elevated in diabetic retina leading to aberrant insulin signaling and neovascularization. We tested this hypothesis in animal models of STZ-induced diabetes and hypoxia-induced proliferative retinopathy. Moreover, we identified the mechanism of TRIB3-mediated progression of DR in primary cultured retinal Muller cells. Our data demonstrated that TRIB3 controls major molecular events in early diabetic retinas. It specifically, regulates a retinal glucose flux, and alters expression of inflammatory and metabolic markers. In addition, we found that

TRIB3 ablation leads to significant retinal ganglion cell (RGC) survival and functional restoration accompanied by a dramatic reduction in pericyte loss and acellular capillary formation in hyperglycemic retinas. In hypoxic conditions, TRIB3 KO retinas significantly diminished GFAP and VEGF expression, thus regulating gliosis and aberrant vascularization, and preserving retinal integrity. In conclusion, overexpression of TRIB3 in hyperglycemic and hypoxic retinas may accelerate the onset and progression of DR to proliferative stages indicating TRIB3 as a potential therapeutic target.

Keywords: Diabetic retinopathy, Glucose metabolism in the retina, Neovascularization, Retinal ganglion cells, and Muller gliosis

DEDICATION

I dedicate my thesis first, in the loving memory of my beloved grandmother, Mrs. *Malini Madhukar Pitale (1925-2010), (Bachelors in English literature)* who was one of the first females in our family to get college education. She had a long career in teaching for almost 35 years. When I was growing up, she was my teacher and best friend. It was her encouragement that I chose medicine and research as my career. Second, I want to dedicate my work to my amazing husband Dr. Ninad Chaudhary and our handsome son, *Avneesh*. They have been truly patient and supportive throughout my dissertation.

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CHAPTER 1

INTRODUCTION

Glucose Metabolism and Transport in the Retina

Retina is a multi-layered structure lining the posterior segment of the eye. These layers are composed of several functionally different neuronal and glial cells (Figure-1). From the outermost to innermost neural layer, these cells are photoreceptor cells, bipolar cells (BP), horizontal cells (HC), amacrine cells (AC), muller cells, retinal ganglion cells (RGC) and the astrocytes.

The retinal cell functions and glucose metabolism have a co-dependent relationship.¹ The preferred energy substrate for any mammalian cell is glucose. Glucose is catabolized to lactate and pyruvate with generation of the energy molecule adenosine triphosphate (ATP). Pyruvate then moves to the mitochondria where it is converted to Acetyl-coenzyme A. Acetyl-coenzyme A is further utilized in the citric acid cycle to produce energy precursor nicotinamide adenine dinucleotide/hydrogen (NAD⁺/NADH) which is required for oxidative phosphorylation (OXPHOS).^{1,2} Glycolysis and OXPHOS co-exist in the mammalian cells and fulfills the cellular energy demands but their relationship can be competitive. For example, most cancer cells utilize glycolysis as a major pathway to produce energy independent of normoxia or hypoxia. This phenomenon is called as aerobic glycolysis, also known as a Warburg effect.³ One of the tissues with the greatest metabolic

rate and the energy demand in the human body using aerobic glycosylation is the retina.¹ Among the retinal cells, photoreceptors are the cells with highest metabolic rate and the light dependent energy utilization. The energy demand is prioritized for utilization in the outer segments (OS) of photoreceptors in the light and in the inner segments (IS), and synapses in the dark.² Additionally, glucose metabolism is compartmentalized depending on the presence of glycolytic enzymes like Hexokinase (HK) II and PK-M2.¹ For example, pyruvate kinase -isoenzyme M2 (PK-M2) is present only in the outer retina, implicating photoreceptors as a primary site of glycolysis in the retina.² In addition to photoreceptors, RPE cells in the outer retina also catabolize glucose obtained from choroidal blood circulation by glycolysis.⁴ Further, glial Muller cells metabolize glucose to lactate even in presence of oxygen.⁵ Recent studies have reported that glucose oxidation in Muller cells is limited to its external limiting membrane (ELM), the cell compartment with a high number of mitochondria.¹ However, the activity of OXPHOS enzymes is reported to be weak in the Muller cells.¹ Researchers have proposed that Muller cells spare oxygen for the retinal neurons,⁶ and produce lactate for utilization by surrounding neurons to meet their high energy requirements.⁵

Glucose influx into the retina is mediated by a facilitated transport via glucose transporters (GLUT) particularly, GLUT-1. GLUT-1 transporter is expressed in the IS of photoreceptors where it transports glucose from the RPE.^{2,4} The apical and basolateral membranes of the RPE, Muller, and the retinal endothelial cells also express GLUT1 transporters.^{2,4,7} GLUT-1 has a high affinity for glucose which explains why the retinal cells are more prone to hyperglycemic damage.^{8,9} Further the monocarboxylate transporters (MCT) 1 and 3 are important for lactate efflux from the retinal cells. For

example, lactate generated from glycolysis is carried out from IS of the photoreceptors and secreted by monocarboxylate transporter-1 (MCT-1) into the subretinal space.⁴ The excess of lactate produced by photoreceptors is taken by the RPE cells via MCT-1 transporter. Lactate efflux from the RPE cells to transpithelial choroidal circulation is carried out by MCT-3 transporter.⁴ A balance between GLUTs and MCTs in the retina is required to achieve homeostasis of glucose and lactate shuttle (Figure-2).

Pathophysiology of Diabetic Retinopathy

The American Academy of Ophthalmology defines that diabetic retinopathy (DR) is a neurovascular eye complication associated with damage of retinal vasculature and growth of abnormal new vessels in the retina in response to high glucose levels. The number of cases with diabetes-associated visual impairments has increased by 64 percent over the past twenty years.¹⁰ DR is classified into two stages based on the pathological changes in affected retinas observed over the period of the disease progression. An early stage of DR is known as a non-proliferative diabetic retinopathy (NPDR) and the later advanced stage is known as a proliferative diabetic retinopathy (PDR). NPDR is characterized by presence of microaneurysms, retinal hemorrhages, intraretinal microvascular abnormalities (IRMA), and venous beading. In the advanced proliferative stage, there occurs aberrant vascular signaling, ischemia, neovascularization and finally diabetic macular edema (DME).¹¹ DME is clinically significant manifestation of DR which results from the disruption of the blood retinal barrier (BRB) leading to leakage of blood and cellular fluids in the neural retina.^{12,13} Chronic hyperglycemia is one of the severe

crucial risk factors for DR and glycemic control has shown to delay the progression of DR.^{14,15}

Both hyperglycemia and hyperglycemia-induced hypoxia are the hallmarks of retinal pathobiology in diabetes that trigger development and progression of DR.¹⁶ In the streptozotocin(STZ) -induced diabetic rats as early as 2 weeks post induction of diabetes, the GLUT-1 expression is reduced by 50 % in the neural retinal cells and retinal microvasculature. However, isolated RPE cells did not show a sign of reduction in GLUT-1 expression at 8 weeks post hyperglycemia induction.⁹ This fact is particular interesting since RPE is the major cell that supplies glucose from the choroidal blood vessels to the photoreceptors.

In normal conditions, the utilization of glucose in the retina occurs primarily via glycolysis. In the diabetic retina, excessive glucose is subsequently diverted to activation of polyol, hexosamine monophosphate (HMP), advanced end glycosylation products (AGE), and protein kinase--c (PKC) pathways.¹⁶ In presence of excess glucose flux, the retinal response is a complex interplay between these signaling pathways. Thereby, the enzyme aldose reductase utilizes nicotinamide adenine dinucleotide phosphate (NADPH) cofactor for activating the polyol (sorbitol) pathway. The NADPH cofactor levels subsequently drops resulting in its deficiency to perform other physiological cellular functions such as maintenance of the antioxidants. Thus, the polyol pathway makes cells vulnerable to oxidative damage.¹⁷ Secondly, in the PKC pathway, increased activation of diacylglycerol (DAG) leads to upregulation of protein kinase C. Activation of PKC signaling in turn affects gene expression of cytokines, enzymes, and growth factors that modulate vascular health.^{18,19} In addition to mentioned above signaling, upregulation of

AGEs in the cells modifies the extracellular matrix components and disrupts the regulation of gene transcription.²⁰

Hyperglycemia reduces the levels of glucose-6-phosphate dehydrogenase (G-6PD) enzyme which is crucial for pentose pathway shunt, HMP, and N-acetyl glucosamine. Cappai et al. reported that patients with deficiency of G-6PD are vulnerable to proliferative retinal damage.²¹ ²² Activation of HMP signaling in turn affects gene transcription profile and post-translational modification. In particular, the expression of the proteins that play a critical role in the maintenance of vascular health were reduced in these studies.^{23,24} In addition to mentioned above cellular pathways, the unfolded protein response is activated in diabetic retina which is an important contributing factor in diabetic retinal pathobiology and is therefore, an aspect of my dissertation.

Unfolded Protein Response and PERK/ATF4 Pathway in DR

"The unfolded protein response (UPR) or the endoplasmic reticulum (ER) stress response is a series of evolutionarily conserved signaling pathways aimed at restoring homeostasis under conditions of ER stress.²⁵ Cellular stress, particularly excess of glucose and hypoxia, can trigger UPR activation in diabetic retina.²⁶ The protein kinase RNA-like endoplasmic reticulum kinase (PERK) signaling is the one of the UPR pathways. One of the downstream targets of this pathway is activating transcription factor 4 (ATF4). During the stress conditions, auto-phosphorylation of PERK leads to phosphorylation of the eukaryotic initiation factor 2α (eIF2 α) causing the selective translation of ATF4 even though peIF2 α causes global attenuation of protein translation. Upregulation of

PERK/ATF4 pathway results in activation of inflammation in response to hyperglycemia as well as hypoxic stress.²⁵ Li et al. reported that both the vascular endothelial growth factor (VEGF) and the tumor necrosis factor (TNF)-alpha cytokines were upregulated in Akita mouse retinas as early as 12- weeks of age. The authors also demonstrated that the elevation of TNF-alpha along with activation of ATF4 and p-eIF2a in oxygen induced retinopathy (OIR) mouse retinas occurs at postnatal day (p) 15-16.²⁷ Experiments with retinal endothelial cells cultured in high glucose medium also revealed the activation of ATF4. Moreover, the inhibition of ATF4 in these cells resulted in downregulation of VEGF and intercellular adhesion molecule (ICAM-1), and the inflammatory cytokine promoting leukostasis and leucocyte adhesion.²⁸ In contrast, Zhong et al. considered consistent hyperglycemia as not a powerful stress trigger of the UPR activation in the retinal pericytes even though the intermittent high glucose incubation of human retinal pericyte cells results in upregulation of ATF/CHOP pathway and downregulation of VEGF cytokine expression.²⁹ However, other researcher emphasized that these intermittent hyperglycemic surges compared to other types of the stress cause even more damage to the cells in patients with DM who often have blood glucose fluctuations.³⁰

PERK pathway plays a critical role in triggering angiogenesis. In cancer cells, the activation of PERK pathway in response to hypoxia has been associated with angiogenic stimulation , upregulation of VEGF, and cell adhesion cytokines.³¹ Similarly, in proliferative retinopathies, PERK signaling activates the nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB) transcription factor, an endogenous controller of angiogenesis and inflammatory signaling.^{32,33} In addition, ATF4 , a PERK pathway mediator, plays a key role in promoting proliferative retinopathy. Our lab's prior work

demonstrated that heterozygous knockdown of ATF4 in OIR mice results in significant reduction of neovascularization at p17. Moreover, expression of VEGF and UPR were significantly reduced in ATF4 (+/-) OIR retinas as early as p12.³⁴

In summary, upon the ER stress activation, the PERK/ATF4 pathway regulates the cytokine activation, modulates angiogenesis, and controls vascular health. However, the question of how ATF4 governs these downstream signaling is still not answered. It is also unknown whether the downstream signaling of ATF4 regulates glucose metabolism in diabetic retinas. Therefore, we hypothesis that the ATF4 promotes angiogenesis and cytokine activation through its downstream targets, one of which is tribbles homologous 3 (TRIB3) also known as Tribbles Pseudokinase 3.

Tribbles Pseudokinase 3 (TRIB3)

TRIB3 is a pseudokinase originally found in Drosophila and was first designated as the gene associated with programmed cell death.^{35,36} There are three mammalian homologs of Drosophila tribbles TRIB1,2 & 3. The human *TRIB* 1 and 2 share homology 71%, 1 and 3 shares 53%, and 2 and 3 share 54%. Past research has described *TRIB3* as apoptotic and hypoxia responsive gene,³⁷ and gene associated with autophagy.³⁸ TRIB3 protein is a pseudo-kinase protein with a kinase like domain that is functionally inactive due to the numerous amino acid substitutions at catalytically important positions.³⁸ *TRIB3* gene overexpression leading to the UPR activation has been investigated in response to different stress triggers. Some of the known cellular stressors are hypoxia, excess glucose, upregulation of fatty acids and certain chemicals causing cell toxicity.³⁸ For example, AFT4/ CHOP (C/EBP Homologous Protein) upregulation induces TRIB3 expression in response to metabolic stress. Interestingly, increased TRIB3 in turn can subsequently downregulate ATF4/CHOP signaling via negative feedback loop mechanism maintaining homeostasis.³⁵

TRIB3 is most exclusively investigated protein amongst tribbles family members in glucose metabolism. The foremost work on TRIB3 in liver cells reported that TRIB3 binds to protein kinase- B (AKT) at the threonine (THR³⁰⁸) and serine (Ser⁴⁷³) residues and inhibits the AKT phosphorylation. Inhibition of AKT phosphorylation causes disruption in downstream signaling.³⁹ One such AKT downstream signaling is the production of Nitric Oxide (NO). Studies on human umbilical vein endothelial cell culture (HUVEC) have reported that phosphorylation of AKT leads to activation of endothelial NO synthetase enzyme (eNOS) which releases the NO and regulates vascular health.⁴⁰ NO in turn functions as a vasodilator and an antithrombotic preventing ischemia. Additionally, NO also regulates the inflammation and cell proliferation. In human endothelial cells, TRIB3 overexpression has been shown to attenuate the insulin-mediated eNOS Ser1177 activation.⁴¹

Role of TRIB3 has been also investigated in insulin resistance signaling in muscular cells. Using skeletal muscles of the STZ injected rodents, Dr. Garvey's group demonstrated that TRIB3 overexpression impairs the insulin mediated glucose transport via inhibition of GLUT-4 transporter. They also highlighted the role of TRIB3 in glucose toxicity by characterizing the glucose dependent TRIB3 upregulation in L6 cells.⁴² Glucose induced overexpression of TRIB3 was also reported to be reliant on the activation of hexosamine biosynthetic pathway (HBP).⁴³ In the mice with muscle cell specific TRIB3 ablation, the STZ injection resulted in attenuation of glucose induced insulin resistance. Alternatively,

the overexpression of TRIB3 in these cells augmented the inflammatory and oxidative damage via downregulation of the AKT phosphorylation upon the treatment with STZ.⁴⁴

TRIB3 gene is also described as a controller of the NOTCH-EGFR (epidermal growth factor receptor) pathway in the cancer studies. TRIB3 upregulation promotes cell proliferation, decreases survival of healthy cells, and activates JAG-1 ligand of NOTCH signaling.⁴⁵ Given the JAG-1 ligand possess pro-angiogenic properties similar to ones reported in tumor cells with aberrant neovascularization,^{46,47} it is no wonder that overexpression of TRIB3 serves as a marker of a poor prognosis for patients with breast cancer. Moreover, TRIB3 overexpression is strongly associated with abnormal angiogenesis and pathological neovascularization through TRIB3-NOTCH-VEGF signaling in the human endothelial cells.⁴⁸ These data provide rationale to further explore role of TRIB3 in PDR pathology.

Overall, our attention towards TRIB3 stems from the published studies on the role of TRIB3 in glucose-induced insulin resistance and the association of TRIB3 with ischemic damage causing endothelial cell apoptosis. Although the role of TRIB3 in cell survival or death has been previously highlighted in diabetes and cancer research, the role of TRIB3 in retinal pathology has not been studied yet.³⁸ To address this gap in the literature, we proposed to understand whether TRIB3 overexpression plays pathological role in diabetic retinas.

Our central hypothesis is that TRIB3 is involved in aberrant insulin signaling and neovascularization. Thus, we aimed to characterize the role of TRIB3 in DR pathology.

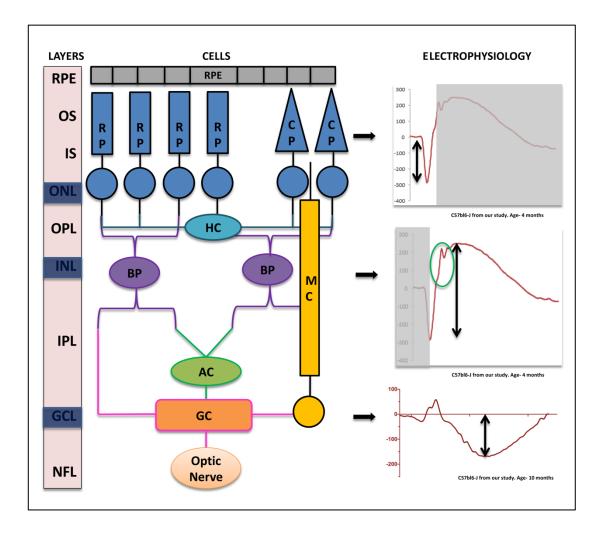


Figure 1. Distribution of retinal cells throughout ten layers of the retina. From outer to inner retina, the retinal layers are RPE layer, ONL composed of photoreceptors' nuclei, INL composed of bipolar cells' nuclei, OPL and IPL composed of horizontal cells and amacrine cells respectively, RGC layer, and finally NFL comprised of the RGC axons as shown in the left panel. The retinal cells' electrophysiological function is measured by ERG. Waveforms representative for different retinal cells in the C57BL6 retina are shown in the right panel. Top- a wave and middle- b wave represent scotopic responses of photoreceptor and BP cells respectively. The bottom right panel depicts amplitudes measured with photopic ERG known as PhNR. These amplitudes reflect the function of RGCs. (RPE: retinal pigment epithelial cell, OS: outer segment, IS: inner segment, ONL: outer nuclear layer, INL: inner nuclear layer, OPL: outer plexiform layer, IPL: inner plexiform layer, RGC: retinal ganglion cell, NFL: nerve fibre layer, RP: rod photoreceptor, CP: cone photoreceptor, BP: bipolar cell, AC: amacrine cell, HC: horizontal cell, MC: muller cell, ERG: electroretinogram, PhNR: photopic negative response)

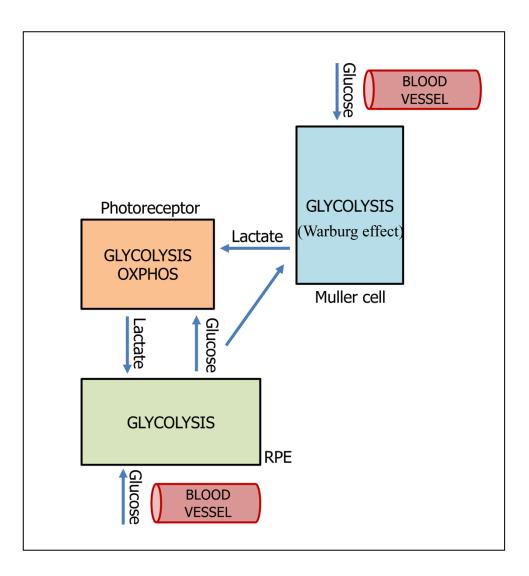


Figure 2. Schematic representation of glucose transport and metabolism in retinal cells. Photoreceptors have high energy demand and have competitive glycolysis and oxidative phosphorylation pathways. Muller cells primarily produce lactate from glycolysis and aerobic glycolysis and release this lactate in subretinal space for photoreceptors. RPE cells not only provide glucose and lactate to photoreceptors but also clear out excess of lactate via transepithelial transport to choroidal blood vessels from subretinal space. Overall, a balance between glucose and lactate transporters is required for the health of the cell.

CHAPTER 2

CHOOSING APPROPRIATE ANIMAL MODEL EXHIBITING HUMAN PATHOLOGICAL CHANGES IN DIABETIC RETINA

Pathobiology of Diabetic Retina

Current clinical trials using *in vivo* imaging techniques have demonstrated retinal morphological changes associated with diabetes. A study with 124 human subjects using spectral domain optical coherence tomography (SD-OCT) demonstrated decreased thickness of the retinal nerve fiber layer (NFL) with no changes in the outer neural retina in patients with early stages DR. In addition to changes detected in NFL, the thickening of the inner nuclear layer (INL) in the diabetic retina has been also reported as a sign of Müller cell hypertrophy and activation.⁴⁹ Furthermore, retinal microangiopathy associated with diabetes is suggested to be a result of increased retinal blood circulation.⁵⁰ A prospective clinical study of 35 subjects showed that multifocal electroretinogram (ERG) analysis employed in patients with severe ocular diabetic complications can be used to successfully evaluate the progression of DME.⁵¹ In another study conducted with 29 individuals with NDPR, the multifocal ERG procedure was used to register the N1 amplitude measured as a peak from a baseline, P1 amplitude measured as a peak from N1 trough, and an N2 amplitude measured as a peak from P1 peak to N2 trough. The amplitudes of these ERG modules were significantly decreased along with delayed implicit time as compared to normal subjects, demonstrating the dysfunction of

the outer neural retina. These functional changes are also associated with the changes in the visual acuity and central macular thickness in the individuals with DME.⁵² In addition, Pardue and colleagues reported results from a cross sectional clinical study involving 16 patients with type II diabetes (T2D) that the follow up ERG examination accompanied with low intensity stimuli may provide earlier detection of neuronal pathology in the diabetic retina.⁵³

While clinical trials concentrate on risk factors and early detection and evaluations of the progression of DR in vivo, access to human donor eye tissue provides a great opportunity to study early molecular changes in the diabetic retina to further understand pathological markers. Multiple studies with postmortem donor eye reported glial cell dysfunction as a primarily manifestation of diabetic retinopathy. A recent study on four postmortem diabetic eyeballs, applied immunolabelling technique for detection of carbonic anhydrase (II) and glial fibrillary acidic protein (GFAP).⁵⁴ The authors revealed the Müller cell nuclei migration and the presence of glial cells in the pre-retinal membranes.⁵⁴ The reported data are in agreement with another study involving 14 donor eyes which demonstrated that the GFAP levels are increased in diabetic eyes as compared to non-diabetic eyes.⁵⁵

In addition to structural and morphological changes occurring in diabetic retinas, there are molecular changes of equal prominence. Studies on post-mortem diabetic eyes have shown presence of inflammation as well as increase in the immunolabelled signal for caspase 3, Fas, and Bax in RGCs and GFAP in the inner retina. ⁵⁶.^{57,58} Studies on vitreous from PDR patients have demonstrated that levels of interleukin-8, monocyte chemotactic protein-1, and macrophage-colony stimulating factor, platelet-derived

growth factor (PDGF) and vascular endothelial growth factor (VEGF) are elevated as compared to non-diabetic individuals.⁵⁹⁻⁶² Microarray analysis of gene expression of fibrovascular membranes extracted from PDR patients during vitrectomy have identified extracellular matrix proteins and elevated expression of genes associated with angiogenesis and apoptosis.⁶³ These analyses have also helped to identify potential therapeutic targets.

Molecular and structural changes observed in the studies using human tissue or postmortem eyeballs have indicated neuronal cell loss in diabetic retinas.⁶⁴ For example, it has been reported that VEGF plays an important role in the development of aberrant neovascularization in diabetic retinas and it's increase can serve as a biomarker of microangiopathy in PDR.⁶⁵ They also found that, in addition to VEGF, increases in the number of apoptotic cells – as measured by a terminal deoxynucleotidyl transferasemediated dUTP nick end labeling (TUNEL) assay - and pericyte and endothelial cell loss are associated with diabetic microvascular complications.⁶⁶ Overall, the studies with postmortem human retinal tissues have informed us on major structural, morphological and molecular changes occurring human diabetic retinas.

Though studies with human donor tissues are an excellent asset for improving our understanding of molecular signaling in diabetic retinas, they cannot provide a complete picture of the mechanism of retinopathy related to diabetic complications. Moreover, the tissues may not be readily available. The use of genetic animal model addresses these limitations as well as, it is an excellent approach for comprehensive understanding of the cellular pathways associated with DR. While choice of the

appropriate animal model mimicking all aspects of human DR pathology is a challenging, several models can capture key cellular and physiological events of diabetic retinopathy in humans. Here in this literature review, we attempted to summarize current status on the developed animal models used in the research focusing on diabetic retinopathy.

Experimental models of DR

One of the key regulators for homeostatic balance of the glucose metabolism in the body is insulin produced by the beta cells of the pancreas. The insulin receptorsignaling pathway facilitates the glucose entry into the cell through the protein kinase B (AKT)-mediated glucose transporter (GLUT1) activation. In humans, fasting blood glucose level is maintained in the range of 92-126 mg/dl while the postprandial blood glucose level is in the range of 97-140 mg/dl. It is well-accepted that under fasting and postprandial conditions, the blood glucose level (BGL) above 126 mg/dl and 180 mg/dl, respectively are considered sustained hyperglycemia.⁶⁷

NPDR is primarily driven by hyperglycemia while PDR is primarily caused by hypoxia. Thus, animal models mimicking events of DR are developed using these two prime inducers. There are several species which have been used for development of DR, including rat, mouse, rabbit, monkey, zebrafish, dog, pig, cat and tree shrew. Accordingly, the experimental diabetes is achieved by induction of a) hyperglycemia using pharmaceutical agents, pancreatectomy, or genetically modified animals.⁶⁸⁻⁷²; b) hypoxia leading to neovascularization (Table.1).

Induction of hyperglycemia

Pharmacological induction of Hyperglycemia. STZ is most frequently used approach to develop T1D model. STZ is an antibiotic produced by the bacterium Streptomyces achromogens and possesses a broad spectrum of antibacterial properties. It has highly reactive methylnitrosourea moiety that exerts the cytotoxic effects resulting in pancreatic β cell necrosis, and the glucose moiety that transports the chemical to the pancreatic β cells. STZ acts via GLUT2 receptor that is abundant on β cell plasma membranes. Therefore, pancreatic β cells are a specific target of STZ.⁶⁸ STZ can be administered either for five consecutive days or as a single dose.⁷³ For example, the range of STZ dosage for multiple doses format is 40-80 mg/kg body weight (bw) injected intraperitoneally (IP) in mouse. The single dose is administered within the range of 150- 200mg/kg bw in mice or 30-100 mg/kg bw in rats by IP injections.⁷⁴⁻⁷⁷ In rabbits, a single dose of 110 mg/kg bw STZ by intravenous administration has been used. In contrast, recently developed tree shrew model of DR requires a single dose of 300mg/kg bw of STZ administrated by IP injections.^{78,79} The maintenance of fasting or non-fasting conditions before STZ injection does not change the post-induction hyperglycemic effect of STZ.⁸⁰ The hyperglycemia after STZ injection is usually seen within 1-4 weeks in most of the species. In some cases, the insulin injections are given to mice and rats to control the extreme fluctuations in the blood glucose levels, although this is not mandatory for STZ models.73

An alternative to STZ, Alloxan, can also induce hyperglycemia and is commonly used in mice, rat, rabbits and pigs. Alloxan is a pyrimidine derivative that directly targets beta cells of the pancreas causing apoptosis by inhibition of glucokinase enzyme and subsequent increase in blood sugar levels due to lack of insulin production.^{70,81} In rodent models, Alloxan-induced hyperglycemia can be developed within 1 week of administration, while it takes less than a day in dogs.⁸²

Surgical and Diet induced Hyperglycemia. Another method to experimentally induce hyperglycemia is surgical removal of pancreas (pancreatectomy) resulting in T1D. Pancreatectomies can be used together with pharmacological agents to get faster induction. For example, in canine models, hyperglycemia develops in 3-4 weeks after the surgery.^{81,83}

In addition to the above methods, dietary modifications can cause changes in the blood glucose levels. The high glucose/galactose diet method is one such approach. Engerman et al. proposed the high galactose diet to induce T2D in dogs.⁷¹ Diet-induced diabetes leading to development of DR has been investigated in several species.⁸¹ However, in dogs and monkeys, this approach can take years to develop changes in the retina secondary to increased blood glucose levels compared to.^{71,84} Rajagopal et. al proposed a high fat diet – induced T2D mouse model. In this model, hyperglycemia was reported to develop at 6 months of age.⁸⁵

Spontaneous (Genetic) model of hyperglycemia

Rodents and zebrafish are the most well-studied genetic models that carry an endogenous mutation leading to spontaneous hyperglycemia. It is relatively easy and economical to develop these models, inbreed them with control backgrounds, and generate a statistically required number of animals for the experiments.

Rat models. There are several spontaneous hyperglycemic rat models: biobreeding (BB) T1D rats, Wistar Bonn/Kobori (WBN/Kob) T2D rats, Zucker diabetic fatty (ZDF) T2D rats, Otsuka Long- Evan Tokushima fatty (OLETF) T2D rats and spontaneous diabetic Torii (SDT) T2D rats. The BB rats are autoimmune DM model which develops hyperglycemia at 3 months of age and shows retinal vascular changes by the 8-11 months.^{86,87} In WBN/Kob rats, the onset of hyperglycemia occurs at 9 months of age, and the male rats are gender preference is.⁸⁸ In contrast, the ZDF rats develop hyperglycemia earlier, between 5-10 weeks of age. These animals are considered a non-insulin-dependent DM model. They are obese and carry missense mutation known as fatty/fa in the leptin receptor gene (Lepr). Originally, these rats are derived from the Zucker rats which are obesity disease model.^{89,90} OLETF male rats develop high blood sugar levels starting from 5 months.⁹¹ In SDT rat model, detection of glucose in urine was reported at 20 weeks of age in males and at 45 weeks of age in females.⁹²

Mouse models. Ins2^{Akita} (T1D) mice, non-obese (NOD) (T1D), db/db mice (T2D), Kimba and Akimba mice are the most popular genetic DM models. The Ins2^{Akita} mice have a point mutation in *insulin2* (earlier reported locus *Mody4*) which causes abnormal insulin production by pancreatic cells leading to pancreatic cell death. The heterozygous

Ins2^{Akita} males are progressively hyperglycemic starting at 4 weeks of age while females exhibit mild symptoms of DM. They have an average life span of 305 days and are primarily a model of early retinal complications.^{93,94} Another T1D model with DR is the NOD mice which mimic the human autoimmune insulin-dependent DM. These mice exhibit CD4 and CD8 cell mediated autoimmune process which destroys the pancreatic cells.^{95,96} Interestingly, there is a gender-based variability in the timeline for the development of hyperglycemia in these mice. Eighty percent of the NOD females develop hyperglycemia at the age of 12 weeks, while males develop hyperglycemia later around 20 weeks of age.⁹⁶ Db/db mice developing T2D model are homozygous for the mutation (Lepr^{db}) leading to spontaneous diabetes and exhibit hyperglycemia at the age of 8-10 weeks (300mg/dl). The db/db (Lepr^{db}) females become hyperglycemic as early as 3-4 weeks of age. In addition to hyperglycemia, these mice are also widely used as a model of obesity and metabolic diseases.⁹⁷ Recently developed Kimba mice are a transgenic line (tr029VEGF) that mimics NPDR and mild PDR.⁹⁸ Another genetically modified Akimba mice are the newly generated model for comprehensive study of the mechanism of DR. These mice are generated by crossing the Ins2^{Akita} and the Kimba mice resulting in the Ins2^{Akita}/VEGF^{+/-} genotype.⁹⁹ They mimic early molecular and advanced microvascular changes in the pathology of DR.

In addition to rodent genetic DR models, *vhl* mutant zebrafish and transgenic zebrafish (Fli-EGFP-Tg) exposed to hypoxic conditions are also the novel genetic models of DM.^{100,101}

Neovascularization model

The most critical pathologic findings of PDR are neovascularization, hemorrhage, and fibro-vascular proliferation observed in the vitreous and the retina, leading to traction retinal detachment and vitreous hemorrhage.¹⁰² The oxygen-induced retinopathy (OIR) in rodents is an accurate and reproducible model of vascular proliferative changes in the retina.¹⁰³ In this model, hypoxia-driven vascular proliferative changes are similar to those seen in retinopathy of prematurity, age-related macular degeneration, and diabetic retinopathy. The OIR in canine and rat models were the first described in early 1950s. In canine models, Arnall Patz and colleagues investigated the effects of hyperoxia on the retinal vessel development to study proliferative retinopathy.^{104,105} To develop a canine model, one-day-old pups were exposed to hyperoxia for 4 consecutive days. In early 1990s, this approach was reintroduced in rodents by Dr. Smith and her colleagues and has gained more popularity. In addition to OIR canine and rodents, the aberrant angiogenesis has also been reported in zebrafish, rabbit, and monkey models (Table1.).

The rodent OIR model is the most common approach to investigate the effect of hypoxia on the retina because it resembles the characteristics of human retinal proliferative changes.^{103,106,107} As the rodent retinal vasculature develops in the first two weeks of birth, researchers can leverage this opportunity to analyze the aberrant vascular development triggered by hypoxia. In this model, hypoxia is induced at p7 after the regression of hyaloid vessels to avoid the development of mixed hyaloidopathy. The rodent pups were then exposed to to hyperoxia (75% oxygen) for five consecutive days from p7 to p12, and then observed at room air from p13 to p17.¹⁰³ The peak changes of neovascularization were seen at p17 which were resolved by p25. The C57BL/6 mice or

the Sprague Dawley (SD) rats are the common strains employed in this model due to their neovascular susceptibility to hypoxia.¹⁰⁶⁻¹⁰⁸

Cellular signaling changes in the pathobiology of DR

Insulin signaling in diabetic retina

Rat models. The basal insulin receptor (IR) signaling has been extensively studied in the STZ-induced diabetic SD rat retina. It was observed that phosphorylation of insulin receptor (IR) in the hyperglycemic retinas remained unchanged up to 8 weeks post-injection, but its kinase activity was reduced by 25% as compared to controls. At 12 weeks post-STZ injection, both kinase activity and auto-phosphorylation of the IR were significantly decreased suggesting that the basal IR activity is diminished in diabetic retina. It was also demonstrated that IRS-2, PI3Kinase expression were significantly reduced at 12 weeks after the STZ injection.¹⁰⁹

Mouse models. Kondo and colleagues observed important differences in insulin signaling between STZ-induced hyperglycemic models and db/db mouse models of DR. Specifically, the IR expression and the tyrosine phosphorylation were upregulated in the first week post STZ treatment in mouse retinas, but no changes were observed in 8 to10-week-old db/db mice. The IRS-1 expression was unaltered while the IRS-2 expression was increased in both db/db and STZ treated mouse retinas. In contrast, few studies reported reduction in IR phosphorylation and increase in the protein tyrosine phosphatase-1B activity (PTP1B) activity in the rod's inner segments 1 week post STZ

injection.¹¹⁰ Analysis of phosphorylated PTP1B in these mouse retinas pointed out PTP1B as a promising therapeutic target to delay the neurodegeneration in diabetic retinas.¹¹¹ Reduced IR kinase activity as early as 12 weeks of hyperglycemia was also reported in study with Ins2^{Akita} mice.⁹⁴

Endoplasmic Reticulum Stress Signaling and Inflammation in the Diabetic Retina

Endoplasmic reticulum (ER) stress is one of the important components of the pathophysiology of the DR. All three branches of the unfolded protein response (UPR), specifically PKR-like ER kinase (PERK)–eukaryotic translation initiation factor 2α (eIF2 α), inositol-requiring protein 1α (IRE1 α)–X-box binding protein 1 (XBP1) and activating transcription factor 6 (ATF6) are activated in the diabetic retina.²⁶ Cellular stresses such as hypoxia and the glucose imbalance can trigger the UPR in diabetic retina.

Rat models. The ER stress markers are upregulated as early as 8 weeks after the onset of diabetes in the SD rats. For example, increase of apoptotic protein caspase 12, C/ERB homologous protein (CHOP), and phosphorylated c-Jun N-terminal kinase 1 (MAPK) was seen at 8 weeks post STZ-induced diabetes in rat retinas. Further, the expression of caspase 12 and MAPK were observed in ganglion cells, while the CHOP levels together with increased GFAP activity were detected in these STZ-treated retinas.¹¹² In another study with SD rats, there was increased expression of *atf4* and *grp78* genes and their related proteins were not detected. These findings suggest that AFT4 might not be the only a signaling molecule responsible for the increased VEGF level in

diabetic retinas.¹¹³ In contrast, immune-histochemical detection of HIF-1α, ATF6, XBP1 and CHOP in STZ -induced diabetic rat retinas showed elevated protein levels at 2 and 4 months. This elevation was accompanied by a decrease in autophagy marker LC3B-II levels indicating potential reduction in autophagy in diabetic retina of mice with 4 months of hyperglycemia.¹¹⁴ The ZFD rat retinas demonstrated apoptotic protein BAX at 6 weeks of age.¹¹⁵

There were also important inflammatory changes across rodent models for DR. For example, the 6 week-old ZFD rat retinas showed increase in the levels of TNF- α and NF-kB.¹¹⁵ Inflammatory proteins like clusterin, tissue inhibitor of metalloproteinase (TIMP)-1, β -2 microglobulin and von Willebrand factor were increased at 4 weeks and significantly overexpressed at 3 months post-STZ injection in the SD rat retinas. In addition, the fibroblast growth factor-2 overexpression was detected in ONL of the diabetic rat retinas at 3 months post-STZ. This group also showed that inflammatory changes were strain dependent in the rats. Thus, as compared to Long Evan and Brown Norway rats, the SD rats have inflammatory changes similar to ones found in human diabetic retinopathy.¹¹⁶ In the OIR SD rat retinas, the activation of inflammatory markers was reported as early as p16.^{106,107}

Mouse models. Using STZ-induced DR mouse retinas, Chung et al. and Zhong et al. have also reported interesting findings on ER stress activation and inflammation. The diabetic mouse retinas had increased expression of GRP78, pPERK, CHOP, VEGF and peIF2 α 4 weeks after STZ-induced hyperglycemia.^{117,118} The inflammatory gene expression is altered in the diabetic retinas deficient in ATF4.¹¹⁸ In addition, the MCP-1

and TNF- α were simultaneously increased in these retinas during the 4-week period.¹¹⁷ The study also highlighted that the ER stress markers, though peaked at 4-weeks, were lower at 6 weeks post-STZ. The Akita mice had increased levels of p-elF2 α and GRP78 proteins in addition to elevated IRE-1 and PERK pathways and the increase in TNF α at 12 weeks of age .^{27,119} Elevated levels of GRP78, ATF4 and peIF2 α were also found in the OIR model at p15.^{27,34} Fifteen-month-old db/db (Lepr ^{db}) mice had increased expression of GRP78, p-IRE-1 α , CHOP, Caspase-3 & ATF4 in the retina along with microglial activation and elevated HIF-1 α / VEGF levels.¹²⁰ The retinal autophagy was also reported to be increased at 40 weeks in the other diabetic models of hypercholesterolemic mice.¹²¹ Overall, these studies emphasized that alterations in the cellular molecular signaling often preceding the retinal pathophysiological events. These findings suggest that dysfunctional insulin signaling, ER stress response, and inflammation are involved in the pathological progression of DR and can be targeted to develop novel cellular therapies for DR.

Retinal cell death and gliosis

Rat models. GFAP activation is often demonstrated in the rodent diabetic retinas. An increase in GFAP immunoreactivity was observed in STZ induced hyperglycemic rat retinas early at 6-7 weeks,¹²² and later at 8-16 weeks post injection.^{114,122} The retinal cell loss and functional changes have also been reported in the STZ induced diabetic rat retinas as early as 2 weeks and later after 24 weeks post STZ injection. In the OIR rats, thinning of the outer segments (OS) of photoreceptors ,and reduction in the INL and IPL thickness have been reported at p18.^{123,124} Moreover, the diabetic rat retinas showed increased apoptotic cells in the ONL, INL, and RGC layer between 12-16 weeks post STZ injections resulted in decrease in a total retinal thickness.^{114,125} At 4 weeks post STZ treatment , ribbon synapses and post synaptic terminals of photoreceptors showed degenerated mitochondria while RGCs developed necrosis.¹²⁵ The WBN/ Kob rat retinas have shown photoreceptor degeneration at 4 weeks ⁸⁸ while STZ-induced diabetic rats manifest severe loss of photoreceptors at 12 and 24 weeks.¹²⁵ Interestingly, the RPE degeneration was also detected in diabetic retinas. For example, 4-month old diabetic BB rat demonstrated the hyperglycemia-induced RPE degeneration with focal necrosis.¹²⁶

The SD rat retinas showed increase in the thickness of the retinal inner limiting membrane (ILM), and degenerating photoreceptors and the OPL at 28 months after hyperglycemia induction.¹²⁶ Extensive glial activation, photoreceptor outer segment (OS) degeneration and decreased RPE-65 were also reported in 32 week-old ZFD retinas.¹²⁷ Further, 9 months after hyperglycemia, decrease in the thickness of the INL, ONL , and the RPE layer have been reported in the retinas of OLETF rats.⁹¹ The hyperglycemic spontaneously diabetic Torii (SDT) rats showed retinal detachment and fibrous proliferation after 50 weeks post hyperglycemia induction.⁹² Recently, in a pre-diabetic rat model, the reversible changes in retinal pathology such as thinning of the inner retina without vascular changes was reported.¹²⁸ Finally, in the OIR SD rat retinas, multiple studies have demonstrated Müller cell reactivity and the thinning of the INL and IPL at p18.^{123,124,129}

Mouse models. STZ-induced diabetic mice have shown the RGCs loss as early as 6 to 12 12 weeks.¹³⁰ Another study conducted with C57BL6/J mice showed the apoptotic

RG cell death at 14 weeks post STZ induced diabetes while decrease in the thickness of total retina was observed earlier at 10 weeks post STZ treatment. They also observed that the number of RGC apoptotic positive cells measured by TUNEL were 25% higher as compared to control retinas.¹³¹ However, the RGC density across the retina varied at 20 weeks post STZ treatment.¹³² Few studies on Ins2^{Akita} mice detected early cone photoreceptor cell loss at 3 months along with reduction in IPL, INL, and RGCs at 22 weeks and reduction in photoreception pre-synaptic and post-synaptic ribbons at 36 weeks of hyperglycemia.^{94,133} Similarly, the OCT analysis of 16-week or 28-week old diabetic db/db mice retinas showed the thinning in NFL and RGC layer at the rate of 0.104 μ m per week resulted in reduction of the total retinal thickness by 28-week.^{132,134} The 28-week old diabetic db/db mice also showed TUNEL positive photoreceptor cells and reduction in the ONL. STZ-induced hyperglycemic mouse retinas manifested GFAP overexpression in retinal astrocytes as early as 5 weeks post STZ treatment while Muller cell gliosis was not seen even after 15 months of DM.^{135,136} In the OIR mouse model, there is a reduction in the total retinal, INL and IPL thicknesses as well as distorted photoreceptor OS, neuronal loss, hyperactivity of Müller cells, microglial activation at p18.¹³⁷

Functional changes in the neural retina

Rat models. Several studies have reported ERG findings in diabetic SD rats. First, there is a delay in the implicit time detected at 4-7 weeks post STZ. Second, a decrease in the a- wave of the scotopic ERG amplitude was detected at 10 weeks while the b- wave amplitude was found to be reduced only at 25 weeks after induction of

hyperglycemia.^{53,122,138,139} Similar ERG findings were observed in SDT rats at 44 weeks post STZ treatment.^{140,141} The OIR rats also demonstrated decreases in the a and b wave amplitudes at p18.^{123,124,129}

Mouse models. The ERG experiments in STZ injected mice have shown decrease in the implicit time for OP at 6 weeks and reduction in b-wave amplitude at 6 months after induction of hyperglycemia.^{115,142,143} Moreover, development of T2D models like db/db (Lepr ^{db}), Ins2^{Akita}, and high fat diet induced DM mice have also reported a chronic timeline of 6, 9 and 12 months, respectively to observe similar ERG changes.^{85,133,144,145} On the other hand, the OIR mice showed reduced a and b wave to OIR rats at p18 similar.¹³⁷

Microvascular changes and neovascularization

Rat models. In albino rats (Wistar–Kyoto), the BRB disruption occurs as early as 2 weeks post STZ injection. Several studies have reported early neovascular changes such as adherent leukocytosis and thickened basement membrane that occur at 8 and 12 weeks, respectively.^{57,146,147} Gong et. al reported that neovascularization occurring in the STZ injected SD can be observed at 3-4 months after induction of hyperglycemia. The neovascularization in these animals correlated with the increase in VEGFR1 and VEGFR2 expression levels.¹⁴⁸ These findings were also observed in the Alloxan induced diabetic rat retinas where leukocytosis and neovascularization were reported at 2 and 9 months after induction of hyperglycemia, respectively. Later at 2 months of sustained hyperglycemia the authors observed pericyte loss, formation of acellular capillaries and the basement membrane thickening.^{149,150} In contrast to previous data, several studies

reported that the BB rat retinas manifested these changes as early as 4 months, while BB, ZDF and OLETF rat retinas demonstrated BRB breakdown and the pericyte loss at 6- 8 months.^{86,87,91,151,152} Downie and colleagues reported increase in extraretinal neovascularization and impaired pericyte distribution in the OIR SD rat retinas as early as p18.¹²⁴

Mouse models. The STZ mice developed microvascular changes early in the course of diabetes than the STZ-induced hyperglycemic rats. For example, vascular permeability measured by imaging of the distribution of fluorescein-conjugated dextran, was compromised as early as 8 days post STZ injection.¹⁵³ However, decrease in arteriolar diameter and velocity were reported at 4 weeks and 8 weeks post STZ injection respectively.⁷⁵ The pericyte loss and acellular capillaries were observed in STZ injected mice from 6 to 9 months later in the course of DR.¹³⁵ The Ins2^{Akita} mice demonstrated increased leukocytosis at 8 weeks, compromised vascular permeability at 12 weeks, microaneurysms at 6 months and neovascularization at 9 months of hyperglycemia^{94,144} In Kimba mice, abnormal blood vessel development was seen early at p28 while increase in vascular permeability and adherent leukocytes was observed at 6 weeks of age. Additionally, loss of retinal capillaries, neovascularization, increased avascular area and alteration in the vessel length and pericyte loss were reported from 9 weeks to advanced age of 24 weeks.^{98,154} Akimba mice were specifically developed to study the microvascular changes of DR and showed these changes at the early age of 8 weeks old.⁹⁹

The db/db mice resembling T2D showed increase in vascular permeability and basement thickness at 13-14 weeks of hyperglycemia.¹⁵⁵⁻¹⁵⁷ Further, high fat diet-

induced T2D mouse model demonstrated pericyte loss, blood retinal barrier disruption and vascular leakage at 12 months of age.⁸⁵ Finally, the OIR mice developed irregular blood vessel and reduction in the retinal inner and deep plexuses at p18, mimicking retinal proliferative events triggered by hypoxia in patients with diabetic complications.¹⁵⁸

Pathological changes in the non-rodent animal models of DR

Apart from rodents, several other species have been used to develop DR models. Characteristics of some of the non-rodent DR models reported over the past two decades have been discussed in this section.

Rabbit

Forty percent of diabetic rabbits with average blood glucose levels of 200mg/dl develop retinopathies after 135 days of initial STZ injections. These retinopathies were classified as degree "3" of retinal pathology representing serious vasculopathy with retinal hemorrhages, vascular lesions and venous thrombosis while 50% were classified as degree "4" corresponding to proliferative retinopathy.⁷⁸ In another rabbit angiogenesis model created by the implantation of human recombinant VEGF at dose of 30 µg, the vitreous showed abnormal tortuous blood vessels followed by vascular leakage at 14 days, and the neovascularization at 21 days of transplantation.¹⁵⁹

Zebrafish

After 12 days of exposure to hypoxia, the zebrafish retina demonstrated new sprouts in the optic capillary plexus and formation of capillary tips.¹⁰⁰ In the *vhl* mutant zebrafish, the increase in aberrant blood vessel formation and the VEGF mRNA expression were observed at 2 days post-fertilization.¹⁰¹ Alternatively, immersions of zebrafish in a high-glucose solution over a period of 30 days caused irreversible reduction in the IPL and INL of zebrafish retina.¹⁶⁰ It has been emphasized that the genetic manipulation and easily achievable hypoxic conditions with this model favors the use of zebrafish as a model for DR.

Monkey

Primate retina is the closest animal model for mimicking the normal morphology and DR pathology of the human retina. The implantation of 100 µg of human recombinant VEGF in the rhesus monkey resulted in increase in vascular permeability, breakdown of BRB and the tortuosity of the blood vessels in the vitreous at 2-3 weeks post-treatment.¹⁵⁹ The STZ induced diabetic monkey model demonstrated retinal pathology after 6-years of hyperglycemia duration. The primary retinal findings in these diabetic animals were cotton-wool spots, macular atrophy, arteriolar occlusion, focal intraretinal capillary leakage and capillary dilatation.¹⁶¹ In addition to STZ induced hyperglycemia, the markers similar to diabetic retinal pathology were also observed in mild hypertensive rhesus monkey and were categorized into three stage of retinopathy: 1) Stage I- microvascular abnormality with capillary dropout; 2) Stage II- vascular leakage, intraretinal exudates and cystoid degeneration with cotton wool spots; and 3) Stage IIIvascular occlusion and retinal atrophy.¹⁶²The above stated retinal findings slowly

developed in the T2D obese monkey over the period of 1.25 to 15 years.¹⁶³ The functional changes in the retina such as decline in the mERG recording, reduced a-wave and oscillatory potential were observed in the aged diabetic rhesus monkey with T2D over 5 years.¹⁶⁴

A recent diet-induced primate model of marmoset monkey developed DR within 2.5 years as compared to other models developing retinal phenotype over the period of 15 years. The marmoset monkeys are smaller than the rhesus macaques, and easier to be maintained. In addition to manifesting a retinal pathological phenotype make these animals as an excellent model of DR.⁸⁴

Tree shrew

Tree shrews are closely related to the primates and have cone-dominant retinas. Tree shrew model generated with either 300mg/kg bw single dose,⁷⁹ or two doses of 80 mg/ kg bw,¹⁶⁵ have resulted in hyperglycemia within a week post STZ injection thus mimicking T1D. Unfortunately, the validation of tree shrews as a model of DR is still under investigation. The only reported findings with the T1D tree shrew retina is the presence of corneal autofluorescence within 7 days as a results of a single STZ dose injection of 300mg/kg bw.⁷⁹

Dog

Induction of hyperglycemia with high-galactose diet is the most common method for development of DR in dogs. Dogs fed with high-galactose diet develop DR changes similar to ones found in human DR pathology and were reported as the first animal model to develop both NPDR and PDR.¹⁶⁶ For example, the pericyte loss, an NPDR marker, is developed in these animals at 9 months post STZ injections. The PDR complications such as hemorrhages, microaneurysms, basement membrane (BM) thickening, vitreous detachment, and neovascularization can be developed in this model within 28 to 68 months post STZ treatment.^{166,167}

Swine

Diabetic pigs are another model of DR. Their retinas have many similarities with human retinal tissues. Alloxan-induced T1D and diet-induced T2D pig model are frequently used models of DR. Alloxan treated pig retinas developed pericyte loss and BRB breakdown at 20 weeks after hyperglycemia.¹⁶⁸ In contrast, another study found that Alloxan induced DM resulted in the Müller cell contraction-promoting activities in the vitreous at 30 days after induction of hyperglycemia. ¹⁶⁹ These findings suggested that swine Müller cell contraction-promoting activity and resultant retinal detachment at advanced stages of diabetes are similar to the changes seen in human retinopathy. In the combination model of DM and hypercholesteremia (DMHC), at there was increased BRB permeability, gliosis, microglial activation, and decreased retinal thickness at 24 weeks.¹⁷⁰ Kleinwort et al. reported that intraretinal microvascular abnormalities and central retinal edema were observed in swine DR model 2 years after the onset of hyperglycemia.¹⁷¹ Lim et al. recently reported an innovative approach where the 4month-old Ossabaw pigs developed chronic DM by feeding the with the western (highfat/high-fructose corn syrup/high-calories) diet for 10 weeks. These animals developed

retinal INL disruption, thickened BM, pericyte ghosts, acellular capillaries and increased fibronectin expression at the age of 6 months.¹⁷²

Cat

The major drawback of DR studies in large animals is the delay in development of histological features, making large animals less attractive species. Hatchell and colleagues were first who developed the feline diabetic model using pancreatectomy and monitored them for 9 years with regular checks for hyperglycemia every 6 months. They subsequently reported presence of thickened BM at 3 months, microaneurysms at 5 years, and neovascularization around 6.5- 8 years post-surgery in the diabetic cat retina.¹⁷³

Limitation of DR models

In this review, we identified the discrepancies in the rodent models of DR induced by STZ injections. The cellular pathology upon long-term development of diabetes was not uniform across different research groups. For example, Martin and colleagues reported retinal cell loss while Feit-Leichman and co-authors observed no changes in the number of retinal cells in STZ treated mouse retinas.^{131,135} These findings could be attributed to different strains used for the development of STZ-induced hyperglycemia. For example, some rodents have consistent high blood glucose level across the studies, while others require additional dosages of STZ over time. Interestingly, STZ induced hyperglycemia is gender dependent. Male mice are more prone to STZ induced pancreatic damage as compared to females, in which the STZ effect is inhibited by the female hormone estrogen.¹⁷⁴ Saadane and colleagues recently reported that increasing

the STZ dose by almost 36 % yields hyperglycemia levels and subsequent retinal pathological changes in females similar to ones found in males.¹⁷⁵

Moreover, in our experience, 20% of the STZ injected males developed normal blood glucose after 20 weeks post STZ, while 45% did not develop hyperglycemia even after 6 weeks with a single high dose of STZ injection. Despite these limitations, the chemical induction of hyperglycemia is still the fastest method to induce diabetes as compared to spontaneous T1D and T2D models. The OIR mouse model is particularly strain dependent model and the maintenance of genetic background in experimental mice is of utmost importance.¹⁰⁸ The rodent models, in comparison to large mammals, provide better accessibility for molecular changes, genetic manipulation, shorter duration to develop diabetes and easy handling. The only advantage of using primates and other large animals is their close resemblance to human retinal physiology. Thus, choosing appropriate animal model for DR research is a challenging process and requires a careful evaluation of all available models and related research resources.

Conclusion

In conclusion, the current review emphasizes the needs f for considering several factors before deciding on the use of DR model. We highlighted the variability in the duration of the study, methods of induction, molecular techniques, and the markers of diabetic retinopathy across different animal models. Because not all animal models accurately mimic the human retinal pathology, it is important to evaluate the strength and weakness of each animal model. (Figure 1).

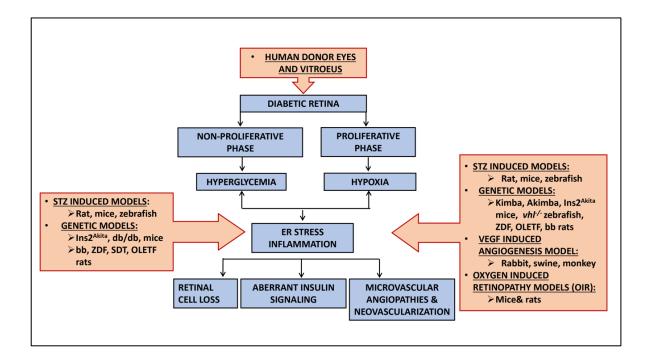


Figure 1. Summary of animal models of diabetic retinopathy. Diabetes – associated ocular complications are divided into non- proliferative and proliferative phases. These phases are governed by hyperglycemia and eventually hyperglycemia-driven hypoxia. The diabetic retinopathy models include pharmacological models, spontaneously diabetic models and finally hypoxia-induced models.

Table 1: Summary of the animal models of DR

A	Hyperglycemia I					
	Method	Species	Dosage	Hyperglycemia	References	
1.	Streptozotocin (STZ)	mouse, rat, rabbit, tree shrew, monkey, cat	mouse and rat- IP 40- 80mg/kg (5 days), mouse- IP 150- 200mg/kg (single dose), rat- IP 30-80mg/kg (single dose), rabbit- IV 110mg/kg (single dose), tree shrew- IP 80 mg twice a week apart and IP 300mg/kg (single dose).	mouse & rat approx. 1-week post STZ	[73-79, 160,164]	
2.	Alloxan	mouse, rat, rabbit, swine, dog			[68,76,82,167 168]	
3.	Pancreatectomy	cat, monkey, dog			[81,83,172]	
4.	High galactose /fat type 2 diet	mouse, rat, dog, swine, zebrafish, monkey			[71,81,84- 85,159,162]	
В	Spontaneous Hy	perglycemia		<u> </u>		
	Mouse			Hyperglycemia	References	
1.	Ins2Akita mouse: Type I DM, mutation in insulin			4 weeks	[93-94]	
2.	Non-obese mouse (NOD): Type I DM, autoimmune model			12 weeks	[95-96]	
З.	db/db (Leprdb) mouse: Type II DM			8-10 weeks	[97]	
4.	Kimba mouse: Tr	ransgenic mo	use (tr029VEGF)		[98]	
5.		Akimba mouse: Ins2 ^{Akita} / VEGF (+/-)			[99]	
	Rat		Hyperglycemia	References		
1.			utoimmune model	3 months	[86-87]	
2.	Wistar Bonn/Kobori (WBN/Kob) rats: Type II DM			9 months	[88]	
З.	Zuker diabetic fat	ty (ZDF) rats:	Type II DM	5-10 weeks	[89-90]	
4.	Otsuka Long- Evan Tokushima fatty (OLETF) rats: Type II DM			5 months	[91]	
5.			T) rats: Type II DM	5 months	[92]	
С	Neovascularization					
	Mouse			References		
1.	Oxygen induced r	etinopathy (C	DIR)		[103]	
2.	Kimba mouse				[98]	
3.	Akimba mouse				[99]	
	Rat, Canine			References		
1.	Oxygen induced retinopathy (OIR)				[104-105,107]	
	Rabbit			References		
1.	Implantation of hu	ıman recomb		[158]		
	Zebrafish			References		
	Angiaganagia		[100,101]			
1.	Angiogenesis					
1.	Monkey				References	

Table 2: Summary of the pathological changes in different models of DR

		A. Molecular Signaling			
	Model	Changes	Duration of hyperglycemia	References	
1.	STZ Rat	Elevated CHOP, Caspase 12, MAPK retinal cytokines	8 weeks	[112-114, 116,109]	
		Reduced IR kinase activity	8 weeks		
		Elevated retinal cytokines	3 months		
		Reduced IR kinase activity & auto phosphorylation and downregulation of IRS-2 & PI3K	3 months		
		Upregulation of HIF-A, ATF-6, XBP1	4 months		
2.	ZFD Rat	Elevated Bax, TNF-α and NF-kappaB	6 weeks	[115]	
З.	OIR Rat	Elevated VEGF, PDEG & TNF-α	P16	[106-107]	
4.	STZ Mouse	Upregulation of GRP78, pPERK, CHOP, VEGF, pEIF2α, retinal cytokine & TNF-α	4 weeks	[27,110-111 , 117-119]	
		Elevated IR expression and tyrosine phosphorylation; upregulated IRS-2 and reduced PDK1/AKT protein levels & phosphorylation	1 week		
		Reduced IR phosphorylation	1 week		
5.	Ins2Akita Mouse	VEGF and TNF-α elevation, increased mRNA expression; protein expression of GRP78 and elevated peIF2α and ATF4 and reduced IR kinase activity	12 weeks	[27,94,119]	
6.	Leprdb (db/db) Mouse	Increased IRS-2 expression and reduced PDK1/ AKT protein levels and phosphorylation	10 weeks	[120,121]	
		GFAP activation, increased expression of HIF-A, VEGF, GRP78, p-IRE-1, CHOP, Casapase-3 and ATF4	15 months		
		B. Microangiopathy			
	Model	Changes	Duration of Hyperglycemia	References	
1.	STZ Rat	Blood retinal barrier disruption	2 weeks	[57,146-148]	
١.	1312 1101		2 100010		
Ι.	1312 Hai	Adherent leukocytes	8 weeks		
1.	512 Hai	Adherent leukocytes Thickened Basement Membrane (BM)	8 weeks 12 weeks		
		Adherent leukocytes Thickened Basement Membrane (BM) Neovascularization	8 weeks		
2.	Alloxan Rat	Adherent leukocytes Thickened Basement Membrane (BM)	8 weeks 12 weeks	[149-150]	
		Adherent leukocytes Thickened Basement Membrane (BM) Neovascularization Leukocytosis Neovascularization	8 weeks 12 weeks 3-4 months		
		Adherent leukocytes Thickened Basement Membrane (BM) Neovascularization Leukocytosis Neovascularization Pericyte loss, acellular capillaries and BM thickening	8 weeks 12 weeks 3-4 months 2 months		
		Adherent leukocytes Thickened Basement Membrane (BM) Neovascularization Leukocytosis Neovascularization Pericyte loss, acellular capillaries and	8 weeks 12 weeks 3-4 months 2 months 9 months		

		Pericyte loss	8 months	
4.	ZDF Rat	BM thickening, pericyte loss and acellular capillaries	6 months	[91]
5.	OLETF Rat	BM thickening, pericyte loss and acellular capillaries	9 months	[152]
6.	OIR SD Rat	Increased extra retinal neovascularization and impaired pericyte distribution	P18	[124]
7.	STZ Mouse	Increased vascular permeability Decreased arteriolar diameter and velocity BM thickening Pericyte loss, acellular capillaries and	8 days 8 weeks 4 to 15 months 6-9 months	[75,135,153]
8.	Ins2Akita	pericyte ghost Leukocytosis	8 weeks	[94,144
5.	Mouse	Increased vascular permeability Blood vessels in the outer plexiform layer (OPL) and microaneurysms Acellular capillaries, BM thickening	12 weeks 6 months 9 months	_
9.	Kimba Mouse	and neovascularization. Abnormal blood vessel development around photoreceptor Increased vascular permeability and	P28 6 weeks	[98,154]
		adherent leukocytes Loss of retinal capillaries, neovascularization, increased avascular area and alteration in the vessel length	9 weeks	-
10.	Akimba Mouse	Pericyte loss Microaneurysms, neovascularization, blood vessel constriction, beading, vessel edema, capillary dropout and new vessel formation it the ONL	24 weeks 8 weeks	[99]
11.	OIR Mouse	Irregular blood vessel development and reduced inner retinal plexus and deep plexus	P18	[158]
12.	Db/db Mouse	Increased vascular permeability and BM thickening13-14 weeksPericyte loss18 weeksAcellular capillaries26 weeks		[155-157]
13.	High Fat diet Mouse	Pericyte loss, blood retinal barrier disruption and vascular leakage	12 months	[85]
		C. Retinal Cell Biology		
	Model	Changes	Duration of Hyperglycemia	References

1.	STZ Rat	Decreased pre and post synaptic photoreceptor ribbon synapses	4 weeks	[114,122, 125]
		Increased GFAP reactivity	6-7 weeks	
		Loss of ONL, INL, GCL	12-16 weeks	
		Severe photoreceptor cell loss	24 weeks	
2.	WBN/Kob Rat	Photoreceptor degeneration	4 weeks	[88]
		Severe OS and ONL degeneration	5 months- 14	
		Ŭ,	months	
3.	BB Rat	RPE degeneration	4 months	[126]
4.	ZDF Rat	Decreased OS, damage to amacrine cells and RPE with gliosis	32 weeks	[127]
5.	OLETF Rat	Decreased INL and photorecptor cells	9 months	[127]
6.	OIR Rat	Reduction in OS, INL, IPL, total retinal thickness, astrocytes and increased muller activity	P18	[123-124]
7.	High galactose fed Rat	gh Increased gliosis and reduced INL and lactose fed OPL		[126]
8.	STZ Mouse	GFAP hyperactivity	5 weeks	[130,132,
		Reduced ONL, INL thickness	6-14 weeks	135-136]
		Total retinal thickness reduced	20 weeks	1
		No retinal cell loss and gliosis	8-12 months	1
9.	Ins2Akita	GFAP hyperactivity	8 weeks	[94,133]
	Mouse	Reduced IPL, INL and cone photoreceptors	3 months	
		Reduced RGCs	22 weeks	
		Decreased presynaptic and post synaptic photoreceptor ribbons	36 weeks	
10.	db/db Mouse	GFAP (8 weeks), (10-16 weeks) and	8 weeks	[132,134]
		Reduced NFL and RGCs	10-16 weeks	
		Reduced total retinal thickness	28 weeks	-
11.	Akimba Mouse	Photoreceptor cell death	25 weeks	[99]
12.	OIR Mouse	Total retinal thickness reduction, distorted photoreceptor OS, neuronal loss, hyperactivity of Müller cells, microglial activation and disrupted INL and IPL D. Electrophysiology	P18	[137]
	Model		Duration of	References
<u> </u>		Changes	Hyperglycemia	
1.	STZ Rat	Decrease in OP amplitude	2-7 weeks	[122,153,
		Decrease in OP implicit time	7 weeks	140-141]
		Decreased a- b wave amplitude	10-12 weeks &	
			44 weeks	
2.	OIR Rat	Decreased a and b wave amplitude	P18	[123-124 ,129]
3.	STZ Mouse	Reduced OP amplitude and implicit time	4-6 weeks	[115,142- 143]
		a and b wave amplitude reduced	6 months	
4.	Ins2Akita Mouse	Decreased OP amplitude, delay in the OP and decreased b wave	9 months	[133-134]
5.	Db/db Mouse	Delay in the b wave, delay in the OP implicit time and decreased amplitude	16-24 weeks	[145]
6.	OIR Mouse			[137]
7.	of a and b wave High fat diet decreased OP amplitude Mouse		12 months	[85]

CHAPTER 3

ROLE OF TRIB3 IN PROGRESSION AND PATHOLOGY OF DR

Introduction

Treatments of severe NPDR with intraretinal hemorrhages and PDR are often focused on vascular abnormalities presented as micro aneurism, vascular leakage, capillary blockade and dropouts, acellular capillaries, and aberrant angiogenesis. The mild and moderate forms of NPDR in patients are controlled by BGL and blood pressure (BP). Although clinical evidence clearly demonstrates that the management of BGL implemented early in the course of diabetes may reduce the development and progression of NPDR,¹¹ a breakthrough treatment that interferes with early DR stages, thus preserving retinal integrity against subsequent damage and preventing progression of DR, remains an unmet critical need.

The tribbles homolog 3 (TRIB3) protein has been proposed as a regulator of insulin signaling (IS) in diabetes. TRIB3 is a metabolic stress indicator and is a critical "stress adjusting switcher," endorsing cell homeostasis shift to metabolic dysfunction. It functions by regulating AKT and AKT/mTOR phosphorylation through pseudokinase activity and controlling autophagy flux and protein degradation.³⁸ TRIB3 overexpression has been linked to aberrant angiogenesis. In a study that employed human umbilical vein endothelial cells (HUVECs) treated with oxidized phospholipids, the induction of VEGF strongly correlated with the highest level of TRIB3.⁴⁸ Mechanistically, TRIB3 reportedly

regulates the nuclear factor kappa B (NF-kB) pathway,¹⁷⁶ promotes activation of TGF-β1 signaling,¹⁷⁷ controls cytokine expression,¹⁷⁸ and governs macrophage health.¹⁷⁹

Recent studies conducted in patients with T2D revealed that the expression of a single nucleotide polymorphism, Q84R in TRIB3 can increase the risk of diabetes and the likelihood of carotid atherosclerosis in part through the effects of abdominal obesity, hypertriglyceridemia, and insulin resistance in the body.¹⁸⁰ These studies have also emphasized that the variant with arginine at position 84 manifests a "gain-of-function" effect due to TRIB3's superior pseudokinase activity in binding AKT and inhibiting insulin-stimulated AKT phosphorylation (Thr308, Ser473).^{41,181-183} TRIB3 mRNA and protein are significantly elevated in human T2D islets, and a substantial reduction of C-peptide has been observed in the plasma of Q84R allele carriers.¹⁸⁴ Q84R polymorphism frequently manifests not only as aberrant insulin signaling, but also with vascular dysfunction and metabolic abnormalities.¹⁸⁵ All of these events occur in the diabetic retina as well.

Furthermore, clinical research conducted with human fibrovascular membranes excised from patients with PDR showed elevated levels of TRIB3.⁶³ However, the exact mechanistic link between TRIB3 overexpression and disease development and progression in DR has not been established. In this study, we explored the effects of abrogated TRIB3 expression in the survival of retinal ganglion, endothelial cells, and pericytes in diabetic mice. Our data reveal distinct underlying mechanism that involves reduced hypoxia regulated GLUT1 and EGFR signaling, which contributes to diminishing neovascularization in DR. Using mouse models of pharmacologically induced diabetes and hypoxia-driven proliferative retinopathies, we demonstrated for the

first time that TRIB3 is a potentially novel therapeutic target that may accelerate the onset and progression of DR to proliferative stages in humans.

We planned to test out hypothesis in three independent specific aims.

1. To investigate role of TRIB3 in early metabolic and inflammatory changes in diabetic retina.

The objective of this aim was to identify whether hyperglycemia can induce the expression of TRIB3, the UPR mediator in diabetic retina. Though the link between the UPR activation and hyperglycemia is known for diabetic retinopathy, the question of whether TRIB3 is upregulated and what the consequences of its upregulation in diabetic retinas are still remained unanswered. Therefore, in aim#1, we hypothesized that TRIB3 expression is elevated in diabetic retinas and its ablation can attenuate early metabolic changes in diabetic retina. First, using TRIB3KO mice, we investigated whether TRIB3 affects glucose transport and metabolism in diabetic retina. Then we examined whether TRIB3 ablation changes the inflammatory profile of retinal cells in diabetic environment.

2. To investigate role of TRIB3 in the retinal neuronal and vascular health in mice with NPDR.

The objective for this aim was to understand the role of TRIB3 in the promoting a neurovascular deficit in diabetic retina. Prior literature has shown that neurodegeneration preceded the vascular damage in diabetic retinopathy.¹³² Therefore, we decided to examine whether TRIB3 ablation can rescue the neural cell loss, in general and the RGC, in particular. We further examined whether rescued RGCs retained the physiological function. Given that one of the hallmarks of early vascular damage in diabetic retina is pericyte loss and increase in acellular capillaries, in the aim #2, we next investigated whether TRIB3 plays role in pericyte apoptosis and whether its ablation can support pericyte and endothelial cell survival in diabetic retinas.

3. To investigate the role of TRIB3 in aberrant neovascularization and activation of gliosis in hypoxic retinas.

The proliferative phase of DR is driven by hypoxia. The oxidative stress and ischemic damage occur in response to hypoxia-mediated VEGF expression resulting in aberrant vascular growth and neovascular tufts formation. Muller cells in the retina are also sensitive to hypoxic changes. Hypoxic stress promotes gliosis and VEGF upregulation in these cells. Therefore, in aim #3, we hypothesized that TRIB3 ablation attenuates aberrant neovascularization and gliosis in hypoxic retinas. In addition, using the in-vitro model of human Muller cells, we revealed HIF1 α as the downstream target of TRIB3 that mediates the TRIB3 –induced VEGF upregulation.

Development of mice hyperglycemic model using STZ: (Aim #1& 2)

Mice were housed at the University of Alabama (UAB) animal facility in a 12hour light-dark cycle condition with unlimited access to food and water as per the UAB-IACUC protocol #09793 and the Association for Research in Vision and Ophthalmology guidelines on the use of animals in ophthalmic and vision research. TRIB3KO mice were generated as previously described,¹⁸⁶ and C57BL6J mice were purchased from the Jackson laboratory. We intraperitoneally injected 8-week-old male TRIB3KO and C57BL6J mice with STZ for both Aims 1& 2. We first developed an acute DM mouse model to observe early metabolic and pro-inflammation retinal changes at 4-weeks. For the proposed Aim-1, we injected mice with 150 mg/kg bw single dose of STZ or vehicle

(0.1 M citrate buffer, pH 4.5). We extracted retinas at 4-weeks after injection to conduct RNA analysis as previously described.^{109,111,118} We next developed non-proliferative chronic DM mouse model to explore cellular and neural damage 8 months post STZ. Therefore, for aim #2, we injected mice with five consecutive doses of 50 mg/kg bw of STZ or vehicle over 5 days.⁴⁴ Electroretinogram and histological analysis were conducted in these mice 32 weeks after injection.

Blood was initially collected via a tail vein, and fasting glucose levels were measured at 1-week post STZ to confirm hyperglycemia in both the STZ models using glucometer. The final measurement of fasting BGL was later conducted at the end of 4th week for the mice used in aim #1. We also measured retinal glucose levels (RGLs) using a calorimetric assay kit in the fourth week. To this end, mice used in aim #2 were analyzed for the fasting BGL every 3 weeks until 32 weeks. Additionally, HbA1c measurements using HbA1c kit were conducted at 15-weeks and 30-weeks post STZ (Supplemental Table 2). Hyperglycemia was considered achieved when the fasting BGL was above 250 mg/dl and >6.5% HbA1c. The animals did not require insulin injections.

Limitation of the STZ mouse model. As previously mentioned, the effect of STZ to induce hyperglycemia is sustained in the male mice only. Estrogens in females inhibit STZ effect eventually giving no rise in BGL .¹⁷⁴ The hyperglycemia in mice also seems to fluctuate over time. The mice might develop acute toxicity initially resulting in transient hyperglycemia and further recovery of BGL. The recovery from hyperglycemia raises the question of whether we should re-inject the mice with STZ after initial

recovery. To overcome these limitations, we used a) male mice only and b) the mice with sustained hyperglycemia throughout the experimental timeline.

Development of OIR mouse model and MIO-M1cells in the hypoxic conditions: (Aim

#3)

The OIR model is one of the most reproducible models of retinal vascular proliferative changes occurring in retinopathy of prematurity, age-related macular degeneration and diabetic retinopathy.¹⁰³ Therefore, we employed the OIR mouse model to generate oxygen-induced retinopathy in C57BL6 and TRIB3 KO pups to study neovascular changes in addition to activated gliosis. The murine OIR model is a wellaccepted model to investigate the effect of hypoxia. Therefore, we hypothesized that neovascularization resulting from hypoxia is delayed and revascularization occurs in the OIR mice deficient in TRIB3 and that direct cytoprotective mechanism is associated with regulation of VEGF expression. A hyperoxia chamber for oxygen exposure was used in the study. Pups were exposed to hyperoxia (75% oxygen) at p7 as described.¹⁰³ At p12, the pups were placed at room air until p17. The choice of time points was determined by the peak of neovascularization and the appearance of revascularization at p17 and p25 respectively.¹⁰³ Thus, it was important to terminate our experiments at p17. Two sets of pups with nursing mothers were placed in the chamber while control mother and pups stayed at room air.

MIO-M1 cells (immortalized human Müller glia cell line)¹⁸⁷ were used to develop a cellular model mimicking hypoxic events in the retinas of mice used aim #3. The MIO-M1 cells were cultured at 37°C in an atmosphere of 5% CO₂ in DMEM optimal

containing 25mM glucose supplemented with 10% fetal bovine serum (FBS) and 1% antibiotic solution.¹⁸⁷ MIO-M1 cells were incubated in the ANAEROgen (W-Zip compact) bags for 48 hours to maintain hypoxic conditions as previously reported.¹⁸⁸

Limitations of the OIR model. OIR animal model is a widely used model to study the impact of hypoxia on angiogenesis, endothelial proliferation, and the role of VEGF in proliferative retinal vasculopathy and inflammation.¹⁰⁸ Although popular, this model has certain limitations. There is a variability in the generation of OIR model depending on the mouse strain.^{103,108} So it is very important to maintain same genetic background in control and experimental animal groups. Therefore, both the experimental and control mice had the same C57BL6 background. Another limitation is the lethal effect of hyperoxia that may occur sometimes in adult mice.¹⁰³ Therefore we used surrogate mothers if needed.

Materials and Methods

The reagents and their catalog numbers and the suppliers are indicated in the Supplemental Table 1.

Quantitative real-time polymerase chain reaction (qRT-PCR) and western blot analysis

Four weeks post STZ injection, mouse retinas were isolated, and RNA was extracted using Trizol. cDNA was synthesized using the Bio-Rad iScriptTM Reverse Transcription Supermix kit. A Thermo Fisher Quantstudio-3 machine was used to perform quantitative real-time polymerase chain reaction (qRT-PCR) using Thermo Fisher TaqMan primers (Supplemental Table 1). Results were normalized to housekeeping Gusb and Gapdh genes and expressed as ratios of mRNA fold changes obtained in diabetic retinas normalized to own nondiabetic controls. For protein analysis, retinal protein extracts were isolated from the OIR and control pups at P13 and separated by polyacrylamide gel electrophoresis, as previously described.¹⁸⁹ Detection of proteins was performed using anti-VEGF, anti-glia fibrillary acidic protein (GFAP), and anti-b-Actin antibodies detecting 21 kDa, 50 kDa, and 42 kDa bands, respectively. Horseradish peroxidase (HRP) goat anti-mouse and infrared (IR) goat anti-mouse antibodies were used as a secondary antibody (Supplemental Table 1). Immunoblots were imaged and analyzed using the LICOR imager system. Differences in VEGF and GFAP levels in diabetic retinas are presented as ratios of band intensities normalized to nondiabetic controls.

Histological and immunohistochemical (IHC) analyses

Human diabetic (84-year-old) and normal (82-year-old) retinal tissues were obtained from the National Disease Research Interchange (IDRI). The tissues were processed for IHC analysis using anti-TRIB3 and anti-CD31 antibodies. Before IHC procedure sections were deparaffinized and rehydrated by immersing the slides through xylene and alcohol of descending concentrations.

Diabetic and the OIR mice were subjected to enucleation at 32-weeks post STZ injection and at P17, respectively. The eyes were fixed for 3.5 hours in 4% paraformaldehyde (PFA) and stored in 30% sucrose before cryopreservation in optimal cutting temperature (OCT) compound. Twelve-micron retinal sections were stained with

hematoxylin and eosin (H&E). A blinded-to-results investigator counted the numbers of retinal ganglion cells (RGCs) and the thicknesses of the inner nuclear layer (INL) and the outer and inner plexiform layer (OPL and IPL). The number of the RGC in 32-week post STZ retinal sections was counted as described in previous studies.^{94,135,190,191} To evaluate the pericyte loss and formation of acellular capillaries, fixed eyes were washed in 1x phosphate-buffered saline (PBS) and the retina cups were excised, as previously described.¹⁹² After digestion, separated retinal vascular tissues were placed on a slide, dried out at room air temperature, and stained with periodic acid-Schiff (PAS) reagent. IHC analysis was performed on retinal sections using anti-TRIB3, anti-GFAP, BRN3A, CD-31/PECAM-1 and Vimentin antibodies (Supplemental Table 1). Co-localization of individual TRIB3 and CD-31/PECAM-1, TRIB3 and BRN3A or GFAP and vimentin proteins in the retinal sections were detected using fluorescent confocal microscopy. To count IBA1 positive microglial cells at P30, mice were injected intraperitoneally (IP)with 10 μ l/g (stock solution of 1 mg/ml) of lipopolysaccharide (LPS). The eyes were enucleated 24 hours after the injection, cryosectioned, and subjected to immunostaining with anti-IBA1 antibody. Blinded-to-results investigators counted the numbers of IBA1 positive cells in the retinal sections.

To detect neovascularization, the eyes of the OIR pups were enucleated at P17, fixed for 1.5 hours and incubated with Isolectin1B-4 diluted in 1x phosphate buffered saline (PBS) overnight. After washing with PBST (PBS +0.02% Triton), the retinal cups were dissected to obtain flat mounts. The flat mount images were used to run a computer program developed by Xiao et al. to analyze areas of neovascularization (NV) and vaso-

obliteration (VO). ¹⁹³ The retinal NV and VO were expressed as ratios of the area with NV or VO to the whole mouse retina.

Retinal physiology

At 32 weeks post STZ injection, the mice were anesthetized to perform electroretinography (ERG) using an UTAS BigShot LKC machine to measure RGC function. Mice were exposed to 15 sweeps (1 sec apart) of white flash of 2.5 cd·s/m2 with a white background of 25 cd·s/m2 intensity and bright fixation to measure the photopic negative response (PhNR) amplitude. The waveforms were measured using LKC EM software.

Seahorse measurement

In Seahorse analysis, the extracellular acidification rate (ECAR) and oxygen consumption rate (OCR) were measured at baseline, post-glucose, and post-oligomycin injections as described in the study.^{194,195} The eyes were collected 4 weeks post STZ injection in the cold 1X PBS. The retinal cup without RPE was dissected in Seahorse XF DMEM medium containing 5 mM HEPES and 2 mM glutamine with no glucose. Retina was placed with RGC layer facing down (Supplemental Figure 1) and secured by the capture screen using an insert tool in each well of a Seahorse XF24 Islet Capture microplate containing 50 μ L of Seahorse XF DMEM. After securing retina in the microplate, 400 μ L of the same dissecting medium were added to each well for final volume of 450 μ L. The microplate was placed in the incubator at 37 °C for one-hour non-CO2 incubation before starting the stress test assay to achieve an even CO2 distribution.

The ports of the Seahorse XF24 Sensor Cartridge were loaded for a final well concentration of 10 mM glucose and 3.0μ M Oligomycin for measuring pH and O₂. Seahorse Wave Desktop Software was used to analyze the data.

Cell culture experiments

MIO-M1 cells (immortalized human Müller glia cell line developed by Limb and colleagues in 2002¹⁸⁷) were cultured at 37°C in an atmosphere of 5% CO₂ in DMEM optimal containing 25mM glucose supplemented with 10% fetal bovine serum (FBS) and 1% antibiotic solution. Cells were exposed to hypoxia or treated with advanced glycosylation end products (AGE-BSA) at dose of 500 mM for 48-72 hours. Protein was extracted using RIPA lysis buffer. Western blot analysis was performed using the anti-TRIB3 antibody and horseradish peroxidase conjugated (HRP) goat anti-mouse secondary antibody.

Experiments were also performed with MIO-M1 cells overexpressing mouse TRIB3-DDK for 72 hours. The cells were harvested, lysed in the IP buffer, and subjected to incubation with TRIB3 anti-body for immunoprecipitation (IP) using agarose G-Plus beads. After verifying expression of TRIB3 by western blot (Supplemental Figure 2), the pull-down samples were submitted to UAB & CCC Mass Spectrometry & Proteomics Shared Facility for protein mass spectrometry and to identify the binding proteins partners of TRIB3. To perform cell toxicity assay, the MIO-M1 TRIB3 overexpressing cells were incubated in MTT solution (0.5 mg/ml/well) for 4 hours at 5% CO2. Following MTT incubation; the cells were incubated with DMSO solution for 15 minutes. The cell survival was measured by estimating optical density at 570 nm using a microplate reader.

To generate hypoxic conditions, 24 hours after transfection, MIO-M1 TRIB3 overexpressing cells were transferred to ANAEROgen (W-Zip compact) bags for 48 hours to maintain hypoxic conditions as previously reported, which generated 1% oxygenation, thus inducing hypoxia.¹⁸⁸ Proteins were extracted using RIPA buffer and TRIB3 (45 kDa), HIF1 α (120 kDa), EGFR1 (170 kDa), GLUT1 (60 kDa), GFAP (50 kDa) and VEGF (21 kDa) were detected, with bands normalized intensities to Actin (42 kDa). Secondary HRP goat anti-mouse and anti-rabbit antibodies and infrared (IR) goat anti-mouse and anti-rabbit antibodies were used for detection. Immunoblots were imaged and analyzed using the LICOR imager system. Experiments with downregulation of HIF1 α and EGFR in MIO-M1 hypoxic cells were conducted using respective siRNAs at concentration of 100 nm.

Glucose uptake assay was performed in MIO-M1cells transfected with siRNA Hif1a and exposed to hypoxia 24 hours later. At 72 hours post transfection, the cells were incubated at 5% CO₂ in glucose-free medium with 0.5% FBS for 45 minutes. Cells were then treated with florescent 2-NBDG glucose for 30 minutes, as recommended by manufacturer, washed with 1x analysis buffer, and resuspended in 400 µl of 1x analysis buffer. The glucose uptake was measured by flow cytometry using Fortessa at UAB Comprehensive Flow Cytometry Core (CFCC). Confocal microscopy was used to obtain florescent images of cellular glucose uptake at 488 nm and florescent signal was quantified using the ImageJ program.

Statistics

Data were analyzed using either two-tailed unpaired Student's t test or oneway/two-way Analysis of variance (ANOVA). Statistical significance was set at p <0.05. The results were plotted in graphs using GraphPad Prism 8 software and data were presented as mean \pm SEM.

Results

In this study, we investigated the role of TRIB3 in the development and progression of human diabetic retinopathy and leveraged STZ-injected and hypoxia driven mouse models mimicking early molecular^{109,111,118} and advanced neurovascular pathophysiological complications of DR, respectively. First, we assessed the TRIB3 level in human diabetic retinas.

Human diabetic retinas overexpress TRIB3 protein

Recent studies conducted with diabetic animal models have clearly demonstrated associations between TRIB3 and numerous diabetes-associated complications, such as hyper-homocysteinemia,^{41,196} non-alcoholic fatty liver disease (NAFLD),^{197,198} diabetic nephropathy (DN),^{199,200} and visceral obesity.²⁰¹ Therefore, we tested the hypothesis that TRIB3 expression is also altered in hyperglycemic retinas. To this end, we analyzed TRIB3 immunoreactivity in the human control and diabetic retinas (Figure 1A). The diabetic retinas of 84-year-old man showed a strong signal for TRIB3 immunoreactivity as compared to the control, the retinas of an 82-year-old man control. Increased TRIB3 staining was observed in retinal endothelial, ganglion, and photoreceptor cells (ONL).

IHC analysis with antibody against CD31, a known endothelial cell marker²⁰², revealed the colocalization (in yellow) of TRIB3 (in red) and CD31 (in green) signals suggesting expression of TRIB3 in the human fibrovascular membrane of diabetic retina (Figure 1A).

STZ-induced mouse diabetic retina and hypoxic muller cells demonstrate overexpression of TRIB3 protein

To proceed further, we took advantage of the TRIB3 KO mice and employed both a pharmacological approach to induce hyperglycemia based on injection with STZ and a hypoxia-induced vascularization approach using an OIR model to mimic the NPDR and proliferative retinopathy, respectively. RNA analysis of four-week post-STZ injected retinas demonstrated about four-fold upregulation of TRIB3 mRNA in hyperglycemic retinas, suggesting that the TRIB3 overexpression occurs early (Figure 1B). TRIB3 expression level was also significantly upregulated in the hypoxic Muller MIO-M1 cells, compared to normoxia, based on western blot analysis (Figure 1C), suggesting that hypoxia in human diabetic retinopathy could upregulate TRIB3 expression level.

The results of the RNA and protein analyses were confirmed in mouse hyperglycemic and hypoxic retinas, which showed a strong immunoreactivity for TRIB3 in IHC (Figure 1D, E). To verify that hyperglycemic RGC overexpresses TRIB3, we further performed an IHC analysis with antibody recognizing the RGC marker, BRN3A, and TRIB3 in control and STZ-treated retinas (Figure 1F). A colocalization shown in yellow was detected from TRIB3 (in red) and BRN3A (in green) signals, suggesting TRIB3 expression in hyperglycemic RGC.

Given that TRIB3 reportedly controls insulin signaling and resistance in cells in general⁴², we next tested the hypothesis that TRIB3 regulates glucose uptake and alters metabolic equilibrium in hyperglycemic retinas.

Hyperglycemic C57BL6 retinas demonstrate raised retinal glucose level, whereas hyperglycemic TRIB3 KO retinas manifest diminished level of glucose

Previously, it was reported that conventional TRIB3 KO in mice does not cause significant difference in the BGL as a response to STZ injection, compared to C57BL6 mice²⁰³. We confirmed this by detecting similar BGLs in both normal and diabetic C57BL6 and TRIB3 KO mouse groups (Supplemental Table 2). Based on this, we asked whether hyperglycemia also increased RGL. Using a calorimetric assay, we determined that four-week hyperglycemic mice have a higher RGL than their non-diabetic controls. Interestingly, TRIB3-ablated hyperglycemic retinas demonstrated significant reduction in RGL, despite the well-matched BGL (Figure 2A, Supplemental Table 2). Given that the retina is extremely sensitive to glucose and an excess of glucose in the retina may modify cellular metabolism; thus, we analyzed early metabolic changes in hyperglycemic retinas.

Hyperglycemic retinas with TRIB3 ablation show altered metabolic responses

TRIB3 is a known metabolic switcher responsible for moving cells from the homeostatic stage to metabolic dysfunction. Therefore, we analyzed the energy metabolism in four-week diabetic retinas of mice mimicking early diabetic changes.^{109,111,118} We monitored both the cellular OCRs and the ECARs in a real-time experiment to measure the mitochondrial respiration and glycolysis rate in the retina

using a Seahorse Extracellular Flux Analyzer (Figure 2B-H). Extracellular acidification is derived from both lactate, produced by anaerobic glycolysis, and CO₂, produced in the citric acid cycle during respiration. However, a previous study revealed that retinas, manifesting the Warburg effect, convert 80–96% of glucose into lactate rather than fully oxidizing it to CO₂ in their mitochondria.²⁰⁴ Here, we found that the ECAR baseline was significantly higher in C57BL6 diabetic retinas, whereas TRIB3 KO hyperglycemic retinas showed no differences compared to both controls (Figure 2B). These results indicate a dramatic difference in acidification rate between C57BL6 and TRIB3 KO diabetic retinas. In fact, the ECAR in the TRIB3 KO diabetic retinas dropped significantly as early as the first ten minutes of the real-time experiment, and this decline continued further over time. However, when we treated the retinal tissue with glucose, dramatic changes were observed in the ECAR rate of TRIB3 KO diabetic retinas. Although both C57BL6 and TRIB3 KO diabetic retinas responded immediately to glucose, the TRIB3 KO hyperglycemic retinas demonstrated a more robust ECAR response at 45 minutes compared to ECAR at 40 minutes (~5-fold TRIB3 KO vs 2-fold in C57BL5, p < 0.01) (Figure 2C). Further analysis of the post-glycolysis ECAR in C57BL6 and TRIB3 KO diabetic retinas revealed that these tissues manifested a similar glycolysis rate (Figure 2D). We next added oligomycin, a known inhibitor of mitochondrial ATP synthase and respiration, to the retinal preparations; this caused the overall glycolytic capacity to drop in a time-dependent manner in both C57BL6 and TRIB3 KO diabetic groups (Figure 2E). Although vehicle- and STZ-injected retinas responded similarly within a single mouse strain, a significant difference was observed

between C57BL6 and TRIB3 KO diabetic retinas: the glycolytic capacity in diabetic TRIB3 KO retina was lower (p < 0.01).

The mitochondrial respiratory rate in the mice was measured with a mitochondrial stress test and the OCR was recorded (Figure 2F). The basal OCR was significantly lower in TRIB3 KO diabetic retinas than in C57BL6 (p<0.0001) suggesting that under hyperglycemic conditions a basal mitochondrial respiration in hyperglycemic retinas could be controlled by TRIB3. In retinal samples stimulated by adding glucose, we observed a dramatic increase of OCR in TRIB3 KO (Figure 2G, ~1.5-fold in TRIB3 KO vs 1.0-fold in C56BL6, p<0.001). Time dependent examination of the retinal tissue responses to glucose demonstrated that hyperglycemic TRIB3 KO retinas uniquely manifested significantly low OCR (p<0.05) (Figure 2H).

Thus, our data demonstrated that TRIB3 ablation in the diabetic retinas manifest altered metabolic responses, which prompts a further exploration of whether other retinal cellular signaling systems are affected at early stages of DR development by TRIB3.

TRIB3 ablation reduces retina inflammatory in diabetic mice

There is a growing literature suggesting that TRIB3 controls inflammatory response in cells.²⁰⁵ To verify the ability of TRIB3 to control the inflammatory response in the retina, mice injected with LPS were sacrificed after 24 hours to analyze IBA1, a microglia/macrophage marker (Figure 3A). IHC analysis demonstrated dramatically reduced number of IBA1 positive cells in the retinas of LPS-injected TRIB3 KO mice, suggesting that TRIB3 ablation could be responsible for the modified pro-inflammatory response in the retina.

Given the early metabolic changes in diabetic retinas at four weeks, we analyzed the expression profile of the genes known to associate with diabetic retinopathy, including *Hifla*, *Aif*1, *Cox*2, *Icam*1, *Nf-kb*, *Rc3h*1 (Roquin), *Zc3h12a* (Regnase or Mcp-1-induced protein-1), and *Vegf*. We found that TRIB3 ablation in diabetic retinas tremendously diminished expression levels of these genes. For example, expression of the intracellular adhesion molecule *Icam*1, responsible for communication between retinal endothelial cells and leucocytes in the diabetic eye, and COX2, a critical player in retinal ganglion cell survival,²⁰⁶ were dramatically reduced in TRIB3 KO diabetic retinas (Figure 3B). These data indicate that TRIB3 ablation results in altered inflammatory response in early diabetic retinas in addition to energy metabolism.

TRIB3 KO results in prevention of retinal ganglion, endothelial, and pericyte cell loss

RGC loss in the diabetic retina has been reported in the literature.^{131,207} At 32weeks of hyperglycemia, we assessed the RGC function and viability in the retinas and found that although the C57BL6 diabetic retinas demonstrated significant loss of RGC, TRIB3 ablation protected hyperglycemic retinas from RGC death. The numbers of RGCs calculated within 100 µm distance in the retinas of TRIB3 KO diabetic mice were comparable to those found in control C57BL6 mice (Figure 4A, B). Therefore, we next recorded the PhNR amplitudes to measure the RGC function known to detect early neuronal damage in the human and mouse retinas²⁰⁷⁻²⁰⁹. Interestingly, the RGC loss in C57BL6 hyperglycemic retinas correlated with dramatic functional loss as measured by PhNR recording, whereas decline in PhNR was remarkably prevented in TRIB3 diabetic retinas (Figure 4C, D).

The RGC layer in the eye is traversed by the retinal capillary beds at varying levels that provide nourishment for both the RGCs and the RGC axons.²¹⁰ Therefore, given that DR is a vascular disease, we next assessed the role of TRIB3 in the retinal endothelial cell health in diabetes. In the hyperglycemic retinas, we observed a dramatic cytoprotective effect of TRIB3 ablation. The pericyte loss observed in C57BL6 retinas was reversed and the increase in acellular capillaries was averted in TRIB3 KO diabetic retinas (Figure 4E-G). Cumulatively, these results indicate that TRIB3 ablation not only has a neuroprotective effect on RGC, but it also maintains retinal vascular homeostasis in diabetic mice.

TRIB3 ablation manifest reduced vascular dysfunction, gliosis, and overall retinal integrity loss in hypoxic retinas

Previously, we reported that ATF4 downregulation is a therapeutic for hypoxic retinas and leads to dramatic reduction in neovascularization and increase in the acellular area of OIR mice.³⁴ TRIB3 is a downstream target of ATF4, expression of which is upregulated during endoplasmic reticulum stress.²¹¹ It has also been suggested that hypoxic conditions upregulate Trib3 expression in the retinal Muller cells.³⁷ Thus, we explored hypoxic TRIB3-ablated retinas employing the OIR model.

At P13, one day after the removal of pups from the oxygen chamber, the VEGF expression was high in the C57BL6 retinas, as expected. Surprisingly, however, TRIB3 ablation in hypoxic retinas resulted in reduced VEGF level (Figure 5A). For example, C57BL6 hypoxic retina normalized to the control demonstrated a 1.3-fold increase in VEGF, whereas normalized TRIB3 KO OIR showed substantial decline in VEGF

expression. Therefore, learning about TRIB3-mediated control of VEGF in the wild type retina, we further analyzed the VEGF-inducted neovascularization in OIR mice.

Analyses of the areas of neovascularization and vaso-obliteration in the OIR retinas demonstrated that TRIB3 ablation significantly reduced formation of neovascular tufts (Figure 5B). Hypoxic TRIB3 KO retinas manifested levels of vaso-obliteration similar to those found in C57BL6 tissues. This difference in neovascularization prompted the question of whether TRIB3-mediated vascularization causes the compromise of retinal integrity and morphology observed in C57BL6 OIR retinas.

Interestingly, TRIB3 KO in hypoxic retinas preserved overall retinal integrity. Although the C57BL6 OIR retinas demonstrated reduced INL, IPL, and OPL thicknesses, TRIB3 KO dramatically reversed these structural changes; no differences were observed between TRIB3 KO OIR and normoxic C57BL6 or TRIB3 retinas (Figure 6A).

Muller cells are known to play a pivotal role in the development of DR and respond promptly to hyperglycemia and hypoxia.²¹² Therefore, we examined Muller cells in hypoxic retinas by detecting the co-localization of GFAP and vimentin, known glial cell markers. ^{213,214} We found that the TRIB3 KO hypoxic mice demonstrated a significantly lower Muller cell immunoreactivity as compared to C57BL6 OIR retinas (Figure 6B). Moreover, detection of the GFAP level in TRIB3 KO OIR mice by western blot also indicated a dramatic reduction in the overall expression, supporting the observation of diminished gliosis in hypoxic TRIB3 KO retinas (Figure 5C).

Together, these results show that TRIB3 is not only involved in early metabolic and inflammatory changes but may also determine the progression of diabetic retinopathy.

TRIB3 controls glucose signaling and VEGF expression, overall impacting the retinal cell viability

The molecular mechanisms of DR are complex and multifactorial. Early molecular changes in diabetic retinas have been reported to occur through a compromised advanced glycation end products (AGE) pathway. Therefore, using Muller cells, we asked whether the treatment of MIO-M1 cells with 500 mM AGE induces TRIB3 overexpression. The results demonstrated significant upregulation of TRIB3 in 48 hours (Figure 7A). These data are in the concord with the results obtained in STZ-induced hyperglycemic mice and the hypoxia treated MIO-M1 cells, indicating that in diabetic hypoxic retina TRIB3 could be overexpressed earlier in the course of disease. Besides overexpression of TRIB3 in hypoxic MIO-M1 cells, we also identified occurrence of overexpression of HIF1 α and EGFR (Figure 7B). The choice of the validated proteins was determined by the literature, and the results obtained in hyperglycemic mouse retina and MIO-M1 cells overexpressing TRIB3 (Supplemental Figure 2). Thus, similar to hyperglycemic retinas, hypoxia in MIO-M1 cells induced both TRIB3 and HIF1 α .

Interestingly, TRIB3 by itself can compromise MIO-M1 cell viability (Figure 7C); more prominent cell death was found in MIO-M1 transfected with TRIB3-DDK plasmid than in control cells. These results are comparable with the RGC cell loss observed in hypoxic diabetic retinas overexpressing TRIB3. Given that HIF1α and EGFR along with TRIB3 were overexpressed in cultured Muller cells, we investigated whether TRIB3 is upstream mediator of these proteins by using TRIB3 overexpression. In hypoxic Muller cells overexpressing TRIB3-DDK cDNA, HIF1α and EGFR were

upregulated along with increased GFAP, VEGF and GLUT1 proteins, indicating that TRIB3 controls these protein expressions as well (Figure 7D). In hypoxic MIO-M1 cells treated with siRNA targeting *Hif1a*, we found that the levels of EGFR, GLUT1 and VEGF were significantly lower, whereas GFAP was unresponsive to the knockdown of *Hif1a*. This suggests that GFAP may be directly regulated by TRIB3 and its expression is not HIF1a-dependent (Figure 7E). Targeting *Egfr* mRNA in hypoxic MIO-M1 cells demonstrated EGFR-dependent HIF1a upregulation. In contrast to experiments with *Hif1a* silencing, the VEGF expression was not responsive to the treatment of cells with *Egfr* siRNA (Figure 7F).

Given that TRIB3 expression is hypoxia-induced and TRIB3, in turn, regulates HIF1 α expression leading to reduction in GLUT1, we asked whether *Hif1\alpha* detected in C57BL6 hyperglycemic retinas (Figure 3) could be responsible for the glucose uptake in Muller cells. Particularly, the observation of higher RGL in hyperglycemic C57BL6 retinas overexpressing *Hif1\alpha* mRNA was worth further investigation. Therefore, we treated hypoxic Muller cells transfected with control or *Hif1\alpha* siRNAs with glucose analog 2-NBDG, known to inhibit glycolysis via its action on hexokinase, and analyzed levels of fluorescence using flow cytometry and confocal microscopy (Figure 8 A and B, respectively). We observed significant reduction in glucose uptake that occurred in HIF1 α -dependent manner, suggesting that the glucose uptake in these cells and the metabolic changes in hyperglycemic retinas could be regulated through a TRIB3 \rightarrow HIF1 α axis. Altogether, these findings indicate that TRIB3 is upstream of HIF1 α , EGFR and GFAP, which allows this molecule to control glycose metabolism, cytokine expression, angiogenesis and gliosis in retinal cells.

Discussion

Overall, the molecular mechanisms of DR are complex and multifactorial. Hyperglycemic conditions affect the intracellular glycose level leading to the retinal endothelial and neuronal cells failing to properly regulate cellular metabolism. Early molecular changes in diabetic retinas have been reported to occur through a compromised polyol pathway, protein kinase C pathway, advanced glycation end products (AGE) pathway, and hexosamine biosynthetic pathway (HBP) in retinal cells.^{16,18,19} Besides the dysregulation of these cellular pathways, hypoxia-induced hypoxia-inducible factor- α (HIF1 α) and vascular endothelial growth factor (VEGF) overexpression play key roles in the progression of NDPR to PDR. In contrast, little is known about how early molecular changes occurring in diabetic retinas lead to the subsequent progression of the disease. Here, we present evidences confirming the critical role of TRIB3 overexpression during the development and progression of diabetic retinopathy. First, we found that TRIB3 was overexpressed in human diabetic retinas and retinas of mice mimicking NPDR and proliferative retinopathy in patients, which provided a rationale to further investigate the role of TRIB3 in diabetic retinas. We discovered that TRIB3 overexpression induced HIF1a-mediated changes in GLUT1 level, glucose uptake, and VEGF expression. In addition to HIF1a, we observed that TRIB3 controlled EGFR expression. Lastly, we determined that TRIB3 ablation in hyperglycemic and hypoxic retinas dramatically prevented retinal neuronal cell functional deficit and loss, as well as VEGF-induced neovascularization.

Retinal vascular homeostasis become compromised with the disease progression. PDR is characterized by uncontrolled growth of fragile blood vessels that can protrude and leak into the vitreous causing blurry vision. This phenomenon is triggered by hypoxia in the retina, one of the known molecular inducers of neovascularization. One of the interesting finding of our study is that both hypoxia and the AGE conditions found in diabetic patients caused the upregulation of TRIB3, which in turn control glycose metabolism, vascular function, and gliosis. Thus, our *in vitro* study indicates that HIF1 α upregulation in diabetic retinas occurs in TRIB3-dependant manner (Figure 3 and 7). A previous study demonstrated that HIF1 α induced TRIB3 overexpression.²¹⁵ In fact, the HIF1 α -TRIB3 relationship could be more delicate and the regulation of their expressions may be interactive. For example, one of such regulation is mediated through ATF4 transcriptional factor, which reportedly cooperates with both proteins by inducing TRIB3 expression²¹¹ and binding to HIF1 α .²¹⁶

Recently conducted study in cultured L6 myocytes^{42,217} indicated that TRIB3 overexpression is induced in glucose–dependent manner resulting in reduced mitochondrial glucose oxidation and the maximal uncoupled oxygen consumption rate. Hence, it noteworthy to find that TRIB3 can regulate glucose uptake in diabetic retina and that this regulation occurs via HIF1α-mediated GLUT1 expression. Indeed, in the whole retinal samples, we found that TRIB3 ablation may modify glucose flux and define glycolytic capacity of diabetic retina. C57BL6 diabetic retinas presented higher RGL, confirming a previous study that reported increase in glucose uptake in diabetic rat retinas.²¹⁸ Changes in glucose flux suggest a plausible explanation for the high RGL in C57BL6 diabetic retinas—stimulation of glucose transporter (GLUT)1–4 activity,

proposed as a target that reduces retinal glucose in diabetic retinas^{219,220}, could be one reason.

TRIB3 KO in diabetic retinas may cause a metabolic shift that allows this retina to function under a "safe operating mode." Previously, it has been demonstrated that diabetic retinas may have higher lactate-pyruvate ratios, suggesting that elevated glucose level mimics the effects of hypoxia on glycolysis and cytosolic free NADH/NAD+.^{221,222} These results were very similar to those observed in this study. C57BL6 diabetic retinas had a higher baseline rate of ECAR derived from lactate produced by anaerobic glycolysis. Although overall glycolytic capacity in the TRIB3 KO diabetic retinas was lower, the response of the retinas to glucose treatment was significantly higher in these mice. Interestingly, the TRIB3 KO diabetic retinas also displayed a lower OCR baseline and revealed a higher sensitivity of the mitochondrial respiration rate in response to glucose treatment. Altogether, these results imply that TRIB3 KO hyperglycemic retinas might respond to glucose elevation similarly to C57BL6 hyperglycemic retinas but operate at limited metabolic mode by reducing glucose flux, instead.

EGFR reportedly promotes angiogenesis, cell proliferation, metastasis, and apoptosis in DR and the inhibition of EGFR signaling protects diabetic retina from insulin-induced vascular leakage.^{223,224} Although we did not assess the level of EGFR in diabetic retinas, we found that TRIB3 overexpression observed in human diabetic retinas induced EGFR in hypoxic MIO-M1 cells. However, unlike the previous studies, we found no direct correlation between EGFR and VEGF levels in Muller cells.²²⁵ Given the fact that diminished HIF1 α level resulted in reduced VEGF, these data indicate that in hyperglycemic and hypoxic retinas, the observed correlation between TRIB3 KO and

VEGF is promoted mostly through a TRIB3 \rightarrow HIF1 α signaling. Future experiments that investigate the TRIB3 \rightarrow EGFR–mediated gene expression profile in diabetic retinas should be conducted; nevertheless, the current study clearly demonstrated that EGFR by itself inhibits HIF1 α expression. Overall, these findings indicate that TRIB3, HIF1 α and EGFR act as pyramid trio set to regulate VEGF expression with TRIB3 upstream and the reciprocal regulation of HIF1 α and EGFR downstream of the pyramid (Figure 9).

Another example of how TRIB3 could control vascular health is through the regulation of an intravascular leukocyte cell adhesion.²²⁶ Although our study did not assess leukostasis in the retinas, we did find that similar to diabetic patients with microalbuminuria and DR ^{227,228}, *Icam1*, a cell surface glycoprotein expressing on endothelial cells and leukocytes, was significantly upregulated in C57BL6 mouse diabetic retinas. TRIB3 ablation led to dramatic reduction of *Icam1* expression. In addition to *Icam1*, we found that TRIB3 controlled the endothelial cell death in diabetic retina and ablation of TRIB3 significantly preserved the formation of acellular capillaries and subsequent pericyte loss. These data provide evidence that TRIB3 plays substantial role in endothelial cell homeostasis in DR.

Inflammation is one of the major factors induced during diabetes, and leads to macular edema, ischemia and neovascularization.²²⁹ In diabetic TRIB3 KO retinas, proinflammatory gene expression profile was altered. Moreover, the increase in retinal microglia and VEGF cytokine production was TRIB3-mediated. TRIB3 could directly or indirectly affect expression of NF-kB and other cytokines, thus modifying early inflammatory response to hyperglycemia. In support of this hypothesis, TRIB3 ablation prevented both neuronal and vascular dysfunction serving as hallmarks of DR in

humans.²³⁰ Therefore, it is of particularly importance that TRIB3 controls RGC cell death. In hyperglycemic TRIB3 KO mouse retinas, RGC preservation correlated with significant improvement of the recorded PhNR amplitudes, known to be diminished in diabetic patients.²³¹ Moreover, in concord with these results, we revealed that in hypoxic retinas, TRIB3 controls retinal integrity too. This latest observation is also supported by the fact that Muller cells overexpressing TRIB3 manifested reduced cell viability.

The current study did not access a cell specific role of TRIB3 in diabetic retinas due to unviability of the mouse model. Therefore, future experiments should identify sources of TRIB3-induced cellular damage, "signaling" neurovascular degeneration in diabetic retinas. Nevertheless, regardless of the sources, we found that hypoxic TRIB3 KO retina manifested less gliotic activity of Muller cells as determined by co-localization of vimentin and GFAP. TRIB3-mediated control of GFAP *in vivo* was confirmed in cultured Muller cells indicating that TRIB3-mediated GFAP upregulation is independent of HIF1α protein regulation. This finding pinpoints TRIB3 as a novel potential regulator of GFAP biogenesis during diabetes.

Overall, our study indicated that TRIB3 mediates the expression of major sets of hypoxia-induced genes, HIF1 α , EGFR, VEGF and GFAP, effects that were observed both *in vitro* and *in vivo*. Our data align with previous studies conducted in human cancer cells, in which TRIB3 was proposed as a molecular regulator of angiogenesis mediated via HIF1 α and VEGF upregulation.²³² In addition to angiogenesis, our investigation indicated that TRIB3 is a master regulator of early metabolic and inflammatory events in diabetic retinas, which affect the overall development and progression of disease to proliferative stages (Figure 9). Given the limitations and side effects of current treatments

for DR and the continuing efforts to understand its complex molecular mechanisms, our findings show TRIB3 is a major game player of diabetic ocular pathophysiology in humans and is a novel therapeutic target, the potential of which should be further evaluated in the clinic.

Acknowledgements

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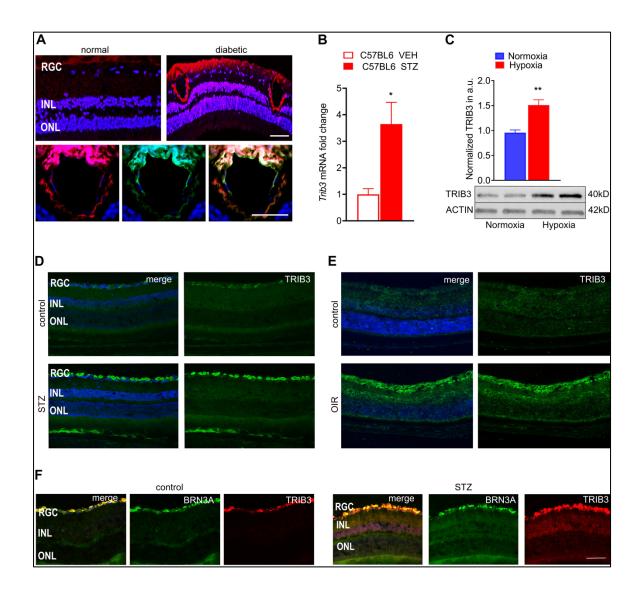


Figure 1. Human and mouse diabetic retinas overexpress TRIB3 protein. A: TRIB3 expression in the human diabetic retinas detected using IHC with anti-TRIB3 primary antibody is shown in red. Strong TRIB3 immunoreactivity is detected in the ONL, INL, RGC and the fibrovascular membrane (FVM) in the diabetic human retina. The TRIB3 immunoreactivity also co-localizes (in yellow) with CD31 positive cells (in green), suggesting that CD31 positive endothelial cells of the FVM also express TRIB3. The scale is 50 µm. B: Detection of TRIB3 expression in mouse diabetic retinas at 4 weeks of hyperglycemia by qRT-PCR (n = 5; *p<0.05). C: TRIB3 overexpression

detected in hypoxic Muller MIO-M1 cells at 72 hours. D: Detection of TRIB3 in the mouse normal and diabetic retinas at 32 weeks after STZ-induced hyperglycemia (in green). DAPI-stained nuclei are shown in blue. Merged images are on the left. Robust TRIB3 expression is detected in the RGC layer. E: Detection of TRIB3 in the normoxic and hypoxic mouse retina at P17. Robust TRIB3 expression is detected in the RGC, IPL, INL and ONL (in green). DAPI-stained nuclei are shown in blue. Merged images are on the left. F: Expression of TRIB3 is co-localized with the RGC marker, BRN3A in the control and diabetic retinas at 32 weeks of hyperglycemia. TRIB3 is shown in red, BRN3A is shown in green, co-localization is shown in yellow (indicated with white arrows). DAPI stained nuclei are shown in blue. Robust TRIB3 expression is detected in the RGC layer. (ONL: outer nuclear layer; INL: inner nuclear layer; RGC layer: retinal ganglion cell layer). The scale is 100 µm.

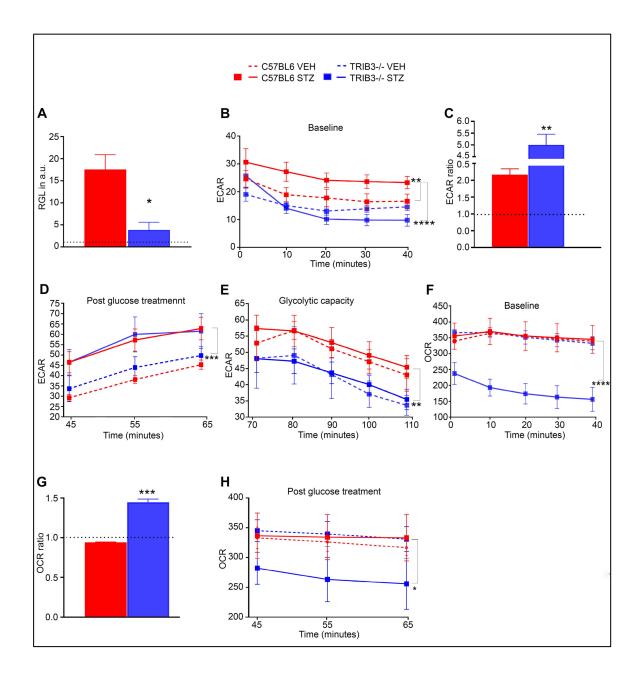


Figure 2. TRIB3 reprograms glucose metabolism in diabetic retinas. A: RGL measured in diabetic retinas (mg/dl) was normalized through own control (dotted line) at 4 weeks of hyperglycemia. Significant reduction of RGL is observed in TRIB3 diabetic retinas (C57BL6 diabetic and control; n = 8-9 and TRIB3 KO diabetic and control; n = 5-11). Please also see Supplemental Table 2. B, C, D, E, F, and H: Results of the metabolic stress experiments (n = 4-6). B: The ECAR baseline of diabetic and control

C57BL6 and TRIB3 KO groups at 4 weeks post injection (mpH/min). Significant increase in the ECAR baseline in C57BL6 diabetic retinas is shown during the first 40 minutes. C: Addition of glucose results in dramatic jumps in the ECAR rate. Thence, the TRIB3 KO diabetic retinas manifest more sensitivity than C57BL6 hyperglycemic tissue. The presented ECAR rates are expressed as the ratios of the normalized ECARs at 45 min over the normalized ECARs measured at 40 min (dotted line). D: Rates of glycolysis in the four experimental groups measured after glucose treatment in a real-time experiment. Both diabetic retinas demonstrate comparable levels of glycolysis. E: Measurement of glycolytic capacity in diabetic retinas after suppression of the mitochondrial respiration rate by using oligomycin. Reduced glycolytic capacity of TRIB3 KO diabetic retinas (about 18%) vs. C57BL6 hyperglycemic retinas (mpH/min). F: The OCR baseline measurements in normal and diabetic mouse groups. Significant reduction in the OCR rate is detected in TRIB3 KO hyperglycemic retinas. G: Addition of glycose at 40 minutes results in a high sensitivity of the OCR rate in TRIB3 diabetic retinas registered at 45 minutes (dotted line). H: However, this jump does not cause an increase in the OCR rate as compared to C57BL6 diabetic retinas in a timely manner. The OCR rate continues to decline in TRIB3 diabetic retinas as measured in pmol/min. (p < 0.05); **(p < 0.01); ***(p < 0.001); ****(p < 0.001));

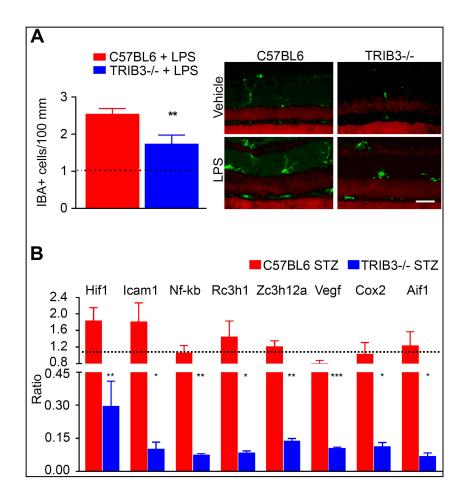


Figure 3. TRIB3 controls immunoresponse in stressed and diabetic retinas. A: Mice were IP injected with 10 µl/g LPS and sacrificed 24 hours later (n = 5). The cryostat retinal sections were subjected by IHC analysis with anti-IBA1 antibody and IBA positive cells per 100 µm were counted (in green). The results are presented as ratios of cells detected in LPS-injected retinas normalized to own controls (dotted lines). B: TRIB3 controls expression of pro-inflammatory genes at 4 weeks of hyperglycemia (n = 3–5). Fold changes for each diabetic group were normalized to own controls. Thus, we observe a dramatic decline in *Hif1a*, *Icam1*, *Nf-kb1*, *Rc3h1*, *Zc3h12a*, *Vegf*, *Cox2* and *Aif1* expression in TRIB3 KO diabetic retinas. *(p<0.05); **(p<0.01); ***(p<0.001). The scale is 100 µm.

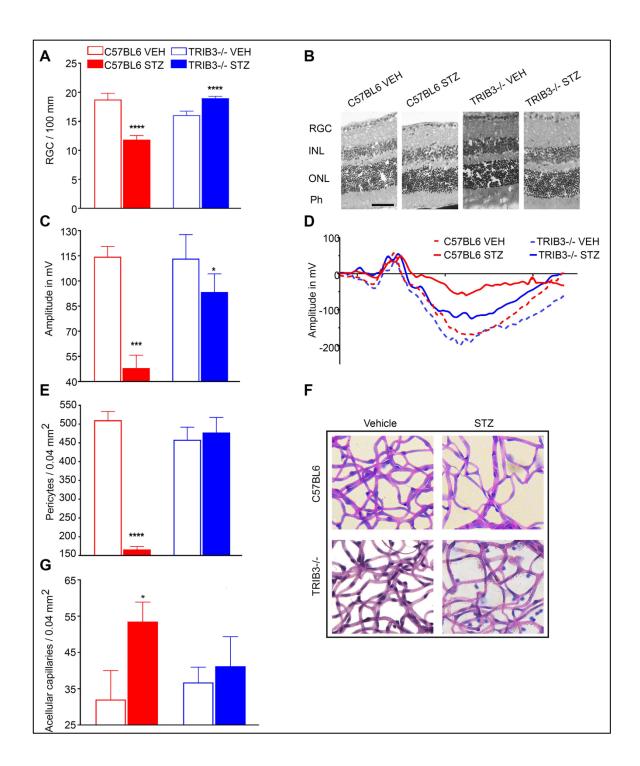


Figure 4. TRIB3 controls neuronal and endothelial health in diabetic retinas. A: The number of ganglion cells/100 μ m is dramatically increased in the diabetic TRIB3 KO retina as compared to other groups. No changes were observed between control or diabetic TRIB3 KO retinas; n = 6. B: Representative images of the H&E-stained control

and diabetic retinas. C: The PhNR amplitudes characterizing the RGC function as recorded by the photopic ERG procedure in the C57BL6 control and diabetic (n = 9 for both) and TRIB3 KO control and diabetic (n = 9-12) mouse groups. Significant diminishing of the RGC function is detected in the C57BL6 diabetic mice whereas dramatic elevation of photopic negative response (PhNR) is observed in TRIB3 ablated diabetic retinas. D: Representative images of PhNR registered in all four groups. E: The number of pericytes detected in 0.04 mm² area of the retina is markedly reduced in the diabetic C57BL6 retinas as compared to other mouse groups; n = 6 for C57Bl6 control and diabetic and n = 5 for TRIB3 KO control and diabetic mice. TRIB3 ablation slows down the pericyte loss in diabetic retinas. F: TRIB3 ablation in diabetic retina prevents formation of acellular capillaries detected in 0.04 mm² area of the retina. G: Representative images for PAS-stained retinal pericytes and acellular capillaries in the retinas of the four mouse groups. *(p<0.05); ****(p<0.0001). The scale is 100 µm.

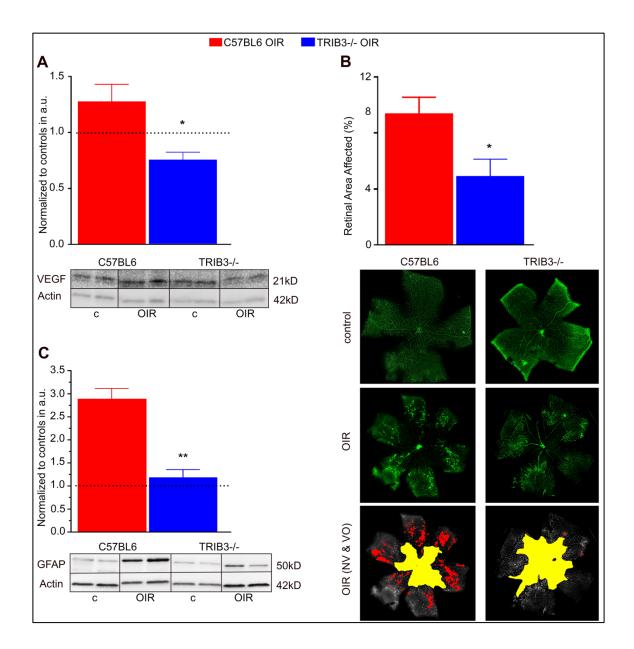


Figure 5. TRIB3 promotes retinal vascularization during hypoxia. A: Expression of VEGF normalized to own controls (dotted line) in two hypoxic groups at P13 is shown (n = 4). Significant reduction of VEGF in TRIB3 KO hypoxic retinas is observed. Representative images of the western blots probed with anti-VEGF antibody are shown at the bottom. B: Ratios of the neovascular (NV) areas in the hypoxic C57BL6 and TRIB3 KO retinas (n = 6-8) to the whole retinal areas at P17 are depicted. The flat mount

images were used to run a computer program developed by Xiao et al. to analyze areas of neovascularization (NV) and vaso-obliteration (VO). ¹⁹³ Significant reduction in NV area is observed in hypoxic TRIB3 KO retinas of pups. Representative images for neovascularization and avascular area in the flat mount retina of pups stained with Isolectin (in green). C: Expression of GFAP in two diabetic C57BL6 and TRIB3 KO groups normalized to own control (dotted line) at P13 is presented (n = 4). A dramatic reduction in the GFAP level in diabetic retinas with TRIB3 ablation was detected. Representative images of the western blots probed with anti-GFAP antibody are shown at the bottom. *(p<0.05); **(p<0.01).

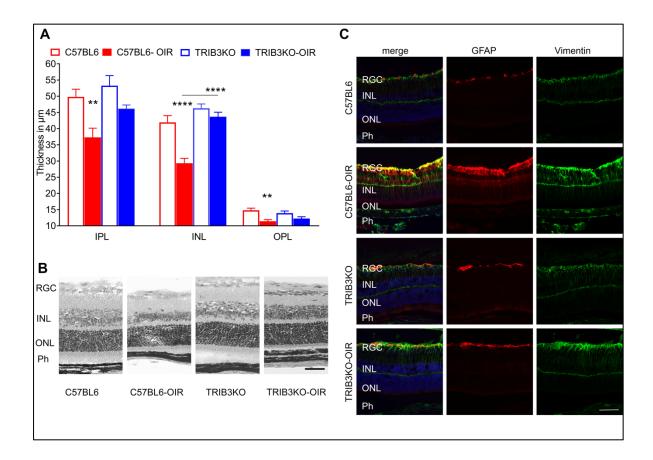
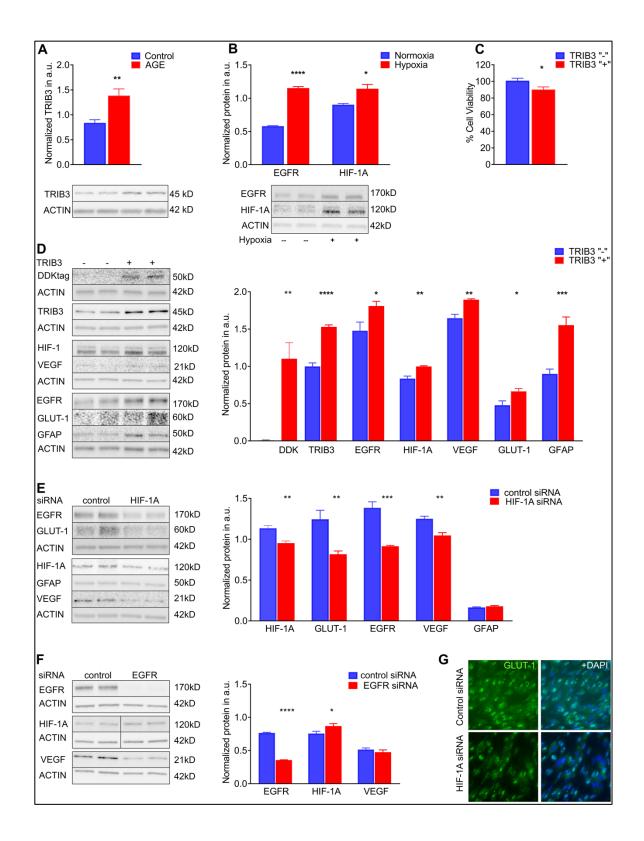
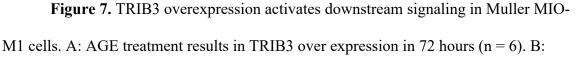


Figure 6. TRIB3 promotes retinal integrity loss and activates gliosis during hypoxia. A: TRIB3 KO significantly preserves the retinal integrity in hyperglycemic mice (n = 5). The IPL, INL, and OPL thicknesses are diminished in C57BL6 diabetic retinas, whereas protection is observed in TRIB3 KO diabetic retinas, demonstrating no difference compared to control retinas in both genetic groups. B: Representative images of the H&E-stained retinas in the four mouse groups. C: Representative images of the retinal sections probed with anti-GFAP (red) and anti-Vimentin (green) antibodies and DAPI (blue) in the four mouse groups taken with fluorescent microscopy. Merged images are shown on the left. (ONL: outer nuclear layer; INL: inner nuclear layer; RGC layer: retinal ganglion cell layer; Ph: photoreceptors). **(p<0.01); ****(p<0.0001). The scale is 100 μ m.





Hypoxia induces HIF1 and EGFR protein in addition to TRIB3 (Figure 1) (n = 4). C: Muller cells overexpressing TRIB3 demonstrate compromised cell viability (n = 4). D: Overexpression of TRIB3 in hypoxic Muller cells results in overexpression of EGFR1, HIF1 α , VEGF, GFAP, and GLUT1 (n = 4–5). E: Knockdown of *Hif1a* mRNA in hypoxic Muller cells results in downregulation of GLUT1, EGFR1, and VEGF1, whereas GFA expression does not respond to the treatment with siRNA (n = 5). F: Knockdown of *Egfr1* mRNA in hypoxic Muller cells results in downregulation of HIF1, whereas VEGF expression does not respond to the siRNA treatment (n = 6). G: Representative images of MIO-M1 cells transfected with control and *Hif1*a siRNA. Detection of GLUT1 (in green) was achieved performing IHC analysis and confocal microscopy. Staining with DAPI (in blue) was used to detect nuclei. *(p<0.05); **(p<0.01); ***(p<0.001).

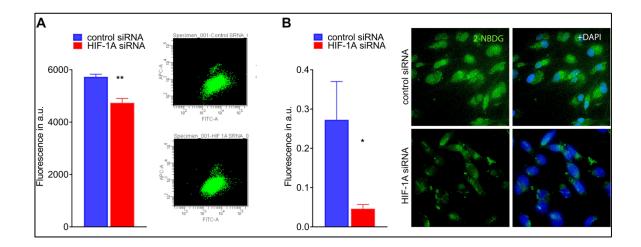


Figure 8. Treatment of hypoxic Muller cells with siRNA targeting *HIF1a* results in reduction of fluorescent signal from 2-NBDG cellular uptake. A: Reduction of fluorescent signal from 2-NBDG uptake measured by flow cytometry (n = 4). B: Reduction of fluorescent signal from 2-NBDG cellular uptake registered in fixed cultured MIO-M1 cells using microscopy (n = 3–4). The calculation of fluorescent signal was performed using the ImageJ program. *(p<0.05); **(p<0.01).

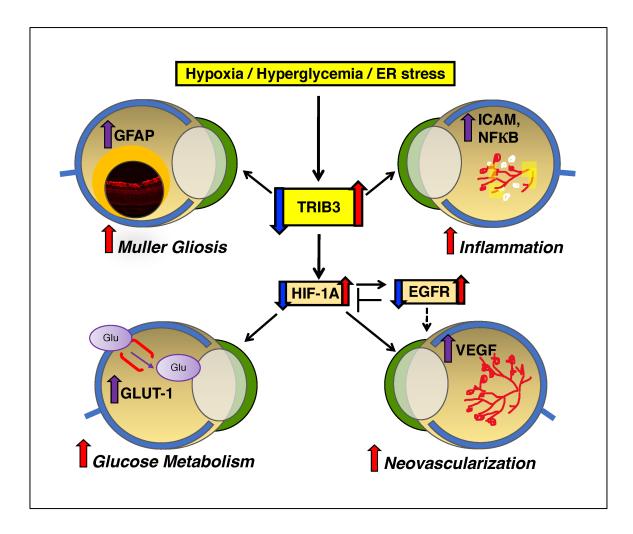


Figure 9. Schematic presentation of the proposed molecular mechanism of diabetic retinopathy controlled by TRIB3. Under hyperglycemic and hypoxic conditions, TRIB3 overexpression leads to upregulation of HIF1 α , EGFR, and GFAP. HIF1 α overexpression results in GLUT1 activation followed by increase in retinal glucose flux, overall affecting retinal metabolism. Additionally, HIF1 α mediates VEGF expression, which compromises vascular cell integrity and triggers angiogenesis. EGFR is upregulated as a result of TRIB3 upregulation as well. This protein reportedly induces

VEGF and cytokines leading to vascular dysfunction. TRIB3 promotes expression of GFAP and reactivation of gliosis in hypoxic retinas. Red and purple arrows indicate upregulation; blue arrows indicate downregulation signaling validated in our study, respectively. Solid lines represent data of the current study. Dashed lines denote the regulation proposed in the literature.

	Manufacturer Catalog Number	
C57BL6J Mice	Jackson Labs# 000664	
Streptozotocin (STZ)	Sigma# S0130	
Glucometer	CVS True Metrix# RE4007-01	
Hba1c Kit	PTS Diagnostics# 3021	
Glucose Estimation Assay Kit	Cell Biolabs, Inc# STA-680	
Trizol	Ambion# 15596	
cDNA Synthesis Kit	Biorad# 1708841	
4% PFA	EMS# 50-980-487	
OCT Compound	EMS# 62550-01	
H& E Kit	EMS# 26754-1A, 26762-01	
Periodic Acid Schiff Reagents	Sigma# 395B-1KT	
Elastase Solution	Calbiochem# 324682	
Lipopolysacchrides (LPS)- E. Coli	Sigma# L2880	
Seahorse XF DMEM Medium	AT# 103575-100	
Seahorse XF24 Islet Capture Microplate	AT# 101122-100	
Seahorse Microplate Insert Tool	AT# 101135-100	
Isolectin1B-4	Vector laboratories # FL-1201	
Anti- Vimentin (IHC)	Cell Signaling# 5741S	
Anti- TRIB3 Antibody (IHC)	Abcam# 73547	
Anti- TRIB3 Antibody (WB)	Santa Cruz# (B-2): sc-390242	
Anti- VEGF Antibody (WB-RETINA)	Santa Cruz # 7269 HRP	
Anti- VEGF Antibody (WB-MULLER CELL)	Invitrogen JH121# MA5-13182	
Anti- GFAP Antibody (IHC &WB)	Sigma# C9205	
Anti- B-Actin Antibody (WB)	Sigma# A2066	
Anti- CD 31/PECAM-1 (IHC)	Novus Biologicals# NB600-562	
Anti- IBA-1 Antibody (IHC)	Wako Chemicals# 019-19741	
Anti- BRN-3A Antibody (IHC)	Santa Cruz # 8429 AF488	
Anti- Glut-1 Antibody (WB& IHC)	Cell Signaling# Rabbit mAb12939	
Anti- HIF-1A Antibody (WB)	Cell Signaling# Rabbit mAb14179	
Anti- EGFR Antibody (WB)	Santa Cruz # sc-03-G	
Peroxidase AffiniPure Goat Anti-Mouse	JacksonImmunoresearch#115-035-206	
lgG, Fcγ subclass 2a specific (WB)		
HRP Goat Anti-Mouse (WB)	LICOR# 926-80010	
IR Goat Anti-Mouse (WB)	LICOR#926-32210	
Donkey anti-Rabbit IgG Secondary	Invitrogen Catalog # A32794	
Antibody, Alexa Fluor 555 (IHC)		
Donkey anti-Rabbit IgG Secondary	Invitrogen Catalog # A-21206	
Antibody, Alexa Fluor 488 (IHC) VECTASHIELD® Antifade Mounting	Venter leberaterient 11,1000,10	
Medium with DAPI	Vector laboratories# H-1200-10	
Taqman Primer- TRIB3	Mm00454879 m1	
Tagman Primer- AIF-1	Mm00479862_g1	
Taqman Primer- COX-2	Mm03294838_g1	
Taqman Primer- ICAM-1	Mm00516023 m1	
	Minio 66 16626_iiii	
Taqman Primer- HIF-1A	Mm00468869_m1	
Taqman Primer- NFKB-1	Mm00476361_m1	
Taqman Primer- RC3HI	Mm01284492_m1	
Taqman Primer- VEGF-A	Mm00437304_m1	
Taqman Primer- ZC3HI2A	Mm00462533_m1	
Taqman Primer- GAPDH	Mm999999915_g1	
Taqman Primer- GUSB	Mm00446953_m1	
Trib3 (BC012955) Mouse Tagged ORF	Origene- Product number MR204806	
Clone	VID (Drof Limb)	
MIO-M1 cells- Müller glia cell line	XIP (Prof Limb)	
AGE-BSA	Abcam# ab51995	
Oxoid™ AnaeroGen™ W-Zip Compact	Thermo Scientific™ AN0010W	
Gas Generator System MTT Solution	Sigma Aldrich# MESSE	
	Sigma-Aldrich# M5655	
HIF-1A siRNA EGFR siRNA	Thermofisher Scientific Assay ID# 42840 Thermofisher Scientific Assay ID# 646	
Neg. Control siRNA	Thermofisher Scientific# 4390844	
2-NBDG Glucose Uptake Assay Kit	BioVision# K68250	

Table: 1. Reagents, antibodies and primers in the study.

Table: 2. Blood and retinal glucose levels in the control and experimental mice.

Experimental groups of		50mg /kg STZ-	5 days dosage			
mice	Mean HbA1c levels (%HbA1c; mmol/mol)					
	15-weeks	P value	30-weeks	P value		
C57BL6 (n=9)	4.7; 28		4.7; 28			
C57BL6 STZ (n=9)	10.8; 95	<0.0001	11.7; 104_	<0.0001		
TRIB3KO (n=10)	5.3; 34		5.1; 32			
TRIB3KO STZ (n=13)	10.0; 86	<0.0001	10.2; 88	<0.0001		
	1		····			
Experimental groups of mice	150mg /kg STZ- single dosage					
mice	Mean Blood glucose levels (mg/dl)					
	4-weeks	P value	6-weeks	P value		
C57BL6 (n=8)	147.5		139.5			
C57BL6 STZ (n=12)	285	<0.0001	291.8	0.0006		
TRIB3KO (n=9)	184		138.5			
TRIB3KO STZ (n=12)	391.5	0.0205	351.0	0.0133		
			<u> </u>			
Experimental groups of		150mg /kg STZ-	single dosage			
mice	Mean Blood glucose levels		Mean Retinal glucose levels			
	(mg/dl)		(mg/dl)			
	4-weeks	P value	4-weeks	P value		
C57BL6 (n=9)	139.9		0.3			
C57BL6 STZ (n=9)	470.6	<0.0001	4.4	<0.0001		
TRIB3KO (n=11)	152.2		0.3			
TRIB3KO STZ (n=5)	434.4	NS	1.3	0.0009		

Chapter 4

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APPENDIX A

IACUC APPROVAL LETTER



MEMORANDUM

DATE: 03-Nov-2017

TO: Gorbatyuk, Marina

FROM:

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Robert A. Kesterson, Ph.D., Chair

Institutional Animal Care and Use Committee (IACUC)

SUBJECT: NOTICE OF APPROVAL

The following application was approved by the University of Alabama at Birmingham Institutional Animal Care and Use Committee (IACUC) on 03-Nov-2017.

Protocol PI:	Gorbatyuk, Marina
Title:	Unfolded Protein Response as a Therapeutic Target for ADRP Animal Models
Sponsor:	UAB DEPARTMENT
Animal Project Number (APN):	IACUC-09793

This institution has an Animal Welfare Assurance on file with the Office of Laboratory Animal Welfare (OLAW), is registered as a Research Facility with the USDA, and is accredited by the Association for Assessment and Accreditation of Laboratory Animal Care International (AAALAC).

This protocol is due for full review by 02-Nov-2020.

Institutional Animal Care and Use Committee (IACUC)		Mailing Address:
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933 19th Street South	Ι	1530 3rd Ave S
(205) 934-7692		Birmingham, AL 35294-0019
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