Diabetes-related adduct formation and retinopathy

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Abstract The pathogenesis of diabetic retinopathy is complex, reflecting the array of systemic and tissue-specific metabolic abnormalities. A range of pathogenic pathways are directly linked to hyperglycaemia and dyslipidaemia, and the retina appears to be exquisitely sensitive to damage. Establishing the biochemical and molecular basis for this pathology remains an important research focus. This review concentrates on the formation of a range of protein adducts that form after exposure to modifying intermediates known to be elevated during diabetes. These so-called advanced glycation end products (AGEs) and advanced lipoxidation end products (ALEs) are thought to play an important role in the initiation and progression of diabetic retinopathy, and mechanisms leading to dysfunction and death of various retinal cells are becoming understood. Perspective is provided on AGE/ALE formation in the retina and the impact that such adducts have on retinal cell function. There will be emphasis placed on the role of the receptor for AGEs and how this may modulate retinal pathology, especially in relation to oxidative stress and inflammation. The review will conclude by discussion of strategies to inhibit AGE/ALE formation or harmful receptor interactions in order to prevent disease progression from the point of diabetes diagnosis to sight-threatening proliferative diabetic retinopathy and diabetic macular oedema.

Keywords Diabetic retinopathy · Advanced glycation · Lipoxidation

A. W. Stitt (⊠) • T. M. Curtis Centre for Vision and Vascular Science, Queen's University Belfast, Royal Victoria Hospital, Grosvenor Road, Belfast BT12 6BA Northern Ireland, UK e-mail: a.stitt@qub.ac.uk Diabetic retinopathy is a leading cause of blindness among people of working age in developed countries [2]. Despite its importance, there are currently few effective means to prevent or treat the disease. Beyond maintenance of tight glycaemic control, laser photocoagulation remains the principal therapy for sight-threatening diabetic retinopathy, but this is always at the expense of functional retina and visual performance [26]. With diabetes rapidly emerging as a global health care problem [78], there remains a genuine and urgent need to develop new, effective therapies for diabetic retinopathy that may be used to augment strict glucose management.

Most studies concerned with the pathogenesis of diabetic retinopathy have focused on alterations in the functional integrity of the intra-retinal blood vessels. The vasodegenerative phase of diabetic retinopathy is characterized by a breakdown of the blood-retinal barrier, a thickening of capillary basement membranes, pericyte and microvascular smooth muscle dropout, microaneurysms and capillary closure [26]. Persistent vascular leakage leads to macular oedema with or without cystoid degenerative changes, photoreceptor atrophy and an irreversible loss of central vision. In the proliferative phase of the disease, the ischaemic retina begins to secrete various growth factors, including vascular endothelial growth factor (VEGF), which stimulates neovascularisation. The new vessels that form during proliferative diabetic retinopathy are unable to replace the flow of necessary nutrients and instead are associated with an increased risk of severe visual loss through vitreal haemorrhage, retinal fibrogliosis and tractional retinal detachment.

While there is no argument that the retinal vasculature is central to the development of diabetic retinopathy, there is accumulating evidence that neuroretinal function is also compromised during this disease [4]. For instance, deficits in visual functioning, such as loss of colour vision [59], contrast sensitivity [63] and abnormalities in the electroretinogram [84] have been documented in patients shortly after the diagnosis of diabetes and before detection of clinically evident vascular retinopathy. Early neuronal changes are also apparent in the retinas of experimental rodent models of diabetes, including neurophysiological deficits similar to those described in human diabetes [57]. Because neuroretinal alterations occur at an early stage of the disease process, it has been proposed that they may a play a causative or contributory role in the initiation and progression of the vascular pathology associated with diabetic retinopathy [4].

Epidemiological signposts for AGE/ALE involvement in diabetic retinopathy

The Diabetes Control and Complications Trial (DCCT) was a major epidemiological study conducted from 1983 to 1993 in type 1 diabetic patients and established the relationship between hyperglycaemia and progression of retinopathy [19]. A similar long-term evaluation of type 2 diabetic patients called the UK Prospective Diabetes Study also highlighted hyperglycaemia as being critical for pathogenesis although it indicated that this occurs in unison with dyslipidaemia and hypertension [1]. Such population-based evidence and a wealth of follow-up clinical data provide a strong basis for current, ongoing research which is seeking to identify the cellular and molecular mechanisms that underpin diabetic retinopathy. Although the formation of advanced glycation end products (AGEs) and advanced lipoxidation end products (ALEs) and activation of receptors for AGEs in the diabetic retina is the focus for this review, hyperglycaemia can evoke many other important pathogenic pathways in the retina, so each should not necessarily be viewed as an independent phenomenon. Indeed, many of these share common biochemistry (such as free radical formation) and evoke common pathological events in various retinal cells. The so-called "unifying concept" whereby hyperglycaemia increases superoxide production (via the mitochondrial electron transport chain) links interrelated pathogenic responses [10]. This hypothesis suggests that enhanced flux through the hexosamine pathway. diacylglycerol-mediated activation of PKC-B and intracellular AGE formation can coalesce to cause cell damage. Evidence for this concept and its involvement in diabetic retinopathy has been provided through the use of the vitamin B₁ thiamine derivative, benfotiamine, which stimulates transketolase activity and shunts excess triose phosphates toward the reductive pentose phosphate pathway which is impaired in high glucose/diabetes [70]. Importantly, benfotiamine can prevent AGE formation and retinal capillary degeneration in diabetic animals [35].

Biochemistry of AGE and ALE formation

Reactions between sugars and free amino groups on proteins, lipids and DNA are an inevitable consequence of aldehyde reactivity, and many molecules in our body manifest chemically attached carbohydrate. These reactions begin with the formation of Schiff bases and ε -amino groups that rearrange to Amadori adducts. These intermediates can undergo further oxidation and dehydration reactions to form irreversible protein-bound compounds collectively termed "AGEs". During diabetes even modest hyperglycaemia can result in significant adduct accumulation on long-lived macromolecules [50].

Glucose is much less reactive than α -oxaloaldehydes which can form AGEs much more rapidly [72]. For example, glyoxal (GO) reacts with arginine residues to form carboxymethyl–arginine [28] while methylglyoxal (MGO) is a precursor for the AGEs N ε -(carboxyethyl)lysine and argininehydroimidazolone [71]. The concentrations of these reactive carbonyls rise in high glucose-exposed cells, occur at elevated levels in diabetic serum and are important precursors for AGEs in the body [40]. Other AGE adducts that form in the body are N $^{\varepsilon}$ -(carboxy-methyl) lysine (CML), crossline, pentosidine, furoyl-furanyl imidazole, hydroimidazolone, argpyrimidine, glyoxal lysine dimer and methylglyoxal lysine dimer [72].

It is important to appreciate that proteins can be modified by lipids as well as carbohydrates. During oxidative stress, reactive oxygen species attack polyunsaturated fatty acids (PUFAs) either in the cell membrane or circulating lipoprotein molecules [3]. This oxidative decomposition of PUFAs initiates chain reactions that lead to the formation of a variety of reactive carbonyl species. Among them are 4-hydroxy-trans-2-nonenal (4-HNE), acrolein (ACR), malondialdehyde (MDA), 4-hydroxy-2-hexenal (4-HHE) and crotonaldehyde [75] (Fig. 1). These compounds subsequently react by the socalled Michael addition mechanism with cysteine, histidine and lysine residues in proteins, generating relatively stable adducts known as ALEs. ALEs chemically characterised to date include N^{ε}-(2-propenal)lysine and dihydropyridine-type adducts (DHP-lysine, MDA-derived), hemiacetal and pyrrole adducts (HNE- and HHE-derived) and β-substituted propanal adducts and N^{ε} -(3-formyl-3,4-dehydropiperidino)lysine (FDP-lysine; ACR-derived) [3] (Fig. 1). In addition to their formation through lipid peroxidation, ALEs can also be formed endogenously by carbohydrate or ascorbate autoxidation, amine-oxidases, cytochrome P450 enzymes and the myeloperoxidase system of neutrophils [54]. While ALEs are often grouped together with AGEs, they should be regarded as a distinct class of adducts due to their differing reaction chemistry. Lipid peroxidation is enhanced in diabetes, and diabetic patients have higher serum levels of MDA than non-diabetic subjects [30]. It has also been reported by our

A. Lipid-derived reactive carbonyls

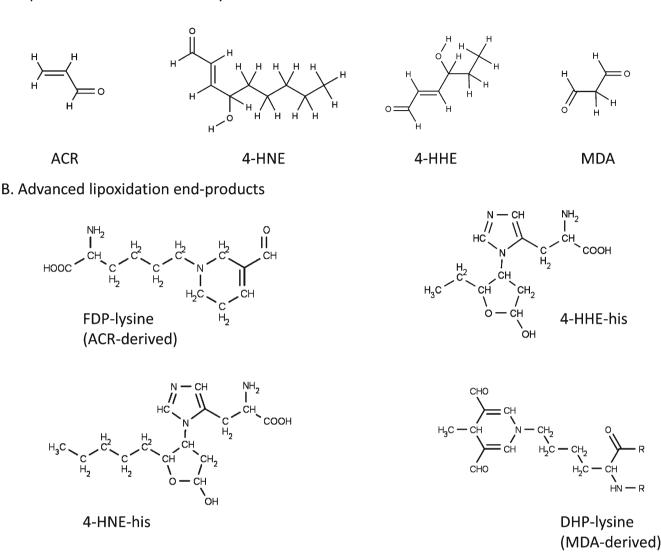


Fig. 1 The structures of lipoxidation-derived reactive carbonyl species and the principal advanced lipoxidation adducts formed following their reaction with proteins

own group and others that FDP-lysine- and HNE-modified proteins accumulate in serum of patients with diabetes compared to control individuals [74, 88]. At present, however, our understanding of the role of ALEs in the pathogenesis of diabetic complications, particularly retinopathy, lags far behind that which is currently known about AGEs.

Most AGE/ALE adducts are highly stable at physiological pH, and their rate of accumulation in tissues depends on factors such as availability of metal ions, redox balances and longevity of the modified protein. Unlike other AGEs/ ALEs, FDP-lysine is not a stable end product but a reactive intermediate that covalently binds to thiols, thereby exacerbating oxidative stress via depletion of the endogenous antioxidant glutathione [24]. Irrespective of their chemical derivation, AGE/ALE modifications can have a significant impact on protein structure and function by mediating protein–protein cross-linking reactions, changing tertiary structure and normal molecular function, conferring resistance to digestion or impairing receptor recognition.

Natural defence against AGE/ALEs

Cells have evolved natural defence mechanisms to protect against reactive carbonyl species. The capacity of cells to metabolise and detoxify carbonyls is dependent not only on the carbonyl but also on the cellular content of carbonylmetabolising enzymes [54]. In general, aldehyde precursors of AGEs/ALEs are detoxified through oxidation reactions mediated by nicotinamide adenine dinucleotide (NAD+) and enzymes of the aldehyde dehydrogenase superfamily (ALDH). At least 19 ALDH isozymes have been identified localized to several cellular compartments and displaying

varying affinities to a wide range of aldehyde substrates [90]. In addition to ALDH, other enzymes with high selectivity towards specific aldehydes have also been shown to play a detoxification role. For example, the glyoxalase complex (formed from glyoxalase I (GLO1) and glyoxalase II (GLO2) components) is an effective detoxification system for GO and MGO [42]. This has been shown in endothelial cells transfected to overexpress GLO1 which accumulate less MGO and subsequently contain less AGEs [62, 83]. Nematode worms engineered to overexpress GLO1 also contain fewer MGO-derived AGE adducts, and the worms demonstrate increased life-span [52, 61]. More recently it has been shown that overexpression of GLO1 in diabetic rats is significantly protective against key retinopathy lesions including Müller glia damage and formation of acellular capillaries [91].

AGES/ALES in diabetic retinopathy

Clinical correlates with diabetic retinopathy

In recent years, numerous studies have examined the correlation between serum levels of AGEs and the severity of retinopathy in patients with diabetes although the data have often been confusing and somewhat contradictory. For example, it has been reported that serum AGEs associate with progression of diabetic retinopathy in patients [55]. Conversely, other studies have demonstrated no significant correlation between AGE levels and retinopathy [69, 77]. Many of these reports are ELISA based using non-adduct-specific antibodies. Beyond this, investigation of defined AGEs such as CML, pentosidine or hydroimidazolone shows correlation of these adducts with retinopathy in diabetic patients [23, 69, 81].

From the ALE perspective, comparatively few studies have been undertaken to examine the association of these lipid-derived adducts and diabetic retinopathy. One study by Losado et al. reported that concentrations of the ALE precursor, MDA, are elevated in type 1 patients with retinopathy when compared to diabetics without retinopathy and healthy controls [45]. More recently, we have demonstrated that haemoglobin levels of the ACR-derived ALE, FDPlysine, are associated with the severity of diabetic retinopathy in type 1 and type 2 diabetic patients [88]. Importantly, the relationship of FDP-lysine with retinopathy was unaltered after adjustment for HbA1c, or other clinical parameters. Thus, haemoglobin FDP-lysine could prove to be a very useful marker for stratifying patients at high risk of retinopathy not evident from simply measuring their longterm glycaemic control.

A number of studies have used skin collagen to investigate the correlations between AGEs and the severity of retinopathy in patient-based studies [49]. The most comprehensive work to date, conducted by the DCCT skin collagen ancillary study group, demonstrated a strong association between the glycooxidation product, CML, and the progression of diabetic retinopathy [48]. The Epidemiology of Diabetes Interventions and Complications trial further followed these patients and demonstrated that retinopathy was less frequent in the group initially maintained under "tight" glycaemic control. Importantly, these benefits extended far beyond the period of intensive insulin therapy [27]. The patients under "conventional" control for the first 10 years maintained a "hyperglycaemic memory" and showed considerable retinopathy progression. The Genuth and Monnier groups have shown that CMLmodified skin collagen predicted the progression of retinopathy (and nephropathy) even after initiation of intensive insulin therapy [27]. Furthermore, they suggest that AGE accumulation on skin collagen is a robust biomarker for retinopathy risk and could offer a molecular-based explanation for hyperglycaemic memory.

AGEs/ALEs in the diabetic retina

AGEs have been extensively quantified in various ocular tissues and are often elevated during ageing and in diabetics when compared to non-diabetic controls [64]. This includes vitreous collagen [68], where the AGE levels correlate with diabetic retinopathy [22]. In the diabetic retina, AGEs and/ or late Amadori products have been localized to vascular cells, neurons and glia [25, 33, 34, 53, 60, 66]. This would be expected to have pathogenic implications for the individual cells and retinal function. Although differential accumulation of AGEs exists in the retina over the course of life, diabetes significantly enhances the occurrence of these adducts in the vascular and neural tissue components [66].

Recent studies by our group have focused on the distribution pattern of ALEs in the diabetic retina [85]. During early experimental diabetes, there is a selective accumulation of FDP-lysine in Müller glia (Fig. 2), whereas no significant differences in the levels of MDA-, 4-HNE- and 4-HHE-derived ALEs were apparent between control and diabetic retinas. Spatiotemporal studies revealed that FDPlysine initially accumulated within the Müller glia end feet after only a few months of diabetes and thereafter spread distally throughout their inner radial processes. The observation that FDP-lysine selectively accumulates in Müller glia during diabetes is intriguing, and the underlying pathobiochemical mechanisms warrant further investigation. In this regard, since other ALE adducts do not appear to be elevated in diabetic Müller glia, it seems unlikely that the accumulation of FDP-lysine can simply be attributed to an increase in the levels of lipid peroxidation. A more likely explanation may relate to the fact that Müller glia are known to function as the main polyamine storage cells of the retina

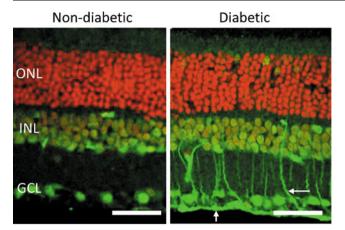


Fig. 2 FDP-lysine immunoreactivity (*green channel*) in retinal sections from non-diabetic and diabetic rats of 4-month disease duration. Cell nuclei are stained *red* by propidium iodide. In diabetes, prominent immunolabelling appeared in Müller glia end-feet and radial processes (arrows). *GCL* ganglion cell layer, *INL* inner nuclear layer, *ONL* outer nuclear layer. *Scale bars*=50µm

[7]. ACR, the precursor for FDP-lysine, can be formed from spermine by spermine oxidase and spermidine by spermidine/ spermine N1-acetyltransferase and acetylpolyamine oxidase [73]. As such, the selective accumulation of FDP-lysine in Müller glia could reflect changes in polyamine catabolism in the diabetic retina, a possibility that is currently being addressed by ongoing studies within our laboratory.

AGEs/ALEs evoke retinal cell dysfunction and death

Retinal capillary degeneration remains a hallmark of retinopathy in diabetic animal models and patients [26]. Retinal capillaries appear to be important targets for AGE-induced toxicity. For example, AGEs induce toxic effects on retinal pericytes by inducing oxidative stress and subsequent apoptosis [16]. In addition, some studies have indicated that AGEs cause osteoblastic differentiation and calcification in retinal pericytes by the activation of alkaline phosphatases [80]. Pericytes growing on AGE-modified basement membrane show acute attenuation of endothelin-1 (ETA receptor mediated) contraction, suggesting that AGE-cross links in a surrounding matrix significantly influence pericyte physiology [36]. Indeed, longer exposure times to these substrate-AGEs induce loss of integrin signalling and apoptosis [44]. The GLO1 detoxification system is also critical for retinal pericyte survival, but this may be insufficient during diabetes since these cells undergo apoptosis as a direct result of MGO-derived AGE formation [47]. Strategies to enhance carbonyl detoxification in retinal cells could be an important future therapeutic strategy.

Retinal microvascular endothelial cells show proangiogenic responses to AGEs at lower concentrations by the involvement of MAPK, PKC and NF-κB signalling pathways [46], although at higher concentrations, these adducts are toxic to endothelial cells [17] and in vivo may eventually lead to enhanced microvascular closure [67]. Under hyperglycaemic conditions, retinal microvascular endothelial cells accumulate MGO and MGO-derived AGE adducts (such as hydroimidazolone and argpyrimidine) which in vivo contribute to premature closure of capillaries [56]. AGEs cause upregulation of ICAM which mediates retinal capillary leukocyte adherence and inner blood retinal barrier breakdown [51]. Independent of the complexities of the diabetic milieu, non-diabetic mice exposed to "diabeticlike" levels of injected AGE-albumin show increased retinal expression of VEGF concomitant with blood retinal barrier dysfunction [15]. Similar treatments may cause loss of pericytes [79], and taken together, this suggests that high serum levels of AGE-modified proteins (as particularly evident in diabetic patients with renal dysfunction) induce lesions that are comparable to those occurring during diabetic retinopathy.

Whilst the detrimental effects of AGEs/ALEs on retinal vascular cells are well recognised, their potential role in mediating neuroretinal changes in diabetes has only recently begun to be explored. In retinal explants obtained from nondiabetic animals, incubation with AGE-modified albumin directly triggers neural cell apoptosis through a caspasedependent pathway [43]. To better understand the possible pathogenic effects of FDP-lysine in Müller glia, we have recently developed an experimental in vitro model of FDPlysine accumulation in human Müller cells [85]. Accumulation of FDP-lysine causes Müller glia dysfunction consistent with that observed in the diabetic retina, including the induction of oxidative stress, upregulation of pro-inflammatory cytokines and VEGF, and dysregulation of K⁺ channel protein expression [85]. Diabetes is one of the few pathological conditions where Müller cell apoptosis has been shown to occur [9]. Exposure of Müller cells to high concentrations of FDPlysine causes extensive cell death (Fig. 3) through apoptosis induction [85], suggesting that FDP-lysine accumulation could be a major factor contributing to Müller cell death during long-term diabetes.

AGE/ALE inhibition and prevention of retinopathy

The first AGE inhibitor was aminoguanidine (Pimagedine) [11]. This small nucleophilic hydrazine drug was shown to prevent AGE formation and attenuate diabetic retinopathy in animal models [25, 32, 38]. In a subsequent clinical trial, aminoguanidine showed positive signs towards slowing the progression of retinopathy, but it ultimately failed [8]. This hydrazine is not a specific inhibitor of AGE formation and is also a potent iNOS inhibitor [58]. A class of drugs called "Amadorins" have an ability to scavenge reactive carbonyls and prevent conversion of Amadori intermediates to AGEs

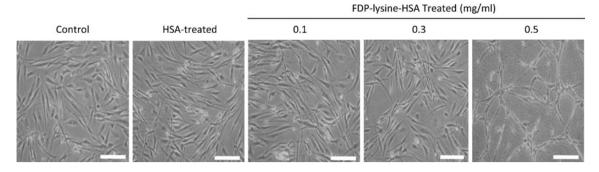


Fig. 3 Phase-contrast micrographs of cultured human MIO-M1 Müller glia exposed to increasing concentrations of FDP-lysine-modified human serum albumin (FDP-lysine-HSA). High concentrations of FDP-lysine-HSA (0.5 mg/mL) induce cell death. Scale bars = $100 \mu m$

and ALEs [39]. The derivative of vitamin B_6 , pyridoxamine (PyridorinTM), is an efficacious and specific post-Amadori inhibitor [76] which reduces retinal AGE accumulation and also attenuates capillary acellularity in the diabetic rat retina [65]. Likewise, a drug called LR-90 has been developed as an effective multi-stage inhibitor of both AGE/ALE formation and prevents diabetic retinopathy [6].

It is noteworthy that drugs such as aminoguanidine and pyridoxamine which have been used widely to prevent advanced glycation reactions in diabetes are relatively poor scavengers of ACR [13]. Indeed, we have found that pyridoxamine is only moderately effective in preventing FDPlysine accumulation in the diabetic retina, although it is still capable of attenuating Müller glial dysfunction and lesion formation [18]. On the other hand, hydrazino compounds such as hydralazine and dihydralazine react readily with ACR and have also been found to target protein-bound FDP-lysine adducts [12, 13]. Hydralazine and dihydralazine have been approved clinically as antihypertensive drugs for many years, but these cardiovascular actions complicate their potential use for the treatment of diabetic retinopathy. Since the hydrazine groups on hydralazine and dihydralazine are known to be responsible for their ACR scavenging abilities [31], but not their anti-hypertensive effects [21], it is evident that other compounds containing hydrazino groups could be highly suitable as therapeutic agents for diabetic retinopathy. Indeed, the non-vasoactive hydralazine analogue, 1hydrazinoisoquinolone, has recently been proposed to protect against tobacco smoke-related lung toxicity through its ability to suppress ACR levels [14]. Future studies are clearly warranted to identify the most appropriate hydrazino compounds for preventing FDP-lysine accumulation in the diabetic retina.

The AGE-RAGE axis in diabetic retinopathy

The receptor for AGEs (RAGE) is the most-established AGE-binding protein and acts as a signalling receptor for the common AGE CML [41] and hydroimidazolone adducts [29]. RAGE is a multi-ligand receptor and binds to many molecules including S100B, high-mobility group box-1,

amyloid- β and Mac-1 [82] and regulates signalling in a range of cells and tissues. Ligand-binding and signal transduction activate transcription of NF κ B and induction of adhesion molecules, cytokines and/or oxidative stress. In the context of the diabetic retina, this is significant since it is now widely appreciated that the retinal expression of proinflammatory cytokines and elaboration of adhesion molecules on the microvasculature are linked to leukostasis and possible occlusion of capillaries [37]. In the diabetic retina, there is a clear imbalance of cytokines, and it seems likely that factors such as IL-1 α , IL-1 β , IL-6 and TNF- α are linked to inflammatory processes including microglial activation and infiltration of monocytes [86, 87].

RAGE is expressed in many retinal cells, although the highest expression levels seem to be in Müller glia where it is constitutively expressed at low levels. During diabetes there is significant upregulation of this receptor where it co-localises with GFAP and vimentin [87, 88]. It is worth noting that some of the other non-AGE RAGE ligands also occur in the retina, and these may be capable of inducing pro-inflammatory signalling. For example, S100B is found in several retinal cell types including photoreceptors and Müller glia [20]. S100B has neurotrophic role at low levels although upregulation occurs in the Müller glia of diabetic animal models where it can induce inflammatory cytokine expression [89].

Blockade of RAGE may be a useful therapeutic strategy. For example, it has been shown that the soluble RAGE fragment (known as "sRAGE") can prevent Muller cell dysfunction [5] during diabetes and retinal capillary leukostasis in AGE-infused (non-diabetic) mice [51]. RAGE blockade using various agents shows great potential for preventing diabetic retinopathy, and the coming years should bring new efficacious agents that can regulate activation of this receptor in the retina.

Conclusion

There is no question that management of retinopathy in patients will involve precise regulation of glycaemic, vasotensive and lipidemic profiles. As we go forward, there is expectation that new therapeutic agents will become available to prevent key biochemical and metabolic abnormalities that are definitively linked to neuroglial and vascular pathology. AGE/ALE formation and activation of AGE receptors appear to play an important role in diabetic retinopathy, and if we can regulate these pathways, there is hope that initiation and/or progression of this important complication can be prevented.

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