

Targeting Bcl-2 family members with the BH3 mimetic AT-101 markedly enhances the therapeutic effects of chemotherapeutic agents in in vitro and in vivo models of B-cell lymphoma

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Overexpression of antiapoptotic members of the Bcl-2 family are observed in approximately 80% of B-cell lymphomas, contributing to intrinsic and acquired drug resistance. Nullifying antiapoptotic function can potentially overcome this intrinsic and acquired drug resistance. AT-101 is a BH3 mimetic known to be a potent inhibitor of antiapoptotic Bcl-2 family members including Bcl-2, Bcl-X_L, and Mcl-1. In vitro, AT-101 exhibits concentration- and time-dependent cytotoxicity against lymphoma and multiple

myeloma cell lines, enhancing the activity of cytotoxic agents. The IC₅₀ for AT-101 is between 1 and 10 μM for a diverse panel of B-cell lymphomas. AT-101 was synergistic with carfilzomib (C), etoposide (E), doxorubicin (D), and 4-hydroxycyclophosphamide (4-HC) in mantle cell lymphoma (MCL) lines. In a transformed large B-cell lymphoma line (RL), AT-101 was synergistic when sequentially combined with 4-HC, but not when both drugs were added simultaneously. AT-101 also induced potent mitochondrial membrane depolariza-

tion (ΔΨ_m) and apoptosis when combined with carfilzomib, but not with bortezomib in MCL. In severe combined immunodeficient (SCID) beige mouse models of drug-resistant B-cell lymphoma, 35 mg/kg per day of AT-101 was safe and efficacious. The addition of AT-101 to cyclophosphamide (Cy) and rituximab (R) in a schedule-dependent manner enhanced the efficacy of the conventional therapy. (Blood. 2008;111:5350-5358)

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Introduction

Bcl-2 and related proteins are key regulators of apoptosis that are expressed in solid tumors and hematologic malignancies.¹ Bcl-2 is known to be constitutively overexpressed in approximately 80% of follicular lymphomas and 20% of diffuse B-cell lymphomas as a result of the t(14;18) translocation and gene amplification, respectively.^{2,3} Overexpression of antiapoptotic family members is associated with inhibition of apoptosis and chemotherapy resistance, resulting in lower clinical response rates and shortened survivals.⁴⁻⁹ Targeting Bcl-2 family members offers new opportunities to address these survival pathways directly. One important benefit of these drugs relates to their ability to lower the threshold required to induce apoptosis, making them potentially complimentary with conventional chemotherapy approaches.

AT-101, a derivative of the natural product gossypol, is a BH3 mimetic capable of binding to Bcl-2, Bcl-X_L, and Mcl-1, attenuating their antiapoptotic influence.¹⁰ Gossypol is a natural compound extracted from cottonseeds (*Gossypium sp*), which was initially used as an antifertility agent¹¹ and subsequently as a cytotoxic agent.¹²⁻²⁰ In the late 1980s, it was discovered that (–)-gossypol enantiomer exhibited the most potent anticancer activity compared with the racemic mixture or the (+)-gossypol enantiomer. The 3D solution structure of small molecules including (–)-gossypol in complex with Bcl-X_L revealed several crucial interactions accounting for the binding specificity of the negative enantiomer. Given the

importance of Bcl-2 overexpression in many types of lymphomas, we investigated the in vitro and in vivo antitumor activity of AT-101 alone and in combination with different cytotoxic and biologic agents. The major objective of these studies was to define the best strategy for combining AT-101 with other agents based upon an understanding of its biologic effects on the cell.

Methods

Cells and cell lines

RL is an Epstein-Barr virus (EBV)–negative diffuse large B-cell lymphoma (DLBCL) line harboring the t(14;18) translocation; H9 is a cutaneous T-cell lymphoma (CTCL) line obtained from ATCC (Manassas, VA); SKI is a diffuse large B-cell lymphoma line^{21,22}; HBL-2 and Granta are mantle cell lymphoma lines and both stain positive for Bcl-2^{23,24}; JJN-3 is a multiple myeloma cell line.²⁵ All cell lines were grown as previously described.²⁶

Materials

All reagents for Western blotting were obtained from Bio-Rad Laboratories (Hercules, CA) and Pierce Biotechnology (Rockford, IL); dimethyl sulfoxide (DMSO) was obtained from Sigma-Aldrich (St Louis, MO). Drugs were obtained as follows: AT-101, from Ascenta Therapeutics (Malvern, PA);

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4-hydroxycyclophosphamide, from Niomech (Bielefeld, Germany); carfilzomib, from Proteolix (South San Francisco, CA); all other drugs were obtained from the Columbia University research pharmacy.

Cytotoxicity assays

For all in vitro assays, cells were counted and resuspended at an approximate concentration of 3×10^5 cells/well in a 24-well plate (Becton Dickinson Labware, Franklin Lakes, NJ). AT-101 was diluted in DMSO that was maintained at a final concentration of less than 0.5%. Concentrations of AT-101 from 1 nM to 10 μ M were used in most experiments. Following incubation at 37°C in a 5% CO₂ humidified incubator, 100 μ L from each well was transferred to a 96-well opaque-walled plate; cell-Titer-Glo Reagent (Promega, Madison, WI) was added in a 1:1 ratio. Contents were mixed for 2 minutes on an orbital shaker to induce cell lysis. The plates were allowed to incubate at room temperature for 10 minutes before recording luminescence with a Synergy HT Multi-Detection Microplate Reader (Biotek Instruments, Winooski, VT). In the schedule dependency experiments, serial dilutions of each drug were prepared in ratios relative to their IC₅₀. Cells were preincubated with AT-101 for up to 72 hours, while 4-HC was added for a 24-hour period, being added at time 0, 24 hours, and 48 hours from the start of incubation. Each experiment was performed in triplicate and repeated at least twice.

Flow cytometry

RL or HBL-2 cells were seeded at a density of 7×10^5 /mL. For the transmembrane mitochondrial membrane potential ($\Delta\psi_m$) determination, cells were incubated with concentrations of AT-101 from 1 nM to 10 μ M for up to 24 hours, stained with 1.3 μ g/mL JC-1 dye (Invitrogen, Carlsbad, CA) for 30 minutes, and incubated at 37°C in normal growth media, then washed once in PBS, resuspended in 200 μ L media, and analyzed using a FACSCalibur Flow Cytometer (BD, Franklin Lakes, NJ). A minimum of 5×10^4 /mL events were acquired from each sample. The green and red fluorescence intensities were resolved by detection in the FL1 and FL2 channels, respectively. Median values obtained from the gated FL1 and FL2 channels were used to calculate the normalized transmembrane mitochondrial membrane potential ($\Delta\psi_m$). Each experiment was performed in triplicate. Data are presented as the average plus and minus a standard deviation (SD). For detection of apoptosis, Yo-Pro-1 and propidium iodide (PI) were used (Vybrant apoptosis assay; Invitrogen). After incubation with the approximate IC₁₀₋₃₀ of AT-101 alone and in combination with the approximate IC₁₀₋₃₀ of bortezomib or carfilzomib for 24 hours, cells were washed in cold PBS, centrifuged, and resuspended in PBS. Yo-Pro-1 (1 μ L) and PI (1 μ L) were added to each 1 mL cell suspension. The fluorescence signals acquired by a FACSCalibur System were resolved by detection in the conventional FL1 and FL3 channels. Cells were considered early apoptotic if Yo-Pro-1 positive but PI negative, late apoptotic if Yo-Pro-1 and PI positive.

Bcl-2 immunoblotting

RL cells were seeded at a density of 7×10^5 /mL and incubated with AT-101 at 100 nM, 1 μ M, or 10 μ M under normal growth conditions for 8 or 24 hours. Bcl-2 immunoblotting experiments were performed as described previously.²⁶

Confocal microscopy analysis

Cells were seeded at a density of 7×10^5 /mL. After incubation with AT-101 at 10 μ M for 24 hours, cells were exposed to 5 μ g/mL Hoechst 33342, 5 μ g/mL MitoTracker Red, and 1 μ g/mL YO-PRO-1 for 20 minutes at room temperature. The membranes of apoptotic cells, but not the membranes of live cells, are permeable to the YO-PRO-1; Hoechst 33342 is a specific stain for AT-rich regions of double-stranded DNA, while MitoTracker Red is concentrated by active mitochondria in living cells. Fluorescence of stained cells was detected with the use of a laser scanning confocal microscope (Leica TCS AOBS SP2, inverted stand; Leica Microsystems, Heidelberg, Germany). Image acquisition and analysis were performed with a semiautomated and design-based

stereology system (MetaMorph version 6; Universal Imaging, Downingtown, PA). The percentage of apoptotic cells (apoptotic ratio, AR) was calculated using the formula $AR = A/T$ where A = apoptotic cells (YO-PRO-1 positive [green] and Hoechst 33342 positive [blue] but MitoTracker Red negative [red]) and T = total cell (Hoechst positive) count. Up to 20 images for each sample were acquired and analyzed in 2 different experiments.

Mouse xenograft models

In vivo experiments were performed as described previously.²⁶ In brief, 5- to 7-week-old severe combined immunodeficiency (SCID) beige mice (Taconic Laboratories, Germantown, NY) were injected with 10⁷ RL-DLBCL cells on the flank via a subcutaneous route. When tumor volumes approached 50 mm³, mice were separated into treatment groups of 8 to 10 mice each. Tumors were assessed using the 2 largest perpendicular axes (l indicates length; w, width) as measured with standard calipers. Tumor volume was calculated using the formula $4/3 \pi r^3$, where $r = (l + w)/4$. Tumor-bearing mice were assessed for weight loss and tumor volume at least twice weekly. Animals were killed when one-dimensional tumor diameter exceeded 2.0 cm, or after loss of more than 10% body weight in accordance with institutional guidelines. All studies were conducted under an approved institutional animal protocol. AT-101 was given by oral gavage. Single-agent experiments were conducted using AT-101 on the following doses and schedules: 25, 35, 50, and 75, 100 mg/kg per day for up to 4 consecutive weeks; 200, 240, and 280 mg/kg per week for up to 3 weeks; and 100, 120, and 140 mg/kg on days 1, 4, 8, and 11 on a 21-day cycle. In the combination experiments, AT-101 was administered at a dose of 35 mg/kg per day for 10 days or 200 mg/kg on days 0 and 6; rituximab (R, 10 mg/kg) and cyclophosphamide (Cy, 50 mg/kg) were administered by intraperitoneal injection on days 2, 4, 6, and 8. For the in vivo experiments, AT-101 was prepared in a vehicle solution of 0.5% sodium carboxymethylcellulose (CMC; Sigma-Aldrich) dissolved in purified sterile water. Control groups were treated with the vehicle solution administered by oral gavage.

Plasma pharmacokinetic analyses

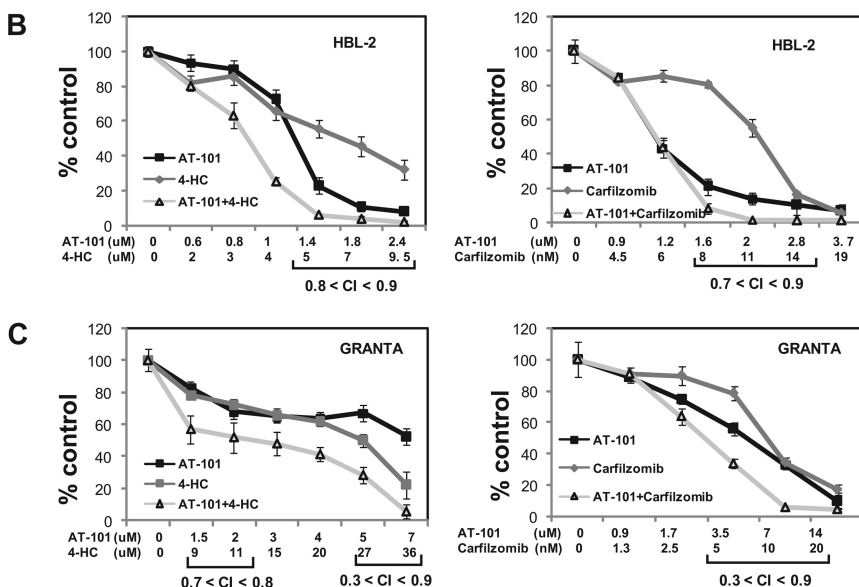
Twenty non-tumor-bearing SCID beige mice received either 35 mg/kg or 200 mg/kg AT-101 in a single dose by oral gavage. Blood collection was performed by intracardiac puncture before the administration of AT-101 and after 30 minutes, 1.5 hours, 3 hours, 6 hours, 12 hours, and 24 hours, with up to 4 mice at each time point. Blood samples were collected in K₂EDTA tubes (BD Vacutainer) and immediately placed in an ice bath. Within a maximum of 30 minutes following blood collection, centrifugation of blood samples at 1800g for 5 minutes at 4°C was applied. Plasma was then transferred into a separate tube. Reduced glutathione (Roche Pharmaceuticals, Nutley, NJ) was added to the whole blood and plasma samples to achieve a concentration between 10 nM and 20 nM. The plasma concentrations of AT-101 in mouse plasma were measured using an electrospray high performance liquid chromatography with tandem mass spectrometry detection (LC-MS/MS) assay (Micromass Quattro-LC triple quadrupole mass spectrometer; Agilent, Palo Alto, CA). The LC-MS/MS assay quantitation range was 20 to 2000 ng/mL based on ex vivo spiking of AT-101 in mouse plasma at an assay volume of 100 μ L. The quantitation of AT-101 in mouse plasma was based on an internal standard approach of spiking internal standard at a constant level to all samples.

Statistical analysis

Tumor volume is presented graphically as the mean at each time point for each treatment group. Each animal's time-tumor volume curve is represented using the area under the curve (AUC), which is interpreted as the total tumor burden of the animal. A logarithmic transformation to normalize the AUC is followed by an analysis of variance for group comparisons with an adjustment for multiple comparisons using resampling. All significance testing was done at the *P* less than .05 level protecting the family-wise error rate. For different in vitro experimental groups, permutation tests were performed to determine whether any of

	HBL-2	GRANTA	H9	RL	SKI	JIN-3
IC ₅₀ (μM) 24h	1.2 (0.5-1.6)	3.5 (1.9-6.3)	3.9 (2.3-6.4)	5.0 (3.0-8.0)	5.6 (3.1-9.8)	7.4 (5.6-9.6)
IC ₅₀ (μM) 48h	0.7 (0.5-0.9)	2.5 (2.2-2.8)	1.7 (0.9-3.2)	1.8 (1.3-2.7)	2.0 (1.2-3.4)	3.9 (2.5-6.0)
IC ₅₀ (μM) 72h	0.3 (0.2-0.5)	1.7 (1.3-2.1)	1.1 (0.4-2.7)	0.9 (0.6-1.4)	0.9 (0.4-1.8)	1.2 (0.8-1.8)

Figure 1. Luminometric assay. (A) AT-101 alone in 6 cell lines of lymphoma and multiple myeloma; IC₅₀s (μM) to AT-101 are shown; estimates from analysis of means and confidence intervals are shown between parentheses. (B) Combination of AT-101 with carfilzomib or 4-HC in HBL-2 after 24 hours. (C) Combination of AT-101 with carfilzomib or 4-HC in Granta after 24 hours. Error bars represent SD.



the experimental groups were superior to a control group. The analysis compared groups using analysis of variance after a normalizing transformation. All *P* values were adjusted using Dunnett method.²⁷ For each cell line, the IC₅₀ (inhibitory concentration of 50% of cells) and the drug to drug interactions in terms of synergism, additivity, or antagonism were computed using the Calcsyn software (Biosoft, Cambridge, United Kingdom; combination index [CI] < 1 defines synergism; CI = 1, additivity; and CI > 1, antagonism).

Results

AT-101 sensitizes mantle cell lymphoma and diffuse large B-cell lymphoma to chemotherapy in vitro

Figures 1-3 present the data for the mantle cell lymphoma cell lines, while Figures 4-6 present in vitro and in vivo data for the large B-cell lymphoma experiments. The IC₅₀ values for AT-101

across a panel of different lymphoproliferative malignancies ranged from 1.2 to 7.4 μM after a 24-hour exposure; 0.7 to 3.9 μM after a 48-hour exposure; and 0.3 to 1.7 μM after 72 hours. In general, the range of IC₅₀ values was relatively restricted, with the mantle cell lymphoma line HBL-2 being the most sensitive, and the multiple myeloma being the most resistant (Figure 1A). Figure 1B,C presents the combination of AT-101 with carfilzomib or 4-HC. In the HBL2 and Granta cell line, the combination of AT-101 plus bortezomib for 24 hours revealed antagonism (CI range: 1.5-2.9), while AT-101 plus carfilzomib showed synergism (CI range: 0.3-0.9). AT-101 plus etoposide, doxorubicin, or 4-HC showed mathematic synergy in both MCL lines (CI range: 0.3-0.9). The combination of AT-101 and 4-HC was synergistic only when the 2 drugs were given in sequence with a 24- or 48-hour preexposure to AT-101 (CI ≤ 0.8, Figure 4D). When both drugs were added simultaneously, antagonism was observed (CI range: 1.2-2.7).

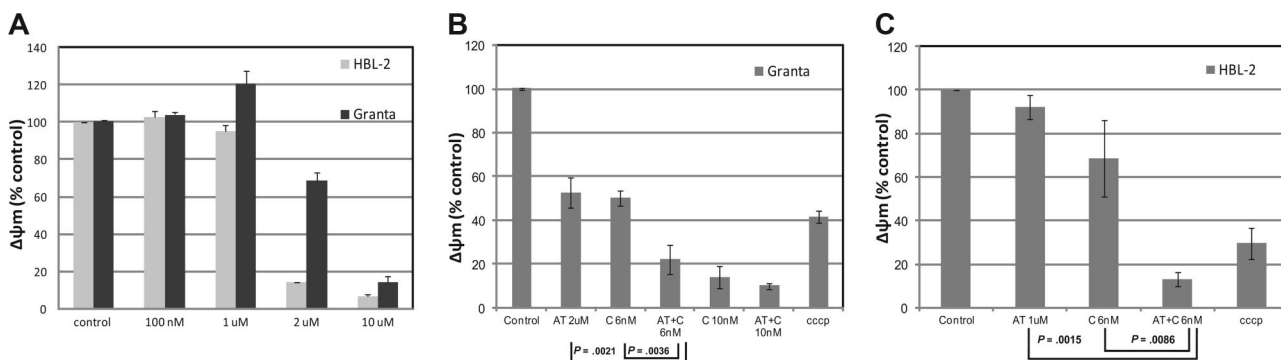


Figure 2. Assessment of the mitochondrial membrane potential ($\Delta\psi_m$) in HBL-2 and Granta (MCL). (A) HBL-2 and Granta cells were incubated with AT-101 from 100 nM to 10 μM for 24 hours. (B) Granta cells were incubated with AT-101 (AT, 2 μM) or carfilzomib (C, 6 or 10 nM) or both for 24 hours. The combination of AT-101 plus C at 6 nM was statistically significant compared to any of the single groups and controls (*P* ≤ .004). (C) HBL-2 cells were incubated with AT-101 (AT, 1 μM), carfilzomib (C, 6 nM) or for 24 hours. Again, the combination of AT-101 plus C was statistically significant compared to any of the single groups and controls (*P* ≤ .009). Error bars represent SD.

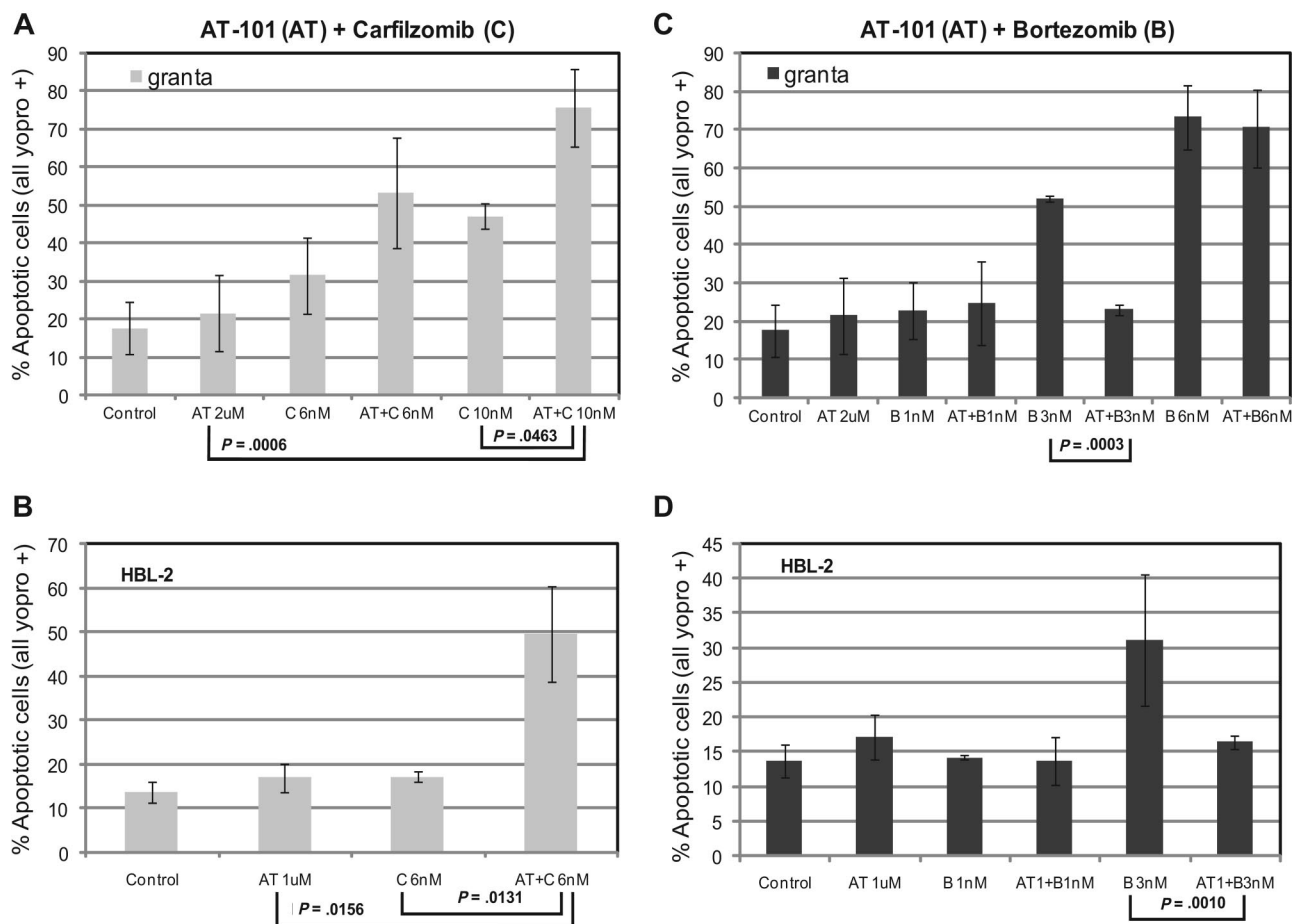


Figure 3. Assessment of apoptosis by Yo-pro-1 and propidium iodide (PI) in HBL-2 and Granta cell lines. (A,B) Treatment with AT-101 (1 μ M or 2 μ M) and carfilzomib (C, 6 nM or 10 nM) induces apoptosis in at least 50% of cells ($P \leq .046$ for Granta and $P \leq .016$ for HBL-2). (C,D) The combination of AT-101 (1 μ M or 2 μ M) and bortezomib (1 nM, 3 nM, or 6 nM) may be antagonistic. Error bars represent SD.

AT-101 disrupts the $\Delta\psi_m$ in a concentration- and time-dependent manner in a diffuse large B-cell and in mantle cell lymphoma lines

$\Delta\psi_m$ antecedes cell death and is a potentially important early biomarker of apoptosis. Figure 4A depicts the time- and dose-dependent changes in the normalized $\Delta\psi$ when cells (RL) were exposed to 1 μ M, 3 μ M, or 10 μ M AT-101 for 1, 4, 8, 12, and 24 hours. After 24 hours, there was a clear concentration-dependent reduction in the $\Delta\psi_m$ (from 1.11 ± 0.0096 at 1 μ M to 0.5 ± 0.0133 for 3 μ M to 0.14 ± 0.0007 for 10 μ M), with more than an 80% change in the mitochondrial membrane potential between the lowest and highest concentrations. Since earlier time points did not elicit a concentration-dependent response, the data in Figure 4A suggest that there may be a critical time of exposure, between 12 and 24 hours, for cells exposed to the higher concentrations of AT-101 (3 and 10 μ M). A multiple comparison analysis revealed that there was a statistically significant difference in $\Delta\psi_m$ that favors 10 μ M for 24 hours compared with all other scenarios ($P < .001$). These changes in membrane potential were not attributable to a reduction in Bcl-2. Figure 4C demonstrates that the levels of Bcl-2 in RL cells were not affected following exposure to AT-101 at 0.1, 1, or 10 μ M. No significant changes in Bcl-2 levels were observed after 8 or 24 hours of exposure compared with control.

In the MCL lines, AT-101 induced a concentration-dependent depolarization of the mitochondrial membrane with a decrease of

the normalized $\Delta\psi_m$ of more than 50% in the range of 1 to 2 μ M for HBL-2 and 2 to 10 μ M for Granta (Figure 2A). The combination of AT-101 and 6 nM carfilzomib produced a significant decrease in $\Delta\psi_m$ for the combinations compared with any single-agent exposure (Granta: $P = .004$ and $P = .002$ for the comparisons to carfilzomib alone and AT-101 alone, respectively; HBL-2: $P = .009$ and $P = .002$ for the same comparisons, Figure 2B,C). The combination of AT-101 and bortezomib did not elicit any significant change compared with bortezomib alone in both cell lines (data not shown).

AT-101 plus carfilzomib induces apoptosis in mantle cell lymphoma

Treatment with AT-101 and carfilzomib for 24 hours induced apoptosis in both HBL-2 and Granta cell lines. Exposure of Granta cells to AT-101 plus carfilzomib induced apoptosis in more than 70% of cells, compared with only 20% for AT-101 alone ($P = .001$) and approximately 45% for carfilzomib alone ($P = .046$, Figure 3A). When HBL-2 cells were treated with AT-101 and a slightly lower dose of carfilzomib, approximately 50% of the cells detected were apoptotic, compared with less than 20% for the individual drugs ($P = .013$ and $P = .016$ for the comparisons to carfilzomib and AT-101 alone, respectively, Figure 3B). When AT-101 was combined with bortezomib, significantly less apoptosis was observed in the combination group compared with bortezomib alone in both cell lines (Figure 3C,D).

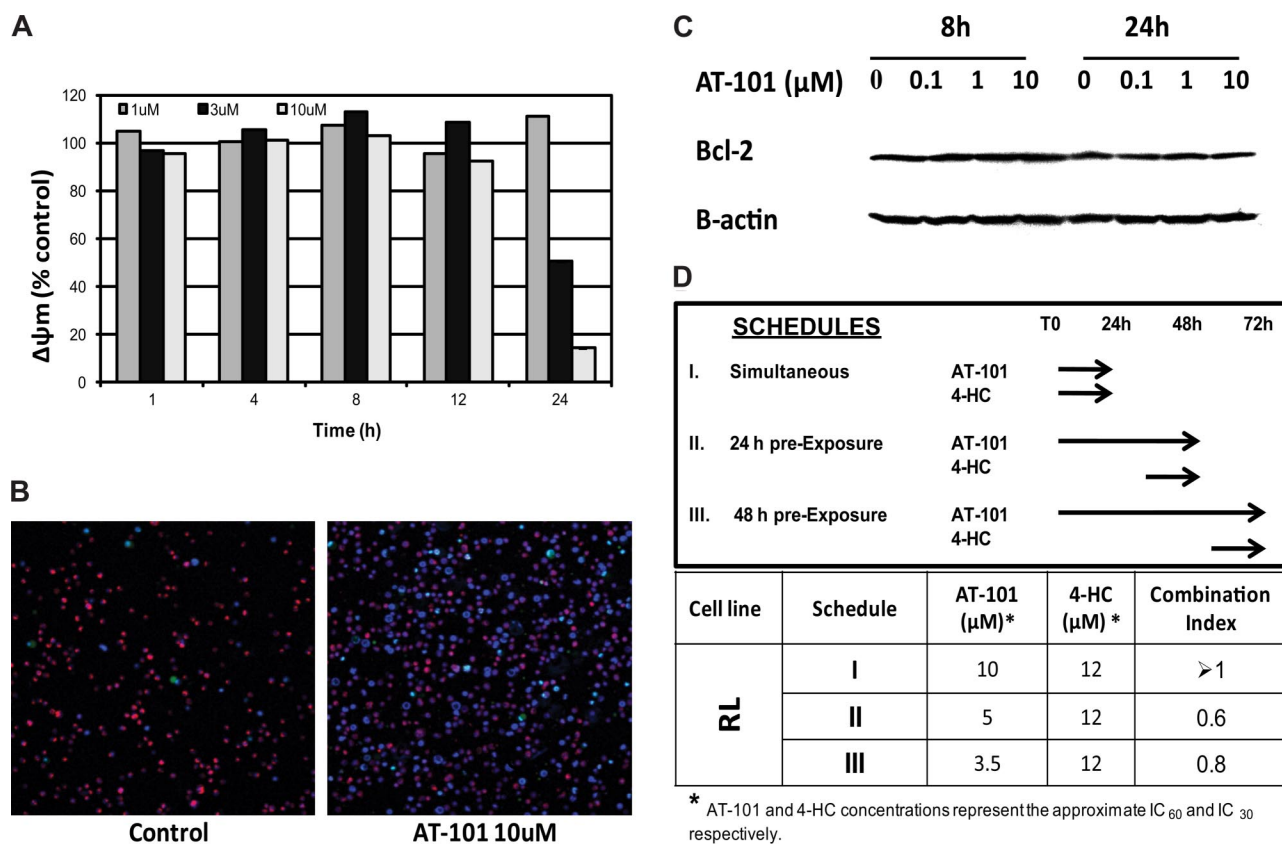


Figure 4. Diffuse large-B cell line (RL). (A) Assessment of the mitochondrial membrane potential ($\Delta\psi_m$). AT-101 (1 μ M, 3 μ M, 10 μ M) from 1 hour to 24 hours of exposure. (B) Confocal microscopic detection of apoptosis in a large B-cell lymphoma line (RL) treated with AT-101 for 24 hours. YO-PRO-1 (nuclei of apoptotic cells, green); Hoechst 33342 (nuclei, blue), MitoTracker Red (active mitochondria). AT-101 induces apoptosis in a large B-cell lymphoma cell line (RL) after exposure for 24 hours. Control, AR=3%; AT-101 10 μ M, AT=38.7%. (C) Immunoblotting for Bcl-2 following treatment with AT-101 in a large B-cell lymphoma line (RL). AT-101 does not affect levels of Bcl-2. (D) Cytotoxicity assay for AT-101 combined with 4-HC in a large B-cell lymphoma line (RL). Model of in vitro exposure to AT-101 and other drugs; a preexposure to AT-101 for up to 48 hours before adding 4-HC for an additional 24 hours revealed a synergistic interaction of the combination.

Confocal microscopy shows induction of apoptosis in a large B-cell lymphoma line

Confocal microscopy was used to directly study changes in the treated cell populations as a function of concentration and duration of exposure. The incubation of RL cells with 10 μ M AT-101 for 24 hours induced apoptosis based upon the YO-PRO-1, Hoechst 33342, and MitoTracker Red staining. Treated cells revealed an apoptotic ratio of 38.7% compared with a 3% in the untreated controls (Figure 4B). Cells exposed to AT-101 for 24 hours revealed a prevalence of both YO-PRO-1- and Hoechst 33342-positive and MitoTracker Red-negative cells, likely representing a later phase of apoptosis. However, treatment of cells at lower concentrations or for shorter durations of exposure produced a minimal change in the YO-PRO-1- and Hoechst 33342-positive and MitoTracker Red-positive cells. These data corroborate the in vitro studies and the $\Delta\psi_m$ data supporting the finding that 10 μ M AT-101 for 24 hours seems to be optimal in inducing apoptosis.

Single-dose AT-101 plasma pharmacokinetics in SCID beige mice

The AT-101 peak plasma concentration was observed after 30 minutes of administration of the drug by oral gavage at 2 dose levels (35 mg/kg and 200 mg/kg). The 200-mg/kg group demonstrated an average plasma concentration (C_{max} and AUC) almost 4 times greater than the 35-mg/kg group (with C_{max} at 27.78 μ M

and 7.88 μ M, respectively). Twenty-four hours after exposure to AT-101, the drug was still detectable in the plasma with average concentrations of 0.49 μ M for the 35-mg/kg group and 0.39 μ M for the 200-mg/kg group (Figure 5A).

In vivo activity of AT-101 reveals marginal single-agent activity but marked potentiation of conventional cytotoxic therapy

Early experiences with AT-101 in the in vivo models were directed toward understanding the toxicity and efficacy of various single-agent schedules. The RL cell line was selected because of its known chemotherapy and AT-101 resistance. A variety of schedules were explored as described in Figure 5B. Overall, increasing the dose of AT-101 from 25 mg/kg to 100 mg/kg per day indefinitely resulted in earlier onset of weight loss equivalent to more than 10% of the pretreatment weight and death in many animals. For example, mice receiving 75 or 100 mg/kg per day lost 10% of their body weight by days 14 and 7, respectively, with almost all animals dying from drug-related toxicity by day 21 (4 of 5) and day 16 (5 of 5), respectively. Mice treated with 50, 75, or 100 mg/kg per day lost a statistically significant amount of weight (ie, > 10% compared with control animals [$P = .029$, $P = .016$, and $P = .016$, respectively]). A statistical analysis of weight changes as a function of time yielded area under the curve values correlating with cumulative weight loss (ie, toxicity) as a function of different treatment groups. These data demonstrated that mice receiving 25 mg/kg per

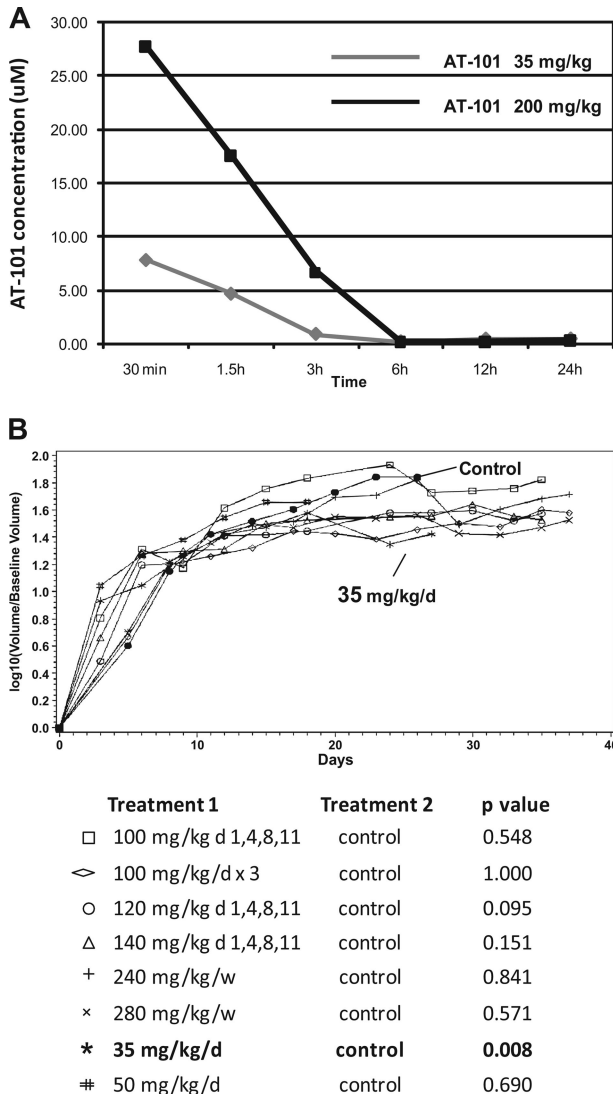


Figure 5. In vivo SCID-beige xenograft model for DLBCL (RL). (A) Pharmacokinetic (PK) modeling of AT-101 after a single administration by oral gavage. AT-101 was administered at a 35 mg/kg or a 200 mg/kg; the AT-101 peak plasma concentration was observed after 30 minutes of administration of the drug in both the dose levels, with the 200 mg/kg group showing a plasma average concentration almost 4 times greater than the 35 mg/kg group (7.88 µM and 27.78 µM respectively). After 24 hours, AT-101 was still detectable in plasma with average concentrations of 0.49 µM for the 35 mg/kg group and 0.39 µM for the 200 mg/kg group. (B) In vivo activity of single AT-101 as a function of schedule. The 35 mg/kg per day for 14 days showed a statistically significant shrinkage of tumor volume compared to controls ($P = .008$); P values are shown; P values for each treatment group compared to control. All significance testing was done at the P less than .05 level. N equals 5 in each group.

day for up to 4 weeks; 35 mg/kg daily for 2 weeks; 200 mg/kg weekly for 3 consecutive weeks; and 100 mg/kg on days 1, 4, 8, and 11 lost some weight relative to control animals, although less than 10% of their pretreatment weight. In another analysis, the time to death from drug-related causes suggested a statistically significant advantage for the 25 mg/kg per day for 4 weeks schedule over any other cohort with a significantly shorter survival and greater toxicity being appreciated for the 240 mg/kg per week \times 3 ($P = .016$), 280 mg/kg per week \times 3 ($P = .016$), 50 mg/kg per day \times 14 days ($P = .029$), 75 mg/kg per day schedules ($P = .016$) and 100 mg/kg per day ($P = .016$) cohorts compared with control.

Interestingly, the assessment of tumor volume across all treatment cohorts revealed no statistically significant tumor shrinkage

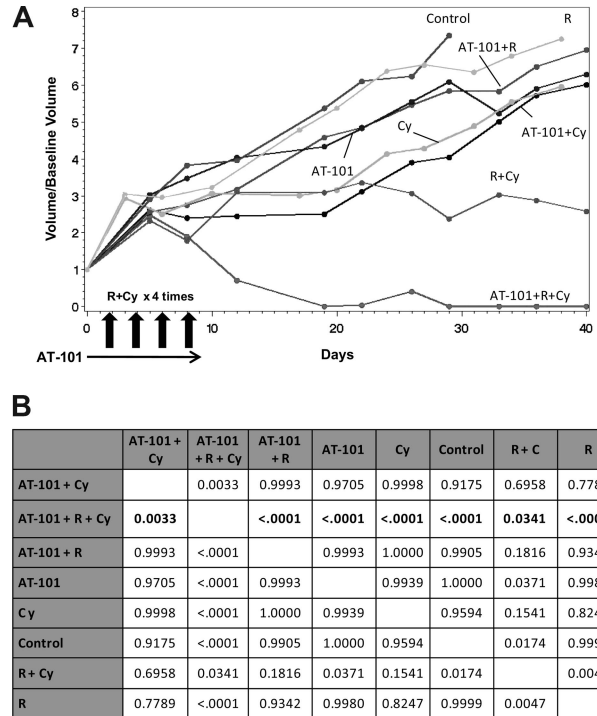


Figure 6. In vivo SCIDbeige xenograft model for DLBCL (RL) Combination experiments. (A) The combination of oral AT-101 35 mg/kg per day for 10 days regimen plus intraperitoneal cyclophosphamide (Cy) and intraperitoneal rituximab (R) in four administrations together on days 2, 4, 6, 8 showed significant tumor volume control compared to any other treatment group. (B) The multiple comparison analysis shows the superiority of the triplet combination to each other group ($P \leq .034$). Data represent P values for the comparisons. All significance testing was done at the P less than .05 level. N equals 7 in each group ($n = 6$ in control group).

for AT-101-treated animals compared with the control, except for the cohort receiving 35 mg/kg per day for 2 weeks ($P = .008$). This latter cohort was the only one that demonstrated a statistically significant reduction in tumor schedule of all the schedules and doses studied. Figure 5B presents the tumor volume growth curves for these different treatment cohorts. Mice receiving 120 mg/m² on days 1, 4, 8, and 11, while trending toward a statistically significant therapeutic outcome, also experienced significant weight loss and toxicity on this schedule. Based on these toxicologic experiments and the efficacy findings, 35 mg/kg per day was chosen for future combination experiments.

Combination experiments focused on integrating AT-101 with cyclophosphamide (C) and rituximab (R). Overall, these regimens were well tolerated. After 8 days of treatment, mice receiving AT-101 alone, AT-101 plus C, and AT-101 plus R plus C showed an average weight loss of between 10% and 15%, which recovered once treatment was stopped. Figure 6A presents the area under the curve of tumor growth as a function of time. These data demonstrate that no single agent displayed a therapeutic advantage versus the control, nor were the differences significantly better compared with each other. In this experiment, intentionally lower doses and less dose-intense schedules were used to identify potentially favorable drug-drug interactions. Among the doublet comparisons, AT-101 plus C and AT-101 plus rituximab were not significantly better than any other doublet ($P = .182$). Interestingly, R plus C was statistically better than AT-101 alone ($P = .037$), rituximab alone ($P = .005$), and control ($P = .017$), but was not superior to C alone ($P = .154$). Despite the lower doses and less dose intensity, AT-101 clearly improved the treatment effects of rituximab and

C ($P = .034$). Results of the multiple comparison analysis are shown in Figure 6B.

An additional combination experiment with AT-101 plus R and C was performed to explore a slightly different schedule of AT-101. This experiment was designed to determine whether higher concentrations of AT-101, leading to more significant changes in $\Delta\psi_m$, would be associated with improved therapeutic outcome in the AT-101 plus C and AT-101 plus R cohorts. In these experiments, 2 single high doses of AT-101 (200 mg/kg) were given on days 0 and 6, while the R and C were given as previously described. Approximately 45 days from treatment, the combination of AT-101 plus R was significantly better compared with AT-101 alone ($P = .044$), rituximab alone ($P = .001$), and control ($P = .004$). The combination of AT-101 plus C was also significantly better than AT-101 alone ($P = .001$), C alone ($P = .002$), the control ($P < .001$), rituximab alone ($P < .001$), and the combination of R plus C ($P = .037$). The multiple comparison analysis trended toward significant ($P = .064$) for the comparison of AT-101 200 mg/kg plus R plus C versus R plus C alone. This experiment appears to corroborate the *in vitro* findings, suggesting that a higher C_{max} -type exposure around the time of the chemotherapy may be the optimal strategy for synergizing with conventional chemotherapy agents.

Discussion

Therapeutic strategies targeting Bcl-2 represent a promising prospect for treating many types of cancers. Given the prominent role Bcl-2 family members play in lymphoproliferative malignancies, there is a strong rationale for targeting these pathways. At present, there are a number of different strategies for targeting both the intrinsic and extrinsic arms of these survival pathways, including small molecules, antisense approaches, and monoclonal antibodies.²⁸⁻³¹ One of the major questions regarding the potential clinical application of these compounds revolves around precisely how to exploit them in combination. Practically, there is little expectation that these compounds will ever emerge as prominent single agents for the treatment of any cancer. Their development path is likely to require their integration into conventional chemotherapy regimens. This approach will require a detailed understanding of how Bcl-2–targeted drugs impact the relevant biology, and how basic pharmacologic considerations affect the activity of the combination. In these experiments, the major objective was to understand the optimal concentrations, doses, and schedules of AT-101, a small-molecule BH3 mimetic that binds to the BH3 domain of Bcl-2, Bcl-X_L, and Mcl-1.^{10,32-34}

Mohammad et al³⁵ demonstrated that (–)-gossypol (AT-101) is capable of overcoming acquired and intrinsic drug resistance to CHOP (cyclophosphamide, doxorubicin, vincristine, and prednisone) chemotherapy. Jazirehi et al³⁶ have demonstrated that rituximab down-regulates Bcl-2 expression and Bcl-X_L transcription in lymphoma cell lines, reinforcing its potential role as a “Bcl-2–directed” therapy. These studies however, did not explore how different doses and schedules of AT-101 affected important pharmacokinetic or pharmacodynamic end points, or how to use this information to establish optimal combination strategies.

Duration of exposure was generally not a critical determinant of cytotoxicity. The cytotoxicity data corroborate the $\Delta\psi_m$ data that similarly established the importance of concentration as major determinant of cytotoxicity. These data support the finding that while AT-101 does not affect levels of Bcl-2, it does affect Bcl-2 function as measured by the $\Delta\psi$. The observed effect on the $\Delta\psi$ was prominent after 24 hours of exposure in a concentration range

between 1 μ M and 10 μ M, suggesting the possible existence of a threshold concentration necessary to trigger apoptosis.

In the mantle cell lines (HBL-2 and Granta), AT-101 showed a synergistic interaction when combined with the proteasome inhibitor carfilzomib,^{37,38} doxorubicin, etoposide, or 4-HC after a 24-hour exposure. The combination with carfilzomib also showed induction of apoptosis and mitochondrial membrane depolarization after 24 hours, with the combination being statistically superior to any single agent in both cell lines. Interestingly, the combination of AT-101 with bortezomib showed antagonism in the cytotoxicity assays and significantly less induction of apoptosis in both cell lines. Kuhn et al recently demonstrated that carfilzomib was more potent than bortezomib in overcoming both primary and secondary resistance to bortezomib in both cell line models and clinical samples.³⁸ The authors speculate that the irreversible effect of carfilzomib on the proteasome and immunoproteasome subunits may partially explain this observation.

Xenograft experiments in SCID beige mice with AT-101 alone showed that a daily dose lower than 50 mg/kg could be safely administered for up to 2 to 4 weeks. The 35 mg/kg per day for a 2-week schedule was safe, and was the only schedule shown to be statistically better than the control or any other treatment cohort ($P = .008$, Figure 5B), although the analysis of time to death was not statistically significantly different between the 35-mg/kg per day and control cohorts ($P = .405$). The single agent *in vivo* dosing experiments suggest that efficacy is not necessarily a function of dose, but rather, schedule as well. For example, 35 mg/kg per day for 2 weeks delivered a cumulative dose of 490 mg/kg, while only approximately 80% of this cumulative dose was delivered with the 100 mg/kg on the 1-, 4-, 8-, and 11-day schedule (400 mg/kg). While both schedules were safe and feasible, the latter schedule was not associated with any therapeutic benefit. A similar comparison can be applied to the 35 mg/kg \times 14-day schedule and the 200 mg/kg \times 3 weekly doses, in which the latter schedule is associated with a greater cumulative dose, although with little to no improvement in tumor shrinkage and survival compared with the daily schedule.

The combination *in vitro* data with AT-101 and 4-HC in RL support the hypothesis that a preexposure to AT-101 for up to 48 hours is necessary to sensitize this relatively drug-resistant cell line to conventional chemotherapy. Using the most effective single-agent schedule of AT-101, a multiple comparison analysis of combination studies with rituximab and cyclophosphamide demonstrated the superiority of the triplet combination with 35 mg/kg per day AT-101 for 10 days, starting 2 days before the administration of the other drugs. Interestingly, complete responses and statistically significant differences in tumor volume shrinkage were observed only in animals receiving rituximab and C, with no additive benefit in those mice receiving AT-101 plus rituximab or AT-101 plus cyclophosphamide. This is in contrast to the *in vivo* experiment in which AT-101 was administered at a very high single exposure (200 mg/kg \times 2 doses on days 0 and 6) in combination with rituximab and C, in which there were significant improvements in the AT-101 plus C and AT-101 plus rituximab combinations, with a trend toward statistical significance in mice treated with AT-101 plus rituximab and C ($P = .064$). These data also demonstrated a statistically significant advantage for those mice treated with AT plus C compared with R plus C, suggesting AT-101 in combination with C exceeded the benefits of adding rituximab. While the contribution of rituximab in these models is likely to be underestimated given the immunocompromised nature of these animals, the

data still support an important role for the addition of this drug to the regimen.

Hence, 2 major observations have emerged from the *in vivo* experiments: (1) a low daily dose of AT-101, in fact the only schedule shown to induce significant responses compared with control, complimented the activity of only the combination of R plus C and not the individual treatments with R or C alone; and (2) higher dose schedules of AT-101 complimented treatments of R and C alone, and trended toward significance with the combination of R plus C. Obviously, each of these 2 different schedules is associated with markedly different pharmacokinetic profiles, with the peak plasma concentrations of 7.88 and 27.78 μM in the 35-mg/kg per day and 200-mg/kg $\times 2$ schedules, respectively, while the plasma concentrations in both cases were less than 1 μM after 24 hours. These data suggest that the average plasma peak concentration for the 200-mg/kg schedule is almost 4 times higher than the low-dose schedule. These PK parameters appear to be in line with the concentrations noted to induce cytotoxicity *in vitro* (IC_{50} values for RL in the range of 3–4 μM) and $\Delta\psi$ in most cell lines (ie, the greatest changes in $\Delta\psi$ occur between 1–10 μM for 12 to 24 hours).

One potential explanation for the differences between the low-dose and high-dose AT-101 schedules with chemotherapy may revolve around the potency of AT-101's ability to influence apoptotic priming. Practically, it is well recognized that combinations of drugs are more effective and efficacious in producing cell death compared with single agents. Assuming there is a relationship between the change in the $\Delta\psi$ and a reduction in the threshold for induction of apoptosis, it is plausible that low-dose AT-101 schedules may result in less change in the $\Delta\psi$ compared with higher dose schedules, therefore requiring a greater "proapoptotic influence" (ie, R + C) to induce cell death. Clearly, AT-101 compliments the combination of R + C when given on the daily schedule. The higher dose schedule produces greater change in the $\Delta\psi$, therefore allowing induction of apoptosis with a relatively less toxic approach, as is seen with the 200-mg/kg dosing schedule with rituximab or C alone. Perhaps most importantly, these data suggest that there may be 2 different strategies to exploit the therapeutic advantage of AT-101, depending upon whether it is to be used as a single agent (daily) or in combination (weekly or pulse high dose). Future strategies for combination studies should focus on achieving

the highest doses possible (ie, greatest changes in $\Delta\psi$) for short periods of time, around the time of cytotoxic therapy administration.

Clearly, future clinical studies will need to explore the appropriate pharmacodynamic end points as a function of various pharmacokinetic parameters to determine the best strategies for integrating these agents into standard chemotherapy regimens. These data have formed the basis for planned clinical studies in lymphoma.

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Authorship

Contribution: L.P. designed and performed all presented experiments, interpreted data, and wrote the paper; M.G. did the statistical analysis; J.R.G. contributed by doing the flow cytometry; J.M. contributed with the cytotoxicity assays; D.Y., J.H., M.S., and L.L. provided expertise regarding the design of the pharmacokinetic and *in vivo* experiments; K.M. provided assistance with the laser confocal microscopy; G.M. provided assistance with Bcl-2 Western blots; M.L.H. provided input regarding experiments measuring the transmembrane mitochondrial potential; O.A.O. provided input regarding experimental design, interpretation of data, and editing of the final paper.

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