CecropinXJ inhibits the proliferation of human gastric cancer BGC823 cells and induces cell death in vitro and in vivo

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Abstract. We have shown that an antimicrobial peptide (AMP) cecropinXJ isolated from the larvae of Bombyx mori selectively inhibits the proliferation of cancer cells. However, the mechanism remains to be determined. In the present study, we examined the antitumor activity of cecropinXJ against human gastric cancer BGC823 cells and explored the mechanism. The results showed that cecropinXJ inhibited the growth of gastric cancer BGC823 cells in vitro and in vivo. MTT and colony formation assays indicated that cecropinXJ suppressed cell proliferation and reduced colony formation of BGC823 cells in a dose- and time-dependent manner, but without inhibitory effect on normal gastric epithelia GES-1 cells. S-phase arrest in BGC823 cells was observed after treatment with cecropinXJ. Annexin V/PI staining suggested that cecropinXJ induced both early and late phases of apoptosis through activation of mitochondrial-mediated caspase pathway, upregulation of Bax expression and downregulation of Bcl-2 expression. Additionally, cecropinXJ treatment increased reactive oxygen species (ROS) and decrease mitochondrial membrane potential (ΔΨm) and led to release of cytochrome c. Importantly, in vivo study showed that cecropinXJ significantly prevented the growth of xenograft tumor in the BGC823-bearing mice, possibly mediated by the induction of apoptosis and inhibition of angiogenesis. These results suggest that cecropinXJ may be a promising therapeutic candidate for the treatment of gastric cancer.

Introduction

Gastric cancer is one of the most common malignant tumors, >70% of new cases and deaths occur in developing countries each year (1). In China, the incidence and mortality rate of gastric cancer are increasing. Although there are distinct discrepancies in diagnosis, prognosis, and treatment efficacy for gastric cancer patients, the 5-year survival rate of gastric cancer is generally <20% (2). At present, the management of gastric cancer includes surgery, radiotherapy, conventional chemotherapy, molecular targeted therapy and biological therapy, but these methods have not reached expected clinical efficacy, which mainly focus on mass cell killing without high specificity and often causes severe side-effects and toxicities. Moreover, recurrences and metastasis frequently occur in the majority of patients (2). Hence, it is urgent to explore safe and effective therapeutic agents for gastric cancer to improve the survival rate and quality of life.

Antimicrobial peptides (AMPs) are evolutionarily-conserved components of the innate immune system (3), which exert their activities against a broad range of microbial pathogens. Compared to conventional treatments, AMPs have many advantages, such as specific cytotoxicity for cancer cells, ability to bypass the multidrug-resistance mechanism, and additive effects in combination therapy (4), suggesting that AMPs has potential as a novel antitumor agent for the treatment of cancer.

AMPs can directly kill cancer cells by a cell membrane-lytic effect (5), or trigger apoptosis in cancer cells via the Fas death-receptor-induced extrinsic pathway (6) and the mitochondria-apoptosome-mediated apoptotic intrinsic pathway (7). AMPs with cationic and amphipathic amino acid composition can penetrate anionic cell membrane and subsequently bind to mitochondrial membrane, which destroy membrane structure, increase the level of reactive oxygen species (ROS) and decrease mitochondrial membrane potential (ΔΨm) (8). Mitochondrial dysfunction elicits the release of cytochrome c from mitochondria to cytosol, which activates the caspase-cascade system and induces cell apoptosis (9). It has been reported that AMPs inhibited the growth or proliferation of cancer cells through regulation of numerous proteins and pathways, including the caspase family and the Bcl-2 family (10,11), the mitogen-activated protein kinase (MAPK) family (12), the nuclear factor-kappa B (NF-κB) (13),

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Abbreviations: AMPs, antimicrobial peptides; ROS, reactive oxygen species; MAPK, mitogen-activated protein kinase; NF-κB, nuclear factor-kappa B; PI3K/Akt, phosphoinositide 3-kinase/protein kinase B

Key words: cecropinXJ, gastric cancer, antitumor, proliferation, apoptosis, mitochondrial-mediated pathway, angiogenesis
phosphoinositide 3-kinase (PI3K)/protein kinase B (Akt) signal transduction pathways (14,15) and ER stress-mediated apoptosis pathway (16,17). Furthermore, certain AMPs are potent inhibitors of blood vessel development (angiogenesis) that is associated with tumor progression (18).

We isolated a 37-AA cationic antimicrobial peptide with specific amphipathic α-helices (named as cecropinXJ) from the larvae of Bombyx mori (19), which has a broad effect spectrum against bacteria (20). Our previous studies have shown that cecropinXJ can inhibit the proliferation of human gastric cancer in vitro such as AGS cells (21), but have no hemolytic activity against human erythrocytes and no toxicity to normal mammalian cells (22,23). Due to difficulty of AGS cells to form a solid tumor in vivo, we investigated the cytotoxicity and the mechanism of cecropinXJ against the human gastric cancer BGC823 cells in the present study. Our results showed that cecropinXJ suppressed the proliferation and induced apoptosis of BGC823 cells in vitro and in vivo through mitochondrial-mediated apoptosis pathways and inhibition of angiogenesis, which could provide experimental evidence for the potential application of cecropinXJ as a new antitumor candidate for treatment of gastric cancer.

Materials and methods

Preparation of antimicrobial peptide cecropinXJ. CecropinXJ of B. mori was prepared through the Saccharomyces cerevisiae eukaryotic expression system and purified with Ni-NTA agarose column as reported (23). The concentration of purified recombinant cecropinXJ protein was detected by a Bradford protein assay kit (KeyGen Biotech, Nanjing, China). The amino acid sequence is WKIFKKIEKMGRNIRDGIVKAGPAIEVLGSAKAIGK. Before use, the peptide was dissolved in Dulbecco's modified Eagle's medium (DMEM; Hyclone, Logan, UT, USA) at a concentration of 1 mg/ml and sterilized by filtration through a 0.2-mm filter.

Cell culture. The human gastric cancer BGC823 cells and normal gastric epithelial GES-1 cells were kindly provided by Professor Youyong Lv (Beijing Cancer Hospital, Beijing, China). Cells were cultured in DMEM medium, supplemented with 10% fetal bovine serum (FBS; Gibco-BRL, Grand Island, NY, USA), 100 µg/ml streptomycin and 100 U/ml penicillin (Beijing Solarbio, Beijing, China) in a humidified atmosphere of 5% CO₂ at 37°C.

Cell viability assay. To evaluate the effects of cecropinXJ on the proliferation of BGC823 cells and GES-1 cells, cell viability was measured by 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) assay (24). The concentration of cecropinXJ (0, 20, 50, 80 and 100 µg/ml) for 24 h. After treatment, the culture medium was removed and 100 µl MTT solution (5 mg/ml) was added into each well, then incubated at 37°C for 4 h. After 4 h, 150 µl of dimethyl sulfoxide (DMSO; Beijing Solarbio) was added into each well and incubated at 37°C for 10 min. Absorbance was measured at OD₄₉₀/₆₅₅ using a 96-well microplate reader (Bio-Rad Laboratories, Hercules, CA, USA). Cell viability (%) = [(ODtreated - ODblank)/(ODnon-treated - ODblank)] x 100%.

Colonies formation assay. One thousand BGC823 cells were resuspended in 4.4 ml mixture of 0.5% agar solution containing cell culture medium (2X DMEM and 20% FBS) at a final concentration of 20, 50, 80 and 100 µg/ml cecropinXJ and layered on the top of 0.7% agar layer in 60-mm petri-dish (Corning). The mixtures without cecropinXJ or with 10 µg/ml Dox served as negative and positive controls, respectively. Dishes were incubated for three weeks at 37°C in a humidified atmosphere of 5% CO₂. Cell colonies were observed by light microscopy (Leica Microsystems, Wetzlar, Germany) and visualized by the treatment of 0.5 ml p-iodonitrotetrazolium violet (Sigma) for 16 h, then photographed and the number of cell colonies was counted. Colonies (%) = the colony number of treatment group/the colony number of control group x 100%.

LDH detection. BGC823 and GES-1 cells at 4x10⁵ cells/well were inoculated into 24-well plates and treated with 0, 20, 50, 80 and 100 µg/ml of cecropinXJ for 24 or 48 h. Medium was collected and the ratio of LDH release was detected according to the manufacturer's instructions.

Cell cycle analysis of cecropinXJ on BGC823 cells by flow cytometry. BGC823 cells at 5x10⁵ cells/ml were inoculated into 100-mm dishes and incubated at 37°C for 24 h. After the medium was removed, fresh medium with different concentrations of cecropinXJ was added and cultured. Medium without cecropinXJ was served as the control. After culture for 24 or 48 h, the cells were collected and washed with cold phosphate-buffered saline (PBS, pH 7.4) and fixed with 75% ethanol overnight at -20°C. After washing, DNA staining solution (Multisciences, Hangzhou, China) was added and incubated at 37°C in darkness for 30 min. The samples were analyzed with a flow cytometry (BD Biosciences, San Jose, CA, USA).

Apoptosis rate determined by flow cytometry. BGC823 cells at 5x10⁵ cells/ml were inoculated into 60-mm dishes and incubated at 37°C for 24 h. After the medium was removed, 2.0 ml of DMEM medium with the final concentrations of cecropinXJ at 20, 50, 80 and 100 µg/ml was added to each dish. No cecropinXJ served as the control. After cultured for 24 or 48 h, cells were collected after digestion with 0.25% trypsin, washed with PBS two times, and suspended in 400 µl binding buffer. Cell suspensions were stained with 5 µl of Annexin V-FITC and 10 µl of propidium iodide (PI) (BestBio Biotechnologies, Shanghai, China) according to the manufac-
Morphological observation by Hoechst 33258 staining and transmission electron microscopy. The BGC823 cells were treated with different concentrations of cecropinXJ for 24 h. For Hoechst 33258 staining, cells were washed with PBS and fixed with 95% ethanol at 4°C for 15 min. After washing with PBS, cells were stained with 0.5 µg/ml Hoechst 33258 (Sigma) for 10 min at 4°C in the dark. Samples were observed under a fluorescence microscope (Leica Microsystems). For transmission electron microscopy, cells were fixed with 2.5% glutaraldehyde in 0.1 M PBS at 4°C overnight, and fixed with 1% osmium tetroxide after washing with PBS. The cells were then dehydrated using graded alcohol and embedded in araldite. Ultrathin sections were stained with uranyl acetate and lead citrate and examined with a Philips CM10 transmission electron microscope.

Detection of reactive oxygen species (ROS) and mitochondrial membrane potential (Δψm). ROS were detected by non-fluorescent 2,7-dichlorodihydrofluorescein (H2DCF-DA) (Sigma) (a fluorogenic freely permeable tracer), which is specific for ROS assessment. Changes in the mitochondrial membrane potential (Δψm) were measured using a lipophilic cationic dye 3,3-dihexyloxacarbocyanine iodide [DiOC6(3)] (Sigma). Briefly, BGC823 cells (2x10⁶ cells/ml) were treated with different concentrations of cecropinXJ for 24 h or treated with 100 µg/ml cecropinXJ for different times in 100-mm dishes. After treatment, cells were incubated with 10 µM H2DCF-DA for 30 min or 40 nM DiOC6(3) for 15 min at 37°C, and washed with PBS. The samples were analyzed by flow cytometry.

Quantitative real-time reverse-transcription PCR (qRT-PCR). Gene expression was analyzed by qRT-PCR. Total RNA was extracted from BGC823 cells and tumors using TRIzol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s protocol. cDNA was synthesized from total RNA by M-MLV reverse transcriptase according to the manufacturer’s protocol. cDNA was then dehybridized using graded alcohol and embedded in araldite. Ultrathin sections were stained with uranyl acetate and lead citrate and examined with a Philips CM10 transmission electron microscope.

Signaling Technology) for 2 h at room temperature. Membrane was then dehydrated using graded alcohol and embedded in araldite. Ultrathin sections were stained with uranyl acetate and lead citrate and examined with a Philips CM10 transmission electron microscope.

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Caspase activity. BGC823 cells (5x10⁶ cells/dish) were seeded in 100-mm dishes and treated with 0, 20, 50, 80 and 100 µg/ml of cecropinXJ for 24 h. After treatment, BGC823 cells were lysed and supernatant was collected. The protein concentration was determined using the bicinchoninic acid (BCA) assay kit. Proteins (40 µg/lane) were separated by 12% SDS-PAGE and transferred onto a nitrocellulose membranes (Millipore, Billerica, MA, USA). After washing with TBS-T buffer (20 mM Tris-HCl, 150 mM NaCl, 0.05% Tween-20), membrane was blocked with 5% skim milk at room temperature for 1 h, and then incubated with the primary antibodies (1:2,000; Cell Signaling Technology, Danvers, MA, USA) overnight at 4°C. After washing with TBS-T, membrane was incubated with the corresponding HRP-conjugated secondary antibodies (Cell Signaling Technology) for 2 h at room temperature. Membrane was washed with TBS-T and exposed using enhanced chemiluminescent (ECL) detection kit (CWBio, Inc., Beijing, China).

Table I. Primers were designed for qRT-PCR.

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<th>Primers</th>
<th>Sequences (5'-3')</th>
<th>Length (bp)</th>
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<tr>
<td>Caspase-3</td>
<td>F: AGACATACTCCTTCCATCA A</td>
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<tr>
<td></td>
<td>R: TCATAGCAGACGACTCATT</td>
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<td>Caspase-8</td>
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<td></td>
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<td>Bcl-2</td>
<td>F: GAATGGGGGAGATTGTGGG</td>
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<tr>
<td></td>
<td>R: GAGTCGCAAGTGTTGCATT</td>
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<tr>
<td>Bax</td>
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<td>267</td>
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<tr>
<td></td>
<td>R: GGCTGAGGGAGCTGAGGAGAG</td>
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<tr>
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<tr>
<td></td>
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<td></td>
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<td>bFGF</td>
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<td></td>
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<td>R: GAGGGATGAAGGAATCTGCC</td>
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<td>GAPDH</td>
<td>F: ACACCCACTCTCTCACCTT</td>
<td>106</td>
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<tr>
<td></td>
<td>R: ACCCTGTGGCTGTAGCCAA</td>
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F, forward; R, reverse.

Western blot analysis. BGC823 cells were treated with cecropinXJ (0, 20, 50, 80 and 100 µg/ml) for 24 h and washing with ice-cold PBS twice, cells were collected and lysed using 100 µl CytoBuster™ protein extraction reagent (Novagen, Madison, WI, USA) on ice for 15 min. After centrifugation at 14,000 rpm 4°C for 10 min, the protein concentration in supernatant was determined by the bicinchoninic acid (BCA) assay kit. Proteins (40 µg/lane) were separated by 12% SDS-PAGE and transferred onto a nitrocellulose membranes (Millipore, Billerica, MA, USA). After washing with TBS-T buffer (20 mM Tris-HCl, 150 mM NaCl, 0.05% Tween-20), membrane was blocked with 5% skim milk at room temperature for 1 h, and then incubated with the primary antibodies (1:2,000; Cell Signaling Technology, Danvers, MA, USA) overnight at 4°C. After washing with TBS-T, membrane was incubated with the corresponding HRP-conjugated secondary antibodies (Cell Signaling Technology) for 2 h at room temperature. Membrane was washed with TBS-T and exposed using enhanced chemiluminescent (ECL) detection kit (CWBio, Inc., Beijing, China).

Caspase activity. BGC823 cells (5x10⁶ cells/dish) were seeded in 100-mm dishes and treated with 0, 20, 50, 80 and 100 µg/ml of cecropinXJ for 24 h. After treatment, BGC823 cells were lysed and supernatant was collected. The protein concentration was determined by Bradford protein assay kit. Cellular extracts (30 µg) were incubated with 20 ng Ac-DEVD-NA and Ac-LEHD-NA for overnight at 37°C and
caspase activity was measured by cleavage of the substrates to pNA, the absorbance was measured at 405 nm. Relative caspase activity was calculated as a ratio OD_{treated}/OD_{untreated}.

Xenograft tumor model. Four-week-old athymic nude female mice (BALB/c nu/nu) were purchased from the Beijing Laboratory Animal Research Center (Beijing, China) and received pathogen-free water and food. The mice were inoculated subcutaneously with BGC823 cells (5x10^6) in 0.1 ml of PBS. All the mice developed tumors at day 5 and were randomized into two groups (n=6). The experiment group was peritumorally injected with 0.1 ml cecropinXJ (100 µg)
five times per week (from Monday to Friday) during a 2-week period and the control group was injected with 0.1 ml of BSA (100 µg). Mouse weight and tumor size were measured every day. After two weeks, mice were sacrificed and tumors were dissected. After weighing, tumors were fixed in 10% formalin and embedded in paraffin. Sections were stained with hematoxylin and eosin.

TUNEL staining. The TUNEL staining assay was performed according to the manufacturer's instructions (Biobox, Nanjing, China). Paraffin-embedded tissue slides were dewaxed and rehydrated by heating at 60˚C, followed by washing in xylene and rehydration through a graded series of ethanol and double-distilled water. Tissue sections were incubated with proteinase K (20 µg/ml) at room temperature for 15 min and with 3% H2O2 for 10 min. Then sections were incubated with the TdT enzyme at 37˚C for 60 min. After incubation with streptavidin-HRP solution in darkness for 30 min, the sections were stained with 3,3-diaminobenzidine (DAB) substrate for 2 min and then counterstained with hematoxylin.

Immunohistochemistry. Paraffin-embedded tissue were cut four-micrometer-thick in consecutive sections and processed for immunohistochemistry for CD31, CD34, basic fibroblast growth factor (bFGF), vascular endothelial growth factor (VEGF) and von Willebrand factor (VWF) (Biosynthesis, Beijing, China). Tissue sections were deparaffinized, dewaxed and rehydrated by heating at 60˚C, followed by washing with PBS/T. Antigen retrieval was carried out in a microwave oven for 10 min. The sections were blocked with 10% goat serum (formulated with 3% BSA) at 37˚C for 15 min and washed thrice with PBS. The primary antibodies against angiogenesis-associated proteins with 1:200 dilution were incubated for 1 h. After washing with TBS-T, slides were incubated with the HRP-conjugated secondary antibodies for 1 h at room temperature. Sections were stained with DAB, counterstained with hematoxylin and mounted. The immunohistochemical staining was observed under light microscope.

Statistical analysis. Data are expressed as the mean ± SD. Statistical analyses were performed using Student's t-test. Values of $P<0.05$ or $0.01$ was considered statistically significant and indicated in the figures with * and **, respectively.

Results

CecropinXJ effectively suppress the proliferation of gastric cancer BGC823 cells. We first examined whether cecropinXJ could suppress the proliferation of gastric adenocarcinoma BGC823 cells via MTT assay. Normal gastric epithelial cell GES-1 was used as control. Compared to GES-1 cells, the proliferation of BGC823 cells was significantly inhibited by cecropinXJ in dose- and time-dependent manner (Fig. 1A and B). Dox suppressed the growth of both types of cells. The inhibitory effects of 100 µg/ml cecropinXJ on BGC823 cells were similar to 10 µg/ml Dox (Fig. 1B). When the concentration of cecropinXJ was <500 µg/ml, the growth of GES-1 cells was not impaired by the cecropinXJ treatment compared to without cecropinXJ treatment (Fig. 1A).

The soft agar colony formation assay showed that cecropinXJ treatment dose-dependently inhibited the colony formation and colony growth of BGC823 cells. The colony formation ratio was only (26.46±4.65%) under 100 µg/ml cecropinXJ treatment (Fig. 1C). These results suggest that cecropinXJ might have a long-term effect on the proliferation of BGC823 cells.
Many AMPs can destroy the lipid membrane and lead to cell lysis and death (24). We investigated whether cecropinXJ could destroy cell membrane through measuring the released cytoplasmic enzyme LDH into the cell culture supernatant. Compared with untreated group, cecropinXJ treatment did not significantly increase LDH release of BGC823 cells either for 24 or 48 h until the concentration of cecropinXJ reached up to 100 µg/ml, suggesting that high concentration of cecropinXJ treatment destroyed the integrity of plasma membrane in BGC823 cells but low concentration of cecropinXJ did not. CecropinXJ did not affect the plasma membrane in GES-1 cells.

CecropinXJ arrests the cell cycle of BGC823 cells at S phase. To determine whether cecropinXJ affects the cell cycle of BGC823 cells, the cell cycle distribution of cecropinXJ-treated BGC823 cells was analyzed by flow cytometry. Upon cecropinXJ treatment, the percentage of cells at G0/G1-phase significantly decreased (56.67 and 57.28% in the 100 µg/ml cecropinXJ treatment for 24 and 48 h, respectively, vs. 68.14 and 68.52% in the control group), whereas the percentage of cells at S-phase significantly increased (35.65 and 35.75% in the 100 µg/ml cecropinXJ treatment for 24 and 48 h, respectively, vs. 23.87 and 23.48% in the control group) (Fig. 2). BGC823 cells with cecropinXJ treatment achieved peak apoptosis. These results indicated that cecropinXJ induced S-phase arrest and apoptosis in a dose-dependent manner.

CecropinXJ dose-dependently induces apoptosis of BGC823 cells. The above results showed that both low (<80 µg/ml) and high concentration of cecropinXJ could inhibit BGC823 cell proliferation with or without damage of plasma membrane. To investigate the mechanism of cecropinXJ-mediated proliferation inhibition of BGC823 cells, BGC823 cells were treated with different concentrations of cecropinXJ for 24 or 48 h and stained with Annexin V-FITC and PI. The samples were analyzed by flow cytometry. CecropinXJ treatment dose-dependently increased apoptosis of BGC823 cells at both early and late stages. The apoptosis of BGC823 cells was increased from (4.88±2.53%) (untreated) to (39.56±4.36%) (100 µg/ml cecropinXJ treated) for 24 h. Compared to 24 h, the apoptosis of BGC823 cells was also increased 48 h after cecropinXJ treatment. Furthermore, 100 µg/ml cecropinXJ treatment induced necrosis of BGC823 cells compared to other groups for both 24 and 48 h (Fig. 3A and B).

Morphological characteristics of apoptosis include chromatin condensation, DNA fragmentation and membrane blebbing (25). Nuclear morphology was observed by Hoechst 33258 staining. As shown in Fig. 3C, the untreated cells were round and homogeneously stained, whereas cecropinXJ-treated cells showed obvious chromatin condensation and fragmentation. Moreover, the numbers of apoptotic nuclei containing condensed chromatin increased significantly with the increased concentration of cecropinXJ.

Ultrastructure of BGC823 cells was observed using transmission electron microscopy and revealed that cecropinXJ induced significantly morphological changes. The untreated BGC823 cells showed intact membrane, normal nucleus and integrated mitochondria. However, cecropinXJ-treated cells showed nuclear heterochromatin and condensed chromatin distribution around the nuclear membrane. Moreover, mitochondria was osteoporosized and swollen and cristae disappeared (Fig. 3D). Taken together, these results suggested that cecropinXJ induced apoptosis of BGC823 cells.

CecropinXJ promotes ROS production and decreases mitochondrial membrane potential (ΔΨm). It has been reported that cancer cell apoptosis induced by cecropins may be due to mitochondrial dysfunction and overproduction of ROS (26). Intracellular ROS production and loss of ΔΨm cause mitochondrial membrane depolarization and trigger a cascade of mitochondrial-dependent apoptotic signaling (27). The ROS level and the ΔΨm were measured by flow cytometry using H2DCF-DA and DiOC6(3), respectively. In this assay, cecropinXJ increased ROS generation in dose- and time-dependent manner. Moreover, cecropinXJ-treated BGC823 cells significantly decreased their ΔΨm (Fig. 4).

CecropinXJ induces apoptosis through activation of caspase pathway. It is well known that proteins in the caspase-family and Bcl-2-family play critical roles in the apoptosis process (28). To investigate whether caspase-family and Bcl-2-family proteins were involved in cecropinXJ-induced apoptosis, we measured the expression and activation of these apoptosis-associated proteins by qRT-PCR and western blot analysis. According to the results of the western blot analysis, these enzymes including caspase-3 (effector caspase), -8 and -9 (initiator caspase) were all cleaved. Moreover, cecropinXJ treatment dose-dependently increased the expression of Bax and cytochrome c at both RNA and protein levels but decreased the expression of anti-apoptotic molecule Bcl-2 (Fig. 5A and B). The Bax/Bcl-2 ratio increased in BGC823 cells, suggesting that these proteins were involved in cecropinXJ-induced apoptosis.

The activities of caspase-3, -8 and -9 were further detected in cecropinXJ-treated cells using the colorimetric tetrapeptide substrates Ac-DEVD-pNA, Ac-IETD-pNA and Ac-LEHD-pNA, which have been shown to be specific for caspase-3-, caspase-8- and caspase-9-like enzymatic activity, respectively (29). After 100 µg/ml cecropinXJ treatment for 24 h, the activities of caspase-3, -8 and -9 were decreased ~2.3-, 1.3- and 1.3-fold compared to untreated control (Fig. 5C). Before cecropinXJ treatment, adding the caspase inhibitors (Ac-VAD-fmk for caspase-3 and -7, Ac-DEVD-fmk for caspase-8 and Ac-LEHD-fmk for caspase-9) significantly reduced cecropinXJ-induced cell death (P<0.01), but caspase-8 inhibitor had no effect (Fig. 5D). These results suggest that cecropinXJ might induce apoptosis of BGC823 cells via activation of caspase-3 and -9.

CecropinXJ inhibits tumor growth in vivo. To investigate whether cecropinXJ could suppress the growth of BGC823 cells in vivo, mice were injected with BGC823 cells and treated with BSA (control group) or cecropinXJ on day 7. Twelve days after treatment, the tumors were removed from these animals and their volumes and weights were measured (Fig. 6). The tumor volume and tumor weight in mice treated with cecropinXJ were markedly reduced compared with BSA. CecropinXJ appeared to be non-toxic to normal cells, and there were no notable side-effects on the body weight of mice (the mice had tumors) at the end of the experiment (Fig. 6D).
Moreover, we observed abundant apoptotic cells in tumors from cecropinXJ-treated group via H&E and TUNEL staining (Fig. 6E).

The expression of caspase-3, -8, -9, Bax, Bcl-2 and cytochrome c in tumors was further detected. We found that cecropinXJ treatment increased the expression of caspase-3,
Figure 4. CecropinXJ promotes ROS production and decreases mitochondrial membrane potential ($\Delta\psi_m$) in BGC823 cells. BGC823 cells were treated with different concentrations (0, 20, 50, 80 and 100 µg/ml) of cecropinXJ for 24 h or 100 µg/ml cecropinXJ for 0, 4, 6, 12 and 24 h. ROS production (A) and $\Delta\psi_m$ (C) were detected by DCFH$_2$-DA and DiOC$_6$(3), respectively, and analyzed by flow cytometry. (B and D) Mean fluorescence intensity (mean ± SD) of ROS and $\Delta\psi_m$ is shown in B and D, respectively. Data are from three independent experiments. *P<0.05, **P<0.01.
Figure 5. CecropinXJ induces caspase-dependent apoptosis in BGC823 cells. BGC823 cells were treated with various concentrations of cecropinXJ (0, 20, 50, 80 and 100 µg/ml) for 24 h and the expression and activity of caspases were detected. (A) The gene expression of apoptosis-associated proteins was detected by qRT-PCR. GAPDH was used as a reference gene. (B) The protein levels of apoptosis-associated proteins were detected by western blot analysis. (C) The activities of caspase-3, caspase-8 and caspase-9 was measured. (D) The effects of caspase inhibitors on cecropinXJ-induced cell death. BGC823 cells were pre-treated with the pan-caspase inhibitor and specific caspase inhibitors and then treated with cecropinXJ. The cell death ratio (%) of BGC823 cells was detected by MTT assay. The data are shown mean ± SD. *P<0.05, **P<0.01.
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-8, -9, Bax and cytochrome c but decreased the expression of Bcl-2, which was consistent with the in vitro data.

CecropinXJ inhibits the expression of angiogenesis-associated proteins in vitro and in vivo. Tumor angiogenesis is initiated by production of angiogenic growth factors from tumor, stromal and infiltrating inflammatory cells (30). Upon cecropinXJ treatment, we examined the expressions of angiogenesis-associated proteins including CD31, CD34, bFGF, VEGF and VWF in BGC823 cells and tumors by qRT-PCR and western blot analysis. As shown in Fig. 7, the expression of all these proteins in BGC823 cells treated with cecropinXJ was downregulated at both RNA and protein levels. Importantly, the expression of all these proteins in tumors treated with cecropinXJ was also decreased by immunohistochemical staining. The results suggest that cecropinXJ could inhibit tumor angiogenesis.

Discussion

Chemotherapy alone shows low efficacy that may be due to drug resistance, and severe side-effects that affect normal mammalian cells (31,32). Since Moore et al (33) first found that Cecropin B possessed potent anticancer activity in mammalian cancer cells, a large body of evidence shows that cecropin families exerted cytotoxic effects on a number of tumor derived cell lines in vitro and exhibited inhibitory effects on tumor growth in vivo, such as human breast adenocarcinoma (34) and hepatocellular carcinoma (35). Cecropins contained amphipathic α-helices that allowed them to target non-polar anionic plasma membranes and formed ion-permeable channels to result in cell depolarization, irreversible cytolysis and final cell death (36). The characteristic of cecropins may avoid drug resistance.

Previous study has determined that the cecropins from Chinese oak silkworm possess effective antitumor activity against the human colon adenocarcinoma cells but without cytotoxicity against normal human gastric epithelial cells (GES-1), suggesting cecropins could specifically kill tumor cells (37). Other evidence also showed that cecropins could suppress the growth of tumor cells such as leukemia cells, but have no hemolytic activity against human erythrocytes and no or little toxicity to normal mammalian cells (38). Consistently, we observed that cecropinXJ inhibited the proliferation of BGC823 cells but did not affect normal GES-1 cells. The inhibitory effects of 100 µg/ml cecropinXJ on the proliferation of BGC823 cells was similar to 10 µg/ml Dox. The specific effects on cancer cells might be attributed to the nature of cecropins contained cationic sequence to interact with non-polar anionic plasma membranes and formed non-polar anionic plasma membranes of cancer cells (39). Wang et al (40) reported that cecropins could disrupt plasma membrane integrity of cancer cells when the concentration was >60.9 or 81.2 µg/ml. Lee et al (41) observed that CopA3 mainly caused necrotic cell death of the gastric cancer cells.
However, in the present study, the LDH release of BGC823 cells was not increased after 80 µg/ml cecropinXJ treatment, and cecropinXJ mainly induced apoptosis of BGC823 cells. Consistent with our results, cecropin of Musca domestica induced DNA fragmentation to induce apoptosis (6). Therefore, cecropins may inhibit tumor cells through different mechanisms via inducing necrosis or apoptosis.

Cerón et al (26) reported that cecropin A was able to induce apoptosis in HL-60 cells through the production of ROS but independent of caspase activation. Huang et al (42) showed that pardxin, an antimicrobial peptide was first isolated from secretions of the Red Sea Moses sole, exerted antitumor effects on human fibrosarcoma HT-1080 cells by increasing intracellular ROS and activating caspase-3/7 activities. Our results showed that cecropinXJ increased ROS level and activated caspase-3/8/9 activity in BGC823 cells. Using caspase inhibitors, we further found that caspase-3/9 were involved in the induced apoptosis, but caspase-8 had no effect.

Bcl-2 family proteins including anti-apoptotic (Bcl-2 and Bcl-xL) and pro-apoptotic members (Bax and Bad) play critical roles in the regulation of mitochondrial-mediated apoptosis or cell survival (43). Bcl-2 protein can bind to the outer membrane of the mitochondrion and prevent apoptosis, while Bax is responsible for permeabilizing the membrane (44,45). It has been reported that melittin could decrease the expression of Bcl-2 and induce the translocation of Bax to the mitochondria with the release of cytochrome c into the cytosol, suggesting that mitochondria are involved in melittin-induced apoptosis (46). In the present study, cecropinXJ treatment increased the release of cytochrome c into the cytoplasm, decreased mitochondria ΔΨm, upregulated the expression of Bax and downregulated the expression of Bcl-2 in vitro, suggesting that cecropinXJ triggered mitochondrion-dependent apoptosis. We further found that cecropinXJ inhibited growth of gastric cancer in mice and induced cancer cell apoptosis in tumors. Consistent with the in vitro results, cecropinXJ treatment could upregulate the expression of Bax and downregulate the expression of Bcl-2. Hence, cecropinXJ were able to induce the BGC823 cells apoptosis both in vitro and in vivo.

Angiogenesis is the formation of new blood vessels by the sprouting of endothelial cells from pre-existing vessels (47) and is essential for tumor growth and metastasis (48). It has been reported that suppression of angiogenesis might be an effective therapeutic approach against gastric cancer (49). A number of growth factors related to tumor angiogenesis have been identified which includes bFGF (50), VWF (51) and members of VEGF family (52). We found the expression of the CD31, CD34, VEGF, bFGF and VWF gene in BGC823 cells and BGC823-xenograft tumors was significantly decreased by cecropinXJ treatment, suggesting that cecropinXJ might inhibit tumor angiogenesis in vivo.

In summary, our results showed that cecropinXJ inhibits the growth of human BGC823 cells in vitro and in vivo through inducing apoptosis and preventing tumor angiogenesis. These results suggested that cecropinXJ has potential as a novel anti-tumor therapeutic strategy for the treatment of gastric cancer.
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