Selinexor overcomes hypoxia-induced drug resistance in multiple myeloma

Barbara Muz
Washington University School of Medicine in St. Louis
Feda Azab
Washington University School of Medicine in St. Louis
Pilar de la Puente
Washington University School of Medicine in St. Louis
Yosef Landesman
Karyopharm Therapeutics
Abdel Kareem Azab
Washington University School of Medicine in St. Louis

Follow this and additional works at: https://digitalcommons.wustl.edu/open_access_pubs

Recommended Citation
Muz, Barbara; Azab, Feda; de la Puente, Pilar; Landesman, Yosef; and Azab, Abdel Kareem, "Selinexor overcomes hypoxia-induced drug resistance in multiple myeloma." Translational Oncology, 10, 4. 632-640. (2017).
https://digitalcommons.wustl.edu/open_access_pubs/6164

This Open Access Publication is brought to you for free and open access by Digital Commons@Becker. It has been accepted for inclusion in Open Access Publications by an authorized administrator of Digital Commons@Becker. For more information, please contact vanam@wustl.edu.
Selinexor Overcomes Hypoxia-Induced Drug Resistance in Multiple Myeloma

Abstract
Increased levels of the nuclear export protein, exportin 1 (XPO1), were demonstrated in multiple myeloma (MM) patients. Targeting XPO1 with selinexor (the selective inhibitor of nuclear export; SINE compound KPT-330) demonstrates broad antitumor activity also in patient cells resistant to bortezomib; hence, it is a promising target in MM patients. Hypoxia is known to mediate tumor progression and drug resistance (including bortezomib resistance) in MM cells. In this study, we tested the effects of selinexor alone or in combination with bortezomib in normoxia and hypoxia on MM cell survival and apoptosis in vitro and in vivo. In vitro, selinexor alone decreased survival and increased apoptosis, resensitizing MM cells to bortezomib. In vivo, we examined the effects of selinexor alone on tumor initiation and tumor progression, as well as selinexor in combination with bortezomib, on tumor growth in a bortezomib-resistant MM xenograft mouse model. Selinexor, used as a single agent, delayed tumor initiation and tumor progression, prolonging mice survival. In bortezomib-resistant xenografts, selinexor overcame drug resistance, significantly decreasing tumor burden and extending mice survival when combined with bortezomib.

Translational Oncology (2017) 10, 632–640

Introduction
Multiple myeloma (MM) is a plasma cell malignancy localized primarily in the bone marrow (BM). Some of the most novel therapies being developed are based on interrupting the interactions between MM cancer cells and the supportive BM microenvironment, which contributes to pathogenesis and facilitates MM progression [1]. The introduction of these therapeutics, including proteasome inhibitors (such as bortezomib, carfilzomib and ixazomib), immunomodulatory drugs, and monoclonal antibodies, substantially improved the survival of MM patients [1–3]; however, more than 90% of MM patients relapse [1,4]. Bortezomib is the first-in-class reversible proteasome inhibitor which has been used as the backbone of frontline therapy and relapsed/refractory settings in combination regimen in MM patients; nevertheless, its use is limited due to the development of drug resistance and side effects [5].

Exportin 1 (XPO1), also known as chromosome region maintenance 1, is the main nuclear exporter of leucine-rich proteins through the nuclear pore complex to the cytoplasm, which is driven by guanosine triphosphate hydrolysis [6]. This tightly controlled protein shuttling is deregulated in cancer cells, where some of the XPO1 cargo proteins, such as tumor suppressors (p53 [7,8], p21, and p27 [9]) and inhibitor of NFkB [10,11], are abnormally localized in the cytosol and thus cannot properly function in suppressing cancer initiation and progression [9]. Overexpression of XPO1 protein is associated with tumor growth and poor patient survival as demonstrated in solid tumors including lung, pancreatic, cervical, and ovarian cancers [8,12–14]; melanoma; glioma; and osteosarcoma [15,16], as well as in hematologic malignancies including acute myeloid leukemia [17], non-Hodgkin lymphoma [18], and MM [11,19,20]. Interestingly, whole-genome sequencing in chronic lymphocytic leukemia patients identified recurring mutations in the highly conserved region of XPO1, suggesting that these oncogenic changes may contribute to the clinical evolution of the disease [21,22]. Inhibition of the XPO1 protein mediated by selective inhibitor of nuclear export (SINE) compounds, developed by Karyopharm Therapeutics, has
shown broad preclinical antitumor activity in many solid and hematologic malignancies independent of p53 status [15,23–25]. SINE is currently in several phase I/II/III clinical trials including more than 20 studies for myeloma patients, both newly diagnosed and relapsed/refractory, in combination with different treatments.

Tai et al. have shown previously that XPO1 protein expression is higher in MM cells in newly diagnosed patients compared to normal plasma cells [20]. Moreover, XPO1 mRNA expression is augmented with MM disease progression and correlates with bone lytic lesions and poor patient survival. It was reported that XPO1 protein plays a crucial role in MM pathophysiology, malignant growth, and cancer cell survival. These findings are based on XPO1 knockdown studies in MM cells which were also mimicked using SINE (KPT-185, KPT-251, KPT-276, and KPT-330). KPT-185 and KPT-330 were the most potent XPO1 inhibitors, which significantly decreased MM cell proliferation, induced cytotoxicity of MM cells, and increased the percentage of apoptotic MM cells that were associated with inhibited export of tumor suppressors such as p53, p27, PP2A, and FOXO3a. In addition, it was shown that KPT-185 and KPT-330 inhibited osteoclastogenesis and bone resorption via reducing NFκB activity; however, they did not affect osteoblastogenesis in vitro [20].

Therapy resistance acquired by MM cells is one of the main problems in MM treatment and tumor recurrence. Interestingly, XPO1 protein and mRNA levels were demonstrated to be increased during MM progression, and MM cells isolated from patients resistant to bortezomib expressed higher XPO1 protein levels [20]. These results suggest that XPO1 could potentially be a biomarker of cancer progression as well as development of drug resistance in MM. Likewise, tumor hypoxia (low oxygenation) is a phenomenon observed in growing malignant tumors, and hypoxia-induced genes have been used in detecting MM tumor growth and drug resistance [26,27]. We and others have shown that tumor hypoxia develops due to uncontrollable MM cell proliferation and correlates with poor prognosis of cancer patients [28–30]. Hypoxia confers treatment resistance of cancer cells by regulating processes such as quiescence via cell cycle arrest [31–33]; inhibiting apoptosis and senescence of cells; controlling autophagy, endoplasmic reticulum (ER) stress, and p53 and mitochondrial activity [26,32]; and maintaining stem cell–like phenotype through induction of dedifferentiated and immature phenotype of cells [34–37]. We have previously tested the role of hypoxia in MM cell proliferation in the presence of proteasome inhibitors, where MM cells were cultured in hypoxic and normoxic conditions and subsequently treated with or without bortezomib and carfilzomib. We have shown that, in hypoxic conditions, cancer cells lose their sensitivity to proteasome inhibitors in vitro [27].

In this study, we examined whether inhibiting XPO1 with the oral, slowly reversible inhibitor selinexor resensitized hypoxia-induced drug resistance to bortezomib in MM. Indeed, selinexor reversed the hypoxia-induced resistance to bortezomib in MM cells both in vitro and in vivo, thus providing evidence for clinical trials combining proteasome inhibitors with XPO1 inhibition.

**Materials and Methods**

**Cell Culture**

The MM cell lines (MM.1S and MM.1S-GFP-Luc) were a kind gift from Dr. Irene Ghobrial (Dana-Farber Cancer Institute in Boston, MA). Cells were cultured in RPMI-1640 (Corning CellGro, Mediatech, Manassas, VA) supplemented with 10% fetal bovine serum (Gibco, Life Technologies, Grand Island, NY), 2 mmol/l of L-glutamine, 100 U/ml of penicillin, and 100 μg/ml of streptomycin (CellGro, Mediatech, Manassas, VA). Cells were cultured at 37°C and 5% CO2 in a humidified tissue culture incubator (21% O2), in the NuAire water jacket incubator (Plymouth, MN), or in hypoxia (1% O2) using the hypoxic chamber purchased from Coy (Grass Lake, MI).

**Selinexor (KPT-330)**

Selinexor (KPT-330) was provided by Karyopharm Therapeutics (Newton, MA). For in vitro experiments, selinexor was solubilized in dimethylsulfoxide, aliquoted, and stored at −80°C. For in vivo studies, the selinexor formulated drug product contained ~70% active ingredient, ~15% Plasdone PVP K-29/32, and ~15% Poloxamer Pluronic F-68 (provided by Karyopharm Therapeutics) and was stored at 4°C for 7 days. Plasdone PVP K-29/32 with Poloxamer Pluronic F-68 served as a vehicle in the mice studies.

**Effects of Selinexor on MM Cell Survival**

Cell survival was assessed using MTT solution (Sigma-Aldrich, St. Louis, MO), followed by absorbance measurement at 570 nm using a spectrophotometer according to the manufacturers’ protocol, where the absorbance is proportional to the number of viable cells. The effects of increasing concentrations of selinexor (KPT-330: 0, 100, 250, and 500 nM), bortezomib (Selleck Chem, Houston, TX; 10 nM), and combination treatment (100 nM of KPT-330 and 10 nM of bortezomib) were tested on MM cell survival/cytotoxicity under normoxic and hypoxic conditions for 24 hours. In addition, the effect of selinexor (0, 50, 100, and 250 nM) and bortezomib (0, 1, 5, and 10 nM) was examined on MM.1S cell survival in normoxic conditions by MTT.

**Effect of Selinexor on MM Cell Apoptosis**

Cell apoptosis was performed using Annexin V–propidium iodide (PI) staining (BD Biosciences, San Jose, CA) according to the manufacturer’s protocol. Briefly, MM.1S cells (1 × 106 cell/ml) were cultured with selinexor (0, 100, 250, and 500 nM) with or without bortezomib (10 nM) for 24 hours. Then, cells were washed and resuspended in 1× Annexin V binding buffer, followed by Annexin V staining for 15 minutes and PI staining for additional 15 minutes, and analyzed with MACSQuant Flow Cytometer (Miltenyi, San Diego, CA). The results were demonstrated as a frequency (%) of viable (Ann–PI–), early apoptotic (Ann+PI−), and late apoptotic/dead (Ann+PI+) MM cells posttreatment.

**Western Blotting**

To test cell signaling involved in proliferation, apoptosis, and cell cycle, MM cells were first treated with selinexor (0, 100, 250, and 500 nM) with or without bortezomib (10 nM) in normoxia or hypoxia for 24 hours. Cells were collected, washed with 1× PBS, and lysed for 30 minutes on ice using 1× lysis buffer (Cell Signaling, Danvers, MA). Protein concentration was assessed by Quick Start Bradford dye reagent (BioRad, Hercules, CA). Sodium dodecyl sulfate–polyacrylamide gel electrophoresis was performed using NuPAGE 4% to 12% Bis-Tris gels (Novex, Life Technologies, Grand Island, NY) and transferred to a nitrocellulose membrane using iBlot (Invitrogen, Life Technologies). Membranes were blocked with 5% skim milk in Tris-buffered saline (TBS) and incubated with primary antibodies overnight at 4°C for apoptosis signaling (cleaved PARP, pPARP; cleaved caspase-3, cCasp3; cCasp8), proliferation signaling (p-p44/42 MAPK; p-S6R), and cell cycle signaling (p-Rb; CDK-6). β-Tubulin was used as a loading control.
control (antibodies were purchased from Cell Signaling). The membrane was washed with Tris-buffered saline/Tween 20, incubated for 1 hour at room temperature with HRP-conjugated secondary antibody, then washed and developed using Novex ECL Chemiluminescent Substrate Reagent Kit (Invitrogen). The density of bands was quantified using ImageJ Software and normalized to α-Tubulin.

**Animal Studies: Tumor Initiation, Tumor Progression, and Survival Study**

SCID-beige mice (females, 8 weeks old) were obtained from Charles Rivers Laboratories (Wilmington, MD). Approval for these studies was obtained from the Ethical Committee for Animal Experiments at Washington University in St. Louis School of Medicine.

For the tumor initiation study, 10 mice were injected intravenously (IV) with MM.1S-GFP-Luc cells at the concentration of $3 \times 10^6$ cells per mouse and were treated instantaneously with vehicle ($n = 5$) or selinexor ($15 \text{ mg/kg, } n = 5$) administered by oral gavage (PO) three times a week (TTW at day 1, 3, and 5). Mice survival was recorded daily, whereas mice weight and tumor growth using bioluminescent imaging (BLI) were monitored at day 0, 28, and 42.

In the tumor progression study, MM.1S-GFP-Luc cells were injected IV into 16 SCID mice at a concentration of $2 \times 10^6$ cells per mouse and allowed to grow for 3 weeks; the tumor growth was determined by BLI. At week 3, mice were randomly allocated into two groups (eight mice per group) which received vehicle ($n = 8$) or selinexor ($15 \text{ mg/kg, } n = 8$). Vehicle and selinexor were administered by oral gavage three times a week (TIW at day 1, 3, and 5). Survival of mice was monitored every day by the investigator, whereas mice weight and tumor growth (BLI) were examined at day 0, 21, and 28.

To test the effect of selinexor on bortezomib-resistant cancer, first we demonstrated development of xenograft mouse model resistant to bortezomib. MM.1S-GFP-Luc cells were injected IV into 16 SCID mice at a concentration of $2 \times 10^6$ cells per mouse and allowed to grow for 3 weeks when the tumor burden was determined by BLI. Mice with established MM tumors were pretreated with vehicle ($n = 8$) or bortezomib ($n = 8; 1.0 \text{ mg/kg}$) administered intraperitoneally (IP) twice a week (BIW) for 2 weeks. A treatment with bortezomib showed initial response in the first week and then relapsed in week 2, demonstrated as ROI normalized to tumor size before pretreatment (Supplementary Figure 1).

Finally, to test the effect of selinexor on bortezomib-resistant cancer, MM.1S-GFP-Luc cells were injected IV into 24 SCID mice at a concentration of $2 \times 10^6$ cells per mouse and allowed to grow for 3 weeks, confirmed by BLI. Then mice were pretreated with bortezomib (1.0 mg/kg) for the next 2 weeks to induce resistance to bortezomib, after which the mice were randomly allocated (day 14) into 3 groups as follow: (1) bortezomib (0.5 mg/kg; $n = 8$) alone, (2) selinexor (10 mg/kg; $n = 8$) alone, and (3) a combination of selinexor (10 mg/kg) and bortezomib (0.5 mg/kg; $n = 8$). Bortezomib was injected IP twice a week (BIW; day 1 and day 4), and selinexor was administered by oral gavage three times a week (TIW; day 1, 3, and 5) along with vehicle controls. Survival of mice was monitored every day by the investigator, whereas mice weight and tumor growth (BLI) were examined at day 0, 14, 21, 35, and 42.

**Statistical Analysis**

The in vitro experiments were performed in quadruples and replicated independently at least three more times. Results are shown as mean ± S.D., and statistical significance was analyzed using Student’s t test. The in vivo data were analyzed using Student’s t test for statistical significance, and results are depicted as mean ± S.E.M. Variation within each group was equally variant and similar between the groups that were statistically compared. Values were considered significantly different for $P$ values less than .05.

**Results**

**Selinexor Inhibits Survival and Induces Apoptosis in a Dose-Dependent Manner in MM.1S Cells**

Within 24 hours, in vitro studies showed that selinexor dose-dependently decreased the survival of both normoxic and hypoxic MM.1S cells (Figure 1A), whereas it increased the percentage (%) of apoptotic and dead cells in normoxic and hypoxic cells (Figure 1B). The mechanism of selinexor-induced cell apoptosis was based on its effects on the expression of the cell signaling proteins (cleaved PARP, Casp3, and Casp8), which was regulated in a dose-dependent manner in both conditions. On the contrary, phosphorylation of ERK1/2 and S6R (proteins involved in cell proliferation) was decreased with increasing concentrations of selinexor, as demonstrated by the intensity of the bands (Figure 1C) and by quantitative densitometry (Figure 1Ci). The effect of selinexor concentrations was more pronounced in hypoxic than normoxic conditions.

**Selinexor Reverses Hypoxia-Induced Resistance to Bortezomib In Vitro**

We have previously demonstrated the development of drug resistance to proteasome inhibitors in hypoxic conditions in MM cell lines [27]. Using the MTT assay, we showed that the combination of selinexor and bortezomib increased the cytotoxic effect of bortezomib and overcome hypoxia-induced resistance to bortezomib (Figure 2A). Similarly, the apoptosis assay showed that the combination of bortezomib and selinexor increased the % of apoptotic and dead cells compared to bortezomib alone, especially in hypoxia (Figure 2B). Mechanistically, the combination of selinexor with bortezomib induced greater cleavage of PARP, Casp3, and Casp8; decreased phosphorylation of Rb; and downregulated the expression of CDK-6 compared to bortezomib alone; there was almost complete abrogation of the latter two proteins, as demonstrated by the intensity of the bands (Figure 2C) and by quantitative densitometry (Figure 2Ci). These results imply halted cell cycle and hence reduced cell proliferation. We tested the effect of combination of increasing concentrations of bortezomib and selinexor on MM.1S cell survival, and we found that the combination had an additive effect, and no synergy was observed (Figure 2D).

**Selinexor as a Single Agent Delays Tumor Initiation and Improves Mice Survival**

Next, we tested the effects of selinexor (15 mg/kg) as a single agent on tumor initiation in vivo. On the first day following MM.1S-GFP-Luc cell injection, we initiated oral gavage treatment of selinexor three times a week (TTW) (see the schematic diagram in Figure 3A). Selinexor delayed tumor initiation, which was detectable in the vehicle-treated group 28 days after treatment (Figure 3B, $P = .034$). At 42 days after treatment, the tumor burden in the selinexor-treated group was nearly half that of the vehicle-treated group ($P = .078$) (Figure 3B). Moreover, the vehicle-treated mice deteriorated rapidly and stayed on the study until day 53, whereas mice treated with selinexor stayed on the study for 10 more days until day 63 ($P = .007$) (Figure 3C).
Selinexor as a Single Agent Delays Tumor Progression in MM-Bearing Mice

Next, we tested the effects of selinexor (15 mg/kg) as a single agent on tumor progression in MM-bearing mice administered by oral gavage three times a week (TIW). Treatment started 3 weeks post-MM cell injection (see the schematic diagram in Figure 4A). Selinexor significantly delayed tumor progression and reduced tumor burden by approximately half when compared to the vehicle-treated group at 21 and 28 days post-treatment initiation ($P = .032$) (Figure 4B). Moreover, selinexor-treated mice survived significantly longer than vehicle-treated mice; the vehicle-treated group stayed on the study until 36 days after treatment initiation, whereas the selinexor-treated group stayed on the study until 42 days after treatment initiation ($P = .014$) (Figure 4C).

Combination of Selinexor with Bortezomib Delays Tumor Growth and Improves Survival in Bortezomib-Resistant MM-Bearing Mice

First, we emulated MM disease refractory to bortezomib by demonstrating that the initial administration of bortezomib alone in established MM mouse model caused bortezomib resistance. Three weeks post-MM cell injection to mice, mice were treated with vehicle ($n = 8$ mice) or bortezomib ($n = 8$; 1.0 mg/kg). Whereas the MM tumor constantly grew over 2 weeks in the vehicle-treated group, a
treatment with bortezomib showed initial response in the first week and then relapse in week 2 (Supplementary Figure 1). We therefore used this mouse model to examine the effect of selinexor, with and without bortezomib, on tumor growth in bortezomib-resistant MM-bearing mice (see the schematic diagram in Figure 5A). Mice with established tumors were first treated with bortezomib IP two times a week (BIW), for 2 weeks, to induce resistance to bortezomib. Mice were then randomized and treated with bortezomib alone, selinexor alone, or the combination of bortezomib and selinexor. At day 35 (21 days after the randomization) and at day 42 (28 days after the randomization), tumor burden in mice treated with the combination of selinexor and bortezomib was 60% smaller than in mice treated with bortezomib alone ($P = .012$ and .036, respectively) (Figure 5B). Similarly, the survival of mice treated with the combination of selinexor and bortezomib was significantly longer than that of mice treated with bortezomib alone, in which the group treated with bortezomib alone all
died at day 50, whereas 50% of the mice in the group treated with the combination of bortezomib and selinexor were still alive at day 52 ($P = .005$) (Figure 5C).

**Discussion**

Despite the implementation of novel therapies (which include proteasome inhibitors), the majority of MM patients relapse due to the development of drug resistance [38]. Low oxygenation (hypoxia) in the BM microenvironment was shown to occur beyond physiologic BM conditions during the progression of MM and, together with cellular and acellular components, plays a critical role in MM dissemination, stem cell–like properties, and drug resistance [27,30,39]. Our research group reported enhanced MM drug resistance to current therapies, including bortezomib and carfilzomib, in hypoxic MM cells [27,40]. Therefore, targeting hypoxic cells to abolish the drug-resistant cancer population is the vital goal to prevent recurrence in MM patients.

Increased levels of XPO1, the nuclear exporter of tumor suppressors, were observed in MM. Moreover, XPO1 was shown to be upregulated in MM cells from patients who became resistant to bortezomib, suggesting that XPO1 could be involved in drug resistance [20]. Because hypoxia has been known to be involved in developing treatment resistance, we hypothesized that inhibition of XPO1 using the novel, orally available inhibitor selinexor (KPT-330) will reverse the hypoxia-induced bortezomib resistance in MM cells both in vitro and in vivo.

In vitro, we have demonstrated that selinexor as a single agent increased apoptosis and decreased survival of MM cells in both normoxia and hypoxia in a dose-dependent manner. In addition, combination treatment of selinexor and bortezomib demonstrated an additive effect on cell survival, and selinexor overcame hypoxia-induced drug resistance to bortezomib in cell culture. In vivo, the tumor initiation study inhibiting XPO1 with selinexor as a single agent delayed tumor progression, decreased tumor growth, and extended mice survival, indicating that XPO1 may be involved in tumorigenesis. In the tumor progression study on MM-bearing mice, which exemplified
newly diagnosed patients, selinexor used as a single agent significantly reduced tumor burden and prolonged mice survival. These results were in accordance with previously published data showing that XPO1 inhibitors decreased tumor size and increased survival in MM-bearing mice as compared to the vehicle-treated group [11,20].

Next, we have mimicked MM patients with refractory disease utilizing a xenograft mouse model resistant to bortezomib achieved by pretreating MM-bearing mice with bortezomib, where bortezomib-treated mice initially responded to chemotherapy but eventually relapsed. In this study, we used the same method to develop the bortezomib-resistant mouse model by pretreating mice with established tumors with bortezomib before adding selinexor to the treatment. Hence, this model emulates the refractory stage of the disease. These in vivo results demonstrate that the combination of selinexor with low-dose bortezomib in bortezomib-resistant MM-bearing mice significantly decreased tumor growth and extended mice survival.

We suggest that the mechanism of the combined effect of bortezomib and selinexor is related to protein homeostasis that each of these drugs (and hypoxia) affects. Protein homeostasis is controlled by several mechanisms such as protein synthesis, nuclear-cytoplasm protein trafficking, protein folding in the ER, and protein recycling by degradation in the proteasome [41]. By blocking XPO1 with SINE, the nuclear-cytoplasm trafficking of the growth-promoting proteins and oncogenic proteins is hampered, causing cell death [9,10]. Hypoxia was shown to induce inhibition of protein synthesis through the mTOR pathway, as well as induction of protein misfolding in the ER due to the lack of recycling of glutathione for production of S-S bonds, ultimately contributing to the ER stress [42,43]. Bortezomib is a proteasome inhibitor that causes inhibition of protein recycling and accumulation of abnormal and misfolded proteins by preventing their degradation in the proteasome, triggering cell death [5]. Hence, we speculate that subsequently the combination of selinexor with bortezomib, especially
in hypoxic conditions, perturbs intracellular protein homeostasis causing protein overload in different cell compartments, induces cell stress, and enhances anticancer activity. A more thorough investigation of the mechanism is warranted.

In conclusion, we report that XPO1 inhibitor selinexor is an effective antitumor agent and can overcome hypoxia-induced drug resistance; hence, it represents a promising novel therapeutic strategy in treating patients with MM. Selinexor decreased survival and increased apoptosis of MM cells as a single agent in vitro and slowed down tumor initiation and tumor progression in the xenograft MM mouse models, significantly prolonging mice survival. In addition, selinexor dramatically reduced tumor growth when combined with low-dose bortezomib in a xenograft-resistant xenograft model. These results provide further support for current and future clinical trials to treat newly diagnosed as well as relapsed/refractory MM patients with selinexor, especially those who are resistant to bortezomib (and possibly other proteasome inhibitors).

Conflict of Interest
Dr. Azab receives research support from Karyopharm, Verastem, Selectys, CellWorks, Cleave Bioscience, Glycomimetics, Abbvie, and Tioma and is the founder and owner of Targeted Therapeutics LLC and Cellatrix LLC. Yosef Landesman is employed by Karyopharm Therapeutics. Other authors state no conflicts of interest.

Supplementary data to this article can be found online at http://dx.doi.org/10.1016/j.tranon.2017.04.010.

Acknowledgement
The study was supported partially by a grant from Karyopharm Therapeutics and partially by the award from the National Center for Advancing Translational Sciences of the National Institutes of Health (NIH) and the National Cancer Institute of the NIH under U54CA199092. The content is solely the responsibility of the authors and does not necessarily represent the official view of the NIH.

References


