Small-Molecule Inhibitors of USP1 Target ID1 Degradation in Leukemic Cells

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Abstract
Inhibitor of DNA binding 1 (ID1) transcription factor is essential for the proliferation and progression of many cancer types, including leukemia. However, the ID1 protein has not yet been therapeutically targeted in leukemia. ID1 is normally polyubiquitinated and degraded by the proteasome. Recently, it has been shown that USP1, a ubiquitin-specific protease, deubiquitinates ID1 and rescues it from proteasome degradation. Inhibition of USP1 therefore offers a new avenue to target ID1 in cancer. Here, using a ubiquitin-rhodamine-based high-throughput screening, we identified small-molecule inhibitors of USP1 and investigated their therapeutic potential for leukemia. These inhibitors blocked the deubiquitinating enzyme activity of USP1 in vitro in a dose-dependent manner with an IC50 in the high nanomolar range. USP1 inhibitors promoted the degradation of ID1 and, concurrently, inhibited the growth of leukemic cell lines in a dose-dependent manner. A known USP1 inhibitor, pimozide, also promoted ID1 degradation and inhibited growth of leukemic cells. In addition, the growth of primary acute myelogenous leukemia (AML) patient-derived leukemic cells was inhibited by a USP1 inhibitor. Collectively, these results indicate that the novel small-molecule inhibitors of USP1 promote ID1 degradation and are cytotoxic to leukemic cells. The identification of USP1 inhibitors therefore opens up a new approach for leukemia therapy. Mol Cancer Ther; 12(12): 2651–62. ©2013 AACR.

Introduction
ID1 (inhibitor of DNA binding 1) protein is a member of the helix-loop-helix (HLH) family of transcriptional regulatory proteins, which consists of 4 members, ID1–ID4 (1, 2). Basic HLH (bHLH) family of transcription factors are key regulators of cellular development and differentiation. The bHLH proteins (e.g., E2A, E2-2, and HEB) form homodimers or heterodimers with other tissue-specific bHLH proteins (e.g., MyoD and SCL/tal1), bind DNA upon dimerization, and activate transcription of differentiation-associated proteins. The ID family of HLH proteins can also form dimers with bHLH proteins (3). However, the bHLH heterodimers are transcriptionally inactive and are unable to bind to the DNA because ID proteins lack a basic DNA binding domain (2). Accordingly, ID proteins antagonize the functions of bHLH proteins and inhibit the bHLH protein–mediated transactivation of genes that promote cellular differentiation. ID proteins, therefore, are also termed inhibitors of differentiation. In addition to binding to the bHLH proteins, ID proteins can also interact with non-bHLH proteins and antagonize their function (4).

ID1, one of the ID proteins, is known to play a role in cellular transformation. It is overexpressed in a variety of human cancer types, including pancreatic, cervical, ovarian, prostate, breast, colon, and brain cancer (5–11). In most of these cancer types, high ID1 expression correlates with poor prognosis and increased chemoresistance (7, 9, 10, 12, 13). Moreover, functional studies have demonstrated that ID1 can promote cell proliferation, inhibit differentiation, delay cellular senescence in primary human cells, and promote cell migration and the metastatic phenotype of cancers (14). ID1 function is also required for maintaining the “stemness” of many normal and malignant tissues (10, 11, 15).

Recent studies indicate that ID1 is also an activator of leukemic cell growth. First, Tang and colleagues (16) observed high levels of ID1 expression in primary acute myelogenous leukemia (AML) samples, and high expression correlated with poor prognosis. Second, ID1 was identified as a common downstream target of constitutively activated oncogenic tyrosine kinases, such as BCR-ABL and TEL-ABL (17). Third, ID1 can immortalize hematopoietic progenitors in vitro and promote a...
Figure 1. High-throughput screening and identification of USP1/UAF1 inhibitors. A, schematic of the USP1/UAF1 constructs. B, Coomassie Blue staining of the purified USP1/UAF1 complex. C, a schematic of the Ub-rhodamine based screening assay. D, chemical structure of a parental USP1 inhibitor compound 527 (C527). E, C527 inhibits USP1 activity in a time-dependent manner. Purified USP1/UAF1 complex was incubated with 1 μmol/L C527 or DMSO for the indicated time, followed by the addition of Ub-AMC at a 0.5 μmol/L final concentration. Relative fluorescence units (RFU) at 535 nanometers was measured to indicate the enzymatic activity of USP1. F, Dose-dependent inhibition of USP1 enzymatic activity by USP1 inhibitor C527. Purified USP1/UAF1 complex was incubated with C527 or DMSO for 3 hours followed by the addition of Ub-AMC at 0.5 μmol/L concentration. (Continued on the following page.)
Targeting ID1 in Leukemia by USP1 Inhibitor

myeloproliferative disease in mice in vivo (18). Moreover, the knockdown of ID1 expression inhibited leukemic cell growth (18). Collectively, these observations suggest that ID1 is a prime therapeutic target for leukemia and other cancer types. However, suitable drugs to therapeutically target ID1 have not been developed to date (14). Protein–protein interactions in the nucleus, such as the interaction of ID1 with HLH factors, are notoriously difficult to inhibit with small molecules (19).

A recent report offers an alternative strategy for knocking down the ID1 protein, namely, through inhibition of the ubiquitin-specific protease, USP1 (20). USP1 is a deubiquitinating (DUB) enzyme, which removes polyubiquitin chains from the ID1 protein (20). ID1 is normally polyubiquitinated and rapidly degraded by the proteasome (21–23). USP1 removes the polyubiquitin chains and rescues ID1 from degradation. Selectively knocking down USP1 using short hairpin RNA (shRNA) results in the rapid degradation of ID1 in osteosarcoma cells. Importantly, USP1 knockdown results in decreased mesenchymal cell proliferation and enhanced differentiation of osteosarcoma cells that overexpress USP1 and ID1 (20), providing a rationale for differentiation therapy of many cancer types including leukemia (e.g., AML), chronic myelogenous leukemia (CML). We, therefore, reasoned that pharmacologic inhibition of USP1 would promote ubiquitin-mediated degradation of the ID1 protein, resulting in differentiation and growth inhibition of immature leukemic cells. Our laboratory has previously shown that human USP1 forms a stable complex with its binding partner, USP1-associated factor 1 (UAF1; ref. 24). USP1 by itself exhibits low DUB activity; however, this activity is significantly enhanced when bound as a USP1/UAF1 complex. Using high-throughput screening, we identified a small-molecule inhibitor of the USP1/UAF1 complex. We describe here a novel small molecule (C527), and multiple derivatives, that inhibit USP1 catalytic activity, promote ID1 degradation, and inhibit leukemic cell growth.

Materials and Methods

High-throughput screening

The USP1/UAF1 complex was prepared as described previously (ref. 24; Fig. 1) and the protein complex was used for high-throughput screening. The fluorogenic ubiquitin-rhodamine (Ub-rhodamine)-based enzyme assay was established in a 384-well format for high-throughput screening. The reaction buffer containing free ubiquitin and USP1/UAF1 enzyme complex was added in 384-well plates using automated liquid handling robot-BioTek Microfill (BioTek Instruments Inc.), followed by the addition of the compounds (in dimethyl sulfoxide, DMSO) from the compound library plates to wells using a pin transfer robotic system at a final concentration of 10 μmol/L. The reactions were then incubated for 30 minutes at room temperature, followed by the addition of Ub-rhodamine to initiate the reactions. The enzyme activity of the USP1/UAF1 complex was determined by measuring the fluorescence of Ub-rhodamine. Of note, 150,000 compounds were screened from the library plates at the Partners Center for Drug Discovery (Cambridge, MA). Details of the screen are provided in the Supplementary Methods.

In vitro deubiquitination assays

Purified USP5 enzyme was purchased from Boston Biochem. UCH-L1 and UCH-L3 were as reported previously (25). USP12/46 was prepared in our laboratory as described previously (26, 27). The in vitro enzymatic assays were performed as described previously (24) using ubiquitin-AMC (Ub-7-amido-4methylcoumarin; Boston Biochem) as a substrate in a reaction buffer containing 20 mmol/L HEPES-KOH (pH 7.8), 20 mmol/L NaCl, 0.1 mg/mL ovalbumin, 0.5 mmol/L EDTA, and 10 mmol/L dithiothreitol. The fluorescence was measured by FLUostar Galaxy Fluorometer (BMG LABTECH). For the Ub-vinylsulfone (VS) assay, the proteins were incubated with Ub-VS (Boston Biochem) at 0.5 μmol/L final concentration for 45 minutes at 30°C, followed by the immunoblotting.

Cells and drug treatments

Leukemic cell lines were grown in RPMI-1640 medium (Invitogen) supplemented with 10% FBS (Invitrogen) and penicillin/streptomycin (Invitrogen). HeLa cells and U2OS cells were grown in Dulbecco’s Modified Eagle Medium (Invitogen) supplemented with 10% FBS (Invitrogen) and penicillin/streptomycin (Invitrogen). K562 cells were provided by Dr. Dipanjan Chowdhury (Dana-Farber Cancer Institute, DFCI, Boston, MA). MOLM14 cells were provided by Dr. A. Thomas Look (DFCI). Luciferase-expressing AML cell lines were provided by Dr. Andrew Kung (DFCI). Mouse osteosarcoma cells were provided by Dr. Stuart Orkin (DFCI). The cells were provided within the last 24 months. No authentication of the cell lines was done by the authors. USP1 inhibitor C527 and its derivatives were synthesized and the purity was validated by high-performance liquid chromatography. Pimozide was purchased from Sigma. Primary human patient samples with AML were collected from DFCI leukemia program under the approval of the appropriate protocols. Cells were treated with DMSO or USP1 inhibitors in appropriate medium for 24 to 72 hours. The viable cell counts were determined using a Countess automated cell counter (Invitrogen). Cell death was measured by Trypan blue exclusion (0.4% Trypan blue in PBS) for HeLa cells and MOLM14 cells or by Annexin V-FITC (BD Biosciences) or PI (BD Biosciences) for K562 cells and U2OS cells. Colony assay was performed as described previously (24) using methotrexate (0.2 μmol/L; Sigma) as a selective agent.
using Trypan Blue staining, CellTiter-Glo reagent (Promega) or MTT assay. The apoptotic cells were detected using Annexin V and 7-amino-actinomycin D (7-AAD; BD Biosciences) staining according to the manufacturer’s instructions using flow cytometry. For benzidine staining, the cells were washed twice with PBS and resuspended in 45 μL of PBS + 5 μL of benzidine stain solution (0.2% in 0.5 M glacial acetic acid, 3% H2O2). After 45 minutes incubation at room temperature, the benzidine-positive cells were detected by light microscopy.

**Antibodies and shRNAs**

Western blotting was performed with the following antibodies: anti-USP1 (A301-699A; Bethyl lab), anti-ID1 (B-8; Santa Cruz Biotechnology), antivinculin (H-10; Santa Cruz Biotechnology), anti-α tubulin (DM1A; Santa Cruz Biotechnology), anti-ID2 (C-20; Santa Cruz Biotechnology), anti-ID3 (C-20; Santa Cruz Biotechnology), anti-FANCD2 (H-30; Santa Cruz Biotechnology), and anti-FANC1 (28). Doxycycline-inducible shRNAs in TRIPZ Lentivirus vectors against human USP1 were purchased from Open Biosystems (Thermo Scientific). Mature antisense sequence for the shRNAs were (i) V2THS_171887: TTATCTTCTCTACAATC, (ii) V2THS_218649: TTAA-GATAGCAAGTATTGC, and (iii) V2THS_171886:TAA-GATAGCAAGTATTGC.

**Mouse xenograft experiments**

Six- to 8-week-old athymic nude mice were used for xenograft studies. The mouse experiments were reviewed and approved by DFCI’s Animal Care and Use Committee. A total of 5 × 10⁸ K562 cells were mixed with Matrigel (BD Biosciences; 1:1 ratio) and injected subcutaneously into the flanks of nude mice. When the tumors reached 2 mm³ size, 15 mg/kg pimozone (dissolved in 2% DMSO + 0.3% tartric acid) or vehicle was administered daily. The tumor growth was monitored by measuring the size.

**Homologous recombination analysis and immunofluorescence**

Homologous recombination activity was analyzed by DRGFP reporter as previously described (29). U2OS-DRGFP cells carrying a chromosomally integrated single copy of homologous recombination repair substrate were used. Double-strand break (DSB)–induced homologous recombination in these cells results in restoration and expression of GFP. Briefly, 24 hours after induction of chromosomal DSBs through the expression of I-SceI, U2OS-DRGFP cells were treated with USP1 inhibitor C527 for 24 hours. Cells were then subjected to fluorescence-activated cell sorting analysis to quantify the percentage of viable GFP-positive cells. The Rad51 foci in cells were detected by immunofluorescence as described previously (30) using anti-RAD51 (Santa Cruz Biotechnology) and Alexa Fluor 488–conjugated secondary antibodies. The quantification of cells with Rad51 foci was performed by counting the number of cells with RAD51 foci.

**Results**

**Identification of a novel USP1 inhibitor using high-throughput screening**

We have previously reported that USP1 by itself exhibits poor DUB activity. However, when bound to its interacting partner, UAF1, the catalytic activity of USP1 is significantly enhanced (24). We therefore used a purified USP1/UAF1 complex and performed a high-throughput screening to identify a USP1 inhibitor (Fig. 1). We used the cDNA encoding a full-length USP1 polypeptide, which had a substitution of GG to AA at amino acid residues 670 to 671 (Fig. 1A). We have previously shown that this amino acid change disrupts an autocleavage site in USP1 but does not alter the interaction of USP1 with UAF1 or the DUB activity of the complex (24, 31). Human USP1 (with GG to AA mutation) and UAF1 proteins were coexpressed in SF9 insect cells to generate a protein complex with elevated DUB activity. The USP1/UAF1 protein complex was purified and the purity was confirmed by coomassie staining (Fig. 1B). The stable USP1/UAF1 enzyme complex was then subjected to a high-throughput inhibitor screen, described schematically in Fig. 1C. A small molecule, C527 (heterocyclic tricyclic 1,4-dihydro-1,4-dioxo-1H-naphthalene; Fig. 1D) was identified, which inhibited the USP1/UAF1 complex.

**USP1 inhibitor C527 has a high affinity for USP1**

We next confirmed the ability of C527 to inhibit the USP1/UAF1 complex in a dose-dependent and time-dependent manner, using Ub-AMC as a substrate (Fig. 1E–G). Pretreatment of USP1/UAF1 with C527 resulted in inhibition of its enzyme activity with an IC₅₀ of 0.88 ± 0.07 μmol/L. To determine the specificity of C527 in USP1/UAF1 inhibition, we next examined its ability to inhibit other DUB enzymes in vitro. UAF1 stimulates not only USP1 by itself but also two other DUB enzymes, USP12 and USP46 (26). We therefore tested C527 for its ability to inhibit the purified USP12/USP46 complex or other DUBs. C527 inhibited the DUB activity of the USP12/USP46 complex and other DUB enzymes in vitro. However, the IC₅₀ of C527 for these DUB enzymes was higher in comparison with USP1/UAF1 complex (Table 1). C527 had considerably less inhibitory effect on UCH-L1 and UCH-L3, a different subclass of DUB enzymes, referred to as the

<table>
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<tr>
<th>DUBs</th>
<th>IC₅₀ (μmol/L)</th>
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<tr>
<td>USP1</td>
<td>0.88 ± 0.07</td>
</tr>
<tr>
<td>USP12/46</td>
<td>5.97 ± 0.37</td>
</tr>
<tr>
<td>USP5</td>
<td>1.65 ± 0.42</td>
</tr>
<tr>
<td>UCH-L1</td>
<td>&gt;10</td>
</tr>
<tr>
<td>UCH-L3</td>
<td>2.18 ± 0.69</td>
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**Table 1. IC₅₀ values of USP1 inhibitor C527 for the indicated DUB enzymes**
ubiquitin C-terminal hydrolases (25), even though they are also cysteine proteases. Taken together, C527 is a pan-DUB enzyme inhibitor in vitro, with a high nanomolar IC₅₀ for the USP1/UAF1 complex.

To generate a USP1 inhibitor with improved cellular potency, we chemically synthesized several analogs/derivatives of C527 and determined their IC₅₀ for USP1/UAF1 inhibition (data not shown). We then selected a derivative, SJB2-043 (IC₅₀ = 0.544 μmol/L), for further studies due to the feasibility of its large-scale chemical synthesis.

**USP1 inhibitor SJB2-043 inhibits the enzyme activity of native USP1**

We next determined whether SJB2-043 inhibits the DUB activity of native USP1 complexes isolated from human cells (Fig. 1H and I). For this purpose, we performed competition assays between SJB2-043 and the ubiquitin active site probe HA-Ub-vinyl sulfone (HA-Ub-VS) reagent (Fig. 1H). This affinity reagent covalently modifies and traps active DUB enzymes in cell extracts (32). Cells were treated with SJB2-043 for 24 hours and cell extracts were incubated with HA-Ub-VS, followed by immunoblotting with anti-USP1 or anti-HA antibodies. Consistent with our previous studies and the most recent studies by Piatkov and colleagues (24, 31, 33), USP1 in the untreated cells was detected on Western blot complex as a doublet; full-length USP1 and an autocleaved USP1 with only N-terminal (Fig. 1I, lane 1). The Ub–USP1 conjugates were generated in the untreated cell lysates and were detected by an increase in molecular weight of USP1 (Fig. 1I, lane 2). However, the formation of the Ub–USP1 conjugate was inhibited by the addition of SJB2-043 (lane 4). Ubiquitin-aldehyde (Ub-aldehyde), a potent known nonspecific DUB inhibitor, caused a complete loss of USP1–Ub-VS conjugates in this competition assay (lane 3). SJB2-043 inhibited the Ub-VS labeling of a limited number of endogenous DUB enzymes (Fig. 1I, anti-HA blot). In addition, SJB2-043 inhibited the labeling of USP1 with Ub-VS in a dose-dependent manner (Supplementary Fig. S1A). Taken together, these data demonstrate that SJB2-043 is an inhibitor of the native USP1/UAF1 complex.

**Novel USP1 inhibitors promote degradation of ID1 in leukemic cells**

shRNA knockdown of USP1 results in enhanced proteasome-mediated degradation of ID1 in osteosarcoma cells (20). ID1 degradation in osteosarcoma cells results in mesenchymal differentiation and decreased cell proliferation. We therefore tested whether small molecule USP1 inhibitors enhance ID1 degradation in cells. Indeed, USP1 inhibitor C527 promoted the dose-dependent degradation of ID1 in human U2OS osteosarcoma cells (data not shown). We next tested USP1 inhibitors for their ability to inhibit USP1 in various cell lines. The K562 cell line (the erythroleukemia cell line derived from a patient with CML) was treated with SJB2-043 (Fig. 2A) for 24 hours, and USP1 and ID1 protein levels were determined by Western blotting. Interestingly, the SJB2-043 caused a dose-dependent decrease in USP1 levels and a concomitant degradation of the ID1 protein in the K562 cells at a micromolar drug concentration (Fig. 2B). The decrease in USP1 and ID1 levels by SJB2-043 exposure resulted from proteasomal degradation because the proteasome inhibitor, MG-132, restored the levels of both proteins (Fig. 2C and Supplementary Fig. S1B).

SJB2-043 also caused a decrease in the levels of other ID proteins, namely ID2 and ID3 in K562 cells (Supplementary Fig. S1C). To determine whether the decrease in ID1 levels was secondary to a decrease in USP1 levels, we next generated a K562 stably transfected with a cDNA encoding doxycycline-induced shRNA to USP1. As expected, doxycycline exposure of these cells resulted in a dose-dependent decrease in USP1 expression (Supplementary Fig. S2A). Moreover, depletion of USP1 by siRNA promoted ID1 degradation in K562 cells (Supplementary Fig. S2B). Importantly, siRNA-mediated knockdown of USP1 in leukemic K562 cells resulted in growth inhibition, increased apoptosis and cell-cycle arrest (Supplementary Fig. S2C–S2F). Collectively, these results indicated that although SJB2-043 may have off-target effects, it exhibits its activity in K562 cells at least in part by promoting degradation of USP1, which results in ID1–ID3 degradation.

**USP1 inhibitors are cytotoxic to leukemic cells**

We next determined whether the USP1 inhibitors are cytotoxic to leukemic cells and whether the cytotoxicity results from ID1 degradation. K562 leukemic cells were treated with SJB2-043, and survival and differentiation were assessed. SJB2-043 caused a dose-dependent decrease in the number of viable K562 cells, with an EC₅₀ of approximately 1.07 μmol/L ± 0.08 (95% confidence interval; Fig. 2D). Moreover, SJB2-043 induced apoptosis of K562 cells in a dose-dependent manner (Fig. 2E). Cytotoxic drug concentrations correlated with the concentrations required for ID1 degradation. In addition, low-dose treatments with SJB2-043 inhibited differentiation of K562 cells into hemoglobin-expressing erythroid cells, as detected by benzidine staining (Fig. 2F).

In addition to the CML cells (K562 cells), we also evaluated the cytotoxicity of SJB2-043 in multiple AML cell lines. We chose AML cell lines that were engineered to express the luciferase enzyme so that these cell lines can be used in future for in vivo optical imaging experiments. The AML cell lines, namely, MOLM14, OCI-AML3, U937, and SK-N01 (Fig. 3) expressed ID1 proteins. Upon treatment with SJB2-043, a dose-dependent cytotoxicity and concomitant ID1 degradation was observed in all of the AML cell lines (Fig. 3) further suggesting the role of ID1 in leukemic cell survival.

To determine whether the cytotoxicity of the USP1 inhibitor was due to USP1/ID1 degradation, we next evaluated analogs of SJB2-043 in leukemic cells (Supplementary Fig. S3). The USP1 inhibitor analogs (e.g., SJB2-127) with no significant activity on USP1 (IC₅₀
> 10 μmol/L) or ID1 degradation at concentrations up to 10 μmol/L, did not exhibit cytotoxic effects on leukemic cells (Supplementary Fig. S3A). The analog SJB2-109 had comparable biochemical inhibitory activity (IC_{50} = 0.416 μmol/L) to our lead inhibitor SJB2-043 and exhibits comparable effects on ID1 degradation and leukemic cell survival. Importantly, a more potent analog SJB3-019A (IC_{50} = 0.0781 μmol/L) was five times more potent than SJB2-043 in promoting ID1 degradation and cytotoxicity in K562 cells.

To determine whether ID1 degradation resulted from USP1 inhibition and was not secondary to apoptosis, we next analyzed USP1/ID1 levels and apoptosis after treating K562 cells with SJB2-043 along with a caspase inhibitor in a time-course manner. USP1/ID1 degradation was observed as early as 7 hours after the 2 μmol/L of SJB2-043 exposure when apoptosis was not significant (Supplementary Fig. S4A). At 18 or 26 hours (Supplementary Fig. S4B and S4C) after the SJB2-043 exposure (2 μmol/L), complete degradation of USP1/ID1 was observed with concurrent robust increase in apoptosis. As expected, caspase inhibitor treatment resulted in decreased apoptosis and inhibition of caspase activity (Supplementary Fig. S4). However, ID1 degradation was not affected by caspase inhibition (Supplementary Fig. S4, left). Even when the caspase activity was significantly abrogated with caspase inhibitor, the apoptosis was not completely blocked. These results suggest that ID1 degradation by SJB2-043 may not be occurring as a result of induction of apoptosis, although apoptosis may play a role in destabilizing ID1 independent of USP1 inhibition.

Collectively, these results indicated that novel USP1 inhibitors promote ID1 degradation by specifically...
targeting native USP1 and cause cytotoxicity in multiple leukemic cell lines.

Pimozide, a known USP1 inhibitor, promotes ID1 degradation and inhibits leukemic cell growth

A known drug, pimozide, is a weak USP1 noncompetitive inhibitor (34). Pimozide may also inhibit STAT5 activation, thereby reducing the survival of BCR/ABL–driven CML cells (35). We examined pimozide for its possible cytotoxic effects on K562 cells (Fig. 4) via destabilization of ID1. Interestingly, pimozide promoted ID1 degradation, decreased USP1 levels, and caused a dose-dependent inhibition of K562 cell growth in vitro with induction of apoptosis (Fig. 4B–D), though at a higher drug concentrations as compared with SJ2-043. The inhibitory drug concentrations correlated with the concentrations required for ID1 degradation. We next examined the effect of pimozide in a mouse xenograft model of leukemia. K562 cells were injected into nude mice and xenografts were established followed by the treatment with pimozide or vehicle. As shown in Fig. 4E, pimozide treatment caused a modest inhibition of tumor growth. Taken together, pimozide, a commercially available weak USP1 inhibitor, promotes ID1 degradation and inhibits leukemic cell growth in vitro and in vivo, suggesting that our more potent USP1 inhibitors may have a stronger antileukemic effect.

USP1 inhibitor promotes ID1 degradation and inhibits growth of primary AML cells from patients

Human AML precursor cells arise from leukemic stem cells (LSC), and LSC differentiation is controlled through coordinate transcriptional regulation including the bHLH proteins. We therefore, tested whether the USP1 inhibitor also promotes ID1 degradation and cytotoxicity in primary human leukemic cells. Primary fresh bone marrow and peripheral blood samples were acquired from patients with AML and mononuclear cells were harvested. The cells were then exposed to the USP1 inhibitor for 48 to 72 hours in culture and ID1 levels as well as cell viability was determined. USP1 inhibitor SJ2-043 exhibited dose-dependent cytotoxicity on the primary bone marrow as well as peripheral blood from patients with AML and mononuclear cells were harvested. The cells were then exposed to the USP1 inhibitor for 48 to 72 hours in culture and ID1 levels as well as cell viability was determined. USP1 inhibitor SJ2-043 exhibited dose-dependent cytotoxicity on the primary bone marrow as well as peripheral blood from patients with AML (Fig. 5A and B). Moreover, SJ2-043 treatment caused inactivation of USP1 and promoted ID1 degradation in primary AML cells (Fig. 5C). The ID1 protein was detected as a doublet, which is likely due to the posttranslational modification of the protein. These results demonstrate that USP1 inhibitor SJ2-043 inactivates native USP1 in primary AML cells and exhibits cytotoxicity. The selectivity of SJ2-043 against normal human hematopoietic cells was then
examined in comparison with pimozide using cord blood CD34<sup>+</sup> cells. Although pimozide was tolerated at high micromolar doses, low micromolar levels of SJB2-043 did result in growth inhibition of primary human cord blood CD34<sup>+</sup> cells (Supplementary Fig. S5).

**USP1 inhibitor increases the levels of Ub-FANCD2 and disrupts the homologous recombination**

Previous studies indicate that USP1 has other substrates besides ID1 (36). For instance, the USP1/UAF1 complex deubiquitinates the cellular monoubiquitinated substrates

![Figure 5](image-url)
Ub-FANCD2 and Ub-PCNA (31, 37). Deubiquitination of FANCD2 is a critical step in the Fanconi anemia (FA)/BRCA DNA repair pathway (38, 39). Knockout or knock-down of USP1/UAF1 results in cellular hypersensitivity to DNA cross-linking chemotherapy agents such as mitomycin C (40, 41). Murine knockout of USP1 results in cellular increase in Ub-FANCD2 and a mouse phenotype resembling Fanconi anemia (41). Knockout of USP1 also results in elevated levels of Ub-PCNA, leading to an increase in DNA translesion synthesis (31).

We next determined whether the USP1 inhibitors can alter cellular levels of Ub-FANCD2 and Ub-FANCI, a binding partner of FANCD2 (Fig. 6). HeLa cells were treated with USP1 inhibitor C527 and levels of FANCD2

Figure 6. USP1 inhibitor increases the levels of Ub-FANCD2, decreases the homologous recombination activity, and sensitizes the cells to chemotherapy agents. A, Western blot analysis of the lysates from HeLa cells treated with DMSO or USP1 inhibitor C527 for 8 hours. B, increased cytotoxicity of C527 on HeLa cells in presence of chemotherapy agents. Cells were treated with C527 (1 μmol/L) for 24 hours, followed by the treatment with mitomycin C (MMC; 0.25 μmol/L), camptothecin (CPT; 0.1 μmol/L), or etoposide (0.5 μmol/L) for 4 days, and cell survival was determined. C, C527 inhibits DRGFP reporter for homologous recombination repair activity. U2OS-DRGFP cells were transfected with I-SCE-I reporter plasmid and then exposed to C527 at the indicated concentration for 24 hours. Cells were then subjected to flow cytometry analysis. The percentage of GFP-positive cells was normalized by solvent vehicle treated group are shown. D, C527 inhibits camptothecin (CPT)-induced RAD51 foci formation. HeLa cells were pretreated with DMSO or C527 at the indicated concentration and then exposed to camptothecin for 1 hour. RAD51 foci were detected using immunofluorescence. Data are represented as mean ± SD from three independent experiments.
and FANCI were determined by Western blotting of the cell lysates. C527 treatments caused an increase in the levels of Ub-FANCD2 and Ub-FANCI (Fig. 6A). SJBJ-043 (another USP1 inhibitor) also increased the level of Ub-FANCD2 (Supplementary Fig. S6A and S6B). An increase in Ub-PCNA was also observed in cells exposed to SJBJ-043 (Supplementary Fig. S6C), suggesting that besides FANCD2, PCNA is also affected by USP1 inhibition. HeLa cells were next pretreated with C527 followed by the treatment with DNA damaging agents including mitomycin C, a DNA cross-linking agent. Pretreatment of cells with the USP1 inhibitor caused an enhancement in the cytotoxicity of mitomycin C and camptothecin (Fig. 6B).

Because disruption of USP1 activity results in decreased homologous recombination (41), we next used a gene conversion assay to examine the effect of the USP1 inhibitor on cellular homologous recombination activity. Interestingly, C527 treatments caused a dose-dependent decrease in gene conversion (Fig. 6C), based on the measurement of cellular GFP in this assay. We also confirmed homologous recombination defect by evaluating Rad51 foci formation, another surrogate marker of cellular homologous recombination activity (Fig. 6D). C527-inhibited camptothecin induced the Rad51 foci in HeLa cells. The more active USP1 inhibitor SJBJ-09A also increased the levels of Ub-FANCD2 and Ub-PCNA, and decreased the homologous recombination activity (Supplementary Fig. S6D–S6F). Taken together, these results suggest that USP1 inhibitor treatments lead to an increase in ubiquitinated forms of FANCD2 and FANCI, cause a decrease in homologous recombination activity, and sensitize cells to DNA damaging agents.

Discussion

ID1 is essential for proliferation and maintenance of many cancer types (14, 42). However, it has not yet been therapeutically targeted with a small molecule in leukemia. Here, we demonstrate that targeting USP1 with a small molecule can promote ID1 degradation and cell death in multiple leukemic cell lines. Our data demonstrate that ID1 can be targeted even in primary human leukemic cells using novel inhibitors of USP1.

The drug of choice for CML is imatinib, although drug resistance remains a problem. Novel therapeutics are also needed for the cure of AML, the most common acute leukemia of adults. USP1 inhibitors represent a novel class of drugs for these cancers. USP1 inhibitors can degrade ID1, promote differentiation, and induce apoptosis in these types of leukemic cells and can be potential therapeutic agents. ID1 blocks cellular differentiation at least in part, through its repressive effect on the expression of cell-cycle inhibitor p21 (20). Consistent with this previous report (20), our USP1 inhibitor C527 promotes the degradation of ID1 and the concurrent upregulation of p21 in mouse osteosarcoma cells (Supplementary Fig. S6B). Cell-cycle arrest through p21 upregulation provides a mechanistic explanation for the increase in erythroid differentiation of leukemic cells. ID proteins have central role in keeping cells in an immature state. Various reports suggest that ID proteins including ID1 play an important role in maintaining cancer-initiating cells in a variety of solid tumors (10, 19, 43, 44). Therefore, therapeutically targeting ID1 may also eradicate leukemic cancer stem cells.

An inhibitor of USP1, pimozide, was recently identified (34). Pimozide, an antipsychotic drug, is currently U.S. Food and Drug Administration approved for treatment of Tourette syndrome. However, its efficacy in promoting ID1 degradation has not been previously reported. In our studies, pimozide promotes ID1 degradation and causes cytotoxicity in leukemic cells. Our in vivo studies using pimozide showed a modest decrease in tumor volume in xenograft models of K562 leukemic cells. This is consistent with the recent report by Nelson and colleagues (45), who observed a modest effect of pimozide in FLT3-AML–driven tumor models. Of note, the EC50 of pimozide on leukemia cells is significantly higher than our novel USP1 inhibitors. Even though pimozide is currently used in humans for the treatment of Tourette syndrome, its efficacy in mouse tumor models is limited because of its toxicity. Accordingly, at the maximum tolerated doses, pimozide may not be an effective anticancer agent, at least in leukemic mouse models. Moreover, in addition to exhibiting an inhibitory activity to USP1, it also abrogates STAT5 activity in leukemic cells (35, 45).

ID1 plays an important role in invasiveness of many cancer types including breast and brain tumors (11, 46, 47). Specifically, ID1 depletion by antisense oligos to ID1 in BCR-ABL–bearing 32D3 cells significantly reduced invasive cell migration in vitro and in vivo severe combined immunodeficient mouse models via MMP9 axis (48). It remains to be investigated whether our small-molecule inhibitors of USP1 can reduce the invasiveness of leukemic cells. Although ID1 has been previously targeted using peptidyl-conjugated antisense oligomers (49), peptide aptamers (50), or cannabidiol (11) as potential ways to inhibit ID1 function in solid tumors, none of these applications have been explored for leukemia.

The USP1 inhibitors reported in this study may have other advantages, because USP1 has other known substrates (31, 37). We have previously shown that USP1, in complex with its stimulatory binding partner, UAF1, deubiquitinates the Fanconi anemia proteins, FANCD2 and FANCI (24). Knockdown of USP1 activity results in elevated cellular levels of Ub-FANCD2, resulting in disruption of the Fanconi anemia pathway (40, 41). Functional Fanconi anemia DNA repair pathway is required for cellular resistance to RNA cross-linking agents and Fanconi anemia pathway deficient cells including USP1−/− cells are hypersensitive to DNA cross-linking agents (39–41). USP1 inhibitors may therefore block the Fanconi anemia pathway, thereby promoting cellular hypersensitivity to DNA cross-linking agents and chromosome instability. In the short run, USP1 inhibitors may therefore have an additional useful function as anticancer agents, by sensitizing tumors to conventional chemotherapeutic
agents, for example, cisplatin (34). In our study, USP1 inhibitors increased the levels of Ub-FANCDD2, decreased the homologous recombination activity, and sensitized cells to chemotherapy agents’ mitomycin C and camptothecin. The increase in Ub-FANCDD2 may result directly from USP1 inhibition or indirectly from the DNA damage incurred by USP1 inhibitor exposure. In either case, USP1 inhibition may have additional anticancer functions that extend beyond the promotion of cellular differentiation. Accordingly, the inhibitors may be useful as radiation or cisplatin sensitizers, based on their ability to disrupt the FA/BRCAl or translesion synthesis pathways.

In summary, we have identified novel small-molecule inhibitors of USP1 that exhibit nanomolar IC50 for USP1 activity. These inhibitors also promote ID1 degradation and cause cytotoxicity in a variety of leukemia cell lines and primary leukemic cells. Further preclinical assessment of these small molecules in mouse models will determine their efficacy as novel anticancer agents. If successful, these newly identified USP1 inhibitors can have broader implications and can be valuable therapeutic agents for not only leukemia but also for other cancer types (e.g., breast cancer and glioblastoma) in which ID1 may be a prime therapeutic target.

Disclosure of Potential Conflicts of Interest
No potential conflicts of interest were disclosed.

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