Abstract. Diabetic retinopathy (DR) is one of the most common causes of blindness in developed countries. Due to its asymptomatic onset and progressive disease course, DR is typically diagnosed at a late stage when treatment options are limited and therefore often results in irreversible blindness. Studies have demonstrated that carnosine may prevent and treat DR. In a previous study, the positive effect of carnosine on DR was determined and it was revealed that there may be an association between carnosine and the mitogen-activated protein kinase (MAPK)/extracellular signal related kinase (ERK) signaling pathway. To assess the interaction between carnosine and the MAPK/ERK signaling pathway, changes in PKC, ERK and p-ERK expression was assessed in diabetic rats following treatment with carnosine, PD98059 or U46619 via reverse transcription-quantitative polymerase chain reaction and western blotting. The results demonstrated that the expression of ERK and p-ERK were significantly suppressed following treatment with carnosine, but no significant effect on the expression of PKC was identified, which indicates that suppressing the activation of the MAPK/ERK signaling pathway may serve an important role in carnosine-induced DR prevention and treatment.

Introduction

Diabetes mellitus (DM) is a systemic metabolic disorder characterized by impaired glucose metabolism, the incidence of which continues to grow at an alarming rate from 10.6% in 1989 to 32.1% in 2009 in Saudi Arabia (1). A severe microvascular complication of DM, diabetic retinopathy (DR), is one of the four major blinding diseases that has become the leading cause of adult blindness, according to surveys conducted in other developed countries, including the United States and Japan (2). DR is typically diagnosed at a late stage when treatment options are limited. Due to its asymptomatic onset and progressive disease course, DR often results in irreversible blindness, which has a shortage of effective treatment due to the complex nature of the disease and limited knowledge of its pathogenic mechanism (3-5). Therefore, finding an effective treatment for DR has important clinical significance.

Poor blood glucose control, hypertension, hyperlipidemia and obesity may be involved in the development of DR, and may also contribute to the development of neuropathy (6). Studies (7,8) assessing the pathogenetic pathways that lead to microangiopathy indicate that many genes may be involved in its development. Known biochemical mechanisms that underlie DR include advanced glycation end-product (AGE) formation, oxidative stress, the activation of protein kinase C (PKC) isoforms and polyl and hexosamine pathway activity (7). As a result of impaired pathway activity, a number of issues may arise, including retinal tissue hypoxia, endothelial cell dysfunction, vasodilation impairment, angiogenic factor hyperactivation and changes to the extracellular matrix (7). At present, the treatment of DR primarily targets its proliferation, but the restoration of lost visual function remains difficult (8).

Carnosine is an endogenous dipeptide consisting of beta-alanine and L-histidine that is produced in skeletal muscle and the nervous system (9,10). It has been reported that many biological effects of carnosine, including anti-inflammation, anti-aging, immune-regulation and anti-diabetes are associated with its anti-oxidative activities (11-13). Carnosine has been revealed to be an effective scavenger for the removal of reactive carbonyl species, as determined in a mouse model of diabetic nephropathy, and the reactive species located in pancreatic β-cells (10,14). Javadi et al (15) demonstrated that carnosine primarily prevents dehydroascorbic acid-induced unfolding and the aggregation of lens proteins and significant lens opacity. Furthermore, other studies have revealed that carnosine exerts a positive effect on the prevention and treatment of DR, but it has no association with anti-oxidation and anti-glycosylation (16,17).

Mitogen-activated protein kinase (MAPK) is a signal transduction pathway that is involved in various physiological processes, including gene expression and the proliferation, differentiation, death and survival of various cells (18,19). Members of the MAPK family are regulated by a cascade
of phosphorylation and are activated by extracellular stimulus (20). In the process of ERK1/2 phosphorylation, the extracellular signal related kinase 1/2 (ERK1/2) cascade is involved in the MAPK pathway, while MAPK signaling is activated by MAPK kinase 1 (21). The activation of the MAPK/ERK signaling pathway has been used frequently for the study of diabetic complications (22). Previous studies have also revealed that the PKC signaling pathway is involved in diabetic wound healing (23) and diabetic myocardial injury (24). The current study assessed the alteration of PKC, ERK, and phosphorylated (p)-ERK expression in diabetic rats following treatment with carnosine, PD98059 (an inhibitor of MAPK/ERK) or U46619 (an activator of MAPK/ERK). Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) and western blotting was performed to assess the association between carnosine and the MAPK/ERK signaling pathway. The results revealed that suppressing the activation of the MAPK/ERK signaling pathway may serve an important role in the prevention and treatment carnosine-induced DR. However, the PKC pathway may not be involved in this process.

Materials and methods

Reagents and animals. Streptozotocin (STZ) was purchased from Sigma-Aldrich (Merck KGaA; Darmstadt, Germany). Cluster of differentiation (CD)31 antibodies were purchased from Abcam (Cambridge, UK), and Goat anti-Mouse Immunoglobulin G (IgG) H&L-cyanine 3 (Cy3) antibodies were purchased from ProteinTech Group, Inc. (Chicago, IL, USA). A total of 25 male sprague dawley rats (age, 7 weeks; weight, 180-185 g) were purchased from the Experimental Animal Center of Southern Medical University (Guangzhou, China). All rats were housed in a specific pathogen free room at a temperature of 25°C and a humidity of 50% under a 12 h light/dark cycle, with ad libitum access to food and water.

Ethics statement. All animal experiments were performed in accordance with the recommendations included in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. Furthermore, the protocol of the current study was approved by the Committee on the Ethics of Animal Experiments of Tangdu Hospital, Air Force Medical University (Xi’an, China).

Establishment of diabetic rat model. Following a week of feeding, the body weight of rats was measured. A total of 10 rats were randomly divided into 2 groups: The DM group (n=5; weight, 200-220 g) and the normal group (n=5; weight, 180-200 g). Rats in the DM group were intraperitoneally injected once with 2% STZ at a dose of 60 mg/kg. Animals were considered diabetic when glucose levels were >16 mM in 24 h following injection. Normal rats were administered the same quantity (100 µl) of citrate buffer solution (0.02 mol/l; pH=4.5). Rats were euthanized by gavage for 10 days) group and a control group. Retinas were fixed in 4% para-formaldehyde solution for 15 min. Retinas were then harvested for cell isolation.

Drug treatment. Rat RVECs from each group were plated in 6-well plates containing coverslips for 24 h at 37°C and then fixed using 4% paraformaldehyde for 10 min at room temperature and washed in triplicate with PBS. Samples were then blocked with normal goat serum (1:50; cat. no. ab7481; Abcam, Cambridge, UK) for 15 min at room temperature. Subsequently cell slides were incubated with primary antibodies (anti-CD31; cat. no. ab28364; 1:200; Abcam) at 37°C for 2 h. Samples were washed with PBS in triplicate and probed with Goat anti-Mouse immunoglobulin G H&L-Cy3 (cat. no. SA00009-1; 1:100; ProteinTech Group, Inc.) at 37°C for 1 h. Hoechst 33258 (Beyotime Institute of Biotechnology, Haimen, China) was used to stain nuclei and images were captured using a fluorescence microscope (magnification, x50; Olympus Corporation, Tokyo, Japan) at room temperature for 15 min. Cell number analysis was performed using Image J 1.45 software (National Institutes of Health, Bethesda, MD) as described previously (25). The number of positive cells (with red coloration) and blue stained nuclei were counted separately. Positive rate was determined to be the number of positive red stained cells/the number of blue stained nuclei.

Establishment of diabetic rat model. Following a week of feeding, the body weight of rats was measured. A total of 10 rats were randomly divided into 2 groups: The DM group (n=5; weight, 200-220 g) and the normal group (n=5; weight, 180-200 g). Rats in the DM group were intraperitoneally injected once with 2% STZ at a dose of 60 mg/kg. Animals were considered diabetic when glucose levels were >16 mM at 72 h following injection. Normal rats were administered the same quantity (100 µl) of citrate buffer solution (0.02 mol/l; pH=4.5). Rats were euthanized by gavage for 10 days) group and a control group. Retinas were then harvested for cell isolation.

Rat retinal vascular endothelial cell (RVEC) isolation and cell culture. To establish a primary cell culture of rat RVECs, retinas isolated from rats of each group were digested with trypsin at 37°C for 30 min. Residual retinas were then minced and fragments were filtered through a 100 µm cell strainer. Following harvest, fragments from the strainer were incubated in 3 ml of mixed collagenase (Sigma-Aldrich) for 30 min at room temperature. Digestion was stopped using complete medium (Dulbecco's Modified Eagles medium supplemented with 10% fetal bovine serum, 100 U/ml penicillin and 100 µg/ml streptomycin, all Thermo Fisher Scientific, Inc., Waltham, MA, USA) at room temperature. Following centrifugation at 252 x g at room temperature for 5 min, cells were suspended in complete medium. The above suspension was transferred to a culture flask precoated with 1% fibronectin (Sigma-Aldrich) and cultured at 37°C in a 5% CO₂ humidified incubator. The medium was changed every 2.5 days until all cells formed confluent monolayers. Cells were passaged at a split ratio of 1:2.

Cell identification. Rat RVECs were seeded (7x10⁵ cells) in 12-well plates containing coverslips for 24 h at 37°C and then fixed using 4% paraformaldehyde for 10 min at room temperature and washed in triplicate with PBS. Samples were then blocked with normal goat serum (1:50; cat. no. ab7481; Abcam, Cambridge, UK) for 15 min at room temperature. Subsequently cell slides were incubated with primary antibodies (anti-CD31; cat. no. ab28364; 1:200; Abcam) at 37°C for 2 h. Samples were washed with PBS in triplicate and probed with Goat anti-Mouse immunoglobulin G H&L-Cy3 (cat. no. SA00009-1; 1:100; ProteinTech Group, Inc.) at 37°C for 1 h. Hoechst 33258 (Beyotime Institute of Biotechnology, Haimen, China) was used to stain nuclei and images were captured using a fluorescence microscope (magnification, x50; Olympus Corporation, Tokyo, Japan) at room temperature for 15 min. Cell number analysis was performed using Image J 1.45 software (National Institutes of Health, Bethesda, MD) as described previously (25). The number of positive cells (with red coloration) and blue stained nuclei were counted separately. Positive rate was determined to be the number of positive red stained cells/the number of blue stained nuclei.
sectioned (0.3 mm thick), stained with CD31 antibodies and Hoechst as aforementioned and examined using fluorescence microscopy (magnification, x50; Olympus Corporation).

**RT-qPCR.** RT-qPCR analysis was performed as described previously (26). Total cellular RNA was extracted from rat retinas using a TRizol reagent (Invitrogen; Thermo Fisher Scientific, Inc.) and cDNA was prepared using the M‑MLV reverse transcriptase kit (Promega Corporation, Madison, WI) with random primers according to the manufacturer's protocol. The thermocycling conditions for PCR were as follows: Pre-denaturation for 10 min at 94˚C, denaturation for 20 sec at 94˚C, annealing for 20 sec at 55˚C, and extension for 20 sec at 72˚C. Each PCR reaction was performed for 40 cycles. RT-qPCR analysis was performed using a Biosystems StepOne™ real-time PCR apparatus (Applied Biosystems; Thermo Fisher Scientific, Inc.) using standard procedures and analyzed using ABI Prism 7500 SDS software (Applied Biosystems; Thermo Fisher Scientific). A Fast SYBR‑Green Master Mix was obtained from Thermo Fisher Scientific, Inc. Data were presented as the relative expression following normalization to the expression of GAPDH using the 2^−∆∆Cq method (27). The PCR primer sequences were as follows GAPDH forward, 5′‑ACA GCA ACA GGG TGG TGG AC‑3′ and reverse, 5′‑TTT GAG GGT GCA GCG AAC TT‑3′; PKC forward (5′‑3′), 5′‑TGG AGT CCT GCT GTA TGA GAT GTT G‑3′ and reverse, 5′‑CGC TTT CCT GGG TGC TTG G‑3′; ERK forward, 5′‑GAA AGC ATT ACC TTG ACC AG‑3′ and reverse, 5′‑CTT TGG AGT CAG CAT TTG G‑3′.

**Western blotting.** The procedure for western blotting was performed as described previously (28). Total cellular protein was extracted using radioimmunoprecipitation assay lysis buffer containing phenylmethylsulfonyl fluoride Protease Inhibitor (Sigma‑Aldrich; Merck KGaA). A BCA Protein Assay kit was utilized for protein determination (cat. no. 23225; Thermo Fisher Scientific). Proteins (30 µg) were resolved in 5 or 12% SDS‑PAGE gels and then transferred onto polyvinylidene difluoride (PVDF) membranes following electrophoresis. Subsequently, PVDF membranes were blocked using 5% skimmed milk in TBST (20 mM Tris, pH 7.6, 150 mM NaCl, 0.1% Tween‑20; wt/vol) for 2 h at room temperature. Membranes were immunoblotted with Mouse anti‑PKC (cat. no. P5704; 1:1,000; Sigma‑Aldrich; Merck KGaA), ERK1/2 Rabbit Polyclonal (cat. no. 16443‑1‑AP; 1:1,000; ProteinTech Group, Inc.) and anti‑p‑ERK antibodies (cat. no. ab65142; 1:1,000; Abcam) at 4˚C overnight. Samples were then incubated with IRDyeTM‑800 conjugated anti‑mouse (cat. no. P/N 925‑32210; 1:10,000; Li‑COR Biosciences, Lincoln, NE) and IRDye® 800CW Goat anti‑Rabbit IgG (H+L)secondary antibodies (cat. no. P/N 925‑32211; 1:10,000; Li‑COR Biosciences) for 30 min at room temperature. The specific protein bands were detected using Odyssey Infrared Imaging System (Li‑COR Biosciences) and analyzed with Image Studio™ Software 4.0 (Li‑COR Biosciences). The quantification of each protein was normalized to β‑actin.

**Statistical analysis.** All data were expressed as the mean ± standard error of the mean and analyzed using GraphPad Prism5 software (GraphPad Software, Inc., La Jolla, CA). One‑way analysis of variance followed by a Tukey's test was used to analyze the significant differences between groups. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**RVEC determination.** To determine whether the isolated and cultivated cells were RVECs, cells were fixed with 4% paraformaldehyde and subsequently stained with CD31, a specific marker of endothelial cells. As presented in Fig. 1, >90% (Data not shown) of the isolated cells stained positive for CD31, thereby confirming that they were RVECs.

**Carosine inhibited DM‑caused retinal overvascularization.** CD31 immunostaining revealed that the density of retinal
microvessels were significantly increased in DM rats compared with control rats and carnosine markedly inhibited this increase of density in DM rats (Fig. 2).

The MAPK/ERK pathway is involved in the prevention and treatment of DR, as caused by carnosine. Changes of PKC, ERK and p-ERK expression in diabetic rats following treatment with carnosine, PD98059 and U46619 were assessed via RT-qPCR and western blotting. The mRNA levels of PKC (Fig. 3A) and ERK (Fig. 3B) of diabetic RVECs were significantly higher than those of normal RVECs. However, PKC mRNA levels in normal or diabetic RVECs were not substantially impacted by PD98059 and U46619 with or without carosine (Fig. 3A). Furthermore, ERK mRNA levels of RVECs was increased by U46619 and reduced by PD98059 and these changes were further enhanced by carosine (Fig. 3B).

The protein levels of PKC, ERK, and p-ERK in RVECs after treatment with PD98059, U46619, and carnosine were assessed via western blotting. Levels of PKC, ERK and phosphorylated ERK protein were significantly increased in diabetic RVECs compared with normal RVECs (Fig. 4A-D). PD98059 and U46619 increased the PKC protein level of normal RVECs but had little effect on the PKC protein level of diabetic RVECs. Furthermore, carosine inhibited PD98059 but augmented the U46619 induced increase of PKC. However,
it did not effect the PKC levels of diabetic RVECs (Fig. 4A and B). ERK and phosphorylated ERK levels were reduced by PD98059 and increased by U46619 in normal and diabetic RVECs. Carosine treatment reduced ERK and phosphorylated ERK levels of normal and diabetic RVECs treated with PD98059 and U46619. Normal RVECs treated with U46619 however, were not affected (Fig. 4A, C and D). Taken together, these data suggest that the MAPK/ERK signaling pathway, rather than the PKC pathway, mediates carnosine-induced microvascular changes in DR.

**Discussion**

Carnosine (β-Ala-l-His) is an imidazole dipeptide that has many biological functions. Numerous studies have demonstrated that the carnosine-carnosinase system serves a key role in the pathogenesis of diabetes and that carnosine may be used as a novel therapeutic agent against type 2 diabetes (10,29,30). Cripps et al (10) revealed that following treatment with carnosine, insulin secretion from isolated mouse islets or insulinoma cell line β-cells was increased. They further revealed that the inhibition of insulin secretion caused by glucolipotoxic was reversed and that glucose uptake into skeletal muscle cells was enhanced. AGEs and oxidative stress serve a role in the development of diabetic complications (31). It has been demonstrated that carnosine can treat diabetic complications by decreasing oxidation and glycation products in the serum and livers of diabetic rats (30).

Visual impairment as a result of DR has a significant negative impact on patient quality of life and their ability to successfully manage their disease (32). Due to its asymptomatic onset and progressive disease course, identifying an effective treatment of DR is clinically significant. Pfister et al (33) determined that treatment with oral carnosine prevents vascular damage in experimental DR, independent of its biochemical function, which includes reactive oxygen species (ROS) or advanced glycation end (AGE) inhibition (33).

The MAPK/ERK signaling pathway has been determined to serve an important role in various pathogenetic pathways and physiological processes, including the regulation of cell cycle entry and proliferation, as well as in the development of DR by upregulating chronic inflammation (34). The MAPK/ERK signaling pathway includes various MAPKs, including ERKs, which integrate multiple biochemical and environmental signals (including Ras activation and the kinase cascade) via phosphorylation cascades (35). Gogg et al (36) demonstrated that the phosphorylation of ERK1/2 was significantly upregulated in Type 2 diabetes. A previous study revealed that after the degradation of the insulin receptor substrate (IRS)-1 protein, p-MAPK levels are upregulated following the increase of ERK1/2 phosphorylation in diabetes (37). However, the basic phosphorylation of ERK1/2 has also been demonstrated to be upregulated in adipose cells in type 2 diabetes (38). Additionally, IRS-1 has also been revealed to be decreased (39).

ERK is strongly activated in primary neuroretinal tissue and has been determined to be a neuroprotective regulator in the neuroretina, which is involved in the maintenance of the neuroretina-retinal pigment epithelium interaction (40,41). It has been reported that the activation of ERK serves an important role in the maintenance of the inner blood-retinal barrier, similar to that of the RPE outer blood-retinal barrier (42). Van Dijk et al (43) demonstrated that the reversible suppression of ERK activity leads to disturbances in the neuroretina-RPE interaction and in the occurrence of SRF, which results in retinopathy.

Figure 4. Carnosine regulates the level of PKC/ERK/p-ERK proteins in RVECs. (A) Western blotting determined the expression of PKC, ERK and p-ERK following treatment with or without PD98059, U46619 and carnosine for 24 h (A). The quantification of (B) PKC, (C) ERK, (D) p-ERK and (E) ERK/p-ERK expression was determined by normalizing to β-actin. Data are presented as the mean ± standard error of the mean from 3 independent experiments. *P<0.05 vs. NC Control; †P<0.05 vs. NC PD98059; ‡P<0.05 vs. U46619; ††P<0.05 vs. DM Control; ‡‡P<0.05 vs. DM PD98059; †††P<0.05 vs. DM U46619. PKC, protein kinase C; ERK, extracellular signal related kinase; p, phosphorylated; RVEC, retinal vascular endothelial cells; NC, negative control; DM, diabetes mellitus.
The glycation of proteins and the formation of AGEs serve an important role in DM and DR. An increase in AGE level has been identified in the retinas of patients with diabetes and has been determined to be positively correlated with the increase of blood glucose and the development of DR (44). Previous studies have demonstrated that the combination of AGE and the AGE receptor causes the activation of the MAPK/ERK signaling pathway, the activation of the oxidative stress response and the overexpression of certain cytokines (including inflammatory cytokines, lymphocyte adhesion molecules, vascular regulators and coagulants), which may lead to retinal lesions and vascular endothelial cell damage in vitro (45,46). The current study revealed that the expression of p-ERK/ERK significantly changed following treatment with PD98059, U46619 and carnosine, respectively or in combination. The current study therefore hypothesized that the protective effect of carnosine in DR may be associated with AGEs and the MAPK/ERK signaling pathway. However, further studies are required to clarify how the interaction between carnosine and the MAPK/ERK signaling pathway improves DR.

A previous study has determined that the activity of PKC, MEK and ERK1/2 is significantly increased in glomerular mesangial cells in STZ-induced and glucose-induced diabetic rats, and that the activation of MEK and ERK1/2 is PKC-dependent (46). The results of the current study revealed that there was almost no effect on the expression of PKC following treatment with PD98059, U46619 and carnosine, respectively or in combination, indicating that the suppression of the PKC-independent MAPK/ERK signaling pathway, rather than the PKC pathway, may serve an important role in the prevention and treatment of carnosine-induced DR.

In summary, the present study demonstrated that the MAPK/ERK signaling pathway may mediate carnosine-induced microvascular changes in DR.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

YG designed the current study, wrote the manuscript and established the diabetic rat model. CG performed western blotting and statistical analysis and RT-qPCR. WH performed retinal immunofluorescence staining, drug treatment and cell identification. ZD performed RVEC isolation and cell culture.

Ethics approval and consent to participate

All animal experiments were performed in accordance with the recommendations included in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. Furthermore, the protocol of the current study was approved by the Committee on the Ethics of Animal Experiments of Tangdu Hospital, Air Force Medical University.

Competing interests

The authors declare that they have no competing interests.

References


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