

# Advanced glycation endproducts enhance proliferation, but not tube formation in choroidal microvascular endothelial cells

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## Abstract

• **AIM:** To investigate the role of advanced glycation endproducts (AGEs) in the pathogenesis of age-related macular degeneration (AMD).

• **METHODS:** Bovine choroidal endothelial cells (CEC) were isolated by the modified protocol using lycopersicon esculentum agglutinin coated Dynabeads, and identified by immunocytochemical staining with anti-Factor VIII antibody and uptake of diI-acetylated low-density lipoprotein (diI-ac-LDL). AGEs were prepared by incubating 50g/L bovine serum albumin and 150g/L glucose at 37°C for 6 weeks, which were characterized by dot blot assay with anti-AGEs antibody. CEC proliferation was evaluated by 3, (4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay, and tube formation in CEC was determined by a Vitrogen system.

• **RESULTS:** More than 90% of the cultured cells were positive to Factor VIII immunostaining and had the ability to uptake diI-ac-LDL, which were the features of endothelial cells. 219 AGEs we prepared were affinitive to anti-AGEs antibody. After treatment with AGEs for a time course of 3 days, CEC proliferation was significantly increased in a dose-dependent manner by AGEs at concentrations between 62.5mg/L and 500mg/L. The cytokine, basic fibroblast growth factor (bFGF), enhanced strongly tube-like structure formation in CEC to 124% ( $P < 0.05$ ) above that of untreated

controls. In this condition, AGEs at the concentrations of 500mg/L and 50mg/L showed no effect on CEC tube formation ( $P > 0.05$ ).

• **CONCLUSION:** The present study demonstrated that CEC proliferation was increased by AGEs. However, there was no statistical effect on CEC tube formation. These findings confirm and extend that AGEs could be a potential initiator in the pathogenesis of choroidal neovascularization in exudative AMD, at least in part, through enhancement of CEC proliferation.

• **KEYWORDS:** advanced glycation endproducts; angiogenesis; choroidal microvascular endothelial cells; proliferation; tube formation; age-related macular degeneration

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## INTRODUCTION

The development of choroidal neovascularization (CNV) or angiogenesis is the major contributor to severe vision loss in patients with age-related macular degeneration (AMD)<sup>[1]</sup>. Although current treatments for CNV, such as laser photocoagulation, photodynamic therapy, pharmacological inhibition, surgical intervention, radiation and transpupillary thermotherapy, have been shown to be therapeutic in some cases, all treatment modalities have limitations<sup>[2, 3]</sup>. Future treatment advances should base on understanding of the pathogenesis of AMD; however, this remains largely unclear to date.

The prevalence of AMD increases dramatically with advancing age<sup>[4]</sup>. Among the current hypotheses about aging process, the accumulation of advanced glycation endproducts (AGEs) has been suggested as an important etiological initiator<sup>[5, 6]</sup>. Moreover, it was demonstrated that cell activation in response to AGEs is associated with angiogenesis<sup>[5, 7-9]</sup>. Recently, immunohistological studies have shown that AGEs accumulate in human Bruch's membrane and choroid

with age<sup>[10,11]</sup>, and were present in CNV of AMD<sup>[12,13]</sup>. These studies have pointed to a causal link between the accumulation of AGEs and the pathogenesis of AMD. However, the interactions of choroidal microvascular cells with AGEs have not been investigated.

AGEs are a group of heterogeneous undefined structures, which are formed by the nonenzymatic glycation of sugars and proteins through the Maillard reaction<sup>[5,8,9,14]</sup>. Several studies have been reported about the effects of AGEs on non-choroidal macro- and microvascular cells<sup>[15-18]</sup>. The relevance of those studies to understand the pathogenesis of CNV is limited because endothelial cells are highly organ-specific and heterogeneous<sup>[19,20]</sup>. Angiogenesis, the formation of new blood vessels from preexisting endothelium, is a complex multistep process which includes proliferation, migration and differentiation of endothelial cells, degradation of the extracellular matrix, tube formation and sprouting of new capillary branches<sup>[21]</sup>. To provide evidence that AGEs are directly implicated in choroidal angiogenesis, we tested the effects of AGEs on proliferation and tube formation in choroidal endothelial cells (CEC), which are crucial components of CNV formation.

## MATERIALS AND METHODS

**Reagents** Lycopersicon esculentum agglutinin (LEA), collagenase/dispase and DNase, rabbit anti-human Factor antibody, fibronectin, bovine serum albumin (BSA, fraction, fatty acid free) and 3, (4,5-dimethylthiazol-2-yl) -2, 5-diphenyl tetrazolium bromide (MTT) were purchased from Sigma, Saint Louis, USA; M-450 Dynabeads, from Dynal, Oslo, Norway; endothelial basal medium (EBM) and endothelial growth medium (EGM) supplements, from Clonetics, San Diego, California, USA; Dil-acetylated low-density lipoprotein (dil-ac-LDL), from Paesel & Lorei, Hanau, Germany; StreptABC/Horseradish peroxidase complex and diaminobenzidine (DAB) kit, from DAKO, Hamburg, Germany; human recombinant basic fibroblast growth factor (bFGF), from Life Technologies, UK; vitrogen was obtained from Cohesion, Palo Alto, California, USA; the AGEs as positive control and anti-mouse carboxymethyllysine (CML) monoclonal antibody (mAb 6D12) were generous gifts from Dr. HP Hammes, Giessen, Germany.

**Cell Culture and Identification** CEC were isolated by the modification of previously described method using LEA (a lectin specific for bovine microvessels)-coated Dynabeads<sup>[22]</sup> from bovine eyes obtained from a local slaughterhouse 12-16 hours after death. In brief, the eye was bisected posterior to the ora serrata and the anterior section was discarded. The vitreous and neural retina were removed from the posterior

eyecup, and then retinal pigment epithelial cells (RPE) were gently scraped out. The choroid was dissected and microvessels were collected with the aid of forceps using a dissecting microscope. The vessels were digested with 1g/L collagenase/dispase and 17.3kU/L DNase at 37°C for 90 minutes. The cell suspension was then filtered through a 70µm mesh filter (Falcon). The LEA-coated beads of 4×10<sup>11</sup> beads/L were mixed with the filtrated cell suspension at a ratio of 1:3, and then were incubated at 4°C for 30 minutes with agitation. The cells attached to the beads were harvested with a magnetic particle concentrator (MCP-1, Dynal, Oslo, Norway), and then were incubated in EGM (EBM supplemented with 12mg/L bovine brain extract, 1mg/L hydrocortisone, 20mL/L fetal calf serum (FCS), and 100mg/L gentamycin) in fibronectin-coated six-well culture plates (Falcon) in 50mL/L CO<sub>2</sub>/950mL/L air at 37°C. Seven to 10 days later, the contaminating cell colonies were scraped out<sup>[23]</sup>, and this procedure was repeated more than 3 times before CEC became confluent. Cells were confirmed by their positive immunostaining for Factor VIII and by their uptaking of dil-ac-LDL. For identification, cells were seeded at density of 1×10<sup>4</sup> cells/well onto fibronectin-coated chamber-slides (Nunc, Wiesbaden, Germany). One part of cells were fixed with 900mL/L ethanol and processed for FITC-immunocytochemical staining with anti-human Factor VIII antibody (dilution 1:200) using routine procedures prior to fluorescence microscopic examination. Another part of cells were incubated with dil-ac-LDL at a final concentration of 10mg/L for 4 hours, and then was fixed in 30g/L buffered formaldehyde at room temperature for 20 minutes. After mounted, they were observed with a fluorescence microscope (Zeiss Axioskop HBO 50/AC, Germany) using standard rhodamine excitation. The parallel experiments were performed using bovine RPE and rat Müller cells. All experiments were carried out in the subconfluent cells at cell passage number 3 to 7.

**Preparation and Characteristics of AGEs** AGEs were prepared as described previously<sup>[17,24,25]</sup> by incubating 50g/L BSA in phosphate buffered saline (PBS) with 150g/L glucose at 37°C for 3 days to 6 weeks under sterile conditions. Control unmodified BSA was incubated under the same conditions except for the absence of glucose. After unincorporated sugars were removed by dialysis against PBS, protein concentration was determined by Bio-Rad protein assay kit (Bio-Rad, Muenchen, Germany) according to the manufacturer's instruction. The resultant preparations were characterized by antibody affinity with mAb 6D12 using dot blot analysis as previously described<sup>[26]</sup>. We loaded 1µL of

positive control AGEs, AGEs preparations and control BSA preparation at protein concentrations of 500, 100 and 50mg/L on nitrocellulose membranes (Biometra, Goettingen, Germany) and dried them at room temperature. Membranes were blocked with 30mL/L skim milk for 30-60 minutes, incubated with 1:5000 mAb 6D12 for 90 minutes, exposed to peroxidase-labelled anti-mouse IgG for 60 minutes, and then with StreptABC/Horseradish peroxidase complex for further 30 minutes. The staining was then developed in DAB solution. Between each step of the procedure, it is necessary to rinse sufficiently the membranes in PBS at room temperature.

**Proliferation Assay** CEC proliferation was evaluated using MTT colorimetric assay of Mosmann<sup>[27]</sup>. Briefly, CEC were trypsinized and plated out at a density of  $4 \times 10^3$  cells in 100 $\mu$ L EGM per well into a fibronectin-coated 96-well plate and allowed to attach overnight. The cells were washed twice with PBS, and then were stimulated with 10 $\mu$ g/L bFGF, 500mg/L BSA control, or AGEs at various concentrations of 500, 250, 125 and 62.5mg/L in EBM with 10mL/L FCS. The cells cultured in EBM with 10mL/L FCS were used as negative control. The cells were incubated at 37°C for 72 hours, and at the last 4 hours 10 $\mu$ L of 5g/L MTT was added in each well. The formazan crystals formed were dissolved in 100 $\mu$ L dimethyl sulfoxide and the optical density was recorded at 570nm on a Molecular Devices Spectra Max 250 Multiscan (BMG, Offenburg, Germany). The impact of AGEs on CEC proliferation was estimated by the percentage (%) compared to negative controls.

**Tube Formation Assay** The tube formation assay was performed as described by Sakamoto *et al*<sup>[28]</sup> with a minor modification. Four hundred  $\mu$ L Vitrogen (2.4g/L) containing 0.001mol/L sodium hydroxide, 200mmol/L HEPES, 5mg/L fibronectin and laminin, and 40 $\mu$ L of 10 $\times$ DMEM medium was put into each well of 24-well plates and incubated at 37°C for 2-3 hours to gel. After polymerization of the gels,  $1.0 \times 10^5$  CEC were seeded onto each gel and incubated with 500 $\mu$ L EGM at 37°C for 24 hours. The medium was then aspirated, and the remaining cells were overlaid with 120 $\mu$ L gel per well, and incubated for 1 hours at 37°C. Next, 50 $\mu$ g/L bFGF, AGE at concentrations of 50mg/L and 500mg/L, 500mg/L BSA in 500 $\mu$ L EBM with 10mL/L FCS, 100kU/L penicillin and 100mg/L streptomycin was added to each well. The incubation was continued to the 3rd day. To evaluate the lengths of tube-like structures in the gels, 4 photographs in different fields around center in each well were randomly taken at 3 days with a phase-contrast microscope ( $\times 5$  objective). The negative films were then

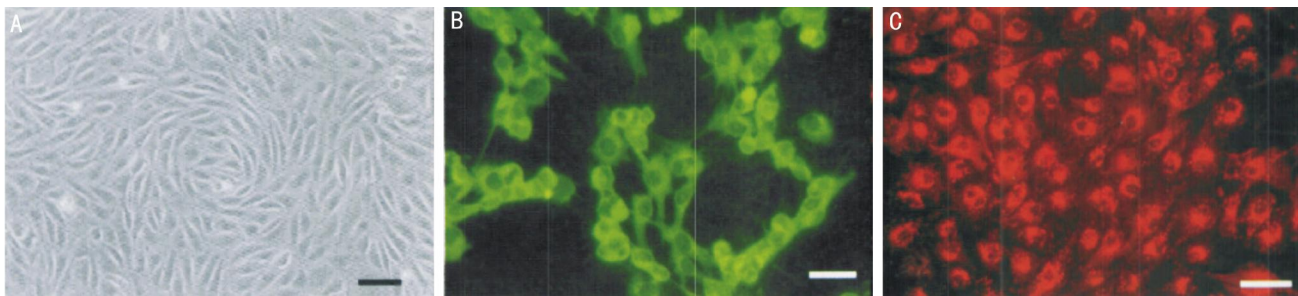
scanned and digitized. All tube structures in the resulting images were traced manually and the lengths of the tubes were automatically measured by the Image Analyzer System KS300 2.0 (Kontron Elektronik GmbH, Eching, Germany). The total additive length of all tube structures in 4 photos was represented as the tube length of the well. Results were shown as percentages compared with the negative controls.

**Statistical Analysis** Proliferation and tube formation assays were done in pentagonal and triplicate respectively; and the experiments were carried out on at least 3 different occasions. Results were shown as mean percentage standard deviation of the mean compared with the negative control of 3 experiments. Mean values were compared using One-way ANOVA and  $P < 0.05$  was considered significant.

## RESULTS

**Bovine CEC** The plated cells became attached and flattened after 3-5 days. In the following 1-3 weeks, cell colonies were formed, and contaminating cells, distinguished from CEC by their appearance, were scraped away. The residual endothelial cells reached confluent in about 3-4 weeks after seeding. At this time, the cultures had the typical 'cobblestone' appearance of endothelium in a confluent monolayer (Figure 1A). Indirect immunocytochemical staining with anti-Factor antibody revealed that positive yellow-green staining was localized in cytoplasm or concentrated in the perinuclear space in the passaged cultures (Figure 1B). The assay for uptake of dil-ac-LDL showed that cytoplasm was filled with red fluorescent granules in the positive cells (Figure 1C). By counting, more than 90% of the cultured cells were confirmed vascular endothelial cells by their positive immunostaining for Factor and their uptake of dil-ac-LDL. The parallel experiments performed in bovine RPE and rat Müller cells had no positive staining (data not shown).

**Dot Blot Analysis** Immuno-dot-blot analysis showed that the highly modified AGEs (positive control) reacted markedly against mAb 6D12 (4-5+). We collected the preparations of BSA and glucose at various time points between 3 days and 6 weeks. Our AGE preparations started to react against this antibody at 2 weeks. The reactivity increased in a time- and concentration-dependent manner, which reached the maximal reactivity (3+) at 6 weeks (Figure 2). No immunoreaction was detected in BSA control at 6 weeks against the CML antibody. CML is a major AGE-epitope<sup>[29]</sup> and the mAb 6D12 is a medium affinity CML-protein-recognizing antibody<sup>[26]</sup>. The results indicated the presence of CML in our AGEs preparations, but not in the control BSA preparation. Based on the above results, AGEs and



**Figure 1 Morphology and identification of CEC** A: The appearance of endothelium in a confluent monolayer under phase-contrast microscope. Bar, 80µm; B: Immunocytochemical staining with anti-Factor antibody (FITC). Bar, 35µm; C: Uptaking assay of dil-ac-LDL (Rhodamine fluorescence). Bar, 50µm

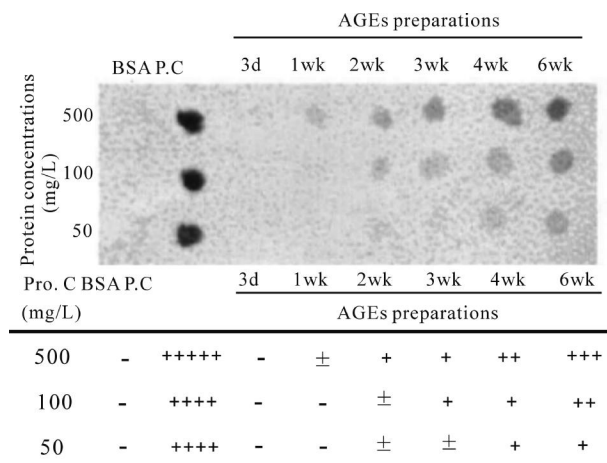
BSA preparations at 6 weeks were used in the following experiments.

**CEC Proliferation** It was observed that CEC proliferation was increased by 10µg/L bFGF by about 2.5 fold over the basal level (EBM group) ( $P < 0.01$ ). Treatment of CEC with AGEs at various concentrations between 62.5mg/L and 500mg/L resulted in a dose-dependent increase in CEC proliferation (Figure 3). CEC proliferation was significantly increased by about 44% ( $P < 0.01$ ) and 33% ( $P < 0.05$ ) by 500mg/L and 250mg/L AGEs respectively. There was no statistical effect of BSA on CEC proliferation ( $P > 0.05$ ).

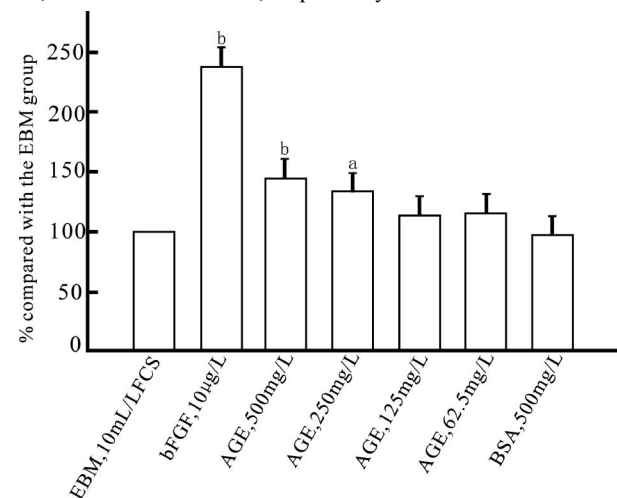
**CEC Tube Formation** Tube formation was evaluated by Vitrogen collagen system. CEC were seeded into solidified Vitrogen and incubated in the presence of 50µg/L bFGF, AGEs or BSA as indicated for 48 hours. Stimulation of CEC with 50µg/L bFGF led significantly to tube-like structure formation ( $P < 0.05$ ). Under the conditions, both AGEs and BSA did not stimulate tube formation in CEC compared with basal level ( $P > 0.05$ ) (Figure 4, 5).

## DISCUSSION

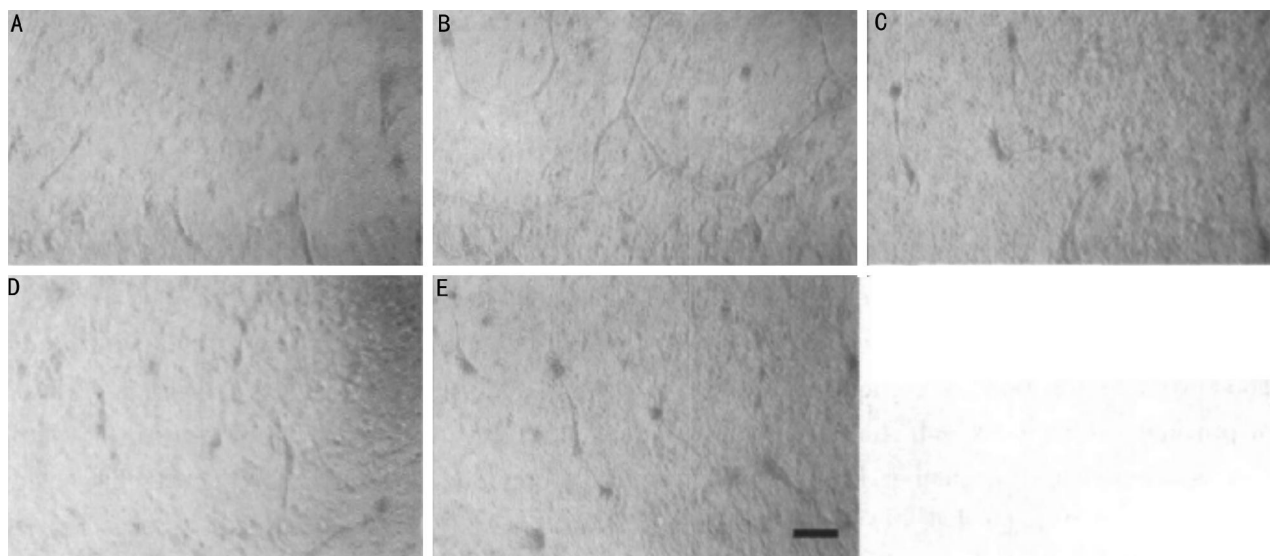
Nonenzymatic glycation, a complex series of reactions between sugars and amino groups of proteins, initiates with the reversible formation of Schiff's base, then undergoes a rearrangement to form stable Amadori product, and after that could end with the formation of irreversible AGEs through multiple chemical rearrangements. This reaction, with subsequent formation of AGEs has been implicated in aging process and accelerated in diabetes [54]. A lot of age-associated diseases, such as cataract, Alzheimer's disease and arteriosclerosis have been proposed to relate to AGEs accumulation [9]. Recently, Handa's group [10, 11] postulated that the formation and accumulation of AGEs in human Bruch's membrane and choroid could promote aging of the RPE-Bruch's membrane-choroid complex. Ishibashi *et al* [12] found that AGEs (CML) accumulated in soft, macular drusen in aged eyes, and in RPE in surgically excised CNV



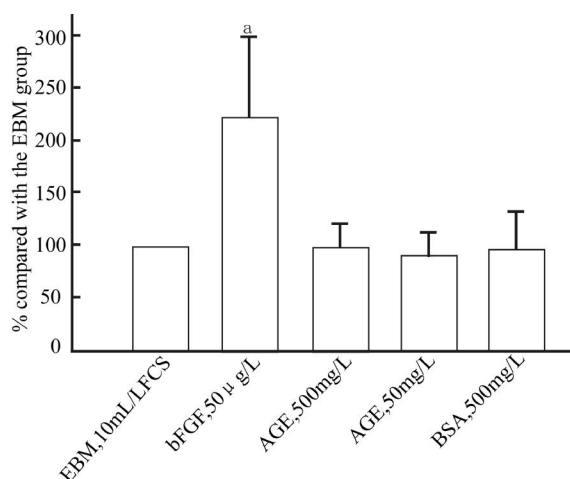
**Figure 2 Dot blot analysis of AGEs using an anti-CML antibody** There were common structures between CML and AGEs preparations. The density of the dots was scored into 7 levels, e.g., negative (-), suspected positive (±), and slightly to strongly positive (1-5+). Pro. C., protein concentrations; P.C., positive control. The numbers and characters, 3 days, 1 week, 2 weeks, 6 weeks, indicate that incubation time for preparation of AGEs are 3 days, 1 week, 2 weeks and 6 weeks, respectively



**Figure 3 Effects of AGEs on CEC proliferation** CEC proliferation was increased by AGEs in a dose-dependent manner. Each column represents mean % compared with EBM group standard deviation of 3 independent experiments. <sup>a</sup> $P < 0.05$ , <sup>b</sup> $P < 0.01$  compared with EBM group



**Figure 4 The representative morphology of tube-like structures in CEC** The tube-like structures in CEC were actively increased by stimulation of bFGF(B). There were no obvious changes in the groups of 500mg/L AGEs (C), 50mg/L AGEs (D) and 500mg/L BSA group (E) compared with EBM group (A). Bar, 250µm



**Figure 5 Effects of AGEs on CEC tube formation** The tube-like structures in CEC were increased by bFGF, but not by AGEs and BSA. Each column represents mean % compared with EBM Group’s standard deviation of 3 independent experiments.<sup>a</sup>*P* < 0.05 compared with EBM group

from patients with AMD. Furthermore, Hammes *et al*<sup>[13]</sup> demonstrated AGEs were adjacent or colocalized with one of the receptors for AGEs (RAGE) in CNV of patients with AMD. Those evidences indicated that AGEs could play a role in the pathogenesis of AMD; however, the precise mechanism should be investigated.

Given that endothelial cells are highly organ specific and heterogeneous<sup>[19, 20]</sup>, it is our interest to evaluate the direct effects of AGEs on the CEC, which is a key cellular component in choroidal angiogenesis. In this study, we showed that

AGEs increased CEC proliferation in a dose-dependent manner. Moreover, there was no statistically significant influence of AGEs on tube formation in CEC. However, the tube-like structures were well formed in the bFGF-stimulated group. These findings are suggestive of an implication of AGEs in choroidal angiogenesis, at least through increasing CEC proliferation directly.

The inconsistent effects of AGEs on the substeps of angiogenesis in different endothelial cell systems have been described, although almost all studies indicated that AGEs exert a potent angiogenic effect on endothelial cells. For example, in human umbilical vein endothelial cells, tube formation and migration but not proliferation were stimulated by AGEs<sup>[15]</sup>. Therefore, the growth (cell number and DNA synthesis) of retinal microvascular endothelial cells treated with AGEs was significantly increased<sup>[16]</sup>. AGEs were found to induce a significant increase in the number of cells as well as their synthesis of DNA, and to stimulate the tube formation in human skin microvascular endothelial cells<sup>[17]</sup>. Exceptionally, AGEs at a higher concentration (500mg/L) reduced the cell number because of cytotoxicity to bovine aortic endothelial cells; however, low concentration (62.5mg/L) of AGEs was significantly mitogenic to the cells<sup>[18]</sup>. One explanation for these discrepant results may be the nature of the cells used in the studies, although other differences in experimental conditions, e.g. AGEs preparation and test methodology, could not be excluded.

Our research was performed using well-characterized cells and AGEs. The CEC we used in this study were isolated by

LEA, a specific lectin for bovine microvessels, and then identified by positive staining to Factor and uptake of LDL, both distinctive markers for endothelial cells [22]. The AGEs were prepared using the classic protocol as previously described by many researchers [17, 24, 25], and immuno-dot blot analysis has demonstrated that there are common structures between CML (a major AGE-epitope [29]) and our AGEs preparations. The current results on proliferation was consistent with our previous research, which showed that CEC proliferation was enhanced by another kind of AGEs made by incubation at 50°C for 4 weeks [30]. With regard to the validity and reliability of the methodology for the tube formation assay, it was morphologically and histologically demonstrated that tube-like structures in gel were formed in cytokine-stimulated CEC (the histological data not shown). In our experience, it seems to need more cytokines to stimulate tube formation than to increase proliferation in CEC. Cytokine bFGF of 3µg/L could increase CEC proliferation by about 150%; however, about 20µg/L bFGF may be necessary for reaching the similar level of increasing tube formation in CEC (unpublished data). Autocrine of cytokines, such as vascular endothelial growth factor (VEGF) by endothelial cells has been considered as a possible angiogenic mechanism of AGEs [17,30]. One possible explanation of our results is that the amount of cytokines in the medium secreted by AGEs-stimulated CEC is sufficient to increase proliferation, but not tube formation in CEC. Of course, we could not exclude other possibilities for explanation of the present results, including the heterogeneity of endothelial cells.

The molecular mechanism used by AGEs to drive angiogenesis is beyond the scope of this paper, but several hypotheses exist. In fact, AGEs have been shown to enhance the following events: ① the expression of VEGF mRNA and/or the production of VEGF protein in cultured Müller cells [24], human RPE and bovine vascular smooth muscle cells (SMC) [31], and microvascular endothelial cells [17,30]; ② the production of platelet-derived growth factor (PDGF)-BB in RPE [32]; and ③ the synthesis of insulin-like growth factor (IGF) and transforming growth factor-β (TGF-β) in mesangial cells [33]. It indicates that the effects of AGEs could be mediated by several growth factors and/or cytokines secreted in paracrine and/or autocrine fashions. Another possibility linking AGEs with angiogenesis is that AGEs could induce expression of vascular cell adhesion molecule-1 (VCAM-1) in endothelial cells, which enhanced adhesivity of circulation monocytes [34]. In addition, AGEs could elicit their cellular effects by modifying the extracellular matrix [8]

or the nitric oxide synthase/ nitric oxide (NOS/NO) pathway in endothelial cells [35]. Most of those effects are supposed to be mediated by the interaction of AGEs with cell surface receptors and other AGEs binding proteins, which have been identified on different cell systems including monocytes, SMC and endothelial cells [7, 8]. The binding of AGEs with their receptors or binding proteins has been shown to induce a cellular-oxidant stress activating the transcription factor NF-κB, which further mediates gene expression of cytokines and VCAM-1 [7,8,36].

However, no matter which of mechanisms may be involved in AGEs function, the present results suggested a quite possibility that their direct impact on CEC may take part in the choroidal angiogenesis. We hypothesized that accumulation of AGEs with age changes the biological behavior of CEC through less well-known mechanisms, at least in part, increases CEC proliferation, and then could contribute to the development of AMD. The immunohistochemical studies confirming the presentation of AGEs and their receptor in CNV of AMD [12,13] are significantly consistent with our results. The current and future research efforts will hopefully lead to a better understanding of the pathogenesis of these changes and ultimately to a cure for AMD.

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## Glycation endproducts enhance proliferation of CEC

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