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Research Paper

Cordycepin attenuates Salivary Hypofunction through the Prevention of Oxidative Stress in Human Submandibular Gland Cells

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Abstract

Xerostomia (dry mouth) is a significant age-related condition. Meanwhile, cordycepin, the natural therapeutic agent, has demonstrated an anti-aging effect. Therefore, the present study aimed to investigate the preventive effects of cordycepin on secretory function in an in vitro model of hydrogen peroxide (H_2O_2) -induced salivary hypofunction. After being exposed to H_2O_2 , human submandibular gland (HSG) cells were treated with various concentrations of cordycepin (6.25-50 μ M) for 24, 48, and 72h. To evaluate cell proliferation and reactive oxygen species (ROS) generation, 3-(4, 5-dimethylthiazol-2-yl)-2, 5-diphenyltetrazolium bromide and 2, 7'-dichlorodihydrofluorescein diacetate assays were performed. The amylase activity was kinetically measured by 2-chloro-p-nitrophenol linked with maltotrioside. The expression of salivary, antioxidant and apoptotic markers at mRNA and protein levels were performed by reverse transcriptase polymerase chain reaction (RT-PCR) and immunofluorescence analysis, respectively. We demonstrated that cordycepin (6.25-25 µM) contributed to significant increases in expression of the salivary marker genes, alpha-amylase I (AMYIA) and aquaporin-5 (AQP5), and in amylase secretion without changes in cell viability. Under oxidative stress, HSG cells showed remarkable dysfunction. Cordycepin rescued the protective effects partially by decreasing ROS generation and restoring the expression of the salivary proteins, AMY and AQP5 via anti-oxidant and anti-apoptotic activity. In addition, the amount of amylase that was secreted from HSG cells cultured in cordycepin was increased. In conclusion, cordycepin demonstrated a protective effect on H₂O₂-induced HSG cells by decreasing ROS generation and upregulating the salivary function markers, AMY1A and AQP5, at both the transcriptional and translational levels.

Key words: cordycepin; salivary gland; xerostomia; saliva; dry mouth

Introduction

Xerostomia, also referred to as dry mouth, is dryness of the oral cavity caused by a low salivary flow rate, lack of saliva and change in the chemical composition of saliva [1, 2]. Therefore, salivary gland dysfunction or salivary hypofunction causes dry mouth, which can lead to several clinical conditions, including mastication problems and a decline in the quality of life, especially for patients who have received radiation as part of cancer therapy and elderly persons [3, 4]. Furthermore, salivary hypofunction is defined as an objective reduction in the salivary flow rate involving physiological atrophy of the salivary gland, which manifests itself as morphological alteration in the gland [5, 6].

Reactive oxygen species (ROS) are one of main causes of oxidative stress, contributing to cell aging, and are strongly associated with age-related diseases [7]. ROS and other free radicals are usually reduced by antioxidants in the body, the so-called oxidation-reduction regulatory mechanism; hence, the accumulation of ROS within the body is repressed [8, 9]. However, antioxidant activity declines with age by interfering with the redox balance, resulting in protein denaturation, DNA damage, lipid peroxidation and apoptosis [10]. Thus, oxidative stress can probably affect the secretory function of salivary glands in the elderly [11]. Nevertheless, the relationship between age-related salivary hypofunction and oxidative stress has not been defined [12].

Cordycepin, also known as 3-deoxyadenosine, is one of the major bioactive compounds of Cordycepes sp., exerting various pharmaceutical properties (e.g. antitumor, anticancer and immunoregulatory effects [13, 14]. Furthermore, the antioxidant activity of cordycepin has been recently studied [15]. In addition, cordycepin could protect cells against oxidative stress, which induces cell damage. Cordycepin has also been demonstrated to inhibit mitochondrial injury and improve immune responses by scavenging ROS [16, 17]. Previous studies have reported that cordycepin inhibits ROS generation and protects several cells (i.e. neuron and mesenchymal stem cells) from oxidative stress [18-20]. Additionally, cordycepin could have antioxidant activity and attenuate oxidative stress in vivo [15].

However, the preventive effect of cordycepin on the dry mouth experimental model is still unclear. This research aimed to examine the effect of cordycepin on improvements in salivary secretion of H_2O_2 -induced salivary hypofunction with xerostomia in human submandibular gland (HSG) cells *in vitro*.

Materials and Methods

Chemicals and reagents

All the chemicals and reagents used in this experiment, including cordycepin (MW=251.2 g/mol, 3-4,5-dimethyl-2-thiazolyl-2,5-diphenyl-2H-tetrazoliu m bromide (MTT), dimethyl sulfoxide (DMSO), 2',7'-dichlorodihydrofluorescein diacetate (DCFHDA), Hydrogen peroxide (H₂O₂) and amylase activity assay kits, were purchased from Sigma-Aldrich Co. (St. Louis, MO). Dulbecco's Modified Eagle Medium (DMEM), non-essential amino acids (NAAs) and foetal bovine serum (FBS) were purchased from HyClone (Logan, UT), whereas 0.5% trypsin-EDTA and L-glutamine were obtained from Gibco (Gibco, CA, USA).

Cell culture

HSG cell line used in this study was obtained from the American Type Culture Collection (ATCC® HTB-41TM, Manassas, VA, USA). Cells were cultured in a DMEM medium supplemented with 10% (v/v) FBS, 1% (v/v) NAAs, 1% (v/v) L-glutamine and 1% (v/v) antibiotics (penicillin-streptomycin). Cells were cultured at 37 °C under an atmosphere of 5% CO_2 with 95% humidity. The culture medium was replaced thrice a week.

Determination of cell viability

Cell viability was evaluated by MTT assay. Briefly, HSG cells were seeded into 96-well tissue culture plates at a density of 1×104 cells/well overnight, then the cells were treated in three different experiments; 1) treated with various concentrations of cordycepin (6.25-50 µM) for 24, 48, and 72 h; 2) treated with different concentrations of H_2O_2 (250-2000 μ M) to induced cell toxicity for 30 min; 3) treated with H₂O₂ for 30 min and cordycepin ranging from 6.25-50 µM for 24 h. After each treatment, the cells were washed with 100 µl of 1× PBS, then 0.5 mg/ml of MTT solution in 1× PBS was added to each well and incubated at 37 °C for 4 h. After removal of the MTT solution, the formazan crystals were dissolved with 100 µl of DMSO, and the purple formazan solution was further measured for absorbance at 570 nm using a microplate reader (BMG Labtech, Ortenbreg). The percentage of cell viability was calculated according to the following equation:

%*Cell viability* =
$$\frac{A}{B} \times 100$$

where A was the absorbance of OD of the treated group, and B was the absorbance of OD of the untreated group.

Determination of ROS generation

Intracellular ROS generation was determined by DCFH-DA assay. Briefly, after the seeding of HSG cells, the cells were incubated with 500 μ M H₂O₂ for 30 min to induce oxidative stress in cells, then washed with 1× PBS and added to cordycepin at various concentrations for 24 h. After being exposed to cordycepin, the cells were stained with 20 μ M DCFH-DA at 37 °C for 30 min in the dark. Afterwards, the fluorescence intensity was measured at an excitation/emission of 485/535 nm using a fluorescence microplate reader (Thermo Scientific Varioskan, USA).

RT-PCR analysis

After 24 h, total RNA was extracted from untreated and treated HSG cells in each well using the NucleoSpin RNA kit (Macherey-Nagel, Dueren, Germany), following the manufacturer's instructions. Complementary DNA was synthesized from 1 μ g of total RNA using the ReverTra Ace qPCR Master Mix kit (Toyobo CO., LTD., Japan), according to the manufacturer's instructions. PCR analysis was performed to determine the expression of salivary-specific genes, alpha-amylase 1 (AMY1A) and aquaporin-5 (AQP5), the expression of antioxidant genes, superoxide dismutase (SOD1), catalase (CAT), glutathione peroxidase (GPX1), and the expression of apoptotic genes, caspase-3 (CASP-3) and B-cell leukemia/lymphoma 2 (BCL-2). The mRNA levels of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were employed as internal controls. The primer sequences and RT-PCR conditions were shown in Table 1. The PCR products were identified by electrophoresis on 1.5% agarose gel and visualized by ethidium bromide staining. The mRNA band density of each gene was analyzed and quantified using densitometer and ImageJ software from the NIH website and shown as the mean ± SD of the results from three independent experiments. Each band image was calculated for the total band density. The relative density of genes of interests and GAPDH was calculated by dividing the density of each gene by the density of GAPDH of the same sample. Lastly, the relative gene expression for the treated group was plotted as a fold-change normalized to the untreated control.

Immunofluorescence staining

Cells were grown on the surface of coverslips in 24-*well plates* until a confluence of 80%. After being treated with cordycepin, cells were fixed with 4% paraformaldehyde (PFA) solution for 30 min at room temperature and rinsed with 1× PBS twice. Cells were permeabilized and blocked in blocking buffer containing 1% bovine serum albumin (BSA), 1% Triton X-100 of 1× PBS at 4 °C for 1 h for amylase and 4% BSA, 0.1% Triton X-100 of 1× PBS at 4 °C for 30 min for AQP5. After cell permeabilization, the cells were stained into each well, then incubated overnight at 4 °C with anti-alpha-amylase (anti-AMY) antibody

Table 1. PCR conditions and primers used in RT-PCR analysis

(1:100 dilution; Santa Cruz Biotechnology Inc.) and ant-aquaporin-5 (anti-AQP5) antibody (1:100 dilution; Sigma-Adrich Co.), then washed with 1× PBS twice. After being washed with PBS, the cells were incubated at room temperature for 1 h with Rhodamine Red-X goat anti-mouse immunoglobulin G (IgG) 1:500 dilution; Invitrogen) for anti-AMY or AlexaFluor 488 goat anti-rabbit IgG (1:1000 dilution; Invitrogen) for anti-AQP5. Lastly, the cells were incubated with 4 µl of 4'6-diamidino-2-phenylindole (DAPI) (Invitrogen) for nuclear staining and observed under a ZOE[™] Fluorescent Cell Imager (Bio-rad, CA, USA). The fluorescence intensity was quantified by ImageJ software. With RGB images, results are calculated using brightness values by the software will convert the RGB pixels to brightness values. To rule out the bias during the determination, we selected two areas of interest with approximately similar cell numbers represented by DAPI nuclear staining. All measurements of each image were set with the same parameters including the subtraction of fluorescence background. The fluorescence intensity was presented as a fold change which was relative to the untreated (control) group.

Amylase secretion

Amylase secretion was measured by biochemical assay. After cell culture periods, the culture medium was collected from each well. Amylase secretion was measured using the amylase activity assay kit (Sigma-Andrich Co.), according to the manufacturer's protocol. Amylase activity was measured using a microplate reader at 405 nm. The kinetic reaction of amylase activity was calculated from the change in absorbance per 30 s for 2 h.

Genes	Direction	Primer sequences $(5' \rightarrow 3')$	Annealing Temperature (°C)	Cycles	Product size (bp)
AMY1A	Forward	AATTGATCTGGGTGGTGAGC	52	35	474
	Reverse	CTTATTTGGCGCCATCGATG			
AQP5	Forward	CCTGTCCATTGGCCTGTCTGTCAC	56	40	249
	Reverse	GGCTCATACGTGCCTTTGATGATG			
CAT	Forward	TCCGGGATCTTTTTAACGCCATTG	55	30	362
	Reverse	TCGAGCACGGTAGGGACAGTTCAC			
SOD1	Forward	CTAGCGAGTTATGGCGAC	52	30	224
	Reverse	CATTGCCCAAGTCTCCCA C			
GPX1	Forward	CGCCAAGAACGAAGAGATTC	54	30	272
	Reverse	CAACATCGTTGAGACACA C			
Caspase-3	Forward	TTTGTTTGTGTGCTTCTGAGCC	54	40	279
	Reverse	ATTCTGTTGCCACCTTTCGG			
BCL - 2	Forward	CGCATCAGGAAGGCTAGAGT	53	40	189
	Reverse	AGCTTCCAGACATTCGGAGA			
GAPDH	Forward	ACCTGACCTGCCGTCTAGAA	54	35	247
	Reverse	TCCACCACCCTGTTGCTGTA			



500 µM H₂O₂+12.5 µM Cor

500 µM H₂O₂+25 µM Cor

500 µM H₂O₂+50 µM Cor

Figure 1. Cordycepin prevented HSG cells from H₂O₂-induced cell toxicity. Cells were treated with various concentrations of cordycepin (6.25-50 μ M) for 24, 48 and 72 h (**A**,**B**) and different concentrations of H₂O₂ (250-2000 μ M) were used to induced cell toxicity for a 30 min culture period (**C**). The post-treatment of HSG cells under 500 μ M H₂O₂ for 30 min and cordycepin ranging from 6.25-50 μ M for 24 h was measured by MTT assay (**D**). Cell morphology of HSG cells exposed to H₂O₂ and cordycepin under an inverted microscope (**E**). The scale bar represents 20 μ m. *p<0.05 and **p<0.01 compared to the untreated group, and ##p<0.01 compared to the H₂O₂ group.

Statistical analysis

Data were presented as the means \pm SDs of three independent measurements (n=3). All analyses were performed using the Statistical Package for the Social Sciences (SPSS) software 23.0 for Windows. For comparison of the treatments, statistical significance was assessed by student's t-test to compare means from two independent sample groups at *P-value* (*p*<0.05).

Results

Cordycepin prevents against H₂O₂-induced toxicity in HSG cells

In this study, we first performed the cytotoxicity test of cordycepin on HSG cells. The HSG cells were treated with various concentrations of cordycepin (6.25-50 μ M) for 24, 48, and 72 h and then measured by MTT assay. As shown in Figure 1A, the HSG cells

exposed to most concentrations of cordvcepin at 24 h did not show their viability alteration. Nevertheless, the highest cordycepin concentration (50 µM) was slightly toxic in HSG cells (19.66% ± 2.45). These suggests that cordycepin especially at low concentrations did not display a toxic effect to the HSG. In addition, the cell viability at different time points (24, 48, and 72 h) showed that % cell viability of HSG remains unchanged at various time points, indicating that cordycepin had no proliferative effects on HSG cells (Figure 1B). The percentage of viable HSG cells was considerably decreased upon exposure to H_2O_2 (Figure 1C), indicating cytotoxicity. Furthermore, we investigated the preventive effects of cordycepin against H₂O₂-induced cell toxicity. The post-treatment group with cordycepin under H₂O₂ revealed significantly increased cell viability when compared to the H₂O₂ group (Figure 1D). Moreover, the post-treatment group with cordycepin in HSG cells showed cell morphology similar to that of the untreated group (Figure 1E).

Cordycepin promotes the expression of salivary genes in H_2O_2 -induced HSG cells

HSG cells were cultured in each concentration of cordycepin treatment for a 24 h period; salivary gene

expression was then comparatively assessed. The salivary-specific gene expression, AMY1A and AQP5, of the cultured HSG cells was determined by RT-PCR analysis. The band intensities of mRNA expression of AMY1A and AQP5 in each cordycepin concentrationtreated HSG cells were demonstrated (Figure 2A). In cordycepin concentrations (6.25, 12.5, 25 µM), the relative expression of AMY1A gradually increased as compared to that found in the untreated group. In particular, 12.5 µM of cordycepcin significantly increased AMY1A expression (Figure 2B). The expression of AQP5 detected in the 12.5 µM of cordycepin group was also higher than that detected in the untreated group (Figure 2C). Interestingly, the increase in salivary-specific gene expression observed among the cells cultured in the cordycepin treatments were much different from one another. In addition, cordycepin had protective effect on H2O2-induced HSG cell dysfunction, the gene expression demonstrated that all cordycepin concentrations significantly increased the levels of AMY1A and AOP5 in H₂O₂-induced HSG cells compared to the induced cells without the cordycepin treatment (Figures 2D-F), suggesting that cordycepin could rescue the salivary function after oxidative stress exposure).



Figure 2. Cordycepin upregulated salivary marker genes in H_2O_2 -induced HSG cells. Cells were treated with cordycepin ranging from 6.25 µM to 50 µM for 24 h. The mRNA expression for AMYIA and AQP5 were analysed by RT-PCR (A-C). Cordycepin promoted AMYIA and AQP5 expression in HSG cells exposed to H₂O₂ for 30 min (D-F). The relative mRNA expression levels of AMYIA (B-E) and AQP5 (C-F) genes were evaluated by image J NIH software and normalized with GAPDH gene. Gel electrophoresis results are from one representative experiment and bar charts are derived from analysis of relative expression from three independent experiments. *p<0.05 and **p<0.01 compared to the untreated group, and ##p<0.01 compared to the H₂O₂ group.



Figure 3. Cordycepin rescued the expression of salivary proteins in H_2O_2 -induced HSG cells. Both AMY (A) and AQP5 (C) protein expression in HSG cells were determined by immunofluorescence. Cells were treated with cordycepin at either 0 μ M (a) or 12.5 μ M (b) for 24 h, treatment with 500 μ M H₂O₂ for 30 min without (c) or with 12.5 μ M cordycepin for 24 h (d). The relative fluorescence intensity of both AMY (B) and AQP5 (D) proteins in cordycepin-treated HSG cells were normalized with the untreated group. Immunofluorescence images are from one representative experiment. Bar charts were expressed as mean ± SD from three independent experiments. The scale bar represents 40 μ m. **p<0.01 compared to the untreated group, and ##p<0.01 compared to the H₂O₂ group.

Cordycepin alters both AMY and AQP5 protein expressions in H₂O₂-induced HSG cells

The expression of AMY and AQP5 proteins in HSG cells under H₂O₂-induced oxidative stress was investigated using immunofluorescence assay. The effect of cordycepin stimulation on AMY protein expression was indicated (Figure 3A). The relative fluorescence intensity revealed that no significant differences were observed in HSG cells treated with either cultured medium or cordycepin. Intriguingly, H₂O₂-induced oxidative stress revealed a significant reduction in AMY protein expression when compared to the untreated group. Compared to the H₂O₂ group, the post-treatment group with cordycepin showed a dramatic increase of AMY protein expression in HSG cells (Figure 3B). The expression of AQP5 protein in HSG cells treated with either cordycepin, H_2O_2 or cordycepin post-treatment was demonstrated (Figure 3C). The relative fluorescence of AQP5 was also significantly increased in HSG cells treated with cordycepin. However, compared to the untreated

group, the AQP5 protein expression of H_2O_2 -induced HSG cells was extremely decreased. Noticeably, the cells post-treated with cordycepin were found to have significantly increased levels of AQP5 protein, indicating that under H_2O_2 -induced oxidative stress, cordycepin could stimulate the expression of both AMY and AQP5 (Figures 3B & D). These results suggest that cordycepin could rescue both AMY and AQP5 protein expressions under H_2O_2 -induced oxidative stress in HSG cells.

Cordycepin attenuates against intracellular ROS generation and affects apoptotic gene expression in H_2O_2 -induced HSG cells

Next, we investigated the effect of cordycepin against H_2O_2 -induced intracellular ROS generation in HSG cells using DCFH-DA fluorescence assay. The fluorescence intensity of DCFH-DA was considerably increased by 500 μ M H_2O_2 when compared to the untreated group, indicating that H_2O_2 enables to generate intracellular ROS. However, the post-treatment with cordycepin effectively reduced

intracellular ROS generation in HSG cells (Figure 4A). To confirm the preventive effects of cordycepin in HSG cells, the expressions of antioxidant genes (i.e. SOD1, CAT and GPX1) and apoptotic genes were evaluated. The band intensities of mRNA expression of these antioxidant genes were upregulated in HSG cells cultured in each concentration of cordycepin post-treatment (Figure 4B & D). The relative expression of CAT and SOD1 were increased significantly in all concentrations of cordycepin whereas that of GPX1 were increased significantly in certain concentrations as compared to that found in the untreated group (Figure 4D). Similarly, we also found that, H₂O₂ induced up-regulation of apoptotic gene, caspase-3 and down-regulated BCL-2 gene expression. Significantly, a decrease in the level of caspase-3 and an increase in BCL-2 in H₂O₂-induced HSG cells after post-incubation with cordycepin were demonstrated (Figure 4C & E). This may indicate the anti-apoptotic activity of cordycepin on H2O2-induced HSG cells.

Cordycepin stimulates the amylase secretion from HSG cells under H_2O_2 -induced oxidative stress

The amounts of amylase secreted from the HSG

cells cultured in each cordycepin concentration were reported (Figure 5). Compared to the untreated specific concentrations of cordycepin group, treatment could stimulate the cells to secrete more amylase after short and long-term culture periods. For the short-term culture periods, cordycepin stimulated amylase secretion, with the high levels during 812 h. The 12.5 µM of cordycepin treated HSG cells showed the highest level of amylase secretion at 8 h. However, at 24 h, the secretion of amylase was dropped in all groups (Figure 5A). It is also noted that amylase secretion from the cells cultured in 6.25, 12.5 and 25 µM of cordycepin (1-5 days) were increased but statistically insignificant (Figure 5B). To further study the effect of cordycepin in amylase secretion of HSG cells, cordycepin was used to treat H₂O₂ -exposed HSG cells. As expected, after 30 min of H₂O₂ treatment, the amylase secretion in the culture medium was suddenly decreased. Surprisingly, the amount of amylase was significantly increased in the culture medium when cordycepin was used for post-treatment (Figure 5C).



Figure 4. Cordycepin attenuated H_2O_2 -induced intracellular ROS generation in HSG cells. Cells were induced with 500 µM H_2O_2 for 30 min and exposed to cordycepin ranging from 6.25-50 µM for 24 h. The relative fluorescence intensity of DCFH-DA was determined by DCFH-DA assay (**A**). The mRNA expression for antioxidant genes, SOD *I*, CAT, and GPX *I* (**B**) and apoptotic genes including *caspase-3* and *BCL-2* (**C**) were analysed by RT-PCR. The relative mRNA expression levels of SOD *I*, CAT and GPX *I* (**D**), *Caspase-3* and *BL-2* (**E**) genes were evaluated by image J NIH software and normalized with GAPDH. Gel electrophoresis are from one representative experiment and bar charts are derived from analysis of relative expression from three independent experiments. **p<0.01 compared to the untreated group; #p<0.01 and ##p<0.01 compared to the H2O2 group.



Figure 5. Cordycepin increased amylase secretion in H₂O₂-induced HSG cells. Cells were treated with various concentrations of cordycepin for short-term culture at 4, 8, 12 and 24 h (**A**) and long-term culture at 1, 3 and 5 days (**B**). Cordycepin increased amylase secretion in HSG cells exposed to H₂O₂ for 30 min (**C**). Amylase activity was kinetically measured in a cultured medium of HSG cells exposed to H₂O₂ without or with cordycepin treatment for 24 h. Data are represented as mean \pm SD (n=3). ***p*<0.01 compared to the untreated group, and ***p*<0.05 and ***p*<0.01 compared to the H₂O₂ group.

Discussion

Xerostomia is a common side effect of numerous medications and other therapies, especially found in patients that have received radiotherapy for the treatment of head and neck cancer and in elderly 22]. individuals [21, Current treatments for xerostomia include salivary substitutes, hyperbaric oxygen and gene therapy as well as salivary stimulants such as pilocarpine (a non-selective muscarinic receptor agonist) [23-25]. However, these treatments still have the potential risk for undesirable adverse effects [26]. The major bioactive compound of Cordyceps sp. is cordycepin, also known as 3deoxyandenosine, which has various pharmaceutical properties including antitumor, anticancer, antiinflammatory, immunomodulatory and antioxidant activities [13, 14, 27]. This study indicated that cordycepin could stimulate the gene and protein expression of salivary markers and amylase secretion, while it reduced intracellular ROS generation in HSG cells via anti-oxidant and anti-apoptotic activities. The protective effect of cordycepin could enhance the H₂O₂-induced HSG cells to survive and revive the functions as a vital saliva-producing cells.

Amylase (AMY) and aquaporin-5 (AQP5) are two important specific markers for functional salivary glands [28]. Amylase, which is encoded by AMY1A gene, is one of the major protein components of saliva and is a digestive enzyme for calcium binding responsible for starch hydrolysis [29-31]. In addition, aquaporin-5, which is encoded by AQP5 gene, is a water-selective channel protein expressed in the apical membrane of epithelial cells of serous acini in the salivary glands [32-35]. The osmolality and the concentration of electrolytes in salivary fluid secretion are directly controlled by AQP5 protein. In this study, cordycepin enhanced the expression of both the AMY1A and AQP5 genes in HSG cells. Although the effective dose to stimulate AMY1A (25 µM) and AQP5 $(12.5 \,\mu\text{M})$ genes was different, it did not show a much difference in a fold-change expression. The concentration at 12.5 µM of cordycepin was chosen for further study since we expected the most effective result with the lower concentration of cordycepin. This concentration showed an increasing effect in AMY1A and AQP5 for both gene and protein levels. This is promising since the previous study showed that AQP5 is one of key proteins to cause xerostoma. In addition, a study has previously reported that high expression of *AQP5* leads to an increase in a salivary amylase level in diabetes induced xerostomia rats [36].

Salivary gland hypofunction and reduced expression of AQP5 are the oxidative stresses leading to xerostomia [2]. Our results showed that cordycepin could promote the expression of AQP5 genes, suggesting that the water channels are opened by cordycepin treatment after administration of H₂O₂. During the study of the expression of protein levels, the reduction in expression of AMY and AQP5 protein levels have been observed in HSG cells exposed to H₂O₂-induced oxidative stress. However, AMY and AQP5 protein levels were significantly increased in cordycepin-treated HSG cells after H₂O₂ administration. These results suggested that the oxidative stress related with xerostomia-induced hyposalivary function can be reversed effectively by treatment with cordycepin. Previous studies have proposed that the mechanism of cordycepin in salivary gland cells might involve several pathways, such as sodium/hydrogen exchanger-1 (NHE1) activity [37], c-Jun N-terminal kinase (JNK) phosphorylation [38] and protein disulphide isomerase (PDI) [39].

ROS generation is a major cause of oxidative stress, which is harmful to the biomolecules inducing the pathology in the body, causing age-related cell damage. Oxidative stress is a main cause of cellular damage in several diseases, including xerostomia [40, 41]. In addition to therapeutic interventions, bioactive compounds with antioxidant activity have been reported to be capable of protecting cells from oxidative stress [42, 43]. The bioactive compound cordycepin has been previously reported to be effective against oxidative and endoplasmic reticulum stress by reducing intracellular ROS generation in vitro [15, 44]. Wang et al. revealed that cordycepin could prevent human bone marrow stem mesenchymal cells (BM-MSCs) from H₂O₂-induced inhibition of osteogenesis [18]. Moreover, cordycepin protected against neuron cell death in PC12 and C6 glial cells from oxidative stress-induced neurotoxicity [19, 20]. In animal models, cordycepin treatment decreased oxidative damage and boosted the differentiation of BM-MSCs [18]. To examine cells under oxidative stress, intracellular ROS generation and the expression of antioxidant genes, which are widely used as markers for oxidative stress, were determined. In this study, we observed that ROS generation was increased by H₂O₂-induced oxidative stress in HSG cells; nevertheless, it was effectively alleviated by cordycepin treatment. To confirm intracellular ROS generation, we determined the effect of cordycepin

through the antioxidant gene expression, SOD1, CAT and GPX1 in the H₂O₂-induced HSG cells. SOD1 is an important endogenous antioxidant enzyme. It catalyzes the dismutation of superoxide anion free radicals to hydrogen peroxide and oxygen and acts as the first enzyme in defense of cell death from ROS [45]. CAT is used in the reduction of hydrogen peroxide to water and oxygen, while GPX1 is the intracellular enzyme used to reduce hydrogen peroxide to water and lipid peroxides [46]. Our results showed that the post-treatment with cordycepin considerably upregulated the expression of the SOD1, CAT and GPX1 genes in HSG cells induced by H₂O₂. Furthermore, we also demonstrated that, posttreatment with cordycepin significantly decreased the expression of caspase-3, but increased BCL-2 expression in H₂O₂-induced HSG cells. Taken together, these results indicated that cordycepin could effectively block ROS generation through the anti-oxidant and anti-apoptotic activities.

The composition of saliva is 99% water and 1% solids and protein, and it produced by the salivary glands consisting of the parotid, sublingual and submandibular glands [47, 48]. Amylase is a major of protein component of saliva, which is synthesized and secreted by acinar cells, a major cell population in the salivary gland. Enzymatic digestion of carbohydrates is the main function of saliva working through the amylase [49]. In this study, we also determined the amylase activity of HSG cells treated with cordycepin. The higher level of amylase secretion in the HSG cells was found during 8-12 h co-culture in cordycepin while long-term culture period (1-5 day) stimulated amylase lesser amount of secretion. The pharmacokinetic research demonstrated that the half-life of cordycepin was short. Cordycepin is rapidly deaminated by adenosine deaminase and promptly metabolized to an inactive metabolite, 3'-deoxy-hypoxanthinosine [50, 51]. It was found that cordycepin permeated the cell membrane of human endothelial-like immortalized cells (EA.hy926) cells within 30 min and was steady for a 3 h culture period [52]. Although, cordycepin are rapidly reduced and metabolized in a short period of time, a cordycepininduced unidentified compounds detected in blood and liver for over 2 h, which might have highly specific and widespread within the body organ or tissue [51, 53, 54]. To overcome the problem of rapid a high concentration might elimination, be administered; otherwise, intracellular concentrations sub-therapeutic. Interestingly, would be we demonstrated that the amylase secretion was recovered after cordycepin treatment in HSG cells induced by H₂O₂ as well as cordycepin significantly enhanced AMY1A and AQP5 expression at both

mRNA and protein levels. It is possible that recovering secretion of amylase under oxidative stress may be due to the ability of cordycepin on AMY1A gene and signaling pathways that are responsible for activating fluid and enzyme secretion [55]. Furthermore, cordycepin probably promotes the effective function of AQP5 gene which is the major pathway for regulating the salivary fluid secretion (water and protein) permeability in acinar cells [56, 57]. Altogether, cordycepin could help cells restoring from oxidative stress by increasing salivary secretion via an increase in AMY and AQP5 protein levels and the resulting amylase secretion. However, despite the fact that we observed significant protein expression in the cordycepin-treated group compared to the untreated group, the enzyme activity was not dramatically different compared to the protein may be described by expression. This two First, explanations. the intracellular protein was analysed through immunoexpression fluorescence staining whereas the protein activity was evaluated from the supernated extracellular protein. Hence, the enzymatic function of secreted amylase may be affected by the post-modification or the formation of complex to function properly. Also, the functional amount of amylase in the supernatant may influence the amylase activity [58]. Second, several previous studies have shown that the 2D in vitro model of salivary gland differentiation can enhance amylase expression and activity. However, the 3D model can better improve the differentiation. This latter model would help salivary gland cells to perform their function and activity better [59]. Therefore, the 2D model that we selected in this study may limit the protein function.

Conclusions

In conclusion, cordycepin could effectively protect against HSG cell death from H_2O_2 -induced toxicity by reducing ROS generation via antioxidant and anti-apoptotic capacities. Furthermore, cordycepin exerts prevention by increasing both the expression of salivary marker genes and amylase secretion. Thus, these results suggest that cordycepin might be used as a new therapeutic agent and an experimental model for oral dryness.

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Competing Interests

The authors have declared that no competing interest exists.

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