Degrasyn Potentiates the Antitumor Effects of Bortezomib in Mantle Cell Lymphoma Cells \textit{In vitro} and \textit{In vivo}: Therapeutic Implications

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Abstract

Mantle cell lymphoma (MCL) is an aggressive histotype of B-cell non–Hodgkin lymphoma that has increased in incidence over the past few decades and is incurable, usually poorly responsive to standard chemotherapy combinations, and associated with poor prognoses. Discovering new therapeutic agents with low toxicity that produce better outcomes in patients with MCL is an ongoing challenge. Recent studies showed that degrasyn, a novel small-molecule inhibitor of the Janus kinase/signal transducer and activation of transcription (JAK/STAT) pathway, exerts antitumor activity in lymphoid tumors by inhibiting key growth and survival signaling (JAK/STAT) pathways. In the present study, we found that treatment of both typical and blastoid-variant MCL cells with degrasyn in combination with bortezomib resulted in synergistic growth inhibition and apoptosis induction \textit{in vitro}. The apoptosis in these cells was correlated with the down-regulation of constitutive NF-κB and phosphorylated STAT3 activation, leading to the inhibition of c-Myc, cyclin D1, and bcl-2 protein expression and the upregulation of bax protein expression. \textit{In vivo}, degrasyn and bortezomib interacted to synergistically prevent tumor development and prolong survival durations in a xenotransplant severe combined immunodeficient mouse model of MCL. These findings suggest that agents such as degrasyn that can pharmacologically target constitutively expressed NF-κB and STAT3 in MCL cells may be useful therapeutic agents for MCL when administered together with bortezomib. 

Introduction

Mantle cell lymphoma (MCL) is an important histotype of aggressive B-cell non–Hodgkin lymphoma (NHL-B) with the poorest long-term outcome among all NHL-B subtypes, the incidence of which has increased over recent decades in the United States and worldwide (1). MCL is characterized by cell cycle abnormalities, including ATM, p53-mediated cell cycle anomalies, DNA repair, apoptosis, in blastoid-variant MCL (BV-MCL), and decreased expression of cyclin-dependent kinase inhibitors such as p16, p21, and p27kip1 (2). Overexpression of the cyclin D1 oncogene and protein are associated with the characteristic nonrandom chromosomal translocation t(11;14)(q13;q32) in MCL as a result of enforced expression by the IgH enhancer on chromosome 14. MCL cases are classified pathologically into at least two subtypes: (a) typical (classic), the less aggressive, small- to medium-cell histotype, and (b) blastoid (BV-MCL), the more aggressive, large-cell histotype (3, 4). The most common clinical course of MCL consists of continual relapses with a median survival duration in most studies of 3 to 4 years (5). Both subtypes of MCL are currently incurable, but some cases show sensitivity and partial or even complete response to dose-intensified regimens of combination chemotherapy (e.g., hyper-CVAD, R-CHOP-CVAD) with occasional extended periods of remission. Overall, however, both MCL and BV-MCL are associated with poor prognoses and relatively short survival (6, 7). Recent approaches to MCL therapy have included the use of humanized monoclonal antibodies (rituximab), various strategies involving allogeneic and autologous bone marrow transplantation (8, 9), and molecular therapeutic targeting with proteasome inhibitors [bortezomib (Velcade; refs. 10, 11), mammalian target of rapamycin antagonists (12, 13), and newer immunomodulatory agents (lenalidomide; Revlimid; refs. 14, 15)]. However, these therapeutic approaches have failed to produce durable remissions in most patients with MCL. Therefore, the need for additional novel approaches to MCL therapy is clearly needed.

Bortezomib (also known as Velcade; PS-341) is one of the first therapeutic proteasome inhibitors demonstrated to be effective in treating hematologic disorders (16, 17).
It is approved in the United States for treating relapsed multiple myeloma and MCL (10). Unfortunately, MCL frequently fails to respond to treatment with bortezomib or subsequently becomes refractory to bortezomib (11). Because single-agent therapy is rarely effective against diseases such as MCL, investigating whether the efficacy of bortezomib can be increased by combining the proteasome inhibitor with other novel type of pharmacologic agents may be necessary. Recent studies, both laboratory and clinical, have indicated that the addition of numerous agents, including thalidomide, lenalidomide, arsenic trioxide, and histone deacetylase inhibitors, to bortezomib may be beneficial in treating MCL (18–20). Recently, WP1066, a novel small-molecular-weight compound, was designed and synthesized by W. Priebe and his coworkers at UT M.D. Anderson. The design of WP1066 and the novel class of Janus kinase/signal transducer and activation of transcription (JAK/STAT) inhibitors to which WP1066 belongs was inspired by the scaffold of the natural product caffeic acid benzyl ester (CABE) and its low potency analog AG490 (originally synthesized as described in ref. 21). WP1066 abilities to suppress the JAK/STAT pathway were documented by suppressing cytokine (e.g., interleukin-6, interleukin-3) activation of STAT3 through downregulating the expression of JAKs upstream from STAT, particularly JAK2 (22–27). Follow-up of design lead optimization of WP1066 led to WP1130, an even more potent inhibitor of both JAK2 and STAT3, possessing additional activities that targeted the degradation of the oncoprotein c-Myc (28). Specifically, degrasyn induces rapid degradation of c-Myc, an oncoprotein found highly expressed in many tumor types that so shown to target Bcr/Abl protein expression that leads to the induction of cell death in chronic myelogenous leukemia cells through inhibition of the transcription factor Oct-1 (Santa Cruz Biotechnology) and an anti–β-actin antibody (Sigma). The secondary antibodies used were peroxidase-conjugated goat anti-mouse and anti-rabbit antibody (Sigma). Peripheral blood mononuclear cells (PBMC) were purified from the donors’ buffy coats by Ficoll gradient.

Antibodies, reagents, and materials

The primary antibodies used in our study included STAT3 and phosphorylated STAT3 (pSTAT3; BD Biosciences Pharmingen), bel-2, bax, cyclin D1, c-Myc, and Oct-1 (Santa Cruz Biotechnology) and an anti–β-actin antibody (Sigma). The secondary antibodies used were peroxidase-conjugated goat anti-mouse and anti-rabbit antibodies (Jackson ImmunoResearch Laboratories). WP1066, WP1129, and degrasyn (WP1130) were synthesized at M. D. Anderson; these compounds were solubilized in DMSO (100 mmol/L) that was further diluted in injectable saline for animal studies. Bortezomib was provided by Millennium.

Gel shift assays and DNA-binding ELISA

Electrophoretic mobility shift assays for nuclear factor NF-κB DNA binding were done according to procedures previously described (35). The DNA binding activity of STAT3 subunits was analyzed using an ELISA according to the manufacturer’s instructions (TransAM STAT Family Transcription Factor Assay kit; Active Motif). Briefly, nuclear extracts were placed in the wells of a 96-well plate that contained an immobilized oligonucleotide carrying a STAT consensus DNA binding site. STAT3 proteins bound to this immobilized oligonucleotide were detected by incubating nuclear extracts with a primary antibody recognizing active STAT3 followed by a horse-radish peroxidase–conjugated secondary antibody and were quantified using spectrophotometry at 450 nm with a reference wavelength of 650 nm.

NF-κB reporter plasmid transfection and luciferase assays

Mino cells were transiently transfected with 5 μg of the 6xNF-κB-luc reporter plasmid according to a nucleofector protocol from Amaxa Biosystems as previously described (31). After transfection, cells were pooled and
separated equally into a 12-well plate. Cells were then treated with specified drug concentrations for 6 and 24 hours. At appropriate time point, cells were harvested and lysed. Whole-cell lysates were used for luciferase assays using the BD Monolight Enhanced Luciferase Assay kit (BD Biosciences) that was normalized according to β-gal activity.

**Cell proliferation assays and synergy calculation**

*In vitro* thymidine incorporation proliferation assays were done for cell growth as previously described (25). Briefly, cells were plated (in triplicate) at 4 × 10^4 cells per well in 200 μL of RPMI 1640 with 10% FCS and the indicated reagents in a 96-well plate, and incubated in 5% CO2 at 37°C. After 24 hours, each well was pulsed with 0.5 μCi/10 μL [3H]thymidine (Amersham) for 16 hours. Cells were harvested, and the radioactivity was measured. The CalcuSyn software program (Biosoft) was used to analyze the result of nonconstant ratio drug combination synergy studies. The combination index and isobologram plots for degrasyn and bortezomib were created using the Chou-Talalay method.

**Immunoblot analysis**

Whole-cell extracts were solubilized in 1% SDS sample buffer and electrophoresed on a 4% to 15% SDS-PAGE gel (Bio-Rad). Proteins were transferred from the gel onto a polyvinylidene difluoride membrane and probed with various specific primary antibodies and appropriate horseradish peroxidase–conjugated secondary antibodies. Proteins were visualized using enhanced chemiluminescence (Amersham).

**Apoptosis and caspase-3 assays**

MCL cells were washed and stained with Annexin V-FITC and propidium iodide in accordance with the manufacturer's protocol recommendation (BD Pharmingen). Apoptosis in the cells was quantified using a fluorescence-activated cell sorter and the CellQuest software program (BD Biosciences). Caspase activity assays were done using a colorimetric substrate, and a protocol and materials supplied by the manufacturer (EMD Chemicals). In brief, cells were lysed, and 50 μg of the resulting cell lysates was added to an assay buffer for a total

Figure 1. Effect of treatment with AG490 congeners on MCL cell proliferation. Mino, DB, Z-138, and JMP-1 cells were treated with WP1066, WP1129, or degrasyn (WP1130) in a dose-dependent manner for 48 h and then analyzed for cell proliferation using thymidine incorporation assays. Points, means and ranges of triplicate cultures of three independent experiments; bars, SD. B, the EC50 of each compound was determined using the GraphPad Prism software program for all four cell lines, and the average EC50 for each compound was calculated. Statistical analysis was determined using the Student’s t test.
volume of 90 μL and incubated at 37°C for 10 minutes. A colorimetric substrate for caspase-3 (final concentration, 200 μmol/L) was added to the mixture. A405 values were recorded for each sample after incubation at 37°C for 2 hours.

RNA isolation and real-time PCR
Total RNA isolation was done by using Trizol LS Reagent (Invitrogen) according to manufactures instructions. Reverse transcription of RNA was carried out with the cDNA archive kit (ABI). Synthesized cDNA was subjected to real-time PCR for the detection of related genes transcripts (CYCLIN D1, C-MYC, 1-OCT, and GAPDH). In brief, 2.5 μL of cDNA was placed in 25 μL of reaction volume containing 12.5 μL of Taqman Universal PCR Master Mix, No AmpErase UNG, 8.75 μL water, and 1.25 μL of primers and probe sets. The primers and probes were purchased from Applied Biosystem (ABI; CYCLIN D1: HS9999904, C-MYC: HS9999903, 1-OCT: HS00427552). Primers and probes were designed to span exon-exon boundaries. Amplification was done in an ABI Fast 7500 Real-time PCR system (Applied Biosciences) using the cycling program: 95°C for 10 minutes; 40 cycles of 95°C for 15 seconds, 60°C for 60 seconds. All samples were analyzed in triplicates. DNA contamination was evaluated by performing PCR on the nonreverse-transcribed control of each sample. The relative expression levels of the genes of interest were normalized to endogenous reference glyceraldehyde-3-phosphate dehydrogenase and relative to control sample as a calibrator using formula: 2−ΔΔCT. The threshold cycle reflects the cycle number at which the fluorescence generated within a reaction crosses the threshold.

Animal studies
Three-week-old female severe combined immunodeficient (SCID) mice were purchased from Taconic. The mice were housed five per cage and maintained under specific pathogen-free conditions at the SCID Mouse Barrier Facility at M. D. Anderson. The experimental protocol was reviewed and approved by the M. D. Anderson Institutional Animal Care and Use Committee. Mino cells

Figure 2. Degrasyn inhibits constitutively activated pSTAT3 and NF-κB in MCL cells. A, cell extracts purified from six MCL cell lines (top), six primary MCL cells (middle), and control normal (unstimulated) and activated B cells (CD154 plus IgM; bottom) were subjected to Western blotting for STAT3, pSTAT3 (ser-727), and actin (loading control). B, mino MCL cells were treated with dose dependently (0–5 μmol/L) with degrasyn for 24 h (top) or time-dependent (0–12 h) with 5 μmol/L of degrasyn (bottom). Cells extracts from the cells were purified and analyzed by Western blot for pSTAT3, STAT3, and Actin. C, Western blots from B were stripped and rebotted for plkBα and IkBα. D, mino cells were cotransfected with the 6xNF-κB-luc reporter plasmid and β-gal reporter plasmid, and then treated with degrasyn or bortezomib (BZ), as a control, for the indicated times. Luciferase activity was measured and normalized according to β-gal activity. Columns, mean of triplicate samples of three independent experiments; bars, SD.
(50 × 10⁶) were injected i.p. into the mice using a 27-gauge needle. One week after inoculation, mice were pooled and randomized into four treatment groups (n = 10/group). Group I received a vehicle control (DMSO) in injection saline (100 μL, twice weekly, i.p.); group II received bortezomib alone (0.25 mg/kg, 100 μL, twice weekly, i.p.); group III received degrasyn alone (20 mg/kg, 100 μL, twice weekly, i.p.); and group IV received a combination of bortezomib (0.25 mg/kg) and degrasyn (20 mg/kg; 100 μL, twice weekly, i.p.). Two additional groups (n = 5/group) that were not inoculated with the tumor cells but received treatment similar to that in groups III and IV were included. Treatment was continued for up to 8 weeks, at which time one mouse in each group was selected at random, euthanized, and subjected to necropsy. The remaining mice were monitored closely and euthanized based on the recommendation of the veterinarians. Three different technicians did the treatment for these mice on different schedules.

Statistical analysis
The IC₅₀ values were calculated by using the sigmoidal dose-response curves. Confidence intervals (CI) are shown in between parenthesis. Data from the in vitro assays were analyzed using a t test with a robust variance estimate. All significance testing was done at the P < 0.05 level. For the animal study, Kaplan-Meier survival curves were created for the mice using the GraphPad Prism software program (version 5.0b; GraphPad Software), and pairwise comparison of survival curves and determination of P values were done using the Log-rank (Mantel-Cox) Test.

Results
Growth inhibition of MCL cell lines with the AG490 congeners
Previous studies in nonlymphoid cells showed that treatment with WP1066, a derivative of the classic JAK2-inhibiting compound AG490, inhibited both JAK2 and STAT3 activation, leading to the inhibition of proliferation
and induction of apoptosis in malignant glioma and melanoma cells (25, 26). Upon further screening and synthesis, we discovered that two additional compounds, WP1129 and degrasyn (WP1130), had superior inhibitory activity in the JAK/STAT pathway when compared with the compound WP1066. We examined the antiproliferative effect of these compounds in MCL cell lines (two typical MCL and two BV-MCL lines). We exposed the four cell lines to each compound at increasing concentrations (0–5 μmol/L) and then analyzed them for cell proliferation using thymidine incorporation assays. These studies showed that cell growth suppression was seen in all three compounds for all four cell lines (Fig. 1A), and the half maximal effective concentration (EC_{50}) of degrasyn was lower than that of the other two compounds in all four MCL cell lines (Fig. 1B), suggesting that degrasyn is a more effective antitumor agent in MCL cells.

### Degrasyn inhibits constitutively activated pSTAT3 and NF-κB in MCL cells

To determine the mechanism of growth suppression by degrasyn, we tested whether degrasyn can target STAT3 and NF-κB activation in MCL cells, as previous studies had shown that both STAT3 and NF-κB were constitutively activated in these cells (31, 36). First, we evaluated the expression of pSTAT3 in MCL cell lines and patient MCL cells by Western blotting and showed that pSTAT3 (serine 727) was constitutively expressed in all of the cell lines and patient MCL cells (Fig. 2A). Constitutive pSTAT3 expression in MCL cells can be downregulated by treatment with degrasyn in a dose- and time-dependent manner in Mino MCL cells (Fig. 2B). Interestingly, degrasyn can also downregulate constitutive cytoplasmic pIκBα protein level in MCL cells within 3 hours of treatment, suggesting that degrasyn targets upstream of the NF-kB pathway (Fig. 2C). Expression of transcription factor NF-kB was also assayed using a reporter gene assay, in which nuclear expression of NF-kB directed the induction of firefly luciferase activity. Degrasyn treatments resulted in the reduced expression of luciferase activity in a dose- and time-dependent manner (Fig. 2D). These results suggest that degrasyn can directly or indirectly target both STAT3 and NF-κB pathways in MCL cells.

### Degrasyn interacts with bortezomib to synergize growth inhibition and apoptosis induction in MCL cells

Bortezomib is a proteasome inhibitor approved for the conventional treatment of MCL. To examine the feasibility of combining degrasyn with bortezomib to treat MCL, we performed in vitro and in vivo experiments assessing the tolerability and antitumor efficacy of these agents in MCL. First, we evaluated the effects of degrasyn in combination with bortezomib in MCL cells in vitro. Figure 3A shows the combined effect of degrasyn and bortezomib in the representative MCL and BV-MCL cell lines Mino and Z-138 on cell growth inhibition. Figure 3B shows that a significant decrease in the cell growth of Mino and Z-138 cell lines was observed in response to treatment with combined low doses of degrasyn (1 μmol/L) and bortezomib (10 nmol/L) than with either agent alone, in a time-dependent manner up to 72 hours. We then

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**Combination index**

<table>
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<tr>
<th>Combination index</th>
<th>Synergism</th>
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<td>0.2–0.3</td>
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<tr>
<td>0.3–0.7</td>
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<tr>
<td>0.7–0.85</td>
<td>slight</td>
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<tr>
<td>0.85–1.0</td>
<td>synergy</td>
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<tr>
<td>&gt;1.0</td>
<td>no synergy</td>
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**NOTE:** Combination index <0.2, synergism; 0.2–0.3 very strong; 0.3–0.7, moderate; 0.7–0.85 slight; 0.85–1 synergy; >1.0 no synergy. The drug combination index for bortezomib and degrasyn was calculated by using the CalcuSyn software program. A combination index value of 1 indicated an additive effect of the two drugs. Combination index values of <1 indicated synergies; the lower the value, the stronger the synergy. In contrast, combination index values of >1 indicated antagonism. Abbreviation: CI, combination index.
evaluated the synergistic effect of the two compounds in four representative MCL cell lines. The concentrations of bortezomib and degrasyn that produced the synergistic effects on cell growth inhibition ranged from 2 to 12 nmol/L and from 0.5 to 2.0 μmol/L, respectively, as shown in a combination index for four representative MCL cell lines (Table 1). Figure 3C shows representative isobologram analysis confirming the synergistic effect of degrasyn and bortezomib in Mino and Z-138 cells. These data show synergistic anti-MCL activity of degrasyn plus bortezomib.

Next, we examined whether degrasyn and bortezomib can interact to synergize apoptosis induction using an Annexin V–based method of early apoptosis detection. At a concentration of 10 nmol/L, bortezomib induced apoptosis in ~15% of Mino cells after 48 hours of treatment, whereas at a concentration of 1 μmol/L, degrasyn induced apoptosis in 5% of Mino cells over the same period. However, the combination of bortezomib and degrasyn administered at similar concentrations induced apoptosis in >50% of MCL of Mino and Z-138 cells (Fig. 4A). These results were similar for both typical MCL and BV-MCL cell lines. In addition, we measured caspase-3 activation, a signature event during apoptosis, in MCL cell lines and primary MCL tumor cells after 48 hours of treatment with degrasyn and bortezomib alone or in combination. The results showed that degrasyn interacted with bortezomib to activate caspase-3 in all four MCL cell lines (both typical MCL and BV-MCL) and four primary MCL tumor cells [two typical MCL (pat # 1 and 2) and two BV-MCL (pat # 3 and 4); Fig. 4B]. Moreover, the apoptotic effect of degrasyn plus bortezomib was evidenced by the induction of bcl-2 protein cleavage and bax protein expression (Fig. 4C). These results suggested the existence of potential synergy between degrasyn and bortezomib in targeting MCL cells through cell growth inhibition and apoptosis induction.

Degrasyn and bortezomib synergy is associated with NF-κB and pSTAT3 inhibition, concurrent with cyclin D1 and c-Myc downregulation in MCL cells

Next, we sought to determine the molecular basis for the mechanism of degrasyn and bortezomib synergy in MCL cells by examining the expression of key growth and survival genes in MCL cells. In these experiments,
we treated Mino cells with degrasyn and bortezomib alone or in combination for 24 and 48 hours and then examined the STAT3 and NF-κB activity in the cells. The results showed that the combination of degrasyn and bortezomib effectively downregulated constitutive STAT3 and NF-κB DNA binding activity after 24 hours; the combination significantly inhibited constitutive STAT3 (Fig. 5A) and NF-κB (Fig. 5B) DNA binding activity after 48 hours of treatment. We observed the downregulation of the protein expression (Fig. 5C) as well as mRNA expression (Fig. 5D) of c-Myc and cyclin D1, target genes of STAT3 and NF-κB, in a similar manner. These results suggested that the combination of degrasyn and bortezomib can target multiple growth and survival signaling pathways as well as targeting genes downstream from these pathways in MCL cells.

**Effects of in vitro treatment of normal human lymphocytes with degrasyn and bortezomib**

To better understand the effects of degrasyn and bortezomib in the normal human B-cell lineage, we also evaluated the cytotoxicity of degrasyn alone and in combination with bortezomib in human PBMCs and activated normal B cells in vitro. We exposed purified PBMCs to degrasyn in dose-response studies at increasing concentrations (0–2.5 μmol/L) alone and with 10 or 15 nmol/L bortezomib. Degrasyn alone had a minimal cytotoxic effect on PBMCs; however, with exposure to the combination of degrasyn and bortezomib, the cytotoxicity of degrasyn and bortezomib in PBMCs increased as the concentration of degrasyn reached >1 μmol/L (Supplementary Fig. S1A). We activated purified human B cells with recombinant CD40L, anti-IgM antibody, and interleukin-4 for 24 hours and then exposed the proliferating B-cell blasts to degrasyn alone or in combination with bortezomib in a manner similar to that used with PBMCs. Unlike PBMCs, the majority of the activated B cells responded to treatment with degrasyn preferentially alone as well as in combination with bortezomib (Supplementary Fig. S1B), suggesting that degrasyn targets highly proliferative lymphocytes.

**Antitumor effects of degrasyn in combination with bortezomib in a xenotransplant SCID mouse model of MCL**

To determine whether degrasyn also enhances the antitumor effects of bortezomib in MCL tumor cells in vivo, we used a xenotransplant SCID mouse model of MCL. We i.p. injected young SCID mice with Mino MCL cells and randomized the animals into four treatment groups as indicated in Fig. 6A. One week after tumor-cell inoculation, we began administering a low-dose combination of degrasyn (20 mg/kg) and bortezomib (0.25 mg/kg) treatment to the mice (i.p.) twice weekly for ~8 weeks. The dose and schedule for both drugs were well tolerated and effective in other tumor models (29, 37). On day 65, when tumors began to appear, we randomly selected one mouse from each treatment group for sacrifice and necropsy. As shown in Fig. 6B, both bortezomib and degrasyn alone had some individual therapeutic effects in MCL tumor inhibition, whereas treatment with a combination of both drugs resulted in complete disappearance.
of tumors on gross pathologic inspection. We closely monitored the remaining mice and sacrificed them at the development of high tumor burden/morbidity. We created a survival curve showing that mice given both degrasyn and bortezomib survived longer than did mice given either agent alone ($P < 0.0001$; Fig. 6C). Bortezomib or degrasyn alone showed modest prolongation of survival compared with vehicle alone ($P < 0.0213$ and $P < 0.105$, respectively; Fig. 6C). In addition, normal control mice ($n = 5$) given treatment with degrasyn alone or in combination with bortezomib remained alive and showed no signs of body weight loss (data not shown) or organ (bone marrow, lungs, kidneys, and liver) toxicity 6 months later (Supplementary Fig. S2). These results suggested that the combination of degrasyn and bortezomib has potential therapeutic efficacy against MCL and warrants further experimental and clinical investigation.

Discussion

Intracellular signaling pathways that control cellular growth and survival mechanisms are complex, interactive, and often cross-talk with each other in various types of human hematologic/lymphoid tumors (38, 39). Frequently, however, similar or identical pathways (NF-$\kappa$B, JAK/STAT, AKT/phosphatidylinositol 3-kinase, etc.) are used and constitutively activated in these tumors (31, 40, 41). Targeting these pathways with new therapeutic agents is likely to provide improved treatment outcomes if adequate agent targeting, molecular specificity, and drug delivery can be achieved. In the present study, we showed that the novel small-molecule compound degrasyn targets multiple signaling pathways in MCL cells, and that treating these cells with degrasyn in combination with the proteasome inhibitor
bortezomib not only synergizes MCL cell growth inhibition and apoptosis induction in vitro but also prolongs host survival in a SCID mouse model of MCL in vivo.

MCL, particularly BV-MCL, remains a poor responder to current combination chemotherapy among the various histotypes of NHL-B and represents a continuing therapeutic challenge (3). In some patients with MCL, however, the disease does respond to treatment, sometimes for extended disease-free durations, with dose-intensification, most recently with combination chemotherapies such as hyper-CVAD and R-CHOP, which are often followed by autologous bone marrow transplantation (2, 7). Our goal in these studies was to continue to develop new, increasingly effective targeted approaches to MCL therapy, particularly for BV-MCL. These studies used experimental therapeutic methodologies, including in vitro techniques with representative MCL cell lines and patient tumor samples, as well as more translational in vivo studies with xenotransplant SCID mouse models of MCL.

Bortezomib is a boronic acid–derived effective reversible inhibitor of the 26S proteasome that, surprisingly, has had relatively minor toxic effects (e.g., minor platelet count decreases, neuropathy) in initial clinical trials for MCL (17). These studies suggested that bortezomib likely would be more effective in combination with conventional chemotherapy and possibly even more effective with molecularly targeted novel therapeutic agents, particularly small-molecule inhibitors such as degrasyn. Although degrasyn is structurally related to the classic JAK2 inhibitor AG490, when tested against aggressive NHL-B MCL cells in vitro, we observed that degrasyn was also an effective inhibitor of constitutive NF-κB activation present in typical MCL and BV-MCL cells. The molecular mechanism of how degrasyn downregulates NF-κB activity in MCL cells remains uncertain. However, recent studies have indicated that degrasyn directly targets c-Myc for proteosomal degradation within hours after treatment (28), suggesting that NF-κB signaling may be a downstream target of c-Myc as previously shown in Drosophila flies (42). Another possible mechanism of NF-κB inhibition exerted by degrasyn occurs through the JAK/STAT signaling pathway, as previous studies have shown that STAT3 maintains constitutive NF-κB activity in tumor cells by prolonging NF-κB nuclear retention through acetyltransferase p300–mediated RelA acetylation (41). Therefore, inhibition of STAT3 activity by degrasyn may have affected the status of NF-κB activity in MCL cells, as seen in the present study. Constitutive STAT3 and NF-κB signaling and interaction are highly displayed in murine and human cancers, and their roles are central to tumor cell growth and survival (39, 41, 43). Studies have shown that concurrently targeting both constitutive STAT3 and NF-κB activity in lymphoid tumor cells is effective both in vitro and in vivo (38, 44). The synergy between degrasyn and bortezomib resulted in the downregulation of STAT3 and NF-κB activity in MCL cells may provide a molecular mechanism accounting for their effect in prolonging the survival of MCL xenotransplanted SCID mice treated with these compounds. However, other still unidentified factors may also contribute to the observed synergistic effects of degrasyn and bortezomib on MCL cell growth and survival.

In terms of cell type specificity, degrasyn is more effective in myeloid and lymphoid tumor cells than normal CD34+ hematopoietic precursor cells, dermal fibroblasts, and endothelial cells (29). The present study also showed that degrasyn has minimal cytotoxicity in normal peripheral blood mononuclear lymphocytes but maximal cytotoxicity in targeted highly proliferative B lymphocytes, suggesting that degrasyn is a potential therapeutic drug candidate for aggressive or relapsed/refractory B-cell lymphomas, which very much need effective new therapies. When combined with bortezomib, degrasyn has antitumor activity in SCID mice with MCL but is not toxic in normal healthy mice, making degrasyn a candidate combinatorial therapeutic agent together with bortezomib in MCL therapy.

In summary, we have shown that MCL cells, both typical MCL and BV-MCL, treated with degrasyn in conjunction with bortezomib resulted in synergistic growth inhibition and apoptosis induction in vitro. Apoptosis in these cells was correlated with the downregulation of constitutive NF-κB and pSTAT3 activation, leading to the inhibition of c-Myc and cyclin D, and the upregulation of cleaved-bcl-2 and bax protein expression. In vivo, degrasyn and bortezomib combined to synergistically prevent MCL tumor development in a xenotransplant SCID mouse model. Agents such as degrasyn that can pharmacologically target constitutively expressed key growth/survival signaling pathways and cell cycle regulators, including NF-κB, STAT3, c-Myc, and cyclin D1, in MCL cells may prove to be useful therapeutic agents for MCL when combined with effective proteasome inhibitors such as bortezomib.

Disclosure of Potential Conflicts of Interest

Dr. Waldemar Priebe is the lead inventor in the patent disclosing WP1130 and related analogs and has financial interest in the company that licensed this patent.

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