# Apoptosis induced by adenosine involves endoplasmic reticulum stress in EC109 cells

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Received June 2, 2012; Accepted July 23, 2012

## DOI: 10.3892/ijmm.2012.1085

Abstract. Apoptosis plays a critical role in the development and homeostasis of multicellular organisms, and endoplasmic reticulum stress (ERS) is one of the intrinsic apoptosis pathways. Previous studies have shown that adenosine induces apoptosis in several cancer cell lines. However, the molecular mechanism remains poorly understood. In this study, we explored whether adenosine triggers apoptosis of EC109 esophageal carcinoma (EC) cells by ERS. The MTT assay was used to determine cell proliferation; cell cycle detection (FCM) and terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling (TUNEL) assay were performed to determine cell apoptosis. The subcellular distribution and expression of the ERS-related proteins GRP78, cleaved caspase-3, cleaved caspase-4, CHOP and NF-KB p65 were detected by western blot techniques. NF-KB activation was measured by electrophoretic mobility shift assay (EMSA). The MTT assay demonstrated that adenosine inhibited EC109 cell proliferation in a dose- and time-dependent manner. FCM and TUNEL assay verified that adenosine caused an apoptotic peak in cell cycle arrest and a higher percentage of apoptotic cells. Western blot analysis confirmed that the expression of GRP78, cleaved caspase-4, CHOP, NF-KB p65 and cleaved caspase-3 were upregulated in a dose-dependent manner after adenosine treatment. EMSA revealed that adenosine activated NF-kB p65. This is the first demonstration that adenosine inhibits cell proliferation, increases GRP78 and NF-kB p65 expression and induces apoptosis by CHOP and caspase-4 pathways. The ERS pathway is involved in adenosine-induced apoptosis in EC109 cells.

# Introduction

Esophageal carcinoma (EC) ranks as the seventh and sixth in terms of cancer incidence and mortality in the world (1), and recent evidence shows that its incidence is gradually increasing. Moreover, nearly 50% of EC cases in the world occurred in China (2) and Shantou is a high-risk region of the disease. Prognosis of this disease is poor, with an overall 5-year survival rate of <10%. Even in resectable stages, the results of standard therapy modalities (surgery or radiotherapy) have been poor, with 5-year survival rates of approximately 20%. Traditional approaches to cancer therapy involve surgery, radiation therapy and chemotherapy. More recent efforts target specific biochemical mechanisms, such as apoptotic pathways.

Adenosine, a metabolite of ATP, is abundantly present inside and outside cells, and exerts diverse biological actions in a wide range of cell types (3). However, high concentrations of nucleotides and their analogues can interfere with the synthesis of nucleic acids and exert cytotoxic activity by incorporating with and altering the DNA or RNA macromolecules, or by modifying the metabolism of physiologic nucleotides (4), which make them exciting candidates for anticancer therapy (5,6). In our previous studies, adenosine induced human hepatoma HepG2 cell apoptosis (7). However, whether adenosine has anticancer effects on human EC is still unknown.

Apoptosis mediated by endoplasmic reticulum stress (ERS) plays a key role in many diseases, such as diabetes and Alzheimer's disease (8). The ER is the primary site for secretory protein synthesis and maturation, Ca2+ storage and lipid biosynthesis. Various stimuli can disturb ER homeostasis and result in the accumulation of unfolded and misfolded proteins and pathological consequences, namely ERS (9). Meanwhile, the accumulating unfolded proteins activate an adaptive signaling cascade known as unfolded protein response (UPR) (10). In response to ERS, ER chaperones such as 78 kDa glucose-regulated protein (GRP78) are upregulated to stabilize protein folding. GRP78 belongs to the heat shock protein 70 (HSP70) family member and generally, it is highly expressed in various types of tumor tissues, resulting in the promotion of tumor cell survival, resistance to chemotherapy, higher incidence of malignancy and poorer prognosis (11). Notably, GRP78 is localized to the plasma membrane of tumor cells where it has receptor-like functions which are associated with increased cellular proliferation and survival (12). The early UPR enhances cell survival by UPR- and ER-associated protein degradation to ensure that the adverse

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*Key words:* adenosine, apoptosis, endoplasmic reticulum stress, glucose-regulated protein 78, C/EBP homologous protein, EC109 cells

effects of ERS are dealt with in a timely and efficient manner. However, prolonged activation of ERS can turn on a cell death pathway through activation of C/EBP homologous protein (CHOP) (13), caspase-12 (14) and/or c-JUN NH2-terminal kinase (JNK) (15). In the present study, we investigated whether adenosine induces EC109 cell apoptosis and if so, the possible ERS-relative molecular mechanisms.

## Materials and methods

Materials. The kit for terminal deoxynucleotidyl transferasemediated dUTP nick end labeling (TUNEL) was from Roche Applied Science (Indianapolis, IN, USA). 4',6-Diamidino-2-phenylindol dihydrochloride (DAPI), propidium iodide (PI), phenylmethylsulfonylfluoride (PMSF), and protease inhibitor cocktail were from Sigma (St. Louis, MO, USA). Primary antibodies against anti-β-actin, caspase-3, cleaved caspase-3, caspase-4, cleaved caspase-4, NF-KB p65, CHOP and all secondary antibodies were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA, USA). Nitrocellulose membrane was from Hybond-C (Amersham Life Science, UK), and an enhanced chemiluminescence detection reagent (ECL kit) was from Thermo Scientific (Rockford, IL, USA). The <sup>32</sup>P-labeled  $\gamma$ -ATP (3,000 Ci/mmol) was from Furui Biotech Co. (Beijing, China). NF-κB oligonucleotide probe (forward, 5'-AGT TGA GGG GAC TTT CCC AGG C-3' and reverse, 3'-TCA ACT CCC CTG AAA GGG TCC G-5'), adenosine, SDS and MTT were from Sangon (Shanghai, China); T4 Polynucleotide Kinase and poly(dI-dC)-poly(dIdC) were from Promega Corporation (Madison, WI, USA). All other chemicals and reagents were of analytical grade.

Cell culture and treatment. EC109 cells were grown in DMEM supplemented with 10% fetal bovine serum, 1% penicillin-streptomycin (final concentration 100 U/ml and 100  $\mu$ g/ml streptomycin), in a humidified atmosphere of 5% CO<sub>2</sub> and 95% air at 37°C. Cells were detached by 0.25% trypsin/0.02% EDTA. For the immunofluorescence assay and TUNEL assay, cells were cultured on coverslips in DMEM for 12 h before addition of adenosine.

Cell proliferation. EC109 cells were seeded at a density of  $2x10^{5}$ /well in a 96-well plate with 200  $\mu$ l culture medium. After the cells were attached, the original medium was replaced with fresh medium containing different concentrations of adenosine (0-4.0 mmol/l) for 36 h or containing 2.0 mmol/l adenosine for 0 to 72 h. The cells were then incubated at 37°C with 20  $\mu$ l MTT (500  $\mu$ g/ml) for another 4 h in a 5% CO<sub>2</sub> incubator. The medium was removed and 150  $\mu$ l DMSO was added to each well, followed by thermal agitation for 5 min protected from light. The optical density of each well was read at 490 nm using a microplate reader (Wellscan K3; KHB Labsystems, Finland). MTT is taken only into viable cells, and the MTT intensity corresponds to the number of viable cells. To assess cell viability, the percentage of the independent basal levels (MTT intensities of cells untreated with adenosine) was calculated.

*Cell cycle analysis*. After EC109 cells were treated with different concentrations of adenosine (0-4.0 mmol/l) for 36 h,

cells in the supernatant were collected by centrifugation and combined with adherent cells that had been trypsinized. Cells were then washed with 5 ml PBS (pH 7.2) and fixed in 70% ethanol at 4°C overnight. Cells were washed twice with PBS to remove the ethanol and passed through a 0.44-mm filter to remove aggregates. The cells were incubated in PBS containing RNase A (2.0 µg/ml) for 1 h at 37°C, followed by staining with PI (5.0  $\mu$ g/ml) for 20 min protected from light on ice. Cells were collected on a nylon mesh filter (pore size, 40  $\mu$ m) and cell cycle phases including the sub-G<sub>1</sub> phase (apoptotic cells) were assayed with a flow cytometer (FACSCalibur; Becton-Dickinson, USA) at an excitation of 488 nm and an emission of 585 nm and analyzed using Lysis II software. A total of  $1 \times 10^4$  cells were counted for each sample. Data were obtained from experiments performed in triplicate.

TUNEL assay. EC109 cells were incubated for 36 h with different concentrations of adenosine (0-4.0 mmol/l), cells were washed with ice-cold PBS and fixed with 10% paraformaldehyde. Cells were subsequently washed 3 times with PBS and permeabilized by 0.2% Triton X-100 in PBS for 5 min at room temperature. After washing twice, cells were equilibrated for 10 min in equilibration buffer [200 mmol/l potassium cacodylate, 25 mmol/l Tris-HC1, pH 6.6, 0.2 mmol/l dithiothreitol (DTT), 0.25 mg/ml BSA, 2.5 mmol/l cobalt chloride] and incubated in TdT reaction mixture (equilibration buffer 98  $\mu$ l, biotinylated nucleotide mix 1  $\mu$ l and TdT enzyme 1  $\mu$ l; total volume 100  $\mu$ l) at 37°C for 60 min in humidified chambers. After immersing the slides in 2X SSC (NaCl, sodium citrate) for 15 min at room temperature and washing the slides twice in PBS, endogenous peroxidases were blocked by immersing the slides in 0.3% hydrogen peroxide for 5 min at room temperature. Slides were then incubated with horseradish-peroxidase-labeled streptavidin (HRP) solution for 30 min and washed twice with PBS. Finally, slides were incubated with diaminobenzidine (DAB) components for 10 min and examined under a light microscope. All experiments were performed 3 times. TUNEL-positive (brown staining) cells were defined as apoptotic cells and 6 randomly selected microscopic fields in each group were used to calculate the percent of TUNEL-positive cells.

Western blot analysis. After adenosine treatment, EC109 cells were washed with ice-cold PBS, harvested and lysed on ice in RIPA buffer [0.25% (w/v) sodium deoxycholate, 0.1% (v/v) Triton X-100, 150 mM NaCl, 0.5% Nonidet P-40, 50 mM Tris-HCl (pH 7.4)] with the addition of protease inhibitor cocktail (Sigma). Cell lysates were centrifuged at 12,000 rpm at 4°C for 10 min. The supernatant [20  $\mu$ g including the sub-G<sub>1</sub> phase (protein/well)] was boiled for 5 min and subjected to 12% sodium dodecylsulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to a nitrocellulose membrane. Membranes were blocked with 5% non-fat milk for 30 min at room temperature, followed by incubation with a primary antibody against  $\beta$ -actin (1:1,000), GRP78 (1:800), caspase-3 (1:800), cleaved caspase-3 (1:800), caspase-4 (1:400), cleaved caspase-4 (1:800), CHOP (1:500) or NF-кB p65 (1:500) at 4°C overnight and with a secondary antibody (HRP-conjugated anti-goat or anti-rabbit IgG) for 1 h. Protein expression was analyzed by NIH image software and normalized to that of  $\beta$ -actin.

Electrophoretic mobility shift assay. The EC109 cells were treated with 2.0 mmol/l adenosine for 36 h. The cells were rinsed with ice-cold PBS and centrifuged. The cell pellets were resuspended with buffer A [10 mmol/l HEPES (pH 7.9), 1.5 mmol/l MgCl<sub>2</sub>, 15 mmol/l KCl, 1 mmol/l DTT, 0.5 mmol/l PMSF, 0.1 mmol/l EDTA, 0.1 mM EGTA, leupeptin, aprotinin and pepstatin (2  $\mu$ g/ml each)] for 10 min on ice. After centrifugation, the pellets were resuspended with buffer A containing 0.5% (v/v) Nonidet P-40 and further incubated for 10 min on ice. The cell suspension was then centrifuged at 4°C for 20 min at 3,000 rpm. The nuclear pellets were resuspended with buffer B [20 mmol/l HEPES (pH 7.9), 420 mmol/l NaCl, 1.5 mmol/l MgCl<sub>2</sub>, 0.2 mmol/l EDTA, 0.5 mmol/l PMSF, 0.5 mmol/l DTT, 25% (v/v) glycerol, leupeptin, aprotinin and pepstatin (2 µg/ml each)]. The resuspended nuclei were then extracted by rocking at 40°C for 1 h, then centrifuging for 20 min at 10,000 rpm. The supernatant (nuclear proteins) was collected and protein concentration was determined using a Bradford protein assay. The EMSA method used was similar to that described previously (16), with slight modifications. A double-stranded NF-kB oligonucleotide probe (forward, 5'-AGT TGA GGG GAC TTT CCC AGG C-3' and reverse, 3'-TCA ACT CCC CTG AAA GGG TCC G-5') was end-labeled with  $[\gamma^{-32}P]$ ATP. The labeling reaction system contained T4 Polynucleotide Kinase buffer and T4 Polynucleotide Kinase (10  $\mu/\mu$ l), nuclease-free water,  $[\gamma^{-32}P]ATP$  (3,000 Ci/mmol at 10 mCi/µl) and unlabeled probe (10 pmol/ $\mu$ l). Ten milligrams of nuclear extraction was incubated for 10 min at room temperature with binding buffer [10 mmol/l HEPES (pH 7.9), 50 mmol/l KCl, 0.2 mmol/l EDTA, 2.5 mmol/l DTT, 0.05% (v/v) Nonidet P-40, 10% (v/v) glycerol and 1  $\mu g/\mu l$  poly(dI-dC)-poly(dI-dC)]. Labeled NF-kB oligo was then added and the reaction was allowed to proceed for a further 20 min at room temperature. In competition experiments, a 100-fold molar excess of the cold unlabeled oligonucleotide was added to the nuclear extracts for 30 min before addition of a radiolabeled probe. The specific antibody against NF-KB p65 protein (final concentration 0.1  $\mu g/\mu l$ ) was pre-incubated with nuclear extracts in binding buffer for 30 min at room temperature before addition of the radiolabeled probe. Samples were then separated on 6% nondenaturing polyacrylamide gel electrophoresis with 0.5X Tris borate EDTA buffer at 4°C. Dried gel was subjected to autoradiography at -80°C.

Statistical analysis. Data are expressed as means  $\pm$  SEM of at least 3 independent experiments. Results were analyzed with the unpaired Student's t-test or one-way ANOVA for different groups in SPSS 16.0. A P<0.05 was considered to indicate a statistically significant result.

#### Results

Adenosine inhibits cell proliferation. EC109 cells were exposed to 2 mmol/l adenosine for different times from 24 to 72 h and cell proliferation was evaluated by MTT assay. The cell relative viabilities were decreased to 57.4, 55.4, 44.3



Figure 1. Time- and dose-dependent cytotoxic effects of adenosine on EC109 cells. (A) Cells were incubated with 2.0 mmol/l adenosine for different durations and the cell relative viabilities were significantly decreased, compared to the control group (0 h). (B) Cells were incubated with adenosine at different concentrations for 36 h and the cell numbers were significantly decreased, compared to the control group (absence of adenosine). Cell numbers were determined by MTT method. Data are presented as the means  $\pm$  SD. Independent experiments were repeated 3 times. \*P<0.05, \*\*P<0.01 vs. control group.

and 28.15%, respectively, compared to the control group (0 h), showing that adenosine inhibited cell proliferation in a time-dependent manner (P<0.01) (Fig. 1A). EC109 cells were exposed to different concentrations of adenosine, 0.5, 1.0, 2.0 and 4.0 mmol/l, for 36 h, and the cell numbers were decreased to 88.2, 67.1, 54.7 and 45.6%, respectively, compared to cell numbers in the absence of the adenosine group, revealing adenosine inhibited cell proliferation in a dose-dependent manner (P<0.05 and P<0.01) (Fig. 1B).

Adenosine induces apoptosis. To explore whether EC109 cell proliferation inhibition caused by adenosine was a result of necrosis or apoptosis, two separate methods were used for these experiments. In one case, we assessed and quantified the presence of apoptotic cells by performing cell cycle assay. After EC109 cells were exposed to various concentrations of adenosine for 36 h, with adenosine concentration increasing, sub-G<sub>1</sub> cell accumulation (hallmark of apoptosis) obviously appeared. The proportion of apoptotic cells in 0.5, 1.0, 2.0 and 4.0 mmol/l adenosine-treated groups was 6.2-, 12.2-, 17.5- and 21.1-fold higher, respectively than that observed in the control group (Fig. 2).

A more specific method for identifying apoptosis was used in a second set of experiments. DNA fragmentation, associated with apoptosis, generates many small nucleosomal-sized DNA fragments, which can be detected by the



Figure 2. Effects of adenosine on EC109 cell cycles. EC109 cells were treated with various concentrations of adenosine (0, 0.5, 1.0, 2.0 and 4.0 mmol/l) for 36 h. Cell cycle phases were determined by propidium iodine staining, followed by flow cytometric analysis. The percentages of apoptotic cells in the sub-G<sub>1</sub> phase (blue peak) were detected and analyzed by the WinMDI 2.9 software.



Figure 3. Effects of adenosine on EC109 cell apoptosis by TUNEL analysis. (A) EC109 cells were treated with various concentrations (0-4.0 mmol/l) of adenosine for 36 h. Apoptotic cells were identified by light microscopy as cells having dark brown nuclei. (B) Apoptotic cells were quantified by counting a minimum of 6 fields of 100 cells/field (magnification, x400), \*P<0.05, \*\*P<0.01 vs. control (absence of adenosine).

TUNEL method. These new DNA fragments increase the number of 3'-OH ends. Incorporating digoxigenin-conjugated nucleotides, via end extension, into these small nucleosomal fragments identifies cells with significant DNA fragmentation. Adenosine treatment caused the appearance of cells with apoptotic (dark brown colored) nuclei (TUNEL-positive cells) (Fig. 3A). Adenosine increased the percentage of TUNEL-positive cells in a dose-dependent manner, as



Figure 4. Effects of adenosine on cleaved caspase-3 by western blot analysis. (A) EC109 cells were exposed to different concentrations of adenosine (0.5-4.0 mmol/l) for 36 h. Cell lysates were subjected to western blot analysis with a specific antibody against caspase-3 or cleaved caspase-3. Blots shown are representative of n=3. (B) Ratio of cleaved caspase-3 to  $\beta$ -actin. \*P<0.05, \*\*P<0.01 vs. control (absence of adenosine).

compared with control cells (Fig. 3B) (P<0.05). The apoptotic cell percentage identified by the TUNEL method was in agreement with the apoptotic peak of the cell cycle assay. Adenosine also significantly increased cleaved caspase-3 expression in a dose-dependent, compared to the control group (Fig. 4) (P<0.05 and P<0.01) and expression of procaspase-3 decreased, which showed that caspase-3 was activated and apoptosis occurred in the EC109 cells.

Adenosine upregulates ERS-related proteins. To assess whether ERS is activated after adenosine treatment, western blot analysis was used to analyze relative protein expression. The results showed that UPR chaperone GRP78 was significantly upregulated in the 1.0-4.0 mmol/l adenosine treatment groups, as compared with the control group (Fig. 5) (all P<0.05), respectively. Caspase-4 and CHOP are two key molecules in the three branches of ERS-associated apoptosis. Both cleaved caspase-4 and CHOP were significantly increased in a concentration-dependent manner, compared to that of the control group (Fig. 5) (P<0.05 and P<0.01); whereas the procaspase-4 was decreased, indicating the increased processing of procaspase-4 to active caspase-4 after adenosine treatment. These results suggested that adenosine activated caspase-4 and CHOP pathways by GRP78.

Adenosine induces NF- $\kappa$ B activation and expression. We performed EMSA on nuclear extracts from EC109 cells using a  $\kappa$ B sequence from  $\kappa$  light chain enhancer in order to observe the effect of different concentrations of adenosine on NF- $\kappa$ B p65 induction. As shown in Fig. 6A, adenosine stimulated DNA binding of NF- $\kappa$ B after 36 h of incubation at concentrations from 0.5 to 4.0 mmol/l. The effects increased with the adenosine dosage; 4.0 mmol/l adenosine appeared to more strongly induce NF- $\kappa$ B activation (Fig. 6B) (P<0.01). Western blot analysis also revealed that NF- $\kappa$ B significantly increased



Figure 5. (A) Expression of ER stress marker-related proteins after adenosine exposure. EC109 cells were exposed to different concentrations of adenosine (0.5-4.0 mmol/l) for 36 h. Cell lysates were subjected to western blot analysis with specific antibodies against GRP78, caspase-4, cleaved caspase-4, or CHOP. Blots shown are representative of n=3. (B) Band intensity analysis of ER stress marker-related protein expression. Band intensity analysis was performed with Quantity one software. The values represent the means  $\pm$  SD of 3 independent experiments. \*P<0.05, \*\*P<0.05 vs. the control group.

in a dose-dependent manner, compared to the control group (Fig. 6C and D) (P<0.05 and P<0.01).

#### Discussion

Adenosine belongs to the cytotoxic nucleotide analogue class of drugs. In physiological conditions, the average concentration of nucleotides in plasma and other extracellular fluids is generally in the range of 0.4-6  $\mu$ mol/l. However, high concentrations of nucleotides and their analogues can interfere with the synthesis of nucleic acids and exert cytotoxic activity by incorporating with and altering the DNA or RNA macromolecules, or by modifying the metabolism of physiologic nucleotides (4), which make it an exciting candidate for anticancer therapy (17).



Figure 6. Effects of adenosine on NF- $\kappa$ B p65 DNA binding and protein expression. (A) EC109 cells (2x10<sup>6</sup> cells) were incubated in culture medium alone (control, lane 2) or in the presence of different concentrations of adenosine (0.5-4.0 mmol/l, lanes 4-7) for 36 h. Nuclear extracts were subjected to EMSA analysis, as described in experimental procedures. The negative control contained the labeled probe and no protein in the binding reactions (lane 1). To demonstrate specificity of induced bands, binding was carried out in the presence of a molar excess (x100) of unlabeled probe containing NF- $\kappa$ B consensus sequence (lane 3). Specific DNA-NF- $\kappa$ B complex and free probes are indicated by the arrows. (B) Results of the densitometric analysis for the specific NF- $\kappa$ B binding. Bands shown are representative of n=3. ("P<0.05, \*\*P<0.01 vs. the control group-lane 2). (C) EC109 cells were exposed to different concentrations of adenosine for 36 h. Cell lysates were subjected to western blot analysis using antibodies against NF- $\kappa$ B p65 and  $\beta$ -actin. Blots shown are representative of n=3. (D) Ratio of NF- $\kappa$ B to  $\beta$ -actin. \*P<0.01 vs. control (absence of adenosine).

Since Rapaport and Fontaine (18) reported that extracellular adenine nucleotides had anticancer activity, adenosine and its analogues (8-Cl-adenosine) have been widely studied in various tumor cell lines, including human thyroid cancer lines (17), colonic cancer Caco-2 (19), breast cancer (20), hepatoma HepG2 (7) and multiple myeloma (21). In this study, EC109 cells were exposed to 2.0 mmol/l adenosine for different times from 24 to 72 h or exposed to different concentrations of adenosine for 36 h. The results showed that adenosine significantly inhibited cell proliferation in a time-dependent and a dose-dependent manner (Fig. 1) and FCM and TUNEL methods verified that the extensive cell death was apoptosis (Figs. 2 and 3). The present experiment indicates that adenosine also has cytotoxic effects on EC109 cells.

In many conditions, the transcription of GRP78 is upregulated as part of a general cellular defense mechanism (UPR) (22-24). To assess whether UPR and ERS are activated after adenosine treatment, western blot analysis was used to analyze relative protein expression. The results showed that UPR chaperone GRP78 was significantly upregulated after adenosine treatment in a dose-dependent manner (Fig. 5), which was in agreement with other studies that chemotherapy drugs can induce GRP78 upregulation and UPR (25-27).

Caspase-4 and CHOP are two key molecules in the three branches of ERS-associated apoptosis: a caspase-4-dependent pathway and/or transcriptional induction of CHOP. CHOP belongs to the C/EBP family of transcription factors. Translocation of CHOP from the cytoplasm to the nucleus regulates the expression of ERS-related genes (28). Overexpression of CHOP promotes cell apoptosis and deficiency or inhibition of CHOP by drugs can protect cells from ERS-induced apoptosis (29). Cascade activation of caspases (such as caspases-3, -8 and -9) play a more important role in cell apoptosis (30). Among 14 known caspases, caspase-12 seems to be involved in signaling pathways specific to ERS-induced apoptosis (14). Cleaved caspase-12 activates caspase-9, followed by activation of caspase-3. Caspase-12knockout mice exhibit resistance to ERS, suggesting that caspase-12 plays a key role in ERS-mediated apoptosis. In humans, caspase-12 protein cannot be produced because the gene is interrupted by a frame shift and a premature stop codon. Caspase-4, another caspase-1 family member, is homologous and functions similarly to mouse caspase-12 (31). In the present study, both cleaved caspase-4 and CHOP were significantly increased in a concentration-dependent manner (Fig. 5). Cleaved caspase-3 was also upregulated in a dose-dependent manner (Fig. 4). Taken together, this study demonstrated that adenosine activated UPR by chaperone GRP78 and UPR triggered CHOP and caspase-4 pathways, finally resulting in caspase-3 activation and cell apoptosis. However, the precise mechanisms need further investigation.

NF- $\kappa$ B is an important mediator of inflammation and carcinogenesis (32). Studies have showed that various agents that induce ERS can activate NF- $\kappa$ B, and the activated NF- $\kappa$ B dimer is rapidly released from the cytoplasm, where it is normally sequestered by the inhibitory unit I $\kappa$ B, and then is translocated to the nucleus, where it activates transcription

of downstream genes (33). Active NF-κB is most commonly composed of the heterodimer DNA binding subunits p50 and p65. The role of NF- $\kappa$ B in regulating cell death is complex. In some cases it has been demonstrated to promote cell survival and in other cases to promote cell death, which may depend on the cell type and the nature of the toxic stimuli (34,35). In this study, we performed EMSA on nuclear extracts from EC109 cells using a  $\kappa$ B sequence from  $\kappa$  light chain enhancer in order to observe the effect of different concentrations of adenosine on NF-kB p65 induction. As shown in Fig. 6A, adenosine stimulated DNA binding of NF-kB after 36 h of incubation at concentrations from 0.5 to 4.0 mmol/l. The effects increased with the adenosine dosage; 4.0 mmol/l adenosine appeared to more strongly induce NF-kB activation (Fig. 6B). Western blot analysis also revealed that NF-KB p65 protein expression significantly increased in a dose-dependent manner (Fig. 6C and D). This is consistent with a previous report that demonstrated that high levels of GRP78 expression may promote cellular proliferation by the NF-κB pathway (23). In our previous study, we observed that NF-κB displayed an anti-apoptotic effect on HepG2 cells (36). In this study, we also found that NF-KB p65 was highly activated and expressed (Fig. 6A and C). We suppose that GRP78-induced activation of UPR both activate the NF-kB pathway and activate ERS-mediated apoptosis pathways by CHOP and cleaved caspase-4. In adenosine-mediated apoptosis, whether the activation of NF-κB p65 represents a cellular defense mechanism and how NF-kB regulates its downstream genes remain to be further investigated.

In summary, the results of this study showed that adenosine inhibited cell proliferation, increased GRP78 and NF- $\kappa$ B p65 expression and induced apoptosis by CHOP and caspase-4 pathways. The ERS pathway may be involved in adenosine-induced EC109 cell apoptosis.

#### Acknowledgements

This study was supported by the National Nature Science Foundation of China (no. 30972925).

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