Cytoprotective effect of amniotic membrane extracts on human corneal epithelial cells exposed to benzalkonium chloride in vitro

Jiayan Liu¹,²#, Huyong Zou¹#, Minzhi Zeng¹#, Amy Michelle Huang³, Yan Chen¹, Elaine Han³, Xiangyin Sha¹, Zhiping Liu¹

¹Ophthalmic Center, the Second Affiliated Hospital of Guangzhou Medical University, Guangzhou 510260, China; ²Department of Ophthalmology, the Sixth Affiliated Hospital of Guangzhou Medical University, Qingyuan 511518, China; ³Department of Ophthalmology, Bascom Palmer Eye Institute, University of Miami Miller School of Medicine, Miami, FL, USA

Contributions: (I) Conception and design: Z Liu, X Sha; (II) Administrative support: Z Liu, X Sha; (III) Provision of study materials or patients: J Liu, Z Liu, X Sha; (IV) Collection and assembly of data: J Liu, H Zou, M Zeng, Y Chen; (V) Data analysis and interpretation: J Liu, H Zou, M Zeng, Z Liu; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

#These three authors contributed equally, and should be considered as co-first authors.

Correspondence to: Zhiping Liu; Xiangyin Sha. Ophthalmic Center, the Second Affiliated Hospital of Guangzhou Medical University, Changgang East Road No. 250, Guangzhou 510260, China. Email: zxiu0001@hotmail.com; shaxiangyin@sina.com.

Background: The goal was to explore the protective effect and potential mechanism of amniotic membrane extracts (AME) on the ocular surface exposed to benzalkonium chloride (BAC).

Methods: The human corneal epithelial cell (HCEC) line SD-HCEC1s was cultured in 5 groups: normal control (NC), NC + AME, BAC, BAC + NC, and BAC + AME. Cell viability analysis, flow cytometry analysis, real-time polymerase chain reaction (PCR), and western blot were employed to measure changes in cell function. Matrix metalloproteinases (MMPs) and inflammatory cytokines were assayed by enzyme-linked immunosorbent assay (ELISA) and activity assays.

Results: Real-time PCR and western blot analysis demonstrated that the expression level of caspase-8 was increased while the levels of Muc1, Muc4, and Muc16 were decreased after treatment with 0.02% BAC for 1 h. When the SD-HCEC1s were withdrawn from the BAC and switched to media containing 10% AME for 2 days, the expression level of caspase-8 was decreased while the levels of Muc1, Muc4, and Muc16 were increased. Real-time PCR and ELISA demonstrated that the mRNA and protein levels of MMP-1, MMP-3, MMP-13, CXCL1, interleukin (IL)-1β, IL-6, and tumor necrosis factor-alpha (TNF-α) were significantly increased after treatment with 0.02% BAC, whereas those of MMP-8 were decreased. When the 0.02% BAC was withdrawn and the SD-HCEC1s were cultured in 10% AME, the mRNA and protein levels of MMP-1, MMP-3, MMP-13, CXCL1, IL-1β, IL-6, and TNF-α were decreased, while those of MMP-8 were increased. MMP-8 activity assays confirmed that IL-1β and TNF-α downregulated the protein levels of MMP-8.

Conclusions: AME protects SD-HCEC1s when stressed in BAC via upregulation of MMP-8 and downregulation of IL-1β and TNF-α. AME may have the potential functions to be employed as a topical adjunctive therapy in eyes chronically exposed to BAC.

Keywords: Amniotic membrane (AM); human corneal epithelial cell (HCEC); benzalkonium chloride (BAC); dry eye

Submitted 20 December, 2019. Accepted for publication 18 February, 2020.
doi: 10.21037/aes.2020.02.06
View this article at: http://dx.doi.org/10.21037/aes.2020.02.06
Introduction

Dry eye disease (DED) is a multifactorial disease of the ocular surface characterized by a loss of homeostasis of the tear film, and accompanied by ocular symptoms, in which tear film instability and hyperosmolarity, ocular surface inflammation and damage, and neurosensory abnormalities play etiological roles (1). Ocular irritation, stinging and foreign body sensation are symptoms frequently reported by patients which are exacerbated by substance such as benzalkonium chloride (BAC) (2).

Most of eye drops contain preservatives to prevent potential contamination, and the most common preservative used is BAC, which is present in eye drops generally ranges from 0.004% to 0.02% (3). BAC is cytotoxic to bacteria and to ocular surface epithelial cells. Therefore, there are a number of reports on the development of ocular surface disorders (OSD) in patients treated with BAC containing eye drops. BAC may induce ocular discomfort, tear film instability, superficial structure and integrity impairment, conjunctival inflammation, subconjunctival fibrosis, and epithelial apoptosis (2,4-8). The toxicity of BAC on cultivated corneal and conjunctival cells have been evaluated in many different species. It induces cytoplasmic damage and apoptosis, and induce DNA strand-breakage and mitochondrial dysfunction (4). BAC has been employed to investigate the pathogenesis of DED in animal models (9). In rabbits, topically applied 0.1% BAC twice daily over 4–14 days results in the development of DED (10,11). In mice, 0.2% BAC applied twice daily triggers clinical signs of DED including loss of conjunctival goblet cells (GCs), increased fluorescein staining, and corneal irregularity (5,12-14).

Amniotic membrane (AM) is avascular and a rich source of biologically active factors with low immunogenicity and as such, promotes healing and acts as an effective material for wound dressing. AME supports epithelialization and exhibits anti-fibrotic, anti-inflammatory, anti-angiogenic and anti-microbial features (5). In ophthalmology, AM and amniotic membrane extract (AME) are widely used for ocular surface reconstruction, including the treatment of persistent epithelial defects and non-healing corneal ulcers, corneal perforations and descemetoceles, bullous keratopathy, as well as corneal disorders with associated limbal stem cell deficiency, pterygium, conjunctival reconstruction, conneoscleral melts and perforations, and glaucoma surgeries (15). Compared to AM, AME can be conveniently applied to the ocular surface without the risk of suture-related complications of amniotic membrane transplantation (AMT) (5). Boiled AME has also been successfully used to ameliorate dry eye induced by BAC in a murine model (5). However, the boiled avenue may destroy some bioactive factors, such as proteins may be destroyed and modified. Although there are numbers of study of BAC toxicity on human corneal epithelial cells (HCECs), there is minimal information regarding the preclinically protective effect of AME on HCECs exposed to BAC.

Based on these beneficial effects of AM on the ocular surface, we hypothesized that AME might protect the corneal epithelium from BAC-induced toxicity. To test the hypothesis, we investigated the effects of 10% AME in cultures of BAC-damaged HCECs and its possible mechanisms.

Methods

Preparation of AME

The human tissues were collected and used after written consent and approval from the Second Affiliated Hospital of Guangzhou Medical University were obtained. The placent AME were obtained from healthy women with no infectious diseases including human immunodeficiency virus (HIV), hepatitis B virus (HBV), hepatitis C virus (HCV), syphilis, and with an uncomplicated pregnancy undergoing an elective cesarean to term (38–40 weeks’ gestation). The placent AME were washed with sterile normal saline and the AM was manually stripped from the chorion. After the AM was separated, it was washed 3 times in phosphate-buffered saline (PBS) with antibiotics (penicillin 50 mg/L, streptomycin 50 mg/L, neomycin 100 mg/L, amphotericin B) under aseptic conditions for approximately 15 minutes each time. After soaking in sterile saline at 2.5 mg/L for 10–20 minutes, the AM was added to an equal volume of PBS solution in the homogenizer, ground, and centrifuged at 1,000 rpm for 5 min to obtain AME. AME was acquired with a 0.22-µm filter and then stored in a −20 °C refrigerator. The protein content of the AME was measured by the Folin-phenol method.

Cell culture

HCEC lines were obtained from a corneal epithelial cell line spontaneously derived from human limbal tissue through serial culture (SD-HCECs) from the research group of Professor Wang Zhichong of Zhongshan Ophthalmic Center, Sun Yat-sen University (16). Thirty cell passages were utilized in this study. The cells were seeded at a density of 500/cm² in a 6-well plate (approximately 5,000 cells/well), and cultured in a 37 °C, 5% CO₂ incubator.

When the cultured cells reached 70% confluence, the
culture media were removed. Cells were incubated with BAC for 1 h at concentrations of 0.02% (the concentration was determined by the inhibition concentration 50 test based on the pre-experiment). To determine whether AME has the protective effects on BAC-damaged HCECs, AME at different concentrations of 50%, 20%, 10%, 2%, 1% were added to the cultures.

The experiment was divided into the following five groups: normal control (NC), NC + AME, BAC treatment untreated group (BAC), BAC treatment withdrawal (BAC + NC), BAC treatment withdrawal + AME group (BAC + AME), and NC + AME group.

**Apoptosis analysis**

Apoptosis and cell death were detected by flow cytometry on floating and adherent cells using an apoptosis detection kit (V13241, Invitrogen, Carlsbad, CA). The cells were collected, washed once with cold PBS, centrifuged at 1,000 rpm for 5 min, resuspended in 1× annexin binding buffer, and then incubated with Annexin V-antibody and propidium iodide (PI). After dark incubation for approximately 1 h, the final suspension was analyzed using a FACS Calibur flow cytometer (BD Biosciences, Shanghai, China) and the results were analyzed using CellQuest Pro software (BD Biosciences, Shanghai, China).

**Gene expression analysis**

Total RNA extraction (n=3) was performed using TRIzol Reagent (Invitrogen). The number of cells used in each group was approximately 10^6. The extraction method was carried out according to the instructions and quantified according to the method described previously (17). The primers were shown in Table 1. First, single-stranded cDNA was synthesized from 1 μg of total RNA using the SYBR Prime Script™ RT-PCR kit (DRR063S, Takara, Dalian, China), following the instructions. The mRNA expressions of caspase-8, Muc1, Muc4, Muc16, Matrix-metalloproteinases (MMPs)-1, -2, -3, -8, -9, -13 CXCL1, IL-1, and IL-6 were detected by the SYBR green system (DRR063S, Takara, Dalian, China) with real-time PCR. The cycling conditions were the following: an initial denaturation temperature of 95 °C for 2 min, then 95 °C for 10 s, then 60 °C for 34 s (fluorescence signal acquisition), then 72 °C for 30 s for a total of 45 cycles, and then lastly, a temperature increase from 60 °C to 98 °C to obtain a melting curve. The results were analyzed by the cycle threshold (CT) method and the housekeeping gene GAPDH was used as a reference.

**Western blot analysis**

According to previous studies (16,17), different groups of cells (approximately 10^5 cells in each group) were collected.

---

<table>
<thead>
<tr>
<th>Gene</th>
<th>Sense primer</th>
<th>Antisense primer</th>
<th>Tm (°C)</th>
<th>Product (bp)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Caspase 8</td>
<td>AGAGTCTGTGCCAAATCAAC</td>
<td>GCTGCTTCTCTTTGCTGAA</td>
<td>58</td>
<td>78</td>
</tr>
<tr>
<td>Muc1</td>
<td>TGCCGGCAAGAAGTAGCT</td>
<td>TGGGGTACTCGCATTAGGAT</td>
<td>60</td>
<td>76</td>
</tr>
<tr>
<td>Muc4</td>
<td>GGAGAGGATCCCGCTGATAG</td>
<td>CCGGTAGCCTGATACTG</td>
<td>59</td>
<td>76</td>
</tr>
<tr>
<td>Muc16</td>
<td>CCAGTCCATACCTCGGTT</td>
<td>AGGGTAGTCTAGAGGGGTT</td>
<td>60</td>
<td>161</td>
</tr>
<tr>
<td>MMP-1</td>
<td>AAAATTACAGCCTGGATTGCC</td>
<td>GGTGTGACATTCCCTAGG</td>
<td>59</td>
<td>82</td>
</tr>
<tr>
<td>MMP-3</td>
<td>CTGACTCCGACACTCTGGA</td>
<td>CAGGAAAGGTTCTGAGATGACC</td>
<td>60</td>
<td>79</td>
</tr>
<tr>
<td>MMP-8</td>
<td>TGCTTACTCCATGTGCAGA</td>
<td>TCCAGGTAGTCCCTGACAGTTT</td>
<td>59</td>
<td>85</td>
</tr>
<tr>
<td>MMP-13</td>
<td>ACTGAGAGGCTCGAGAATG</td>
<td>GAACCCCGGATCTTGCT</td>
<td>60</td>
<td>103</td>
</tr>
<tr>
<td>IL-1β</td>
<td>ATGATGGCTATTACATGAG</td>
<td>GTCCGGAGATCGCTGGA</td>
<td>59</td>
<td>132</td>
</tr>
<tr>
<td>IL-6</td>
<td>CCTGACACTTTCAAGATG</td>
<td>TTTGACGGCAAAGTCTC</td>
<td>61</td>
<td>75</td>
</tr>
<tr>
<td>CXCL1</td>
<td>TGCTTACCAACAAGGGGGTTA</td>
<td>AAAGGGCCCTTGTTCCC</td>
<td>59</td>
<td>110</td>
</tr>
<tr>
<td>TNFα</td>
<td>CAGCCCTCTCTCTCCCTGAT</td>
<td>GCCCAGGCAGTCAGAT</td>
<td>59</td>
<td>139</td>
</tr>
</tbody>
</table>

RT-PCR, real-time polymerase chain reaction.

---

Table 1 Primer sequences used for RT-PCR
for western blot analysis. Cells at passage 30 were used to detect the expressions of the related proteins. The primary antibodies used were mouse anti-human caspase-8 (1:100, Santa Cruz Biotechnology, Santa Cruz, CA), mouse anti-human Muc1 (1:100, Abcam, Cambridge, UK), mouse anti-human Muc4 (1:100, Abcam, Cambridge, UK), and mouse anti-human Muc16 (1:100, Abcam, Cambridge, UK). β-actin (1:3,000, Cell signaling, USA) was used as positive control. The secondary antibodies used were horseradish peroxidase (HRP)-conjugated goat anti-mouse immunoglobulin G (IgG) and goat anti-rabbit IgG (Sigma-Aldrich, St. Louis, MO). Localization of antibodies was detected by a chemical luminescence using a ECL kit (Amersham, USA) following the manufacturer’s instructions. The images were analyzed using the Gel-Pro analyzer (Media Cybernetic, Inc.) for statistical analysis.

The expression of inflammatory cytokines by an enzyme-linked immunosorbent assay (ELISA)

ELISA (Biosource, San Francisco, CA, USA) were employed to detect the expression of MMP-1, -3, -8, -13, CXCL1, IL-1β, IL-6, and TNF-α. ELISA procedures were performed following the manufacturer’s protocol.

The NC + 10% AME group was treated with IL-1β (0.1, 1.0, or 10.0 ng/mL) or TNF-α (0.1, 1.0, or 10.0 ng/mL) for 24 h to analyze the activity levels of MMP-8 protein in the supernatants of SD-HCEC1s cells. After that, the supernatants were collected and centrifuged before assays. The supernatants, pro-MMP-8 standard, and assay buffer were incubated in the 96-well plates which were pre-coated with anti-MMP-8 antibody at 4 °C for 12 hours. Then, the activity levels of MMP-8 protein were assayed using the commercial kits (Biotrak, Amersham).

Statistical analysis

All values are presented as mean ± standard deviation (SD). All analyses were evaluated with the analysis of variance followed by Boferroni’s post hoc test using IBM SPSS Statistics 25 (IBM Corp., Armonk, NY, USA). In cases of nonparametric data distribution, the Kruskal-Wallis test was performed, followed by the Mann-Whitney U test. A probability value of 0.05 (P<0.05) was considered to be statistically significant.

Results

AME enhances cell viability and inhibits apoptosis of SD-HCEC1s treated with BAC

As shown in Figure 1, cell proliferation assays showed that SD-HCEC1s in 10% AME possessed relatively strong growth capacities as compared to those in the 50% AME, 20% AME, 2% AME, and NC groups. These results showed that the SD-HCEC1s possessed a higher proliferative capacity in 10% AME after withdrawal of the 0.02% BAC.

Morphologically, the SD-HCEC1s treated with 0.02% BAC exhibited relatively more vacuolated cells (Figure 2). The SD-HCEC1s in the 0.02% BAC 1 h + 10% AME group exhibited relatively lower vacuolated cells than the cells in the 0.02% BAC group.

Flow cytometry revealed that the percentage of apoptotic cells in the 0.02% BAC, 0.02% BAC 1 h + NC, 0.02% BAC 1 h + 10% AME, NC, NC + 10% AME groups were 74.13%±4.00%, 52.60%±1.68%, 29.52%±0.72%, 20.13%±5.04%, 18.46%±4.73%, respectively. Compared to the continuously cultured 0.02% BAC group, the 0.02% BAC 1 h + 10% AME possessed fewer apoptotic cells (P<0.001, n=3) (Figure 3A,B). Compared to the NC groups, SD-HCEC1s cultured in the NC + 10% AME groups exhibited fewer apoptotic cells (P<0.001, n=3).
AME downregulates the expression of caspase-8 and enhances the maintenance of the functional properties of mucins in SD-HCEC1s

The mRNA expression of caspase-8 was significantly increased after adding 0.02% BAC to SD-HCEC1s for 1 h. However, the level of mRNA transcripts for caspase-8 was detected at lower levels in the 0.02% BAC 1 h + 10% AME group (P<0.001, n=3) (Figure 3C). The mRNA expression levels of Muc1, Muc4, and Muc16 were significantly decreased after adding 0.02% BAC for 1 h (P<0.05, n=3), while the mRNA levels of Muc1, Muc4, and Muc16 were dramatically increased (P<0.01, n=3) in 0.02% BAC 1 h + 10% AME group (Figure 4A). Western blot showed that the protein expression of caspase-8 was significantly increased in 0.02% BAC group (P<0.05, n=3). The protein expression levels of Muc1, Muc4, and Muc16 were significantly decreased after SD-HCEC1s were exposed to BAC, while the protein expression of Muc1, Muc4 and Muc16 were dramatically increased in the 0.02% BAC 1 h + 10% AME group (P<0.01, n=3) (Figure 4B,C).

To evaluate the effect of AME on the regulation of the inflammatory cytokines in SD-HCEC1s exposed to BAC, cytokines MMP-1, MMP-3, MMP-8, MMP-13, CXCL1, IL-1β, IL-6, and TNF-α were examined. Compared to the cells exposed to 0.02% BAC, the mRNA expression levels of MMP-1, -3, -13, IL-1, IL-6, TNF-α were significantly reduced in 0.02% BAC 1 h + 10% AME group (P<0.01, n=3). Interestingly, the expression levels of MMP-8 were increased in 0.02% BAC 1 h + 10% AME group (P<0.01, n=3) (Table 2). The concentrations of MMP-1, MMP-3, MMP-13, CXCL1, IL-1β, IL-6, and TNF-α proteins were significantly higher in the 0.02% BAC group than the cells cultured in both the NC group (P<0.001, n=3) and 0.02% BAC 1 h + 10% AME group (P<0.05, n=3). MMP-8 levels in NC + 10% AME were higher than the cells in NC group (P<0.001, n=3). Compared to NC group, the concentrations...
Figure 3 Changes of apoptosis in SD-HCEC1s in each group. (A) Representative flow cytometry images of SD-HCEC1s in each group; (B) the total percentage (%) of apoptotic cells in each group; (C) quantitative real-time PCR showed the expression levels (relative fold of mRNA) of caspase-8 in each group. Data are expressed as mean ± SE. Q1, death cells; Q2, later apoptotic cells; Q3, early apoptotic cells; Q4, survival cells. PCR, polymerase chain reaction.

of MMP-1, MMP-3, MMP-13, CXCL1, IL-1β, IL-6, and TNF-α proteins in NC + 10% AME were significantly lower (P<0.001, n=3) (Table 3).

MMP-8 activity in the conditioned media was significantly reduced after treatment with 0.1, 1, and 10 ng/mL of IL-1β, respectively. Similarly, MMP-8 activity was inhibited by TNF-α. The concentrations of MMP-8 in the control group, 0.1, 1, and 10 ng/mL of TNF-α were 29.49±0.59, 22.74±1.31, 14.38±1.46, and 7.63±0.23 ng/mL, respectively (Figure 5).
Figure 4 The expression levels of mucins and caspase-8 in each group. (A) Quantitative real-time PCR showed the expression levels (relative fold of mRNA) of Muc1, Muc4, and Muc16 in each group; (B,C) Western blot results showed the changes in levels of caspase-8, Muc1, Muc4, and Muc16 in each group. Data are expressed as mean ± SE. AME, amniotic membrane extracts; BAC, benzalkonium chloride; NC, normal control.

Discussion

The current study demonstrated that BAC decreased HCECs viability and proliferation, and increased cellular apoptosis. Addition of 10% AME remarkably protected HCECs against BAC-induced toxicity. The key finding is that the protective effect of AME on SD-HCEC1s exposed to BAC was mediated partially by upregulation of MMP-8 and downregulation of IL-1β and TNF-α.

DED is well documented that typical pathologic alterations of dry eye involve ocular surface inflammation, conjunctival GC loss, corneal epithelium squamous metaplasia, and epithelial disruption. In DED, the inflammations may play vital roles (18,19). Interleukin (IL)-1β and tumor necrosis factor (TNF)-α both play important roles in the mechanisms of DED (20,21). IL-6 plays a vital role in the induction of acute inflammation which is generated by macrophages and T cells. Bian et al. found that an increase in the expressions and activities of MMPs-
Table 2 The mRNA expression levels of cytokines in SD-HCEC1s in each group

<table>
<thead>
<tr>
<th>Variables</th>
<th>0.02% BAC</th>
<th>0.02% BAC 1 h + NC</th>
<th>0.02% BAC 1 h + 10% AME</th>
<th>NC</th>
<th>NC + 10% AME</th>
</tr>
</thead>
<tbody>
<tr>
<td>MMP-1</td>
<td>1.28±0.10^a</td>
<td>1.12±0.06^d</td>
<td>0.85±0.03^b</td>
<td>1.00±0.01^c</td>
<td>0.73±0.03^c</td>
</tr>
<tr>
<td>MMP-3</td>
<td>1.37±0.11^a</td>
<td>1.13±0.09^d</td>
<td>0.86±0.04^b</td>
<td>1.00±0.01^c</td>
<td>0.85±0.03^c</td>
</tr>
<tr>
<td>MMP-8</td>
<td>0.84±0.02</td>
<td>0.90±0.04^d</td>
<td>1.07±0.02^b</td>
<td>1.00±0.01^c</td>
<td>1.34±0.04^b</td>
</tr>
<tr>
<td>MMP-13</td>
<td>1.32±0.07^a</td>
<td>1.06±0.09</td>
<td>0.91±0.07^b</td>
<td>1.00±0.01^c</td>
<td>0.68±0.03^c</td>
</tr>
<tr>
<td>CXCL1</td>
<td>1.06±0.08^a</td>
<td>0.85±0.07</td>
<td>0.76±0.04^b</td>
<td>1.00±0.01^c</td>
<td>0.53±0.03^c</td>
</tr>
<tr>
<td>IL-1β</td>
<td>1.29±0.06^a</td>
<td>0.94±0.08</td>
<td>0.88±0.09^b</td>
<td>1.00±0.01^c</td>
<td>0.62±0.07^a</td>
</tr>
<tr>
<td>IL-6</td>
<td>1.26±0.07</td>
<td>1.21±0.08^d</td>
<td>1.01±0.05^b</td>
<td>1.00±0.01^c</td>
<td>0.71±0.04^c</td>
</tr>
<tr>
<td>TNF-α</td>
<td>1.19±0.06^a</td>
<td>0.84±0.08</td>
<td>0.78±0.09^b</td>
<td>1.00±0.01^c</td>
<td>0.52±0.07^c</td>
</tr>
</tbody>
</table>

^a, P<0.05 0.02% BAC vs. 0.02% BAC 1 h + NC; ^b, P<0.05 0.02% BAC vs. 0.02% BAC 1 h + 10% AME; ^c, P<0.05 0.02% BAC vs. NC; ^d, P<0.05 0.02% BAC 1 h + NC vs. 0.02% BAC 1 h+10% AME; ^e, P<0.05 NC vs. NC + 10% AME. Data were expressed as mean ± SD. AME, amniotic membrane extracts; BAC, benzalkonium chloride.

Table 3 Comparisons of MMPs and inflammatory cytokines in each group

<table>
<thead>
<tr>
<th>Variables</th>
<th>0.02% BAC</th>
<th>0.02% BAC 1 h + NC</th>
<th>0.02% BAC 1 h + 10% AME</th>
<th>NC</th>
<th>NC + 10% AME</th>
</tr>
</thead>
<tbody>
<tr>
<td>MMP-1 (ng/mL)</td>
<td>77.01±6.00^a</td>
<td>40.32±2.16^d</td>
<td>20.40±0.72^b</td>
<td>19.00±0.19^c</td>
<td>17.21±0.17^c</td>
</tr>
<tr>
<td>MMP-3 (ng/mL)</td>
<td>73.77±5.94^a</td>
<td>36.61±2.92^d</td>
<td>18.58±0.86^b</td>
<td>17.10±0.17^c</td>
<td>15.52±0.15^c</td>
</tr>
<tr>
<td>MMP-8 (ng/mL)</td>
<td>28.50±0.29^a</td>
<td>33.48±1.80^d</td>
<td>58.32±2.59^b</td>
<td>53.81±1.28^c</td>
<td>60.13±1.44^c</td>
</tr>
<tr>
<td>MMP-13 (pg/mL)</td>
<td>75.42±7.80^a</td>
<td>38.16±3.24^d</td>
<td>21.84±1.68^b</td>
<td>19.00±0.19^c</td>
<td>17.21±0.17^c</td>
</tr>
<tr>
<td>CXCL1 (ng/mL)</td>
<td>47.57±3.60^a</td>
<td>23.02±1.89^d</td>
<td>13.74±0.72^b</td>
<td>14.25±0.14^c</td>
<td>12.98±0.13^c</td>
</tr>
<tr>
<td>IL-1β (ng/mL)</td>
<td>77.54±3.60^a</td>
<td>33.84±2.88^d</td>
<td>21.12±2.16^b</td>
<td>19.00±0.19^c</td>
<td>17.21±0.17^c</td>
</tr>
<tr>
<td>IL-6 (ng/mL)</td>
<td>94.50±5.25^a</td>
<td>54.26±3.60^d</td>
<td>30.30±1.50^b</td>
<td>23.75±0.24^c</td>
<td>21.44±0.21^c</td>
</tr>
<tr>
<td>TNF-α (ng/mL)</td>
<td>53.66±2.70^a</td>
<td>22.68±2.16^d</td>
<td>14.04±1.62^b</td>
<td>12.83±0.14^c</td>
<td>11.71±0.13^c</td>
</tr>
</tbody>
</table>

^a, P<0.05 0.02% BAC vs. 0.02% BAC 1 h + NC; ^b, P<0.05 0.02% BAC vs. 0.02% BAC 1 h + 10% AME; ^c, P<0.05 0.02% BAC vs. NC; ^d, P<0.05 0.02% BAC 1 h + NC vs. 0.02% BAC 1 h+10% AME; ^e, P<0.05 NC vs. NC + 10% AME. Data were expressed as mean ± SD. AME, amniotic membrane extracts; BAC, benzalkonium chloride; NC, normal control.

1, -2, -3, -8, -9, and -13 were positively associated with the severity of corneal diseases. Additionally, in corneal wound healing procedures, the generations and activities of MMPs have elevated (22-24).

BAC has been the most common preservative used in eye drops, especially in anti-glaucoma medications (6,7). BAC acts as a surfactant to solubilize ionic components into immiscible solvents, which facilitates effective emulsification and stabilization of medications, and prolongs shelf-life. Further, BAC is superior to other preservatives in inhibiting microbial activity (3,25,26). The half-life of BAC is approximately 20 hours in the corneal epithelium and conjunctiva, which may lead to chronic ocular surface damage in patients using BAC-containing eye drops (3,5,27).

Prolonged exposure to BAC caused toxicity to the ocular surface. Preservative-free eye drops are the ultimate choice for ocular surface disease. However, preservative eye drops had a risk of microbial contamination when used more than once over a period of 10 hours. It is critical to manage BAC-induced ocular surface toxicity. Our study showed that AME had beneficial effects in preventing and treating BAC-induced toxicity in HCECs (27). AM can inhibit the release of inflammatory mediators and their matrix surface can also capture inflammatory effects when covering inflammatory AME had beneficial effects in preventing and treating BAC-induced toxicity in HCECs (27). AM can inhibit the release of inflammatory mediators and their matrix surface can also capture inflammatory effects when covering inflammatory effects when covering...
the suture-related risk. It is generally believed that AME has effective components, including HA-HC complex (hyaluronic and the heavy chains of inter-α-inhibitor, which are responsible for anti-inflammatory, and anti-scarring), fibronectins, peptides, small proteins, and growth factors (15,31-33). Unlike boiled method, we employed homogenized and filtered method to acquire AME, which can possess almost all the effective components of AM maximally. This work demonstrated that AME protected SD-HCEC1s when stressed in BAC via upregulation of MMP-8 and downregulation of IL-1β and TNF-α.

In summary, this study suggested that AME provided protection for SD-HCEC1s exposed to BAC. AME maintained the normal function of SD-HCEC1s by increasing cellular activity, inhibiting apoptosis, and maintaining mucin secretion via upregulation of MMP-8 and downregulation of IL-1β and TNF-α. AME may have the potential to be employed as a topical adjunctive therapy in eyes chronically exposed to BAC.

**Acknowledgments**

**Funding:** This study was supported by the “Yangcheng Scholar” Youth Research Backbone Training Project of Guangzhou Municipal College (No. 1201581612), Guangzhou Science and Technology Project (No. 201804010038), and Guangdong Natural Science Foundation of China (2020A1515010276; No. 2015A030313479).

**Footnote**

**Conflicts of Interest:** The authors have no conflicts of interest to declare.

**Ethical Statement:** The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The human tissues were collected and used after written consent and approval from the Second Affiliated Hospital of Guangzhou Medical University were obtained.

**References**


31. He H, Li W, Tseng DY, et al. Biochemical characterization and function of complexes formed by hyaluronan and the


doi: 10.21037/aes.2020.02.06