Proteasome inhibition by MG132 induces growth inhibition and death of human pulmonary fibroblast cells in a caspase-independent manner

BO RA YOU and WOO HYUN PARK

Department of Physiology, Medical School, Institute for Medical Sciences, Chonbuk National University, JeonJu 561-180, Republic of Korea

Received November 26, 2010; Accepted January 17, 2011

DOI: 10.3892/or.2011.1211

Abstract. MG132 as a proteasome inhibitor that can induce apoptotic cell death in various cell types including lung cancer cells. We investigated the cellular effects of MG132 on human pulmonary fibroblast (HPF) cells in relation to cell growth inhibition and death, and described the molecular mechanisms of MG132 in HPF cell death. This agent dose-dependently inhibited the growth of HPF cells with an IC_{50} of approximately 20 μ M at 24 h and induced cell death accompanied by the loss of mitochondrial membrane potential (MMP; $\Delta \Psi_m$) and an increase in caspase-3 and -8 activities. MG132 increased intracellular ROS levels and GSH-depleted cell numbers. However, all the tested caspase inhibitors intensified HPF growth inhibition by MG132 and caspase-9 inhibitor also enhanced cell death and MMP ($\Delta \Psi_m$) loss. Moreover, the administration of Bcl-2 siRNA augmented HPF cell death by MG132 whereas p53, Bax, caspase-3 and -8 siRNAs did not strongly affect cell death. In addition, each caspase inhibitor and siRNA differently affects ROS levels including O2+ regardless of cell growth inhibition and cell death levels. Caspase-8 and -9 inhibitors

Correspondence to: Dr Woo Hyun Park, Department of Physiology, Medical School, Chonbuk National University, Institute for Medical Sciences, JeonJu 561-180, Republic of Korea E-mail: parkwh71@chonbuk.ac.kr

Abbreviations: HPF, human pulmonary fibroblast; MG132, carbobenzoxy-Leu-Leu-leucinal; ROS, reactive oxygen species; MMP ($\Delta\Psi_m$), mitochondrial membrane potential; Z-VAD-FMK, benzyloxycarbonyl-Val-Ala-Asp-fluoromethylketone; Z-DEVD-FMK, benzyloxycarbonyl-Asp-Glu-Val-Asp-fluoromethylketone; Z-IETD-FMK, benzyloxycarbonyl-Ile-Glu-Thr-Asp-fluoromethylketone; Z-LEHD-FMK, benzyloxycarbonyl-Leu-Glu-His-Asp-fluoromethylketone; FBS, fetal bovine serum; PI, propidium iodide; FITC, fluorescein isothiocyanate; H₂DCFDA, 2',7'-dichlorodihydrofluorescein diacetate; DHE, dihydroethidium; GSH, glutathione; CMFDA, 5-chloromethylfluorescein diacetate; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; siRNA, small interfering RNA

Key words: MG132, proteasome, cell death, human pulmonary fibroblast, reactive oxygen species

increased the number of GSH-depleted cells in MG132-treated HPF cells. In conclusion, MG132 induced growth inhibition and death in HPF cells in a caspase-independent manner. The growth inhibition and death of HPF cells by MG132 and/or each caspase inhibitor or apoptosis-related siRNA were not tightly related to the changes in ROS levels.

Introduction

The ubiquitin-dependent proteasomal system presents the foremost non-lysosomal corridor (1,2). The proteasome consists of large multi-subunit protease complexes; a 20S catalytic and two 19S regulatory subunits. The protease activity works in a channel at the center of the 20S subunit and exhibits 3 distinguishable enzymatic activities: chymotrypsin-like, trypsin-like and caspase-like (3). Many proteins despoiled by proteasome are implicated in crucial processes: for example, cell cycle-regulatory proteins (cyclins A, B, D and E; p21 and p27) and apoptotic-related proteins (p53 and Bcl-2) (4). Transformed cells including cancer cells accumulate more misfolded/mutated/damaged proteins due to the elevated duplication rate of malignant cells, which are disposed of by the proteasome (4). Thus, these cells can be much more vulnerable to proteasome inhibition than normal cells. Apoptosis in cancer cells is closely connected with the activity of ubiquitin/proteasome pathways (5,6). Accordingly, the inhibition of proteasome function has emerged as a useful strategy to maneuver apoptosis.

The peptide aldehyde MG132 (carbobenzoxy-Leu-Leuleucinal) efficiently prevents the proteolytic activity of the proteasome complex (7). Various proteasome inhibitors including MG132 have been demonstrated to stimulate apoptotic cell death through the induction of reactive oxygen species (ROS) (8,9). ROS formation and glutathione (GSH; a major non-protein antioxidant in the cell) depletion by proteasome inhibitors may trigger mitochondrial dysfunction and subsequent cytochrome c release, which can lead to cell death (10,11). The mechanism underlying ROS generation after inhibition of the proteasome is still imprecise. In addition, although cells have antioxidant systems to manage their redox state, excessive production of ROS can be induced by endogenous and/or exogenous sources, which then initiates events that lead to cell death or survival, depending on cell types (12-14).

Lung cancer is a main cause of cancer death in developed countries. Since the clinical use of conventional drugs is limited due to intrinsic or acquired resistance and toxicity, various novel remedial strategies are presently under consideration (15). Studies of the molecular mechanisms of cytotoxic drug action have shed light on the treatment of lung cancer. Novel drugs that aim at specific intracellular pathways related to the distinctive properties of cancer cells continue to be developed. It has been reported that proteasome inhibitor bortezomib (PS-341, Velcade) inhibits lung cancer cells (16,17). However, the role of MG132 in lung cancer cells is not fully understood. We recently demonstrated that MG132 reduced the growth of Calu-6 and A549 lung cancer cells via apoptosis and GSH depletion (18,19). On the other hand, little is known about the cellular effects of MG132 on normal primary lung cells in relation to cell death.

Because cancer cells accumulate more misfolded/mutated/ damaged proteins which are disposed of by the proteasome (4), it is worthy to know the susceptibility of normal primary lung cells to MG132 and to understand its toxicological mechanisms in these cells. We assumed that the susceptibility of normal human pulmonary fibroblast (HPF) cells to MG132 is lower than that of lung cancer cells. In the present study, we investigated the cellular effects of MG132 on HPF cells in relation to cell growth inhibition and death, and describe the molecular mechanisms of MG132 in HPF cell death. In addition we examined the effects of MG132 and/or caspase inhibitors or apoptosis-related siRNAs on ROS and GSH in HPF cells.

Materials and methods

Cell culture. The human pulmonary fibroblast (HPF) cells from PromoCell GmbH (Heidelberg, Germany) were maintained in humidified incubator containing 5% CO_2 at 37°C. HPF cells were cultured in RPMI-1640 supplemented with 10% fetal bovine serum (FBS) and 1% penicillin-streptomycin (Gibco-BRL, Grand Island, NY). HPF cells were used between passages 4 and 8.

Reagents. MG132 was purchased from Calbiochem (San Diego, CA) and was dissolved in dimethyl sulfoxide (DMSO; Sigma-Aldrich, St. Louis, MO) solution buffer. The pancaspase inhibitor (Z-VAD-FMK; benzyloxycarbonyl-Val-Ala-Asp-fluoromethylketone), caspase-3 inhibitor (Z-DEVD-FMK; benzyloxycarbonyl-Asp-Glu-Val-Asp-fluoromethylketone), caspase-8 inhibitor (Z-IETD-FMK; benzyloxycarbonyl-Ile-Glu-Thr-Asp-fluoromethylketone) and caspase-9 inhibitor (Z-LEHD-FMK; benzyloxycarbonyl-Leu-Glu-His-Aspfluoromethylketone) were obtained from R&D Systems, Inc. (Minneapolis, MN) and were dissolved in DMSO. Based on a previous study (20), cells were pretreated with or without each caspase inhibitor for 1 h prior to MG132 treatment. DMSO (0.2%) was used as a control vehicle and it did not appear to affect cell growth or death.

Cell growth inhibition assays. Cell numbers in HPF cells treated with MG132 and/or each caspase inhibitor were

determined by trypan blue cell counting, and cell growth inhibition in these cells was indirectly evaluated by measuring 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) dye absorbance as previously described (21,22). In brief, 2x10⁵ cells per well were seeded in 24-well plates (Nunc, Roskilde, Denmark) for cell counting and 5x10³ cells per well were seeded in 96-well microtiter plates (Nunc) for MTT assays. After exposure to the indicated amounts (0.1-30 μ M) of MG132 with or without 15 μ M of a given caspase inhibitor for 24 h, the cells in the 24-well plates were collected with trypsin digestion for trypan blue cell counting and the cells in the 96-well plates were used for MTT assays. Twenty microliters of MTT solution [2 mg/ml in phosphate-buffered saline (PBS)] was added to each well of the 96-well plates. The plates were incubated for additional 4 h at 37°C. Media in plates were withdrawn by pipetting and 200 μ l of DMSO was added to each well to solubilize the formazan crystals. Optical density was measured at 570 nm using a microplate reader (Spectra MAX 340, Molecular Devices Co., Sunnyvale, CA).

Western blot analysis. The patterns of ubiquitinated proteins were evaluated using Western blot analysis. In brief, 1x106 cells in 60-mm culture dish (Nunc) were incubated with 10 or $30 \,\mu M$ MG132 for 24 h. The cells were then washed in PBS and suspended in 5 volumes of lysis buffer (20 mM HEPES, pH 7.9, 20% glycerol, 200 mM KCl, 0.5 mM EDTA, 0.5% NP40, 0.5 mM DTT, 1% protease inhibitor cocktail). Supernatant protein concentrations were determined using the Bradford method. Samples containing 40 μ g total protein were resolved by 12.5% SDS-PAGE gels, transferred to Immobilon-P PVDF membranes (Millipore, Billerica, MA) by electroblotting and then probed with anti-ubiquitin and anti-\beta-actin antibodies (Santa Cruz Biotechnology, Santa Cruz, CA). Membranes were incubated with horseradish peroxidase-conjugated secondary antibodies. Blots were developed using an ECL kit (Amersham, Arlington Heights, IL).

Cell cycle and sub-G1 analysis. Cell cycle and sub-G1 cells were determined by propidium iodide (PI, Sigma-Aldrich; Ex/Em = 488 nm/617 nm) staining, as previously described (23). In brief, 1x10⁶ cells in 60-mm culture dish (Nunc) were incubated with the indicated amounts (0.1-30 μ M) of MG132 for 24 h. Cells were then washed with PBS and fixed in 70% ethanol. Cells were washed again with PBS, then incubated with PI (10 μ g/ml) with simultaneous RNase treatment at 37°C for 30 min. Cellular DNA content was measured using a FACStar flow cytometer (Becton-Dickinson, Franklin Lakes, NJ).

Annexin V/PI staining for cell death detection. Apoptosis was determined by staining cells with Annexin V-fluorescein isothiocyanate (FITC, PharMingen, San Diego, CA; Ex/Em = 488 nm/519 nm) and propidium iodide (PI; Sigma-Aldrich; Ex/Em = 488 nm/617 nm). In brief, $1x10^6$ cells in 60-mm culture dish (Nunc) were incubated with MG132 with or without a given caspase inhibitor or siRNA duplex for 24 h. Cells were washed twice with cold PBS and then resuspended in 500 μ l of binding buffer (10 mM HEPES/NaOH pH 7.4, 140 mM NaCl, 2.5 mM CaCl₂) at a concentration of $1x10^6$ cells/ml. Five microliters of Annexin V-FITC and PI (1 μ g/ml) were then

added to these cells, which were analyzed with a FACStar flow cytometer (Becton-Dickinson). Viable cells were negative for both PI and Annexin V; apoptotic cells were positive for Annexin V and negative for PI, whereas late apoptotic dead cells displayed both high Annexin V and PI labeling. Non-viable cells, which underwent necrosis, were positive for PI and negative for Annexin V.

Measurement of MMP ($\Delta \Psi_m$). MMP ($\Delta \Psi_m$) levels were measured using a rhodamine 123 fluorescent dye (Sigma-Aldrich; Ex/ Em = 485 nm/535 nm) as previously described (22). In brief, 1x10⁶ cells in 60-mm culture dish (Nunc) were incubated with MG132 with or without 15 μ M of a given caspase inhibitor for 24 h. Cells were washed twice with PBS and incubated with the rhodamine 123 (0.1 μ g/ml) at 37°C for 30 min. Rhodamine 123 staining intensity was determined by flow cytometry (Becton-Dickinson). An absence of rhodamine 123 from cells indicated the loss of MMP ($\Delta \Psi_m$) in HPF cells. The MMP ($\Delta \Psi_m$) levels in the cells, excluding MMP ($\Delta \Psi_m$) loss cells, were expressed as the mean fluorescence intensity (MFI), which was calculated by CellQuest software (Becton-Dickinson).

Quantification of caspase-3 and caspase-8 activities. The activities of caspase-3 and -8 were assessed using the caspase-3 and -8 Colorimetric Assay Kits (R&D Systems, Inc.). In brief, 1x10⁶ cells in 60-mm culture dish (Nunc) were incubated with 10 μ M MG132 for 24 h. The cells were then washed in PBS and suspended in 5 volumes of lysis buffer [20 mM HEPES pH 7.9, 20% glycerol, 200 mM KCl, 0.5 mM EDTA, 0.5% NP40, 0.5 mM DTT, 1% protease inhibitor cocktail (from Sigma)]. Supernatant samples containing 50 μ g of total protein were used for determination of caspase-3 and -8 activities. These samples were added to each well in the 96-well microtiter plates (Nunc) with the DEVD-pNA for the caspase-3 activity assay or IETD-pNA for the caspase-8 activity assay at 37°C for 1 h. The optical density of each well was measured at 405 nm using a microplate reader (Spectra MAX 340, Molecular Devices Co.). The activities of caspase-3 and -8 were expressed in arbitrary absorbance units.

Detection of intracellular ROS and O_2^{\bullet} levels. Intracellular ROS were detected by means of an oxidation-sensitive fluorescent probe dye, 2',7'-dichlorodihydrofluorescein diacetate (H₂DCFDA, Invitrogen Molecular Probes, OR; Ex/Em = 495 nm/529 nm) (24). As H₂DCFDA is poorly selective for O₂[•], dihydroethidium (DHE, Invitrogen Molecular Probes; Ex/Em = 518 nm/605 nm), which is highly selective for O_2^{\bullet} , was used for its detection. In brief, 1x10⁶ cells in 60-mm culture dish (Nunc) were incubated with MG132 with or without a given caspase inhibitor or siRNA duplex for 24 h. Cells were then washed in PBS and incubated with 20 μ M H₂DCFDA or DHE at 37°C for 30 min. DCF and DHE fluorescence was detected using a FACStar flow cytometer (Becton-Dickinson). ROS and O2. levels were expressed as mean fluorescence intensity (MFI), which was calculated by CellQuest software (Becton-Dickinson).

Detection of the intracellular glutathione (GSH). Cellular GSH levels were analyzed using a 5-chloromethylfluorescein diacetate dye (CMFDA, Invitrogen Molecular Probes; Ex/

Em = 522 nm/595 nm) as previously described (24). In brief, 1x10⁶ cells in 60-mm culture dish (Nunc) were incubated with MG132 with or without a given caspase inhibitor for 24 h. Cells were then washed with PBS and incubated with 5 μ M CMFDA at 37°C for 30 min. CMF fluorescence intensity was determined using a FACStar flow cytometer (Becton-Dickinson). Negative CMF staining (GSH depleted) cells were expressed as the percent of (-) CMF cells.

Transfection of cells with apoptosis-related siRNAs. Gene silencing of p53, Bax, Bcl-2, caspase-3 and caspase-8 was performed as previously described (25). A non-specific control siRNA duplex [5'-CCUACGCCACCAAUUUCGU (dTdT)-3'], p53 siRNA duplex [5'-CACUACAACUACAUGU GUA(dTdT)-3'], Bax siRNA duplex [5'-GCUGGACAUUGGA CUUCCU(dTdT)-3'], Bcl-2 siRNA duplex [5'-CAGAAGUCU GGGAAUCGAU(dTdT)-3'], caspase-3 siRNA duplex [5'-AGU AUGCCGACAAGCUUGA(dTdT)-3'] and caspase-8 siRNA duplex [5'-GCUGCUCUUCCGAAUUAAU(dTdT)-3'] were purchased from the Bioneer Corp. (Daejeon, South Korea). In brief, 2.5x10⁵ cells in 6-well plates (Nunc) were incubated in RPMI-1640 supplemented with 10% FBS. The next day, cells (~30-40% confluence) in each well were transfected with the control or each siRNA duplex [80 pmol in Opti-MEM (Gibco-BRL)] using Lipofectamine 2000, according to the manufacturer's instructions (Invitrogen, Branford, CT). One day later, cells were treated with or without 30 μ M MG132 for additional 24 h. The transfected cells were collected and used for the measurement of Annexin V-FITC/PI staining cells and ROS levels.

Statistical analysis. The results represent the mean of at least three independent experiments (mean \pm SD). The data were analyzed using Instat software (GraphPad Prism4, San Diego, CA). The Student's t-test or one-way analysis of variance (ANOVA) with post hoc analysis using Tukey's multiple comparison test was used for parametric data. Statistical significance was defined as p<0.05.

Results

Effects of MG132 on cell growth in HPF cells. We examined the effect of MG132 on the growth of HPF cells at 24 h. Based on an MTT assay, MG132 dose-dependently inhibited the growth of HPF cells with an IC₅₀ of ~20 μ M (Fig. 1A). In addition, treatment with 0.5-30 μ M MG132 significantly decreased the population of viable (trypan blue-negative) HPF cells in a dose-dependent manner whereas the numbers of dead (trypan blue-positive) cells dose-dependently increased in MG132-treated cells (Fig. 1B). The ratio of dead cells to viable cells was increased by MG132 treatment. When the cell cycle distributions in MG132-treated HPF cells were examined, 10 µM MG132 significantly induced S and G2-M phase arrests of the cell cycle at 24 h compared with control cells (data not shown). Moreover, we observed that MG132 increased the level of anonymous ubiquitinated proteins in HPF cells compared with that of the control cells (Fig. 1C).

Effects of MG132 on apoptosis and MMP ($\Delta \Psi_m$) *in HPF cells.* We determined whether MG132 induces cell death via apop-

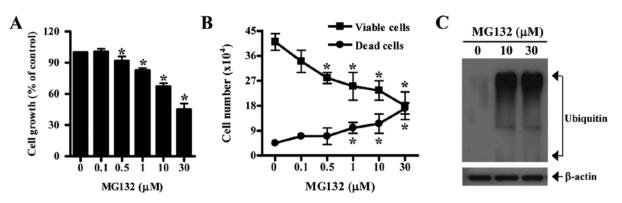


Figure 1. Effects of MG132 on cell growth in HPF cells. Exponentially growing cells were treated with the indicated amounts of MG132 for 24 h. (A) The graph shows cellular growth changes in HPF cells, as assessed by MTT assays. (B) The graphs show the viable (trypan blue-negative) and dead (trypan blue-positive) cell numbers in HPF cells. (C) Samples (40 μ g) of protein extracts were resolved by SDS-PAGE gel, transferred onto PVDF membranes and immunoblotted with the indicated antibodies against ubiquitin and β -actin. *p<0.05 compared with the control group.

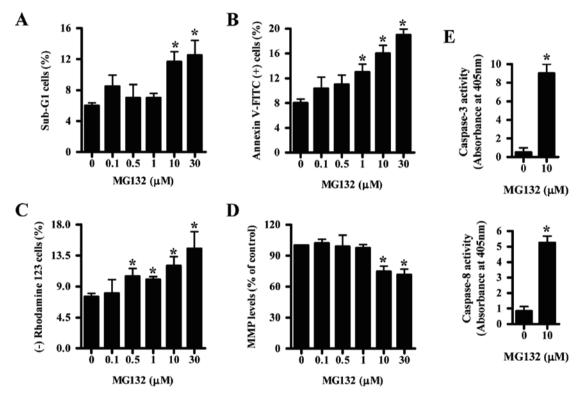


Figure 2. Effects of MG132 on cell death and MMP ($\Delta \Psi_m$) in HPF cells. Exponentially growing cells were treated with the indicated amounts of MG132 for 24 h. (A and B) Sub-Gl cells (A) and Annexin V-FITC cells (B) were measured with a FACStar flow cytometer. (C and D) MMP ($\Delta \Psi_m$) was measured with a FACStar flow cytometer. Graphs show the percent of rhodamine 123-negative [MMP ($\Delta \Psi_m$) loss] cells (C) and MMP ($\Delta \Psi_m$) levels in the cells, excluding MMP ($\Delta \Psi_m$) loss cells (D). (E) The graphs show the activities of caspase-3 and caspase-8 in HPF cells treated with or without 10 μ M MG132. *p<0.05 compared with the control group.

tosis in HPF cells. As shown in Fig. 2A, MG132 increased the number of sub-G1 cells at 24 h. When HPF cells were stained with Annexin V-FITC, the proportion of Annexin V-stained cells following MG132 treatment was significantly increased in a dose-dependent manner (Fig. 2B). In addition, when the loss of MMP ($\Delta \Psi_m$) in MG132-treated HPF cells was determined using a rhodamine 123 dye, MG132 induced the loss of MMP ($\Delta \Psi_m$) in HPF cells in a dose-dependent manner (Fig. 2C). Treatment with 10 or 30 μ M MG132 reduced MMP ($\Delta \Psi_m$) level in the cells excluding MMP ($\Delta \Psi_m$) loss cells (Fig. 2D). Caspase-3 plays an essential role as an executioner in apoptosis

(26). As shown in Fig. 2E, MG132 increased caspase-3 activity in HPF cells. We also observed the disappearance of procaspase-3 protein in MG132-treated HPF cells (data not shown). Furthermore, the activity of caspase-8, which is known to be involved in receptor- or extrinsic-mediated apoptosis (27), was increased in MG132-treated HPF cells (Fig. 2E).

Effects of caspase inhibitors on cell growth, apoptosis and MMP ($\Delta \Psi_m$) in MG132-treated HPF cells. Next, we investigated the effects of caspase inhibitors on the growth and death of MG132-treated HPF cells. For this experiment, 10 μ M

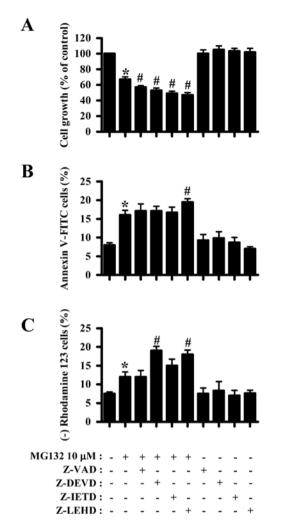


Figure 3. Effects of caspase inhibitors on cell growth, cell death and MMP $(\Delta \Psi_m)$ in MG132-treated HPF cells. Exponentially growing cells were treated with 10 μ M MG132 for 24 h following a 1-h pre-incubation with 15 μ M of a given caspase inhibitor. (A) The graph shows cell growth changes in HPF cells as assessed by MTT assays. (B and C) Annexin V-FITC cells and MMP $(\Delta \Psi_m)$ loss were measured with a FACStar flow cytometer. Graphs show the percent of Annexin V-positive staining cells (B) and rhodamine 123-negative [MMP ($\Delta \Psi_m$) loss] cells (C). *p<0.05 compared with the control group. #p<0.05 compared with cells treated with MG132 only.

MG132 as a suitable dose was used to differentiate the levels of cell growth inhibition and death in the presence or absence of each caspase inhibitor [pan-caspase inhibitor (Z-VAD)], caspase-3 inhibitor (Z-DEVD), caspase-8 inhibitor (Z-IETD), or caspase-9 inhibitor (Z-LEHD). A concentration of 15 μ M was used as an optimal dose of each caspase inhibitor. All of the caspase inhibitors enhanced growth inhibition in MG132-treated HPF cells (Fig. 3A). In relation to cell death, caspase-9 inhibitor among other inhibitors enhanced increased Annexin V-positive cell numbers in MG132-treated HPF cells (Fig. 3B). Moreover, caspase-3, -8 and -9 inhibitors intensified MMP ($\Delta \Psi_m$) loss in these cells (Fig. 3C).

Effects of MG132 and/or caspase inhibitors on ROS and GSH levels in HPF cells. ROS and GSH levels in HPF cells treated with MG132 and/or each caspase inhibitor were assessed at 24 h. As shown in Fig. 4A, the intracellular ROS level (as determined by DCF) was significantly increased in

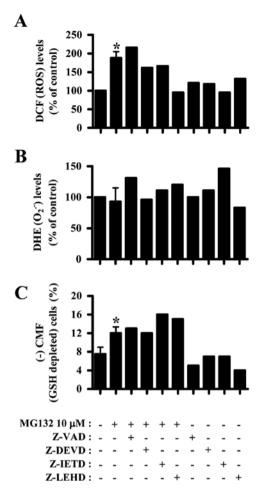


Figure 4. Effects of caspase inhibitors on ROS and GSH levels in MG132treated HPF cells. Exponentially growing cells were treated with 10 μ M MG132 for 24 h following a 1-h pre-incubation with 15 μ M of a given caspase inhibitor. ROS and GSH levels were measured with a FACStar flow cytometer. (A and B) Graphs indicate ROS (as determined by DCF) levels (%) (A) and DHE (O₂⁻) levels (%) compared with control cells (B). (C) Graph shows the percent of (-) CMF (GSH-depleted) cells. *p<0.05 compared with the control group.

MG132-treated HPF cells. While Z-VAD slightly enhanced ROS levels in MG132-treated cells, other caspase inhibitors, especially caspase-9 inhibitor decreased the level in these cells (Fig. 4A). Red fluorescence derived from DHE, which reflected the O_2^{\bullet} accumulation, was not significantly changed in MG132-treated HPF cells (Fig. 4B). Z-VAD and caspase-9 inhibitor seemed to increase O_2^{\bullet} level in MG132-treated HPF cells. Caspase-8 inhibitor alone increased O_2^{\bullet} level in control cells (Fig. 4B). In relation to GSH levels in HPF cells, MG132 increased GSH-depleted cell numbers in HPF cells compared with those of the control cells and caspase-8 and -9 inhibitors relatively increased GSH-depleted cell numbers in MG132-treated HPF cells (Fig. 4C).

Effects of apoptosis-related siRNAs on apoptosis and ROS levels in MG132-treated HPF cells. Furthermore, it was determined whether apoptosis-related siRNAs changed cell death and ROS levels in MG132-treated HPF cells. As shown in Fig. 5A, 30 μ M MG132 increased the proportion of Annexin V-stained cells about 15% compared with that in control siRNA treated HPF cells without MG132. Treatment with 10 μ M MG132 did

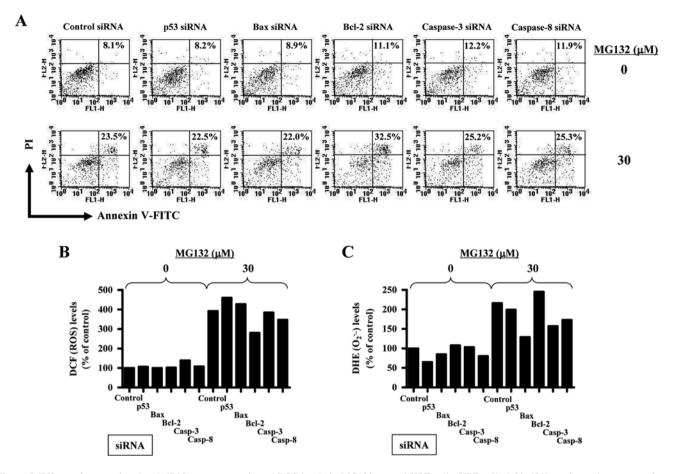


Figure 5. Effects of apoptosis-related siRNAs on apoptosis and ROS levels in MG132-treated HPF cells. HPF cells (~30-40% confluence) were transfected with either non-target control siRNA or each apoptosis-related siRNA. One day later, cells were treated with 30 μ M MG132 for additional 24 h. (A) Annexin V-FITC and PI cells were measured with a FACStar flow cytometer. The number (%) in each figure indicates Annexin V-FITC positive cells regardless of PI negative and positive cells. (B and C) Graphs indicate DCF (ROS) levels (%) (B) and DHE (O₂*) levels (%) compared with MG132-untreated control siRNA cells (C).

not clearly increase Annexin V-stained cell number in this system (data not shown). Probably, Lipofectamine 2000 in the medium attenuated the biological activity of MG132. While p53 and Bax siRNAs did not affect Annexin V-stained cell number in MG132-treated HPF cells, Bcl-2 siRNA definitely increased the number in these cells (Fig. 5A). Caspase-3 and -8 siRNAs also slightly increased Annexin V-stained cell number in MG132-treated HPF cells (Fig. 5A). In relation to ROS levels, p53 and Bax siRNAs slightly increased ROS (DCF) level in MG132-treated cells whereas Bcl-2 and caspase-8 siRNAs relatively decreased the level in these cells (Fig. 5B). Caspase-3 siRNA alone slightly increased ROS level in HPF control cells (Fig. 5B). In view of O₂⁻ level, p53, Bax and caspase-8 siRNAs decreased O_2^{\bullet} level in HPF control cells (Fig. 5C). In this experimental condition, treatment with 30 μ M MG132 clearly increased O₂[•] level in HPF cells (Fig. 5C). Bax, caspase-3 and -8 siRNAs attenuated the increased O₂[•] level in MG132-treated HPF cells whereas Bcl-2 siRNA relatively increased the level (Fig. 5C).

Discussion

In the present study, we focused on investigating the molecular mechanisms of MG132 in HPF cells in relation to cell growth

inhibition and death. MG132 inhibited the growth of HPF cells in a dose-dependent manner. In addition, MG132 dosedependently induced cell death in HPF cells. Furthermore, MG132 blocked the activity of proteasome in HPF cells. Consequently, this inhibition seemed to influence the growth inhibition and death of HPF cells. Based on an MTT assay, an IC_{50} of MG132 in HPF cells was approximately 20 μ M. Our previous reports demonstrated that the doses of IC₅₀ of MG132 in A549 and Calu-6 lung cancer cells were approximately 20 and 5 μ M, respectively (18,19). Therefore, these results did not strongly support the assumption that the susceptibility of HPF cells to MG132 is lower than that of lung cancer cells. Thus, the susceptibility of MG132 seems to depend on the cell type. Our DNA flow cytometric analysis demonstrated that 10 μ M MG132 induced S and G2-M phase arrest of the cell cycle in HPF cells. Since 10 μ M MG132 inhibited the growth of HPF cells approximately 40% but this dose induced cell death at most 6-8% compared with that of control cells. This result implies that S and G2-M phase arrest by MG132 is a pathway to suppress the growth of HPF cells. According to previous reports, 0.5 µM MG132 induced a G2/M phase arrest in Calu-6 cells and 1-30 µM MG132 induced an S phase arrest in these cells (19). MG132 induced a G1 phase arrest of the cell cycle in A549 cells (18). Thus, the deregulation of the ubiquitin-proteasomal system by MG132 can differently affect cell cycle distribution depending on incubation dose and cell type in lung cells.

To gain insight into the molecular mechanism involved in MG132-mediated HPF cell death, sub-G1 cells, Annexin V staining and MMP ($\Delta \Psi_m$) were assessed in HPF cells. MG132 clearly induced HPF cell death which was accompanied by MMP ($\Delta \Psi_m$) loss. In addition, MG132 increased the activities of caspase-3 and -8 in HPF cells. Thus, MG132 seemed to induce apoptosis in HPF cells. However, none of the tested caspase inhibitors prevented HPF cell death by MG132. Instead, caspase-9 inhibitor (Z-LEHD) increased Annexin V-positive cell numbers in MG132-treated HPF cells. In addition, all of the tested caspase inhibitors enhanced cell growth inhibition by MG132, and caspase-3 and -9 inhibitors significantly increased MMP ($\Delta \Psi_m$) loss. Furthermore, caspase-3 and -8 siRNAs slightly increased Annexin V-positive cell number in MG132treated HPF cells. These data imply that MG132-induced HPF cell death occurred in a caspase-independent manner. The slight enhancement of cell death by caspase inhibitors or siRNAs in MG132-treated HPF cells probably resulted from increased necrotic cell death following the blockade of the apoptotic pathway by these agents. In relation to Calu-6 cell death by MG132 (19), Z-VAD (pan-caspase inhibitor) significantly rescues Calu-6 cells from MG132 insults. Therefore, the mechanism of cell death by MG132 can be different in lung cancer and normal lung cells. In addition, when p53, Bax and Bcl-2 siRNAs were administered in HPF cells treated with MG132, p53 and Bax siRNAs did not affect Annexin V-stained cell number in these cells but Bcl-2 siRNA clearly increased the number. These data imply that Bcl-2 protein, but not p53 and Bax proteins, was tightly involved in MG132-induced HPF cell death.

Proteasome inhibitors including MG132 can induce apoptotic cell death through formation of ROS (8-10). According to our results, ROS level (as determined by DCF) was significantly increased in HPF cells treated with MG132. Treatment with 10 μ M MG132 did not significantly affect O₂[•] level but 30 µM MG132 clearly increased O2 · level in control siRNAadministered cells. However, each caspase inhibitor affects differently ROS levels including O2+ regardless of cell growth inhibition and cell death levels. Especially, caspase-9 inhibitor showing a strong decreasing effect on ROS (DCF) level in MG132-treated cells enhanced cell death and MMP ($\Delta \Psi_m$) loss in these cells. In addition, Z-VAD increased O₂[•] level in MG132-treated HPF cells regardless of cell death. Caspase-8 inhibitor alone increased O2 - level in HPF control cells without the induction of cell death. In relation to siRNA-administered HPF cells, Bcl-2 siRNA showing an enhancement in MG132induced HPF cell death decreased the ROS (DCF) level. Bax, caspase-3 and -8 siRNAs clearly reduced O2 - level in MG132treated cells regardless of cell death. Therefore, the ROS level changes in HPF cells treated with MG132 and/or each caspase inhibitor or apoptosis-related siRNA need to be further clarified in view of cell death. The redox status of cellular GSH is a crucial regulatory element in protein ubiquitination system (28). GSH depletion due to proteasome inhibitors can lead to cell death (10,11). Likewise, MG132 treatment increased the number of GSH-depleted cells in the HPF cells. In addition, caspase-8 and -9 inhibitors increased the number of GSH-depleted cells. This result seemed to correlate with the observation that these inhibitors intensified cell growth inhibition or cell death in MG132-treated HPF cells.

In conclusion, MG132 induced growth inhibition and death in HPF cells in a caspase-independent manner. The growth inhibition and death of HPF cells by MG132 and/or each caspase inhibitor or apoptosis-related siRNA seemed to be related to the changes in GSH levels rather than ROS levels. Our present data provide useful information to understand the toxicological and molecular effect of MG132 on normal lung cells.

Acknowledgements

This study was supported by Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education, Science and Technology (2010-0007059).

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