Sorafenib Inhibition of Mcl-1 Accelerates ATRA-Induced Apoptosis in Differentiation-Responsive AML Cells

Rui Wang, Lijuan Xia, Janice Gabrilove, Samuel Waxman, and Yongkui Jing

Abstract

Purpose: All trans-retinoic acid (ATRA) is successful in treating acute promyelocytic leukemia (APL) by inducing terminal differentiation-mediated cell death, but it has limited activity in non-APL acute myeloid leukemia (AML). We aim to improve ATRA therapy of AML by enhancing apoptosis through repression of the antiapoptotic proteins Bcl-2 and Mcl-1.

Experimental Design: APL and AML cell lines, as well as primary AML samples, were used to explore the mechanisms regulating differentiation and apoptosis during ATRA treatment. Stable transfection and gene silencing with siRNA were used to identify the key factors that inhibit apoptosis during induction of differentiation and drugs that accelerate apoptosis.

Results: In differentiation-responsive AML cells, ATRA treatment induces long-lasting repression of Bcl-2 while first upmodulating and then reducing the Mcl-1 level. The Mcl-1 level appears to serve as a gatekeeper between differentiation and apoptosis. During differentiation induction, activation of MEK/ERK and PI3K/Akt pathways by ATRA leads to activation of p90RSK and inactivation of glycogen synthase kinase 3β (GSK3β), which increase Mcl-1 levels by increasing its translation and stability. Sorafenib blocks ATRA-induced Mcl-1 increase by reversing p90RSK activation and GSK3β inactivation, maintains the repressed Bcl-2 level, and enhances ATRA induced apoptosis in non-APL AML cell lines and in primary AML cells.

Conclusions: Inhibition of Mcl-1 is required for apoptosis induction in ATRA differentiation-responsive AML cells. ATRA and sorafenib can be developed as a novel drug combination therapy for AML patients because this drug combination augments apoptosis by inhibiting Bcl-2 and Mcl-1.

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Introduction

All trans-retinoic acid (ATRA) together with chemotherapy used as the first-line treatment for the newly diagnosed acute promyelocytic leukemia (APL) patients achieves approximately 75% cures (1). However, all APL patients treated with ATRA alone relapse, most likely because of the cells that escape terminal differentiation (2, 3) and also escape death. Although ATRA can induce differentiation in most non-APL AML cell lines and primary AML cells in culture, its clinical activity in non-APL AML patients is very limited (4). Past efforts have focused on the search of agents that can enhance ATRA-induced differentiation (4, 5). These have been proven to be unsuccessful in clinical testing (6).

The main pathway through which chemotherapy kills leukemia cells is induction of apoptosis. Apoptosis is mediated through intrinsic (mitochondrial) and extrinsic (death receptor) signaling pathways. The extrinsic pathway affected by TRAIL and death receptor 5 (DR5) is thought to be responsible for apoptosis of terminally differentiated APL cells after ATRA treatment (3). The mitochondrial apoptotic pathway is controlled by Bcl-2 family of antiapoptotic proteins. ATRA treatment was shown to have variable effects on the transcription of the Bcl-2 family of antiapoptotic proteins; it decreases Bcl-2, increases Bfl-1, while showing no effect on the Mcl-1 mRNA level (7, 8). However, we and others have shown that Mcl-1 protein is upregulated by ATRA and that this regulation is time dependent (9, 10). Mcl-1 is a key prosurvival protein that protects mature neutrophils from cell death (11) and its ATRA-induced increase might save AMLs from apoptosis.

We propose that a rapid AML cell death can be induced by blocking Mcl-1 in cells in which Bcl-2 is repressed by ATRA. Mcl-1 has a short half-life which is regulated through translation and protein degradation (12). We explored the mechanism of Mcl-1 upregulation by ATRA in AML cells and found that ATRA increased Mcl-1 levels at translational and posttranslational levels associated with p90RSK activation and GSK3β inactivation. We also studied the role of Mcl-1 in differentiation and apoptosis using cells in which Mcl-1 was either forced expressed or silenced and found that Mcl-1 plays a gate keeper role that protects cells from apoptosis. We then looked for an agent that, by reducing Mcl-1, will eliminate the block to apoptosis. We found that sorafenib, a multikinase inhibitor, counteracts ATRA-induced Mcl-1 increase and augments apoptosis induction in non-APL AML cell lines and in primary cells.

Materials and Methods

Reagents
ATRA, rapamycin, LY290024, SB216763, and cycloheximide were purchased from Sigma Chemical Co. U0126, PD184352,
Translational Relevance

Cytotoxic chemotherapy induces acute myeloid leukemia (AML) cell death by apoptosis through mitochondrial pathway which is blocked in part by Bcl-2 and Mcl-1, the two main antiapoptotic proteins. All trans-retinoic acid (ATRA) induces terminal differentiation-mediated cell death causing complete remission in patients with acute promyelocytic leukemia (APL), but only limited effect in other AML subtypes. In APL cells, Bcl-2 is repressed while Mcl-1 is induced during ATRA induced differentiation process. Inhibition of Mcl-1 accelerates apoptosis of differentiated cells. Increase in the ATRA-induced Mcl-1 protein is due to increased translation and stability associated with RSK activation and RSK3β inactivation. Sorafenib, a multikinase inhibitor, currently in clinical trials in AML patients, counteracts Mcl-1 induction by ATRA, together with ATRA augments apoptosis in cell culture, and in vivo extends the survival time of mice xenografted with human AML cells. This work provides a rationale for utilizing ATRA and sorafenib as a combination therapy for AML patients by repressing both Bcl-2 and Mcl-1 proteins and augmenting apoptosis.

and sorafenib were obtained from LC Laboratories (Woburn). Antibody to Mcl-1, ERK1, Bcl-2, β-actin, and USP9X were purchased from Santa Cruz Biotechnology, Inc.; to p-MEK (Ser217/221), p-ERK (Thr202/Tyr204), Akt, p-Akt (Ser473), p-mTOR, p-p70S6K, p-p70S6K (Thr389), p-p70S6K (Thr422)/Ser404, p-p90RSK (Ser380), p-p90RSK, p-pS6 (Ser240/244), p56, GSK-3β, p-GSK-3β (Ser9), caspase-3, -8, -9, -3 and Trail were from Cell Signaling Technology, Inc.

Cell lines and treatments. APL-derived NB4 and its ATRA resistance clone R4 were obtained from Dr. Willson Miller Jr (McGill University, Montreal, QC, Canada) (13). AML-derived HL-60 and THP-1 were obtained from ATCC, and M1-1 cells were obtained from Dr. Shigeru Fujita (School of Medicine, Ehime University, Japan) (14). MOLM13 cells were obtained from DSMZ. All cell lines were not authenticated by us once received in our laboratory and were cultured in RPMI1640 medium supplemented with 100 U/mL penicillin, 100 µg/mL streptomycin, 1 mmol/L L-glutamine, and 10% (v/v) heat-inactivated FBS (FBS). HL-60/Bcl2 cells transfected with Bcl-2 were obtained from Dr. Michael Cleary (Stanford University School of Medicine, Stanford, CA) (15). Mcl-1-expressing HL-60/M15 cells were generated in our laboratory (16). These cells were seeded at 10^7/mL and treated with ATRA alone or in combination with other agents for 2 days. For the time course studies, the cell number was adjusted and new medium with ATRA was changed every 2 days.

Detection of apoptosis, differentiation, protein levels, and silencing of Mcl-1. Apoptotic cells were determined by the Annexin V assay, according to the manufacturer’s instructions [Annexin V–FITC Apoptosis Detection Kit (BD Biosciences)]. The nitroblue tetrazolium (NBT) reduction assay and CD11b expression were used to evaluate differentiation using FACS. Western blot analysis was used to measure protein levels. To silence Mcl-1 we used siRNA. All these methods were done as we previously reported (8, 17).

Isolation of patient-derived leukemic blasts. Leukemic blasts were obtained from a bone marrow and three peripheral blood samples of AML patients with FAB subtypes of M2 and M4. These studies have been sanctioned by the Investigation Review Board of Mount Sinai School of Medicine (New York, NY) and all patients provided informed consent. Whole blood or bone marrow collected in heparinized tubes were subjected to Ficoll-Hypaque density gradient separation, the blasts were washed and resuspended in RPMI1640 with 20% FBS at 1 × 10^7 cells/mL.

Animal survival study. Twenty NOD/SCID/IL-2Rγ (NSG) mice (age, 8 weeks) were injected with 5 × 10^6 log-phase MOLM13 cells via the lateral tail vein. Five days after leukemia cell injection, the mice were divided randomly into 4 groups. A Cremophor EL/ethanol/water solution (12.5% Cremophor EL/12.5% ethanol/75%water) containing sorafenib was prepared and 20 mg/kg was administered by oral gavage. ATRA was prepared in DMSO and diluted with PBS to containing 10% DMSO and 10 mg/kg was administered intraperitoneally. Both drugs were given alone or as a combination daily for five continuous days a week until the mouse died. Control mice received the vehicle through the same route and on the same schedule. Survival times were compared with Kaplan–Meier survival analysis.

Statistical analysis

Data were analyzed for statistical significance using the Student t test (Microsoft Excel, Microsoft Corp.). A P value of <0.05 was considered statistically significant.

Results

Mcl-1 protein is increased while Bcl-2 protein is decreased in ATRA differentiation-responsive NB4 cells

Treatment of NB4 cells with 1 µmol/L ATRA for 2, 4, and 6 days produced on the respective days approximately 26%, 57%, and 38% of differentiated cells (Fig. 1A) and approximately 8%, 15%, and 40% of apoptotic cells (Fig. 1B). The parallel reduction in the percent of differentiated cells, accompanied by the rise in apoptotic cells, suggests that terminally differentiated cells eventually undergo apoptosis. Both the extrinsic and the intrinsic apoptotic pathways were examined by measuring the cleavage of caspase-8 and caspase-9 during the 2- to 6-day treatment (Fig. 1C). The cleaved fragments of caspase-3, -8, and -9 were not detected on day 2. PARP cleavage, detected on day 4, was associated with the extrinsic pathway caspase-8 cleavage but only minimally with caspase-9 cleavage. Consistent with the previous report, we found that the levels of both TRAIL and DR5 were induced by ATRA on day 2 and 4 (3). Increased PARP cleavage associated with cleaved fragments of caspase-3, -8, and -9 was detected in cells treated with ATRA for 6 days (Fig. 1C), indicating activation of both apoptotic pathways after prolonged treatment.

We also tested the levels of Bcl-2 and Mcl-1 protein in NB4 cells treated with 1 µmol/L ATRA for 2, 4, and 6 days. The Bcl-2 protein level was decreased after 2 days of ATRA treatment and remained repressed throughout the treatment (Fig. 1C). The level of Mcl-1 was increased on day 2 and gradually declined during day 4 and 6, at the time when the proportion of the differentiated cells declined and of the apoptotic cells rose. These data suggest that increased Mcl-1 protein protects differentiated cells from apoptosis and that Mcl-1 protein needs to be reduced for maximal...
apoptosis to occur. To test the role of Mcl-1 protein in ATRA-induced NB4 differentiation process, Mcl-1 was knocked down using siRNA (Fig. 1D). Although silencing of Mcl-1 decreased the ability of ATRA to induce differentiation at day 2 (Fig. 1E) it strongly enhanced ATRA-induced apoptosis as determined by PARP cleavage (Fig. 1D) and Annexin V staining (Fig. 1F). These data suggest that Mcl-1 protein is required for induction of differentiation by ATRA and that it functions as a main prosurvival protein protecting differentiated cells from death.

Increase in Mcl-1 protein levels by ATRA is associated with p90RSK activation and GSK3β inactivation

Utilizing ATRA differentiation-sensitive NB4 and its resistant R4 subclone, the protein levels of Mcl-1 and Bcl-2 were determined during ATRA-induced differentiation defined as the ability to reduce NBT. After 4 days of treatment with 1 μmol/L ATRA there were 52.7% NBT-positive cells in NB4 and only 8.7% in R4 cells (Fig. 2A). Treatment with ATRA for 2 days increased the level of Mcl-1 protein but reduced Bcl-2 in NB4 cells (Fig. 2B). Previously, it has been shown that ATRA represses Bcl-2 at both mRNA and protein levels (7). ATRA did not increase Mcl-1 mRNA level in NB4 cells as measured using RT-PCR (Fig. 2B), suggesting a posttranscriptional regulation. MCL-1 is an unstable protein with a very short-half life. To distinguish between ATRA effect on translation and stability of Mcl-1 protein, NB4 cells were pretreated with or without ATRA for 2 days and then treated with cycloheximide for 1 to 4 hours. ATRA treatment increased the Mcl-1 half-life in presence of cycloheximide from approximately 20 minutes to 80 minutes (Fig. 2C and D), indicating that ATRA stabilizes Mcl-1 protein. In contrast, ATRA effect on Bcl-2 was not influenced by cycloheximide treatment (Fig. 2C).

Raf/MEK/ERK and Akt/mTOR signaling pathways are activated during ATRA-induced differentiation process (18, 19) and ATRA treatment of NB4 cells increases the levels of phosphorylated p70S6K and rpS6 (20). Both Raf/MEK/ERK and Akt/mTOR pathways phosphorylate p70S6K and then rpS6, increasing protein translation (21). ERK phosphorylates p70S6K at Thr389/Ser424.
sites while mTOR phosphorylates it at Thr<sup>389</sup> site (22). The existence of a link between rise in Mcl-1 and these factors was tested in ATRA-treated NB4 cells. NB4 cells treated with 10<sup>-8</sup>–10<sup>-6</sup> mol/L of ATRA had elevated levels of phosphorylated MEK, ERK, Akt, mTOR, p70S6K, and rpS6 as well as an increased level of Mcl-1 protein. Importantly, we found that at the lower (10<sup>-8</sup> mol/L) concentration of ATRA there was an increase in ERK pathway phosphorylated Thr<sup>202</sup>/Tyr<sup>204</sup> p-p90RSK level, while p-p70S6K (Thr<sup>389</sup>; the target of mTOR pathway) was only induced by ATRA at concentrations higher than 10<sup>-7</sup> mol/L (Fig. 2E). Although MAP kinase inhibitor U0126, or PI3K inhibitor LY294002, or mTOR inhibitor rapamycin (40 nmol/L) decreased the level of phosphorylated p70S6K at Thr<sup>389</sup> (Fig. 3C) but neither blocked ATRA-induced phosphorylation of p90RSK, GSK3β, and rpS6 nor Mcl-1 increase. These data suggest that ATRA-activated MEK/ERK and PI3K/Akt increase Mcl-1 protein by both enhanced translation and increased stability involving activation of p90RSK and inactivation of GSK3β (Fig. 3D).

The stability of Mcl-1 protein is regulated by phosphorylation by ERK and GSK3β and by ubiquitination. GSK3β increases Mcl-1 degradation while deubiquitinase USP9X stabilizes Mcl-1 (23, 24). GSK3β has been found to be inactivated by phosphorylation in ATRA-differentiated NB4 cells (25, 26). GSK3β can be phosphorylated by AKT and p90RSK, the downstream target of ERK (27). We found that ATRA treatment of NB4 cells increased the phosphorylation of p90RSK, GSK3β, and ERK, and that those are associated with the increased levels of Mcl-1 (Fig. 2F). Because the level of USP9X was decreased by ATRA treatment (Fig. 2E), we concluded that the ATRA induced increase in Mcl-1 stability proceeds through activation of p90RSK, ERK, and inactivation of GSK3β.

To further identify the specific ATRA upregulated kinase pathways which mediate the Mcl-1 induction, we treated cells with MEK inhibitor U0126, or PI3K inhibitor LY294002, or mTOR inhibitor rapamycin. Inhibition of MEK and PI3K blocked the increase in phosphorylated-p90RSK, -GSK3β, and -rpS6 as well as the level of Mcl-1 protein (Fig. 3A and B). Rapamycin (40 nmol/L) decreased the level of phosphorylated p70S6K at Thr<sup>389</sup> (Fig. 3C) but neither blocked ATRA-induced phosphorylation of p90RSK, GSK3β, and rpS6 nor Mcl-1 increase. These data suggest that ATRA-activated MEK/ERK and PI3K/Akt increase Mcl-1 protein by both enhanced translation and increased stability involving activation of p90RSK and inactivation of GSK3β (Fig. 3D).
Mcl-1 level is crucial for ATRA-induced differentiation and apoptosis in FAB-M2 HL-60 cells

We tested the effect of 1 μmol/L of ATRA for 2, 4, and 6 days in differentiation-responsive FAB-M2 HL-60 cells. We found that similar to NB4 cells the percent of differentiated cells rose from 31% on day 2 to 60% on day 4 (Fig. 4A) and, like in NB4, it dropped to 40% on day 6; this drop was accompanied by a sharp rise (to 43%) in apoptotic cells (Fig. 4B). Moreover, like in NB4, the Mcl-1 protein levels increased on day 2 and dropped on day 6 to the level in untreated cells (Fig. 4C). ATRA treatment of HL60 cells increased the levels of phosphorylated p90RSK, phosphorylated (inactive) GSK3β, and phosphorylated rpS6 (Fig. 4C).

Compared with NB4 cells (Fig. 1), Bcl-2 was repressed on day 2 and 4, but this was not accompanied by substantial degree of apoptosis. Silencing of Mcl-1 with siRNA (Fig. 4D) produced increased PARP cleavage (Fig. 4D), and Annexin V staining (Fig. 4E), which rose to 48.8% in cells treated with ATRA for only 2 days. The fact that this profound apoptosis-inducing effect was not observed in absence of ATRA treatment (Fig. 4D and E), suggests that both, the repressed Bcl-2 and the silenced Mcl-1, were crucial.

To conclusively test the role of Mcl-1 and Bcl-2 in differentiation and apoptosis, we used an Mcl-1-overexpressing clone of HL-60 (HL60/M15 cells) and a clone with overexpressed Bcl-2 (HL-60/Bcl2 cells) and, respectively, compared ATRA-
induced differentiation and apoptosis to those in HL-60 clones transfected with empty vector (HL-60/V5 and HL-60/neo) (Supplementary Fig. S1A and S1D). ATRA treatment of HL-60/M15 cells enhanced differentiation but blocked apoptosis (Supplementary Fig. S1B and S1C). Bcl-2 overexpression had no effect on differentiation (Supplementary Fig. S1E), but protected HL-60 cells from apoptotic cell death (Supplementary Fig. S1F). These data show that although both Bcl-2 and Mcl-1 block apoptosis, probably only Mcl-1 participates in ATRA-induced differentiation.

Enhancement of ATRA-induced apoptosis by MEK inhibitors is associated with blockade of Mcl-1 induction in FAB-M2 HL-60 cells.

MEK inhibitors can enhance ATRA-induced apoptosis in AML cells but the mechanism has not been elucidated (28). The combined effects of ATRA with either of the two MEK inhibitors, U0126 and PD184352, were tested in HL-60 cells. ATRA combined with either U0126 or PD184352 induced more than 50% of the treated cells to undergo apoptosis (Supplementary Fig. S2A). The increased levels of p-ERK, p-p90RSK, p-rpS6, p-GSK3β, and Mcl-1 proteins by ATRA were blocked by either U0126 or PD184352 (Supplementary Fig. S2B). ATRA-mediated repression of Bcl-2 protein was not reversed either by U0126 or by PD184352. At the concentrations used, these agents alone did not induce apoptosis in HL-60 cells, even with repressed Bcl-2 or Mcl-1, respectively (Supplementary Fig. S2B). These data suggest that ATRA-induced Mcl-1 could be counteracted by inhibiting MEK/ERK signaling.

Sorafenib augments ATRA-induced apoptosis in AML cell lines and primary cells.

Sorafenib, an inhibitor of B-raf, the upstream activator of the MEK/ERK pathway, was approved for clinical use (29) and it...

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**Figure 4.** Mcl-1 plays an important role in ATRA-induced differentiation and apoptosis in HL-60 cells. HL-60 cells were treated with 1 μmol/L ATRA for the indicated times and the percent of differentiated cells was determined by measuring CD11b expression (A) and of apoptotic cells using Annexin-V-FITC (B). HL-60 cells were treated with 1 μmol/L ATRA for the indicated times and the levels of Mcl-1, Bcl-2, cleaved caspase-9, -8, -3, PARP and β-actin were determined using specific antibodies with Western blot (C). HL-60 cells transfected with Mcl-1 siRNA or control siRNA were treated with 1 μmol/L ATRA for 2 days. The protein levels of PARP, Mcl-1, Bcl-2, and β-actin were determined using specific antibodies with Western blot (D) and the percent of apoptotic cells was determined using FACS of cells stained with Annexin V-FITC (E).
has been reported to inhibit Mcl-1 translation in leukemia cells (30, 31). We tested the combined effects of ATRA and sorafenib in FAB-M2 HL-60 cells, additional AML cell lines, FAB-M4 ME-1 cells and FAB-M5 THP-1 cells as well as primary AML cells from four AML patients. Like HL-60 cells (Fig. 4A), both ME-1 and THP-1 cells are responsive to ATRA-induced differentiation (Supplementary Fig. S3). ATRA increased the levels of p-MEK, p-90RSK, p-GSK3β, and p-rpS6 as well as Mcl-1 protein while repressing Bcl-2 protein in HL-60 cells (Fig. 5A), ME-1 cells (Fig. 5C), and THP-1 cells (Fig. 5E). Sorafenib at 2.5 μmol/L blocked the ATRA-induced rise in the levels of p-MEK, p-GSK3β, p-90RSK, and p-rpS6 as well as in Mcl-1 protein (Fig. 5A, C and E) and together with ATRA induced apoptosis in more than 40% to 50% of cells; each agent alone induced only 5% to 20% apoptosis (Fig. 5B, D and F). The four primary samples were isolated from two FAB-M2 AML patients and two FAB-M4 AML patients. Three cases had normal genetic phenotype and one case had inv (16) like ME-1 cells. Neither of the patient’s sample had FLT3 and (or) NPM mutations. Treatment of these samples showed that ATRA or sorafenib alone induced less than 10% apoptosis above the untreated control; the combination of ATRA with sorafenib induced apoptosis in more than 30% to 40% of cells, suggesting a synergistic effect (Fig. 6A). ATRA increased the levels of p-MEK, p-90RSK, p-GSK3β, and p-rpS6 as well as Mcl-1 proteins in one case of tested primary AML cells (Fig. 6B); these effects were blocked by the addition of sorafenib. We found that sorafenib blocking of ATRA-induced Mcl-1 levels always coincided with inhibition of ATRA-induced phosphorylation of p90RSK, GSK3β, and rpS6 (Figs. 5A, 5C, 5E, and 6B). Although the regulation on p-ERK levels by either ATRA or sorafenib in the different cell lines was inconsistent, the combination of ATRA with sorafenib...
consistently augmented apoptosis in all cell lines and primary cells tested and this was always associated with downregulation of Bcl-2 and Mcl-1 protein (Figs. 5A, 5C, 5E, and 6B).

Sorafenib is currently being tested as a FLT3-ITD inhibitor for AML treatment (32) and FLT3-ITD AML cells are responsive to sorafenib-induced apoptosis at lower concentrations (33). Recently, sorafenib together with ATRA was used to treat 3 patients with FLT3-ITD AML, achieving durable responses (34). In MOML13 cells, which have FLT3-ITD, sorafenib, at lower concentrations, was able to induce apoptosis (35). This cell line undergoes differentiation in response to ATRA (36). We found that ATRA-induced differentiation of MOML13 cells (Supplementary Fig. S3) increased the levels of phosphorylated pRSK, phosphorylated GSK3β, and Mcl-1 (Fig. 6D). ATRA combined with lower concentrations of sorafenib was more effective in apoptosis induction than sorafenib alone (Fig. 6C). The effects of ATRA, sorafenib, and their combination were tested in vivo using MOML13 cells xenografted into NSG mice. ATRA alone did not increase survival while, as compared with control, sorafenib prolonged the survival of mice xenografted with MOML13. Importantly, ATRA significantly enhanced the sorafenib-induced survival (Fig. 6E and F). These data show that the antileukemia effect of the two drugs is mediated by sorafenib but greatly improved by the addition of ATRA.

Discussion
We found that, similar to normal neutrophils (37), APL cells terminally differentiated by ATRA treatment, die through apoptosis, and that this process is controlled, at least in part, by Mcl-1 protein. Silencing of Mcl-1 appears to accelerate the apoptosis of the differentiation-responsive AML cells. This
observation provides a rationale for achieving an accelerated apoptosis induction by combining ATRA with an Mcl-1 inhibitor in ATRA differentiation-responsive AML cells. Mcl-1 has been found to be essential for the development and survival of AML (38-40). We found that Mcl-1 protein levels were differently regulated in differentiated and apoptotic APL NB4 cells. The ratio of differentiated to apoptotic cells was approximately 4:1 on day 4 with elevated Mcl-1; while on day 6 of ATRA treatment, the ratio dropped to approximately 1:1 and Mcl-1 was strongly reduced (Fig. 1C). Although it is unclear how Mcl-1 is decreased in the terminally differentiated cells undergoing apoptosis, the observation that Mcl-1 knockdown increased apoptosis of ATRA-treated cells (Figs. 1D and 4D) indicates that Mcl-1 protein is needed for survival of differentiated cells. Knockdown of Mcl-1 in control cells, which had high levels of Bcl-2, increased apoptosis, but only slightly (Figs. 1F and 4E), indicating that both antiapoptotic proteins might play an important role in protecting these cells from death. This is supported by the observations that overexpression of either Bcl-2 or Mcl-1 blocks apoptosis of HL-60 cells treated with ATRA for 6 days (Supplementary Fig. S1C and S1F). However, our results show that it is the upregulated Mcl-1 that is the crucial prosurvival protein in NB4 and HL-60 cells during ATRA treatment as Bcl-2 level is suppressed by this treatment (Figs. 1C and 4C). Mcl-1 is regulated by a large number of pathways at translational and posttranslational levels (12, 41). We found that both MEK/ERK and Akt/mTOR pathways are activated in ATRA-treated APL cells (Fig. 2E) and that this leads to increase in phosphorylation of p70S6K, p90RSK, and p-rps6 levels and in Mcl-1 protein (Fig. 2E). Pathway-specific inhibitors showed that both MEK/ERK/RSK and PI3K/Akt, but not mTOR pathways regulate Mcl-1 protein levels in NB4 cells (Fig. 3A–C). On the basis of these observations, we propose that ATRA increases the rate of Mcl-1 translation by activating p90RSK and rps6 and stabilizes Mcl-1 protein by inactivating GSK3β and/or activating ERK (summarized in Fig. 3D). However, it is possible that other unexplored factors also participate in Mcl-1 protein regulation during ATRA treatment. Could targeting Mcl-1 improve the ATRA therapy for other AMLs? In an earlier publication (28), ATRA/MEK inhibitor-treated AML cells were shown to undergo accelerated apoptosis without signs of differentiation and through an unknown mechanism. We propose that the apoptotic effect of this combination is due to dual inhibition of Bcl-2 and Mcl-1. Treatment of HL-60 cells with either of the two MEK inhibitors, U0126 or PD184352, inhibited both the basal level and the ATRA-induced level of phosphorylated ERK, p90RSK, GSK3β, and rps6 as well as Mcl-1, while also maintaining Bcl-2 repression by ATRA, leading to a profoundly increased apoptosis (Supplementary Fig. S2). As a preclinical approach, we tested a combination of ATRA with sorafenib, a multikinase inhibitor with a transient effect in resistant and relapsed AML patients (29, 42). We showed that in FAB-M2 HL60 cells, a short (2 days) treatment with sorafenib was almost as effective as MEK inhibitors in reversing ATRA-induced Mcl-1 increase while allowing for ATRA repression of Bcl-2 to continue (Fig. 5A) and strongly increasing apoptosis (Fig. 5B). Relevant to our goal of expanding this treatment to other AMLs, sorafenib reversed ATRA-induced Mcl-1 level and maintained the low ATRA-repressed Bcl-2 level in FAB-M4 ME-1 cells and FAB-M5 THP-1 cells, in a sample of primary human AML cells from a FAB-M2 patient and in MOLM15 cells with FLT3-ITD mutation (Figs. 5 and 6). The pathways regulating Mcl-1 by ATRA in these cells were in general similar to those in APL with induced p-p90RSK/p-rps6 and p-GSK3β. Moreover, like in APL, the level of Bcl-2 was reduced by ATRA. The only difference from that observed in APL was the effect on ERK phosphorylation. While in the primary AML and HL-60 cells, ATRA treatment increased p-ERK level and sorafenib reduced it by approximately 50%, the effect in ME-1 and THP-1 was unexpected. ATRA induced pERK in ME-1 cells but sorafenib did not block it, while in THP-1 cells ATRA strongly reduced the level of pERK (Figs. 5 and 6). All the kinases in this pathway, including ERK, are subject to multiple regulatory controls and rapid phosphorylation and dephosphorylation (43, 44). We did not further study the reasons for the differences in phosphorylation status of ERK after treatment with ATRA and/or sorafenib but the kinetics of ERK phosphorylation and dephosphorylation might be different in these cells which would require more detailed kinetic study to prove it (45). We also did not resolve the question of p90RSK upregulation in absence of pERK upregulation but possibly, the active p90RSK persists longer. Importantly, the apoptosis augmenting effect of sorafenib when added to ATRA is present in all 8 samples tested; 4 non-AML APL cell lines and 4 primary AMLs (Fig. 6) and, it is consistently accompanied by reduction of both Mcl-1 and Bcl-2 proteins. Several reports showed that ATRA repressed Bcl-2 mRNA in differentiated AML cells (7, 46) and we found the cell lines used to be responsive to ATRA-induced differentiation (Fig. 4A, Supplementary Fig. S3). Therefore, the downregulation of Bcl-2 by ATRA might be transcriptional and secondary to differentiation. ATRA increases Mcl-1 protein, but not its mRNA (Fig. 2B). The Mcl-1 protein has a short half-life and it can be regulated by many factors such as cytokines, growth factors, and reactive oxygen species as well as activated caspase due to spontaneous apoptosis of cells (11). Therefore, the regulation of Mcl-1 by ATRA may be cell line specific and influenced by the culture conditions. However, differentiated AML cells with repressed Bcl-2, like normal neutrophils without Bcl-2 expression, rely on Mcl-1 for survival (47). Although we found that ATRA-increased phosphorylation of GSK3β and p90RSK, which contributes to Mcl-1 induction by ATRA (Fig. 3D), is inhibited by sorafenib, this inhibitor alone also reduces the basal level of Mcl-1 in the tested cell lines. The inhibition of basal Mcl-1 levels by sorafenib alone might be due to other mechanisms such as inhibition of FLT3-ITD and/or eIF4E phosphorylation, which are known to increase Mcl-1 translation (31, 48–50). Overall, we found that ATRA together with sorafenib augments apoptosis in vitro and prolongs survival of AML-xenografted mice, providing a translational rationale for studying this combination in AML cells and patients.

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

Authors’ Contributions
Conception and design: R. Wang, J. Gabrilove, Y. Jing
Development of methodology: R. Wang, L. Xia, Y. Jing
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): I. Xia, J. Gabrilove, Y. Jing
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): R. Wang, Y. Jing
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