Combination of bortezomib and daunorubicin in the induction of apoptosis in T-cell acute lymphoblastic leukemia

XIN DU1*, JIA TONG1*, HONGYING LU1*, CONG HE1, SHENGHONG DU1, PEIMIN JIA1, WEILI ZHAO1, HANZHANG XU2, JUNMIN LI1, ZHIXIANG SHEN1, YINGLI WU2, JIANHUA TONG1 and LI ZHOU1

1State Key Laboratory of Medical Genomics, Department of Hematology, Faculty of Medical Laboratory Science, Ruijin Hospital, Shanghai Jiaotong University School of Medicine; 2Hongqiao International Institute of Medicine, Shanghai Tongren Hospital/Faculty of Basic Medicine, Department of Pathophysiology, Chemical Biology Division of Shanghai Universities E-Institutes, Key Laboratory of Cell Differentiation and Apoptosis of The Chinese Ministry of Education, Shanghai Jiaotong University School of Medicine, Shanghai 200025, P.R. China

Received March 19, 2016; Accepted February 27, 2017

DOI: 10.3892/mmr.2017.6554

Abstract. Despite advances in the treatment of T-cell acute lymphoblastic leukemia (T-ALL), the outcome of T-ALL treatment remains unsatisfactory, therefore, more effective treatment is urgently required. The present study examined the cytotoxicities of bortezomib in combination with daunorubicin against human Jurkat and Molt-4 T-ALL cells and primary T-ALL cells. Compared with treatment alone, co-exposure of cells to bortezomib and daunorubicin resulted in a significant increase in cell death in the Jurkat cells, as evidenced by the increased percentage of Annexin V-positive cells, the formation of apoptotic bodies. In addition, the administration sequence of bortezomib and daunorubicin had an effect on cell viability. Treatment with bortezomib followed by daunorubicin treatment was more effective, compared with treatment with daunorubicin followed by bortezomib. Co-treatment with bortezomib and daunorubicin markedly enhanced the activation of caspase-3, -8 and -9, which was reversed by the pan-caspase inhibitor, Z-VAD-FMK. In addition, cotreatment with bortezomib and daunorubicin enhanced the collapse of mitochondrial transmembrane potential and upregulated the proapoptotic protein, B-cell lymphoma 2 (Bcl-2)-interacting mediator of cell death (Bim). Taken together, these findings indicated that the combination of bortezomib and daunorubicin significantly enhanced their apoptosis-inducing effect in T-ALL cells, which may warrant further investigation in preclinical and clinical investigations.

Introduction

T-cell acute lymphoblastic leukemia (T-ALL) is an aggressive hematologic disease (1), accounting for ~15 and 25% of ALL in pediatric and adult patients, respectively (2). Despite progress in treatment and children having an approximate cure rate of 90%, the cure rate of adults is <50%, even with hematopoietic stem cell transplantation (3). In adult patients with ALL who have achieved complete remission, the majority relapse, and few patients are cured even when treated using the best therapeutic regimens currently available. Therefore, more effective treatment is urgently required for this disease.

The 26S proteasome is vital in eukaryotic cell function and viability. It is responsible for several integral cellular processes, including the timely degradation of cell cycle regulator proteins and transcription factors, and the maintenance of cellular homeostasis, all of which are essential for cell proliferation, differentiation apoptosis and angiogenesis (4). The proteasome has consequently become an attractive target for therapeutic intervention in cancer chemotherapy. Bortezomib (Velcade; PS-341), a dipeptide boronic acid analog and the first clinically available proteasome inhibitor, has been widely applied to treat multiple myeloma (5). The drug exhibits selectivity towards the proteasome of cancer cells, rendering it with a distinct therapeutic advantage. Preclinical studies have shown that bortezomib also induces apoptosis in acute leukemia and solid tumor cells (6). However, as a single agent, the clinical results of bortezomib in patients with acute myeloid leukemia and pediatric refractory ALL are less encouraging (7). Accumulating clinical studies have indicated that the effect of bortezomib combined with anthracycline drugs, including idarubicin and doxorubicin, is likely to be
active with acceptable toxicity in hematological malignancies, including acute leukemia (8-10). Daunorubicin belongs to the anthracycline family, and is one of the major antitumor agents in the treatment of myeloid leukemia (11). Daunorubicin kills leukemic cells by the inhibition of topoisomerase II, induction of DNA damage and generation of reactive oxygen species (ROS) (12). However, whether the combination of daunorubicin and bortezomib is also effective in T-ALL cells remains to be elucidated.

In the present study, the cytotoxicities against T-ALL cells of bortezomib and daunorubicin, alone and in combination were compared. It was found that cotreatment of daunorubicin and bortezomib was more effective, compared with either agent used alone at inducing T-ALL cell death, and it was demonstrated that the mitochondrial pathway was important in this process.

Materials and methods

Cells and drug treatments. The T-ALL Jurkat and Molt-4 cells and Burkitt lymphoma Daudi cells (both from American Type Culture Collection; ATCC, Manassas, VA, USA) were maintained at 37°C in RPMI-1640 medium (Gibco; Thermo Fisher Scientific, Waltham, MA, USA) supplemented with 10% heat-inactivated fetal bovine serum (FBS; HyClone, Logan, UT, USA) in a humidified atmosphere containing 95% air and 5% CO₂. Primary ALL cells were harvested from bone marrow samples of 5 patients with T-ALL in Ruijin Hospital (Shanghai, China), from October 2013 to July 2014 (patient information is shown in Table I). Bone marrow mononuclear cells were isolated using Ficoll-Paque isolation solution and centrifugation for 30 min at 400 x g at 25°C and re-suspended in RPMI-1640 medium supplemented with 10% FBS. The study protocol was approved by the local institution review board at Shanghai Jiaotong University School of Medicine (Shanghai, China) and informed consent was obtained from the patients. Cells were seeded at 5x10⁵ cells/ml in 6-well plates. Jurkat cells were treated with 10 nM bortezomib (Velcade; Millennium Pharmaceutical, Cambridge, MA, USA) or 1 µM daunorubicin hydrochloride (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany), and Molt-4 cells were treated with 15 nM bortezomib or 15 nM daunorubicin hydrochloride, alone or in combination, for 24 h or 48 h in a humidified atmosphere containing 95% air and 5% CO₂.

Cell proliferation assays. The cells were seeded at 2-3x10⁵ cells/ml in a 6-well plate and were treated with the appropriate drugs. The cells were counted every 12 h using a Beckman Coulter counter (Beckman Coulter, Inc., Fullerton, CA, USA) and cellular morphology was evaluated using light microscopy with Wright's staining. The experiment was performed in duplicate at least three times independently.

Apoptotic assays. For the apoptotic assays, 2x10⁶ cells were washed twice with phosphate-buffered saline (PBS), and then labeled using Annexin V-fluorescein isothiocyanate and propidium iodide (PI) according to the manufacturer's protocol (BD Pharmingen, San Diego, CA, USA). Fluorescence intensity was measured using flow cytometry (BD Biosciences, Franklin Lakes, NJ, USA). Data were analyzed using CellQuest software (version 3.3; BD Biosciences). For each analysis, 10,000 events were recorded. The experiment was performed in duplicate at least three times independently.

Determination of mitochondrial membrane potential (ΔΨm). For measurement of alterations in ΔΨm, the cells (1x10⁶) were incubated with 100 µg/ml Rhodamine123 for 30 min at room temperature with 50 µg/ml PI. Fluorescence intensity was measured using flow cytometry. The experiment was performed at least three times independently in triplicate.

Western blot analysis. The cells were washed with PBS and lysed with lysis buffer containing 62.5 mM Tris-HCl (pH 6.8), 100 mM DTT, 2% SDS and 10% glycerol by boiling at 100°C for 5 min. The cell lysates were centrifuged at 20,000 x g for 10 min at 4°C, and proteins in the supernatants were quantified using a bicinchoninic acid protein assay (Merck KGaA). Protein extracts (10-20 µg) were equally loaded to an 8-14% SDS-polyacrylamide gel, electrophoresed, and transferred onto nitrocellulose membrane (GE Healthcare Life Sciences, Buckinghamshire, UK). The blots were stained with 0.2% Ponceau S red to ensure equal protein loading. Following blocking with 5% nonfat milk in PBS, the membranes were probed with antibodies against caspase-3 (cat. no. 9662), caspase-8 (cat. no. 9746), caspase-9 (cat. no. 9502), Bim (cat. no. 2933), Bcl-2 (cat. no. 4223) and Bcl-xL (cat. no. 2764), overnight at 4°C (all at 1:1,000 dilution; Cell Signaling Technology, Inc., Danvers, MA, USA), then followed by incubation with horseradish peroxidase (HRP)-linked secondary antibodies (1:2,000 dilution; Cell Signaling Technology, Inc.) for 1 h at room temperature. The signals were detected using a chemiluminescence substrate (HRP) kit (Cell Signaling Technology, Inc.) according to the manufacturer's protocol. As necessary, the blots were stripped and re-probed with anti-β-actin antibody (1:5,000 dilution; cat. no. PM053; Medical and Biological Laboratories Co., Ltd., Nagoya, Japan) as an internal control. All experiments were repeated three times.

Statistical analysis. Statistical analysis was performed using SPSS software (version 17.0; SPSS, Inc., Chicago, IL, USA). Student's t-test was used to compare differences between two groups. P<0.05 was considered to indicate a statistically significant difference. Synergistic interactions between bortezomib and daunorubicin were analyzed by the Chou-Talalay method using CalcuSyn software version 2.1 (Biosoft, Cambridge, UK) (13). A combination index (CI) of <1, equal to 1, and >1 indicate synergistic, additive and antagonistic effects, respectively.

Results

Bortezomib and daunorubicin cotreatment promotes late apoptosis of Jurkat and Molt-4 cells. The present study first examined the possible combinatorial effect of bortezomib and daunorubicin against Jurkat and Molt-4 cells. The Jurkat cells were treated with 1 µM daunorubicin in the presence or absence of 10 nM bortezomib. Bortezomib and daunorubicin alone caused a 24.9±0.74 and a 72.6±2.17% reduction in the proliferation of Jurkat cells 48 h post treatment, respectively. The
combined use of the two agents resulted in a marked decrease in the proliferation of the Jurkat cells (90.1±3.23%; Fig. 1A), with similar results observed in the Molt-4 cells (Fig. 1B).

Wright’s staining of the Jurkat cells treated with bortezomib (10 nM) and daunorubicin (1 µM) alone or in combination for 24 h showed that an induction of apoptosis was observed in the combination group, with the appearance of cell shrinkage, nuclear condensation and the formation of apoptotic bodies (Fig. 1C and D). Flow cytometric analysis revealed that, following treatment for 24 h, the percentage of early apoptotic Jurkat cells treated with bortezomib was 24.2±6.35% (P<0.05, vs. control) whereas the percentage of late apoptotic Jurkat cells was 2.8±5.65% (P>0.05, vs. control; Fig. 1E). The percentage of early apoptotic Jurkat

Table I. Information of patients with T-cell acute lymphoblastic leukemia.

| Number | Sex  | Age (years) | Blast (%) | Chromosome                  | White blood cells (x10⁹/l) |
|--------|------|-------------|-----------|-----------------------------|---------------------------|
| 1      | Male | 16          | 95.5      | ND                          | 15.4                      |
| 2      | Female | 37         | 85.0      | ND                          | 2.2                       |
| 3      | Female | 24         | 91.5      | 51-53,XX,3p-, del(6),       | 22.1                      |
| 4      | Female | 14         | 95.0      | 46,XX                       | 15.6                      |
| 5      | Female | 37         | 90.0      | 46,XX,3p+,del(9),           | 32.3                      |

Figure 1. Effects of Bor+DNR on the proliferation of Jurkat and Molt-4 cells. (A) Jurkat cells were treated with 10 nM Bor with or without 1 µM DNR, and (B) Molt-4 cells were treated with 15 nM Bor, with or without 15 nM DNR for 12, 24, 36 and 48 h, respectively, and cell viability was determined. The morphology of the (C) Jurkat and (D) Molt-4 cells treated with Bor, with or without DNR, was examined using Wright’s staining (magnification, x100). Apoptotic cells were determined for (E) Jurkat and (F) Molt-4 cells treated with Bor, with or without DNR, for 24 h, by Annexin V/propidium iodide staining. All values are expressed as the mean ± standard deviation. Each experiment was performed at least 3 times independently. *P<0.05, compared with Bor or DNR treatment. Bor, bortezomib; DNR, daunorubicin.
cells treated with daunorubicin was 12.4±2.75% (P<0.05, vs. control) whereas the percentage of late apoptotic Jurkat cells was 5.8±2.55% (P>0.05, vs. control; Fig. 1E). The combination of daunorubicin and bortezomib caused no apparent increase in the percentage of early apoptotic cells, compared with the cells treated with daunorubicin alone (27.9±4.81%; P>0.05), however, a significant increase in the percentage of late apoptotic cells was observed (23.0±4.51%; P<0.05, vs. daunorubicin or bortezomib alone; Fig. 1E). Similar results were observed in the Molt-4 cells (Fig. 1F). These data indicated that the combination of daunorubicin and bortezomib enhanced the late apoptosis of Jurkat and Molt-4 cells.

To determine whether these two drugs synergistically inhibited the proliferation of leukemia cells, the CI values were calculated using the Chou-Talalay method. As shown in Tables II-IV, daunorubicin and bortezomib cotreatment had a synergistic effect on the Jurkat and Molt-4, but not on the Daudi cells.

To determine whether the administration sequence affected the combination effect of daunorubicin and bortezomib, daunorubicin or bortezomib was added prior to, or following, the other drug. As shown in Fig. 2A and B, bortezomib treatment followed by daunorubicin was more effective, compared with daunorubicin treatment followed by bortezomib, or with the two drugs simultaneously.

Bortezomib promotes cell death by activating the caspase cascade. To investigate the mechanisms underlying bortezomib+daunorubicin-induced cell death in the Jurkat and Molt-4 cells, the present study examined the activation of caspase using western blot analysis. It was found that cotreatment of the Jurkat (Fig. 3A) and Molt-4 (Fig. 3B) cells with daunorubicin and bortezomib significantly increased the levels of cleaved caspase-3, -8 and -9. To determine the role of caspase activation, the Jurkat (Fig. 3C) and Molt-4 (Fig. 3D) cells were

| Table II. Combination index values of bortezomib and daunorubicin in Jurkat cells. |
|---------------------------------|---------------------------------|-----------------|-----------------|-----------------|
| Dose | Dose | Growth | Combination |
| bortezomib (nM) | daunorubicin (µM) | inhibition (%) | index |
| 10.0 | 1.00 | 52 | 0.31398 |
| 10.0 | 1.25 | 58 | 0.33839 |
| 10.0 | 1.50 | 64 | 0.34315 |
| 15.0 | 10.00 | 70 | 0.29281 |
| 15.0 | 10.25 | 84 | 0.12266 |
| 15.0 | 10.50 | 78 | 0.40632 |
| 20.0 | 1.00 | 40 | 0.66699 |
| 20.0 | 1.25 | 62 | 0.47163 |
| 20.0 | 1.50 | 70 | 0.38763 |

| Table III. Combination index values of bortezomib and daunorubicin in Molt-4 cells. |
|---------------------------------|---------------------------------|-----------------|-----------------|-----------------|
| Dose | Dose | Growth | Combination |
| bortezomib (nM) | daunorubicin (nM) | inhibition (%) | index |
| 12.5 | 10.0 | 62 | 0.79070 |
| 12.5 | 12.5 | 72 | 0.62921 |
| 12.5 | 15.0 | 73 | 0.77304 |
| 15.0 | 10.0 | 70 | 0.68773 |
| 15.0 | 12.5 | 73 | 0.69217 |
| 15.0 | 15.0 | 80 | 0.62526 |
| 17.5 | 10.0 | 72 | 0.72619 |
| 17.5 | 12.5 | 76 | 0.68364 |
| 17.5 | 15.0 | 80 | 0.68750 |

| Table IV. Combination index values of bortezomib and daunorubicin in Daudi cells. |
|---------------------------------|---------------------------------|-----------------|-----------------|-----------------|
| Dose | Dose | Growth | Combination |
| bortezomib (nM) | daunorubicin (nM) | inhibition (%) | index |
| 10.0 | 40.0 | 51 | 1.13996 |
| 10.0 | 50.0 | 64 | 0.90496 |
| 10.0 | 60.0 | 67 | 1.03848 |
| 15.0 | 40.0 | 53 | 1.12142 |
| 15.0 | 50.0 | 68 | 0.82208 |
| 15.0 | 60.0 | 63 | 1.40427 |
| 20.0 | 40.0 | 61 | 0.93591 |
| 20.0 | 50.0 | 70 | 0.83512 |
| 20.0 | 60.0 | 71 | 0.99927 |
treated with bortezomib and daunorubicin in the presence or absence of the broad-spectrum caspase inhibitor, Z-VAD-FMK (10 µM). The results of the flow cytometric analysis revealed that Z-VAD-FMK significantly attenuated the cell death induced by cotreatment with bortezomib and daunorubicin (P<0.05). These findings suggested that bortezomib and daunorubicin promoted apoptosis through activating caspase-3, -8 and -9.

Bortezomib+daunorubicin causes marked dissipation of ΔΨm and increases the expression of Bim. The mitochondrial pathway is critical in the execution of apoptotic cell death, and the collapse of ΔΨm is an early event in the mitochondrial cell death pathway. The present study examined alterations in ΔΨm using rhodamine123/PI staining. Marked dissipation of ΔΨm was observed in the Jurkat cells cotreated with bortezomib and daunorubicin at 24 h (35.4±9.3%), which was significantly higher, compared with that in the Jurkat cells treated with bortezomib (11.1±4.6%) or daunorubicin (9.3±2.0%) alone (Fig. 4A). Similar results were observed in Molt-4 cells (Fig. 4B). Alterations in the expression of the Bcl-2 family proteins, Bcl-2, Bcl-xl and Bim, were also examined. In the Jurkat cells, daunorubicin alone had no apparent effect, whereas bortezomib upregulated the expression of Bim. However, the combination of bortezomib and
daunorubicin caused a marked increase in the expression of Bim at 24 h (Fig. 4C). By contrast, the combination regimen exerted no noticeable effect on the expression of Bcl-2 or Bcl-xl (Fig. 4C). Similar results were observed in the Molt-4 cells (Fig. 4D). These data indicated that the combination effect of bortezomib+daunorubicin was associated with the collapse of ΔΨm, in which Bim may be involved.

**Bortezomib and daunorubicin in combination are effective in inducing the apoptosis of primary T-ALL cells.** To determine whether the combination effect of bortezomib and daunorubicin was limited to Jurkat cells, five primary T-ALL cells were treated with daunorubicin (500 nM) in the presence or absence of bortezomib (10 nM) for 24 h. As shown in Fig. 5A, bortezomib+daunorubicin caused a significantly higher apoptotic rate, compared with either agent used alone (P<0.05). In addition, flow cytometric analyses showed that bortezomib or daunorubicin alone caused only modest mitochondrial injury, whereas the combination of bortezomib and daunorubicin resulted in more prominent mitochondrial injury (Fig. 5B). Furthermore, the western blot assays showed that the combination treatment markedly increased...
the expression of Bim in primary ALL cells, compared with either bortezomib or daunorubicin alone (Fig. 5C). Together, these findings indicated that the combination of bortezomib and daunorubicin exerted more marked cytotoxicity against primary ALL cells by disrupting the ΔΨm.

Discussion

T-ALL has a poor prognosis due to its intrinsic chemoresistance and severe immunosuppression. Although traditional chemotherapeutic regimens have shown improved response rate, they have failed to achieve a significant effect on long-term survival rates (2). Therefore, novel strategies are required for the treatment of T-ALL. In the present study, it was demonstrated that bortezomib and daunorubicin cotreatment induced apoptosis in T-ALL cells via disrupting the ΔΨm.

Human leukemic cells express abnormally high levels of proteasomes, compared with normal peripheral blood cells, and they are significantly more sensitive to proteasome inhibition, compared with normal bone marrow progenitor cells or peripheral blood lymphocytes (6,14,15). Clinical trials are underway to assess the efficacy of bortezomib in several human malignancies, including leukemia (10). Available data suggest that bortezomib as a single drug may only yield minor clinical benefits in patients with leukemia (7-16). However, bortezomib has been shown to enhance the efficacy of several conventional therapies and may overcome resistance to conventional anticancer drugs, including belinostat, SAHA, gemcitabine and imatinib, and the effects in combination studies appear promising (10,17-24). In the present study, using human T-ALL Jurkat and Molt-4 cells as a model, it was demonstrated that bortezomib and daunorubicin cotreatment was more effective, compared with either agent used alone at inducing apoptosis, as reflected by the activation of caspase-3, -8 and -9, and the appearance of apoptotic morphology. Notably, it was demonstrated that this combination effect was also present in primary T-ALL cells. It is well known that standard chemotherapy is inhibitory to normal hematopoietic cells and frequently results in severe myelosuppression, and anthracyclines can cause cumulative dose-dependent cardiotoxicity (25). Thus, the combination of bortezomib and daunorubicin at a lower dose may reduce their dose-associated side effect but increasing their efficacy. Similar to the findings obtained in the present study, Koyama et al reported that bortezomib and doxorubicin also induced apoptosis in T-ALL cell lines (26). However, the combination effect of these drugs on primary leukemia cells was not investigated.

The mitochondrial and cell death receptor apoptotic pathways are two major apoptotic cell death pathways. It has been shown that mitochondrial signaling exerts a critical role in bortezomib-induced apoptosis (27-30). The present study found that the combination of these two agents caused extensive loss of ΔΨm, indicating the involvement of the mitochondrial apoptotic pathway. Consistent with this, bortezomib and daunorubicin cotreatment enhanced the collapse of ΔΨm in primary T-ALL leukemia cells. The cell death receptor pathway may also be activated by cotreatment of bortezomib and daunorubicin, as evidenced by the activation of caspase-8.

An important event in the mitochondrial apoptotic pathway is mitochondrial outer membrane permeabilization, which is primarily mediated and controlled by the Bcl-2 family members (31). When mitochondrial outer membrane permeabilization occurs, it precipitates cell death through either the release of molecules involved in apoptosis or the loss of mitochondrial functions essential for cell survival. The present study determined the effect of bortezomib daunorubicin cotreatment on several Bcl-2 family members. The bortezomib daunorubicin cotreatment markedly increased the proapoptotic regulator protein, Bim, in the Jurkat and primary ALL cells, but exerted minimal effect on the expression of Bcl-2 or Bcl-xL. Bim is a member of the BH3-only protein family, which mediates cell death from physiologic stimuli, including cytochrome c deprivation and signals from activated oncogenes. The upregulation of Bim triggers the release of cytochrome c from the mitochondria and the onset of apoptosis (32). The results of the present study indicated that Bim may be important in bortezomib+daunorubicin-induced cell death. Consistent with this, several reports have shown that Bim-targeting contributes to the bortezomib-based combination regime (33-35). However, whether Bim contributed to bortezomib+daunorubicin-induced mitochondria impairment, and how cotreatment with bortezomib and daunorubicin upregulated the expression of Bim required further investigation. BH3-interacting domain death agonist (Bid), another proapoptotic Bcl-2 family member, may also be involved in this process (36,37). As shown in Fig. 2, bortezomib and daunorubicin cotreatment induced the activation of caspase 8. Caspase 8 can cleave Bid into t-Bid, which then causes mitochondrial outer membrane permeabilisation. This leads to the mitochondrial release of apoptogenic proteins, including cytochrome c.

In conclusion, the present demonstrated that bortezomib cooperated with daunorubicin to induce the apoptosis of Jurkat and Molt-4 cells, and primary T-ALL cells, in which the mitochondrial apoptotic pathway was pivotal. These findings provide a rationale for use of the combination of bortezomib and daunorubicin in the treatment T-ALL in future preclinical and clinical investigations.

Acknowledgements

This study was supported by the Shanghai Commission of Science and Technology (grant nos. 10411966900 and 15401901800), the National Natural Science Foundation of China (grant nos. 81100980, 81570118 and 81570112) and the Innovation Program of Shanghai Municipal Education Commission (grant no. 13YZ028).

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