# Review Article

# Inflammatory mechanisms in diabetic retinopathy: pathogenic roles and therapeutic perspectives

Han-Ying Qian, Xiao-Hong Wei, Jin-Ou Huang

Department of Ophthalmology, Shengzhou People's Hospital (Shengzhou Branch of The First Affiliated Hospital of Zhejiang University School of Medicine), Shengzhou 312400, Zhejiang, China

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Abstract: Diabetic retinopathy (DR), a leading cause of global vision impairment, represents one of the most prevalent microvascular complications of diabetes. Numerous studies have confirmed that inflammatory processes and aberrant angiogenesis constitute pivotal pathological mechanisms in DR. Elevated levels of pro-inflammatory mediators - including cytokines, chemokines, and adhesion molecules - have been consistently detected in the serum, ocular fluids (aqueous humor and vitreous), retinal tissue, and tear film of DR patients, forming an intricate molecular network that drives disease progression. Importantly, modulation of these inflammatory components demonstrates potential to attenuate both vascular abnormalities and neurodegeneration in DR. This mechanistic understanding positions inflammation as a promising therapeutic target, highlighting the need for further investigation into anti-inflammatory strategies for DR management.

**Keywords:** Diabetic retinopathy, inflammation, pro-inflammatory mediators, pathological mechanisms, therapeutic target

#### Introduction

Diabetic retinopathy (DR), a prevalent microvascular complication of diabetes mellitus (DM), affects nearly 40% of diabetic patients and is the primary cause of blindness and visual impairment among adults aged under 50 years worldwide [1]. The global burden of DR is escalating in parallel with the rising prevalence of DM, with epidemiological projections estimating 129.84 million affected adults by 2030, potentially reaching 160.50 million by 2045 [2, 3]. Disease progression correlates strongly with diabetes duration, suboptimal glycemic control, concurrent hypertension, dyslipidemia, and inflammatory activation [4]. Clinically, DR manifests initially as non-proliferative diabetic retinopathy (NPDR), characterized primarily by retinal microaneurysms, which may progress to proliferative diabetic retinopathy (PDR), developing hard exudates, cotton-wool spots, pathological neovascularization, and retinal hemorrhages [5].

In the context of hyperglycemia, retinal microglial cells undergo pathological activation and

secrete a wide range of pro-inflammatory cytokines, including tumor necrosis factor-α (TNFα), interleukin-1beta (IL-1β), interleukin-3 (IL-3), interleukin-6 (IL-6), along with other inflammatory mediators such as vascular endothelial growth factor (VEGF), reactive oxygen species, glutamate, matrix metalloproteinases (MMPs), and nitric oxide (NO). This inflammatory cascade orchestrates multiple pathogenic processes through: (1) upregulation of adhesion molecules, (2) induction of apoptotic pathways, (3) facilitation of leukocyte infiltration, and (4) disruption of blood-retinal barrier (BRB) [6]. A substantial body of research has established the synergistic interplay between inflammatory and angiogenic pathways in DR pathogenesis [7-11]. Variations in the levels of various pro-inflammatory and angiogenic mediators have been detected in biological compartments such as serum, ocular fluids (aqueous humor and vitreous), retinal tissue, and tear film of DR patients. Importantly, targeted modulation of these mediators demonstrates therapeutic potential to mitigate both vascular abnormalities and neurodegeneration in DR [12-15], positioning inflammatory pathways as promising intervention targets. This review systematically examines the mechanistic contributions of inflammation to DR progression and evaluates emerging anti-inflammatory strategies with translational potential.

#### Inflammatory mechanisms in DR

Inflammation serves as a central pathogenic driver in DR, with inflammatory responses persisting throughout all disease stages. These processes fundamentally contribute to the two hallmark pathological features of DR-mediated visual impairment in DM: increased vascular permeability and pathological neovascularization, both in the retina [16-18]. DM induces the local or systemic upregulation of numerous inflammatory molecules implicated in DR progression, such as chemokines, cytokines, transcription factors, growth factors, and vascular adhesion molecules (VCAM-1) [19-21]. These elevated inflammatory mediators promote leukocyte activation, endothelial adhesion, and subsequent capillary sequestration [22, 23], leading to capillary occlusion, retinal ischemia, endothelial cell dysfunction, and BRB breakdown. Characteristic clinical manifestations include retinal edema, intraretinal hemorrhages, hard exudation, and microaneurysm formation [24, 25]. Concurrently, activated retinal glial cells amplify local inflammation through the secretion of pro-inflammatory mediators and the recruitment of immune cells to inflammatory foci [26, 27].

Increasing evidence suggests that retinal neurodegeneration serves as a significant pathophysiological mechanism in DR [28]. Retinal glial cells, including astrocytes, Müller cells, and microglial cells, play crucial roles in providing structural support and preserving retinal homeostasis [29, 30]. Importantly, dysfunction of these cells contributes significantly to the initiation and progression of retinal inflammation during early DR [31]. Chronic hyperglycemia potently activates retinal microglia [32]. In the early stages of DR, this activation initiates a cascade of neurodegenerative events, including neuronal apoptosis and progressive thinning of the nerve fiber layer, ultimately leading to measurable visual dysfunction and potential vision loss [33].

#### Inflammatory cytokines pathways in DR

Hyperglycemia induces retinal inflammation accompanied by elevated production and release

of multiple inflammatory mediators [34]. This inflammatory milieu stimulates a cascade of pro-inflammatory factors, including chemokines, inflammatory cytokines, and other related molecules [35], which collectively contribute to leukocyte stasis, cellular apoptosis, and retinal capillary leakage (**Table 1**). The key pathological cascades in DR development are illustrated in **Figure 1**.

#### Inflammatory cytokines

Elevated levels of macrophage-derived IL- $1\beta$  are consistently observed in both serum and vitreous samples from DR patients. Mechanistically, IL- $1\beta$  synergizes with TNF- $\alpha$  to: (1) upregulate endothelial adhesion molecules via NF- $\kappa$ B-mediated transcription, (2) enhance IL-6 and interleukin-8 (IL-8) production, (3) activate caspase-1-dependent inflammatory pathways [36]. Additionally, IL- $1\beta$  promotes: (1) accumulation of ROS through inflammatory cell recruitment, leading to mitochondrial dysfunction and retinal cell apoptosis [37], as well as (2) pathological angiogenesis via the mitogen-activated protein kinase (MAPK) cascade, promoting DR progression [38].

TNF- $\alpha$ , a pleiotropic cytokine, mediates multiple pathological processes in DR. First, it initiates leukocyte-endothelial adhesion and retinal vascular inflammation. Second, TNF- $\alpha$  activates NF- $\kappa$ B signaling pathways [39]. Third, TNF- $\alpha$  contributes to BRB disruption (demonstrated by TNF- $\alpha$ /epidermal growth factor receptor axis inhibition in diabetic mouse models) [40]. Additionally, TNF- $\alpha$  induces retinal microvascular cell loss [41].

Other inflammatory factors, such as IL-6 and IL-8, have been found to have significant elevation in the serum, vitreous, and aqueous humor, with concentrations correlating with DR severity [42, 43]. IL-6 plays significant pathological roles. It activates retinal glial cells, induces TNF- $\alpha$  secretion, and initiates early BRB breakdown [44]. IL-6 and IL-8 stimulate VEGF production in fibroblasts and monocytes, thereby promoting inflammation-associated neovascularization and diabetic macular edema (DME) progression [45, 46].

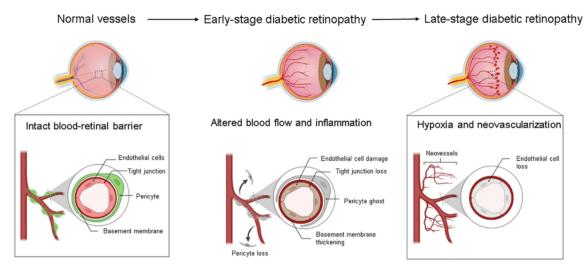
#### Adhesion molecules and integrins

The upregulation of adhesion molecules constitutes a critical step in DR pathogenesis. Key

Table 1. Inflammatory cytokines involved in DR

Category	Mediator	Mechanism	Stage of DR Involved	Therapy	References
Inflammatory Cytokines	IL-1β	Increases ICAM-1 expression, induces reactive oxygen species formation, iNOS expression, NF-kB, MAPK activation	Increased expression in NPDR and correlated with the severity of DR	Anakinra, Canakinumab, Gevokizumab	[36-38]
	TNF-α	Mediates leukocyte adhesion, activates NF-кВ	Increased expression in PDR and correlated with the severity of DR	Infliximab, Adalimumab, Golimumab, Triamcinolone acetonide, Fluocinolone, Dexamethasone	[39-41]
	IL-6	Increases vascular permeability, and disrupts the BRB	Increased expression in PDR, DME	Tocilizumab, Triamcinolone acetonide, Fluocinolone, Dexamethasone	[45, 46]
	IL-8	Promotes VEGF expression, induces angiogenesis	Increased expression in PDR, DME	Risuteganib	[47]
Adhesion Molecules	ICAM-1	Increases leukocyte adhesion, endothelial cell damage, capillary non-perfusion	Increased expression in NPDR and correlated with the severity of DR	Triamcinolone acetonide, Fluocinolone, Dexamethasone	[49-51]
	VCAM-1	Increases leukocyte adhesion and stasis, promotes inflammatory factor release	Increased expression in NPDR and correlated with the severity of DR	Triamcinolone acetonide, Fluocinolone, Dexamethasone	[49, 50]
Growth Factors	VEGF	Increases vascular permeability and neovascularization	Increased expression in PDR, DME	Ranibizumab, Aflibercept, Bevacizumab, Fariximab	[52-55]
	PIGF	Promotes angiogenesis	Increased expression in PDR	Faricimab, Conbercept, Aflibercept, Roveredimab, TB-403	[56-58]
	TGF-β	Regulates cell growth, differentiation, proliferation, and apoptosis	Increased expression in PDR	Faricimab, Aflibercept, Ranibizumab, Ozurdex®, Iluvien®	[59, 60]
Chemotactic Factors	MCP-1/CCL-2	Increases VEGF expression, enhances endothelial cell adhesion, promotes inflammatory damage	Increased expression in PDR	Ozurdex®, Iluvien®, Aflibercept, Faricimab	[61, 63]
	CXCL10	Immune stimulation	Increased expression in PDR	Ozurdex®, Iluvien®, Baricitinib, Faricimab	[60, 62, 63]
Others	iNOS	Oxidative stress, cytotoxicity	Increased expression in PDR	$\alpha\text{-lipoic}$ acid, $Ozurdex^{\oplus},$ dimethyl fumarate, Aflibercept	[64, 65]
	COX-2	Promotes neovascularization	Increased expression in DR	Diclofenac, Bromfenac, Celecoxib, Yutiq, Iluvien	[66, 67]
	MMP-9	Oxidative stress, promotes angiogenesis	Increased expression in DR	Ranibizumab, Aflibercept, Faricimab	[68]
	Ang-2	Promotes angiogenesis	Increased expression in PDR	Faricimab	[69]

Abbreviations: DR, diabetic retinopathy; IL-1β, interleukin-1beta; ICAM-1, intercellular adhesion molecule-1; iNOS, inducible nitric oxide synthase; NF-κB, nuclear factor κB; MAPK, mitogen-activated protein kinase; NPDR, non-proliferative diabetic retinopathy; TNF-α, tumor necrosis factor-α; PDR, proliferative diabetic retinopathy; IL-6, interleukin-6; BRB, blood-retinal barrier; DME, diabetic macular edema; IL-8, interleukin-8; VEGF, vascular endothelial growth factor; VCAM-1, vascular adhesion molecule-1; PIGF, placental growth factor; TGF-β, transforming growth factor-β; MCP-1/CCL-2, monocyte chemoattractant protein-1; CXCL10, interferon-Induced Protein 10; COX-2, cyclooxygenase-2; MMP-9, matrix Metalloproteinase-9; Ang-2, angiopoietin-2.



#### Mechanisms of the pathological cascade of diabetic retinopathy

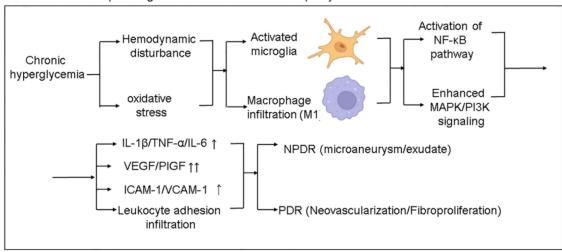


Figure 1. The mechanisms of pathological cascade of DR. Abbreviations: NF- $\kappa$ B, nuclear factor  $\kappa$ B; MAPK, mitogenactivated protein kinase; PI3K, phosphoinositide 3-kinase; IL-1 $\beta$ , interleukin-1beta; TNF- $\alpha$ , tumor necrosis factor- $\alpha$ ; IL-6, interleukin-6; VEGF, vascular endothelial growth factor; PIGF, placental growth factor; ICAM-1, intercellular adhesion molecule-1; VCAM-1, vascular adhesion molecule-1.

molecules include intercellular adhesion molecule-1 (ICAM-1) and VCAM-1, induced by hyperglycemia and oxidative stress during early DR. The elevated expression of these two molecules mediates leukocyte-endothelial interactions through integrins and facilitates multistep leukocyte recruitment: migration, aggregation, and adhesion, resulting in increased retinal capillary permeability and stimulated retinal neovascularization [47, 48]. Capillary occlusion can trigger the release of toxic substances (including nitric oxide synthase [NOS]) and inflammatory factors (e.g., IL-1, TNF-α), which exacerbate retinal ischemia and further stimulate ICAM-1 synthesis, forming a vicious cycle that accelerates DR progression [49].

#### Growth factors

VEGF serves as a potent inducer of retinal vascular permeability and angiogenesis, playing a crucial role in neovascularization in DR [50]. Under conditions of elevated glucose levels, oxidative stress, hemodynamic changes, and inflammatory mediators, VEGF expression is tightly regulated. VEGF enhances capillary permeability by phosphorylating tight junction proteins, leading to increased central subfield thickness, DME, and neovascularization [16]. Inflammatory mediators involved in DR pathogenesis can induce microvascular abnormalities, expand the avascular zone in the central fovea, and exacerbate macular ischemia [51].

These mediators further enhance VEGF expression through the upregulation of ICAM-1 in endothelial cells, which exacerbates the release of cytokines and promotes white blood cell stasis, thereby amplifying the inflammatory response. Additionally, VEGF, as a pro-inflammatory molecule, can stimulate the expression of various inflammatory cytokines such as monocyte chemoattractant protein-1 (MCP-1), macrophage inflammatory protein-1 $\alpha$ , and IL-8. Specific inhibition of VEGF has been shown to reduce the levels of ICAM-1, TNF- $\alpha$ , and NF- $\kappa$ B in diabetic mice, which alleviates retinal leukostasis, and mitigates disruption of the BRB [52].

Placental growth factor (PIGF), a member of the VEGF family, serves as a potent pro-angiogenic factor and plays a crucial role in modulating the inflammatory response. It achieves this by stimulating tissue factor production and promoting the growth, chemotaxis, and survival of monocyte/macrophage through binding to VEGF receptor-1 [20]. Clinical studies have demonstrated elevated levels of PIGF in the vitreous humor of PDR patients [53]. Conversely, the absence of PIGF in diabetic mice has been shown to prevent BRB disruption and inhibit retinal cell apoptosis [54].

Prolonged hyperglycemia in diabetic patients induces abnormal secretion of transforming growth factor-β (TGF-β) by retinal endothelial cells and macrophages. Elevated levels of TGF-β in the bloodstream then stimulate the migration of retinal pigment epithelial cells, fibroblasts, and glial cells into the vitreous humor and subretinal space, thereby promoting the development of PDR [55]. Moreover, the fragile nature of retinal neovascularization, which is prone to rupture and leakage, enables platelets and macrophages from the bloodstream to infiltrate the vitreous humor and retina. This infiltration triggers the release of substantial amounts of TGF-β, creating a vicious cycle that accelerates the formation and progression of PDR [56].

### Chemotactic factors

In inflammatory conditions, chemotactic factors such as Monocyte Chemoattractant Protein-1 (MCP-1, also known as CCL-2) and Interferon-Induced Protein 10 (CXCL10) are expressed at elevated levels in the vitreous

humor of PDR patients. Furthermore, the concentrations of these factors are positively correlated with the DR severity [57, 58].

MCP-1, through signaling pathways such as phosphoinositide 3-kinase (PI3K) and MAPK, upregulates the expression of NO and VEGF, thereby promoting neovascularization in the retina. Additionally, inflammatory factors such as IL-1β, TNF-α, INF-γ, and platelet-derived growth factor (PDGF) can induce and positively regulate the expression of MCP-1. When acting in concert, these factors contribute to inflammatory damage. MCP-1 can activate adhesion molecules, strengthen the adhesion between vascular endothelial cells, and trigger retinal inflammatory responses [59]. CXCL10 plays a crucial role in the development, migration, and adhesion of T cells, and it can also activate monocytes and natural killer cells, contributing to the onset and progression of various autoimmune disorders [60]. These chemotactic factors further promote the activation and recruitment of white blood cells, leading to the leakage of fluid and neutrophils from blood vessels into retinal tissues.

#### Other inflammatory mediators

Inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2) play significant roles in the inflammatory response during the progression of DR [61]. iNOS catalyzes the production of NO, which, under normal physiological conditions, is essential for regulating vasodilation and inhibiting platelet aggregation. However, under high glucose conditions, the overexpression of iNOS leads to excessive NO production. The reaction between NO and superoxide anions generates peroxynitrite, a potent oxidant that induces oxidative stress, impairing retinal cell function and triggering apoptosis [62]. Furthermore, NO enhances vascular permeability and promotes neovascularization through the activation of the cyclic guanosine monophosphate signaling pathway, thereby contributing to the pathological processes of DR [63].

COX-2, as a key enzyme in the inflammatory response, triggers the release of the pro-inflammatory cytokine prostaglandin E2 in the early phases of DR. This, in turn, leads to increased VEGF expression and promotes the formation

of neovascularization in the retina [64]. Research by Wu et al. [65] has shown that salidroside can directly bind to COX-2, inhibiting its expression and thereby mitigating the inflammatory response in ultraviolet-induced mouse epidermal JB6Cl41 cells and human skin keratinocytes.

Matrix metalloproteinase-9 (MMP-9) plays a dual role in DR progression by degrading extracellular matrix components and inducing oxidative stress under hyperglycemic conditions. Additionally, MMP-9 interaction with CD44 triggers TGF-β release, which contributes to neovascularization in PDR [66]. The balance between the angiopoietin (Ang) family members (Ang-1 and Ang-2) and the Ang/tyrosine kinase receptor 2 system determines vascular formation and stability. Clinical evidence demonstrates significantly elevated serum Ang-2 levels in PDR patients compared to NPDR cases. This Ang-2 upregulation drives the formation of immature and leak-prone neovessels, exacerbating retinal edema and inflammatory responses [67].

#### Role of inflammatory cells in DR

Retinal cell dysfunction may exacerbate inflammation in the early phases of DR. Key pathological changes, including microglial activation, Müller cell proliferation, retinal pigment epithelium secretion, endothelial cell proliferation, and pericyte loss, are associated with BRB breakdown and DME progression [68, 69]. The involvement of specific inflammatory cells in DR pathogenesis is shown in **Table 2**.

#### Leukocytes

Leukocyte adhesion and aggregation to the vascular endothelium represent critical early events in retinal inflammation, contributing to microcirculatory dysfunction and BRB breakdown in DR [70]. This process is mediated by mutual recognition between endothelial cells and leukocyte surface adhesion molecules. Under inflammatory conditions, VEGF and ICAM-1 expression in retinal endothelial cells is upregulated. ICAM-1 binds to its receptors, facilitating leukocyte transendothelial migration and subsequent adhesion. VEGF can induce the expression of endothelial NOS, which further promotes leukocyte adhesion.

#### Macrophages

Several studies suggest that macrophages are a primary driving force in PDR pathogenesis [71, 72]. In the early stages of DR, macrophages can polarize into M1 phenotype, exacerbating retinal vascular endothelial damage by secreting inflammatory factors. This increases vascular permeability, disrupts the BRB, and leads to retinal leakage, hemorrhage, microaneurysm formation, and other pathological changes such as vascular occlusion [73]. In the late stages of DR, M2 macrophages become progressively activated, inducing chemotaxis and fibrosis through the secretion of leukotrienes and fibronectin. They also influence cell proliferation by synthesizing VEGF and PDGF, promoting neovascularization and fibrovascular membrane formation [74]. Studies have shown that both in vitro and in vivo, human monocytes and macrophages are activated, with elevated levels of CD11c and iNOS in macrophages treated with high glucose, as well as in circulating monocytes from DR patients [75]. Additionally, activated macrophages release excessive inflammatory factors, including TNF-α, IL-1β, IL-6, and IL-12, through the NF-κB signaling pathway. However, their phagocytic function is impaired, suggesting that macrophage dysfunction may exacerbate DR inflammation [76, 77].

## Astrocytes

Astrocytes are the primary source of VEGF and inflammatory factors in the retina [78]. In DR, retinal astrocytes are closely associated with retinal blood vessels and play a crucial role in maintaining the BRB [79]. Additionally, astrocytes contribute to the inflammatory response by producing pro-inflammatory cytokines such as IL-1 $\beta$ , TNF- $\alpha$ , and iNOS, which exacerbate damage to the retinal neurovascular unit and promote neuroinflammation [80].

#### Müller cells

Müller cells, the primary glial cells of the retina, provide support and nourishment to retinal neurons. They play a crucial role in maintaining retinal integrity and are involved in various pathological processes [81]. In DR, the expression of glial fibrillary acidic protein (GFAP) is increased [82], and the secretion of pro-inflammatory mediators such as IL-1β, IL-6, IL-8, VEGF,

Table 2. The role of inflammatory cells in DR

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Category	Mediator	Mechanism	Stage of DR Involved	Therapy	References
Leukocyte	VEGF/ICAM-1	adhesion, aggregation, and sedimentation	causing BRB damage in early DR	Yutiq, Iluvien, Trivaris, Triesence, Ozurdex, Aflibercept, Faricimab, Ranibizumab	[70]
Macrophage	M1/M2 type	Early destruction of BRB, late promotion of neovascularization and fibrovascular membrane formation, exacerbation of DR inflammation	Increased expression in NPDR and correlated with the severity of DR	Yutiq, Iluvien, Ozurdex, INCB-8765, PLX3397, BLZ945, Pioglitazone, Glycyrrhizin	[73, 74]
Astrocyte	VEGF	Increases vascular permeability and neovascularization	Increased expression in PDR, DME	Ozurdex®, Iluvien®, Aflibercept, Faricimab	[78, 79]
Müller cells	GFAP/VEGF/ICAM-1	It is the main component of retinal astrocytes, providing support and nutrition for retinal neurons	Participate in the inflammatory process of DR	Aflibercept, Faricimab, Ranibizumab, Ozurdex®, Iluvien®, N-acetylcysteine, Epalrestat	[81-85]

Abbreviations: DR, diabetic retinopathy; VEGF, vascular endothelial growth factor; ICAM-1, intercellular adhesion molecule-1; BRB, blood-retinal barrier; NPDR, non-proliferative diabetic retinopathy; GFAP, glial fibrillary acidic protein; DME, diabetic macular edema.

and ICAM-1 is elevated. This suggests that, during the progression of diabetes, Müller cell proliferation and reactive gliosis contribute to the inflammatory response [83-85].

## Therapeutic approaches for DR inflammation

Given the critical role of inflammation in the pathogenesis of DR, targeting or inhibiting inflammation is considered a potential treatment strategy. Glucocorticoids exhibit potent antiinflammatory effects, by suppressing the expression of various inflammatory mediators, reducing retinal edema, and alleviating inflammatory responses. Local application of glucocorticoids can effectively treat vision-threatening DR [86]. Studies [87] have demonstrated that long-acting water-soluble steroid dexamethasone reduces VEGF expression. Further research indicates that dexamethasone reduces VEGF levels by inhibiting p38 MAPK activity, extracellular signal-regulated kinase activity and MMP-9 expression, thereby reducing neovascularization [88].

Anti-VEGF therapy has emerged as a novel and widely adopted approach in clinical practice in recent years. Currently available anti-VEGF drugs include Ranibizumab, Bevacizumab, Conbercept, and Aflibercept, with Conbercept injections as the primary treatment. Conbercept, a VEGF multi-target recombinant fusion protein, effectively blocks the VEGF receptor signaling pathway, thereby reducing retinal edema, promoting the absorption of hemorrhages, decreasing retinal vascular permeability, alleviating inner retinal hypoxia and ischemia, and reducing retinal nerve fiber layer thickness. These effects collectively inhibit neovascularization and retinal microvascular leakage, facilitating visual recovery and enhancing treatment efficacy [89, 90]. Study have demonstrated that 20 weeks of Conbercept intervention in diabetic mice inhibits the NF-κB signaling pathway, reduces the expression of ICAM-1, IL-1 $\beta$ , IL-6, and TNF- $\alpha$ , and improves PDR [91].

Nonsteroidal anti-inflammatory drugs achieve their anti-inflammatory effects by inhibiting COX activity and reducing the formation of prostaglandins. Sodium bromfenac, as a new generation NSAID, is highly effective and safe for anti-inflammatory and analgesic purposes, primarily targeting COX-2 [92]. In a preliminary

study, local application of Sodium bromfenac significantly reduced central foveal retinal thickness in DME patients, although it did not significantly improve vision [93]. Nepafenac is a prodrug that inhibits COX-1 and COX-2 activity through its active metabolite Amfenac [94]. Animal experiments have shown that topical Nepafenac significantly inhibits pro-inflammatory cytokine prostaglandin E2 content in corneal tissues, improves corneal edema symptoms, and decreases corneal neovascularization [95].

Moreover, numerous new anti-inflammatory targets are being developed to address key molecules in the inflammatory pathway, such as TNF- $\alpha$ , IL-1 $\beta$ , and IL-6, representing an emerging area of research in DR treatment. TNF- $\alpha$  inhibitors, including Infliximab, Adalimumab, and Golimumab [96], have demonstrated some degree of improvement in vision and retinal edema in patients [97]. Canakinumab, an IL-1 $\beta$  inhibitor, reduces DME and stabilizes the condition, aiding in the treatment of PDR without affecting neovascular formation [98]. IL-6 inhibitors such as Siltuximab, Sirukumab, Olokizumab, and Clazakizumab can globally block IL-6 signal transduction [99].

#### Conclusion and future perspectives

DR represents a major microvascular complication of diabetes, characterized by complex inflammatory pathogenesis involving multiple interrelated pathways rather than isolated factors. This review elucidates the mechanistic roles of inflammatory cells, cytokines, and associated signaling pathways in DR progression, highlighting that targeted modulation of intra- and intercellular inflammatory signaling may offer promising therapeutic strategies. While substantial advances have been achieved in understanding inflammatory mechanisms in DR, current research remains constrained by several limitations. Most therapeutic studies rely predominantly on animal models and in vitro systems, with a notable paucity of largescale clinical trials to validate long-term efficacy and safety in human populations. In the future, the mechanism of action of multi-target inflammatory factors should be further elucidated, and research into anti-inflammatory mechanisms should be enhanced. Additionally, the development of new formulations for the anti-inflammatory treatment of DR is critical for advancing clinical treatment options.

#### Disclosure of conflict of interest

None.

Address correspondence to: Jin-Ou Huang, Department of Ophthalmology, Shengzhou People's Hospital (Shengzhou Branch of The First Affiliated Hospital of Zhejiang University School of Medicine), No. 666 Dangui Road, Chengnan New District, Shengzhou 312400, Zhejiang, China. E-mail: 13605855780@163.com

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