Synergism of a novel MCL-1 downregulator, acriflavine, with navitoclax (ABT-263) in triple-negative breast cancer, lung adenocarcinoma and glioblastoma multiforme

ABNOK LEE1*, HYEON-OK JIN2*, MD. MASUDUL HAQUE3, HEE YEON KIM1, HANA JUNG3, JIN HEE PARK3, ILWHAN KIM4, JOO YEON SONG5, HYE KYOUNG YOON6, HYOUNG KYU KIM7, JIN HAN7, IN-CHUL PARK8, KWANG SEOK KIM8* and SAE GWANG PARK3,9*

1Department of Surgery, Busan Paik Hospital, College of Medicine, Inje University, Busan 47392; 2KIRAMS Radiation Biobank, Korea Institute of Radiological and Medical Sciences, Seoul 01812; 3Department of Microbiology and Immunology, College of Medicine, Inje University, Busan 47392; 4Department of Internal Medicine, Division of Oncology, Haeundae Paik Hospital, College of Medicine, Inje University, Busan 48108; 5Department of Pathology, Dongnam Institute of Radiological and Medical Sciences, Busan 46033; 6Department of Pathology, Inje University, Busan 47392; 7Cardiovascular and Metabolic Disease Center, Smart Marine Therapeutic Center, College of Medicine, Inje University, Busan 47397; 8Division of Radiation Biomedical Research, Korea Institute of Radiological and Medical Sciences, Seoul 01812; 9Innovative Therapeutic Research Institute, College of Medicine, Inje University, Busan 47397, Republic of Korea

Received April 29, 2021; Accepted September 27, 2021

DOI: 10.3892/ijo.2021.5292

Correspondence to:
Dr Sae Gwang Park, Department of Microbiology and Immunology, College of Medicine, Inje University, 75 Bokji-ro Street, Busan 47392, Republic of Korea
E-mail: micpsg@inje.ac.kr

Dr Kwang Seok Kim, Division of Radiation Biomedical Research, Korea Institute of Radiological and Medical Sciences, 75 Nowon-ro Street, Seoul 01812, Republic of Korea
E-mail: kskim@kirams.re.kr

*Contributed equally

Key words: acriflavine, navitoclax, myeloid cell leukemia sequence 1 downregulation, triple-negative breast cancer, drug resistance

Abstract. Myeloid cell leukemia sequence 1 (MCL-1), an anti-apoptotic B-cell lymphoma 2 (BCL-2) family molecule frequently amplified in various human cancer cells, is known to be critical for cancer cell survival. MCL-1 has been recognized as a target molecule for cancer treatment. While various agents have emerged as potential MCL-1 blockers, the present study presented acriflavine (ACF) as a novel MCL-1 inhibitor in triple-negative breast cancer (TNBC). Further evaluation of its treatment potential on lung adenocarcinoma and glioblastoma multiforme (GBM) was also investigated. The anticancer effect of ACF on TNBC cells was demonstrated when MDA-MB-231 and HS578T cells were treated with ACF. ACF significantly induced typical intrinsic apoptosis in TNBCs in a dose- and time-dependent manner via MCL-1 downregulation. MCL-1 downregulation by ACF treatment was revealed at each phase of protein expression. Initially, transcriptional regulation via reverse transcription-quantitative PCR was validated. Then, post-translational regulation was explained by utilizing an inhibitor against protein biosynthesis and proteasome. Lastly, immunoprecipitation of ubiquitinated MCL-1 confirmed the post-translational downregulation of MCL-1. In addition, the synergistic treatment efficacy of ACF with the well-known MCL-1 inhibitor ABT-263 against the TNBC cells was explored (combination index (CI)<1). Conjointly, the anticancer effect of ACF was assessed in GBM (U87, U343) and lung cancer (A549 and NCI-H69) cell lines as well, using immunoblotting, cytotoxicity assay and FACS. The effect of the combination treatment using ACF and ABT-263 was estimated in GBM (U87, U343 and U251), and non-small cell lung cancer (A549) cells likewise. The present study suggested a novel MCL-1 inhibitory function of ACF and the synergistic antitumor effect with ABT-263.

Introduction

While breast cancer is a preeminent threat to the healthcare of women worldwide, triple-negative breast cancer (TNBC) is the most notable subtype due to its incomparable aggressiveness. Currently, treatment for TNBC primarily relies only on systemic cytotoxic chemotherapy, with a rather unsatisfactory prognostic outcome (I). Therefore, effective therapy for TNBC is urgently required.

Apoptosis, programmed cell death, is an intrinsic phenomenon for normal embryonic development and tissue
homeostasis. Dysregulation of apoptosis is a critical step in cancer pathogenesis and is one of the major barriers to effective cancer treatment (2,3). The B-cell lymphoma 2 (BCL-2) protein family regulates the apoptosis pathway through mitochondrial outer membrane permeabilization, the cytosolic release of mitochondrial protein, cytochrome c, and caspase activation (4,5). Cancer cells resist apoptosis through various strategies, including increased expression of anti-apoptotic BCL-2 family proteins. The BCL-2 family includes anti-apoptotic members such as BCL-2, BCL-XL, BCL-W, as well as myeloid cell leukemia sequence 1 (MCL-1), and pro-apoptotic members such as BAX, BAK, and BH3-only (6).

MCL-1 has unique features among the BCL-2 family, including structure, function, and regulation (4). MCL-1 has a high affinity for pro-apoptotic BH3 peptides of BAX, BAK, BID, PUMA and NOXA compared with other BCL-2 anti-apoptotic family proteins (7). MCL-1 expression is quickly and readily induced by environmental stimuli including cytokines and growth factors unlike other proteins that downregulate, destabilize, or inactivate MCL-1 (12-14). Thus, MCL-1 has recently been recognized as an emerging therapeutic target in cancer.

Acriflavine (ACF), a mixture of 3,6-diamino-10-methylacridinium chloride (trypafavin) and 3,6-diaminoacridine (proflavine) has been used as an antimicrobial agent a century ago (15). Recently, its potential as an anticancer agent has been highlighted. The antitumor activity of ACF has been reported in various cancers, including colorectal, pancreas, and prostate (16-21). Its mechanism is still under investigation; however, HIF-1 inhibition is considered as the main anticancer mechanism.

In the present study, the anticancer activity of ACF on TNBC cells and its cell death mechanism were evaluated. Furthermore, the interaction of MCL-1 protein in TNBC cells during ACF-induced cancer apoptosis was revealed. The potential of MCL-1 downregulation by ACF as a breakthrough strategy against BCL-2 inhibitor resistance was evaluated. Additionally, the therapeutic potential of ACF against non-small cell lung cancer (NSCLC) and glioblastoma multiforme (GBM) was demonstrated.

Materials and methods

Reagents and antibodies. ACF was purchased from Sigma-Aldrich; Merck KGaA. APC Annexin V Apoptosis Detection Kit was purchased from BioLegend, Inc (cat. no. 640930). APC anti-human IgG was purchased from BioLegend, Inc. (cat. no. 366906). ABT-263 was purchased from Santa Cruz Biotechnology, Inc. (cat. no. sc-207241). Monoclonal antibodies to detect MCL-1 (product no. 5453), GSK-3β (product no. 9315), phosphorylated (p)-GSK-3β (product no. 9323), GAPDH (product no. 2118) and polyclonal antibodies to detect p-MCL1 (product no. 4579), BCL-XL (product no. 2762), BCL-2 (product no. 2872), β-catenin (product no. 9562), cleaved PARP (product no. 9541), caspase-8 (product no. 9746), caspase-9 (product no. 9502), ubiquitin (product no. 3933), XAF1 (product no. 138055) and XIAP (product no. 2042) were purchased from Cell Signaling Technology, Inc. Monoclonal antibody against β-actin (cat. no. A5316) was obtained from Sigma-Aldrich; Merck KGaA. Monoclonal antibodies to evaluate caspase-3 activation were purchased from Abcam (product code ab136812). MG-132 (product no. M7449), and cycloheximide (CHX; product no. C859) were purchased from Sigma-Aldrich; Merck KGaA. Protein A/G PLUS-Agarose (cat. no. sc-2003) was purchased from Santa Cruz Biotechnology, Inc.

Cell line culture. The human cancer cell lines MDA-MB-231 (HTB-26), HS578T (HTB-126), HCC-70 (CRL-2315), H460 (CCL-185), and NCI-H69 (HTB-119) cells were purchased from the American Type Culture Collection (ATCC). U87 (glioblastoma of unknown origin), U343 and U251 cells were kindly received as a gift from Professor Jongsun Park (Department of Pharmacology, College of Medicine, Chungnam National University, South Korea). STR profiling was performed on all cell lines. All cells were cultured in Dulbecco's modified Eagle's medium (DMEM) or RPMI-1640 medium with 10% fetal bovine serum (FBS) (all from Cytiva) and 1% penicillin, and cultured at 37˚C with 5% CO2. Cells were routinely controlled to exclude mycoplasma contamination.

Determination of cell viability. The In Vitro Toxicology Assay Kit (product no. TOX6; Sigma-Aldrich; Merck KGaA) was used to determine cellular viability. Briefly, cells (0.5-2x10^4 cells/well) were seeded and attached to 96-well culture plates. The indicated doses (0, 0.5, 1, 2, 10, 50, 100 μM) of ACF for different experiments were administered for 6-48 h. Cells were fixed in 10% trichloroacetic acid for 1 h at 4°C and washed five times with water. Fixed cells were stained with 0.4% sulforhodamine B for 15 min at room temperature and washed five times with 1% acetic acid. The incorporated dye was solubilized with 10 mM Tris Base, pH 8.8. The absorbance was spectrophotometrically measured at 565 nm using an EL800 microplate reader (BioTek Instruments, Inc.).

Cell death assay. Cell death was assayed using the Muse Annexin V & Dead Cell Kit (cat. no. MCH100105; Lumisex Corporation) according to the manufacturer's protocol. Briefly, 5x10^5 cells were collected, washed using cold phosphate-buffered saline (PBS), and resuspended in a medium containing 1% FBS. Following staining with the Muse Annexin V and Dead Cell reagent at room temperature for 20 min, cells were analyzed using the Muse Cell Analyzer (Luminx Corporation). Cell death was detected as the percentage of Annexin V- and/or 7-amino-actinomycin D (7-AAD)-positive cells.

Caspase-3/7 activity assay. Caspase-3/7 activity was analyzed using the Muse Caspase-3/7 kit (cat. no. MCH100108; Luminox Corporation). Briefly, 5x10^5 cells were collected...
and washed using cold PBS. Cells were resuspended in 1X assay buffer BA and mixed with Muse Caspase-3/7 reagent. Following 30 min of incubation at 37°C, Muse Caspase-7-AAD working solution was added to the cells. The cells were detected using the Muse Cell Analyzer and results were analyzed using Muse Cell Analyzer software version 1.6 (both from Luminex Corporation). The results were reported as the percentages of live cells (lower left panel, caspase-3/7-negative and 7-AAD-negative), apoptotic cells expressing caspase-3/7 activity (lower right panel, caspase-3/7 positive and 7-AAD-negative), late apoptotic/dead cells (upper right panel, caspase-3/7-positive and 7-AAD-positive), and necrotic cells (upper left panel, caspase-3/7-negative and 7-AAD-positive).

Western blot analysis. Cells were lysed using lysis buffer (cat. no. 9803S; Cell Signaling Technology, Inc.) supplemented with a protease inhibitor cocktail (cat. no. 11697498001; Roche Diagnostics GmbH) for 30 min at 4°C. Protein concentration was measured by bicinchoninic acid (BCA) assay. A total of 10-50 µg of protein per lane were separated using 8-14% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to nitrocellulose membranes (GE Healthcare Life Sciences; Cytiva). The membranes were blocked using 2% skim milk (product no. 70166; Sigma-Aldrich; Merck KGaA) for 1 h at room temperature followed by immunoblotting with specific primary antibodies (BCL-XL, BCL-2, GAPDH, β-actin, MCL-1, p-MCL1, XIAP, XAF1, cleaved PARP, GSK-3β, and p-GSK-3β; ubiquitin, caspase-8 and caspase-9) at a dilution of 1:2,000 overnight at 4°C. Horseradish peroxidase-conjugated anti-mouse (product no. 7076P2) and anti-rabbit (product no. 7074P2; both from Cell Signaling Technology, Inc.) secondary antibodies were incubated at a dilution of 1:1,000 at room temperature for 1 h. Following three washes with Tris-buffered saline 0.1% Tween-20, immunoreactive bands were visualized using an enhanced chemiluminescence detection system (Cyanagen Sr).

Annexin V and 7-AAD staining. Cells (4-5×10^5 cells/well) were seeded and attached to 6-well culture plates. The indicated doses (0, 1, 2 and 10 µM) of ACF were administered for 6-48 h. Cells were resuspended in binding buffer or PBS and stained using Annexin V-APC and 7-AAD for 15 min at room temperature. Flow cytometry was performed using a FACSCanto II instrument (BD Biosciences) and FlowJo v10.8.0 software (Tree Star, Inc.).

RNA extraction, cDNA synthesis, and reverse transcription-quantitative (RT-q) PCR. The mRNA was isolated using TRIzol (Thermo Fisher Scientific, Inc.). Extracted RNA was subjected to complementary cDNA synthesis using TOPscript RT DryMIX (Enzymics, Inc.) according to the manufacturer's protocol. qPCR was performed using SYBR Green TOPreal qPCR 2X PreMIX (Thermo Fisher Scientific, Inc.). Samples were normalized according to GAPDH (forward, 5'-GGAGCGGATCCCTCCAAAT-3' and reverse, 5'-GGC TGTGTGCATATTCTTCTATGG-3') and compared with assigned expression of untreated cells using the MCL-1 primers (forward, 5'-CGACGGCGTAAACAACT-3' and reverse, 5'-GGAAGAACTCCACAAAACT-3'). Amplification was conducted with the CFX96 Touch Real-Time PCR detection system, a PCR machine (Bio-Rad Laboratories, Inc.). The thermocycling conditions consisted of initial denaturation at 95°C for 30 sec, followed by 40 cycles at 95°C for 5 sec, annealing at 60°C for 30 sec and extension at 72°C for 30 sec. All reactions were examined in technical triplicate. The 2^ΔΔCq method (22) was used to calculate the relative gene expression.

Mitochondrial membrane potential and reactive oxygen species (ROS) measurement. Mitochondrial membrane potential and ROS generation were evaluated using specific fluorescence probe staining including tetramethylrhodamine-ethyl ester-perchlorate (TMRE; Thermo Fisher Scientific, Inc.), and 6-chloromethyl-2',7'-dichlorodihydrofluorescein diacetate acetyl ester (CM-H2DCFDA; Thermo Fisher Scientific, Inc.), respectively, as previously reported (23). Briefly, cells exposed to ACF (2 µM) and ABT-263 (1 µM) for 2, 6 or 24 h were incubated in 200 nM TMRE and 10 µM CM-H2DCFDA probe for 30 min at 37°C. Following washing twice using PBS, the fluorescence signals of TMRE and DCF-DA were detected and analyzed using a confocal microscope (LSM 700; Carl Zeiss AG).

Post-translational regulation and immunoprecipitation. For post-translational regulation of MCL-1, MDA-MB-231 and HS578T, cells were treated with cycloheximide (CHX; 10 µg/ml) and/or ACF (10 µM) and after 0, 1 and 2 h the expression of MCL-1 was determined using western blot analysis. For the proteasomal regulation study, MDA-MB-231 and HS578T cells were treated with MG-132 (10 µM) and/or ACF (10 µM) and after 6 h the expression of MCL-1 was determined using western blot analysis. For ubiquitination study, HS578T cells were treated with MG-132 (10 µM) and/or ACF (10 µM) and after 6 h the cells were harvested and washed twice with PBS. Cells were then resuspended in Pierce® IP Lysis Buffer (25 mM Tris-HCl pH 7.4, 150 mM NaCl, 1% NP-40, 1 mM EDTA, 5% glycerol) mixed with protease inhibitor and EDTA. A total of 100 µg of protein was incubated with anti-MCL-1 antibody for 2 h at 4°C. Subsequently, 20 µl of resuspended Protein A/G PLUS-Agarose (cat. no. sc-2003; Santa Cruz Biotechnology, Inc.) was added and incubated overnight at 4°C. Immunoprecipitates were collected and washed 4 times with PBS by centrifugation at 1,000 x g for 5 min at 4°C. The supernatant was discarded and the beads were resuspended with 40 µl of electrophoresis buffer and analyzed by SDS-PAGE and western blotting.

Combination index (CI) analyses. Using the CompuSyn software (CompuSyn v1.0; CompuSyn Inc.), the CI values were calculated according to the cell viability at corresponding concentrations. The CI value, CI=1 indicates a synergistic effect, CI>1 indicates an additive effect and CI<1 indicates and antagonistic effect (24,25).

Statistical analyses. Differences between groups were analyzed using one-way ANOVA and subsequent Tukey's multiple comparison post hoc test. Error bars represent ± standard deviation from three independent experiments. All statistical analyses were conducted using Graph-Pad Prism.
ACF induces apoptosis in TNBC cells in time and dose-dependent manners. To assess whether ACF has anticancer activity on TNBC cells under normoxic conditions, the effect of ACF on survival and apoptosis in MDA-MB-231 and HS578T cells was examined. ACF (5 µM for 72 h) induced typical apoptosis morphologies in TNBC cells characterized by shrinkage of cells and fragmentation into ultra-structured apoptotic bodies (Fig. 1A). Flow cytometric analysis revealed that ACF-induced cell death was apoptosis via activation of caspase-3/7 (Fig. 1C and D). Cleaved PARP, which is a known cellular substrate of caspases (Fig. 1E), and ROS levels were also increased following treatment with ABT-263, a BCL-2 protein inhibitor. However, ACF markedly increased the ROS levels within 2 h and reduced the mitochondrial membrane potentials at 6 h following treatment (Fig. 2). From a molecular perspective, ACF causes apoptosis via the intrinsic pathway.
point of view, the expression of caspase-8 and caspase-9 in ACF-treated (5 µM) cells was analyzed via western blot analysis. A total of 24 h following ACF treatment, the expression of caspase-9 was decreased while the expression of caspase-8 remained unchanged and the cleaved form of caspase-9 was not detected (Fig. S2). Collectively, these data suggested that ACF induces apoptosis via the intrinsic pathway.

Antitumor effect of ACF is mediated by the downregulation of MCL-1 protein. To evaluate the mechanism of ACF-induced intrinsic apoptosis, the protein levels of anti-apoptotic BCL-2 family proteins were determined using western blotting. ACF downregulated the protein levels of MCL-1, but not of other BCL-2 family proteins (Fig. 3A). The time kinetic study revealed that the MCL-1 protein level rapidly decreased and completely disappeared at 3-6 h following ACF treatment, while the majority of the cleaved PARP was formed 24 h following treatment (Fig. 3B). These results suggested the presence of other molecules between MCL-1 and cleaved PARP during ACF-mediated apoptosis. To determine which molecules linked MCL-1 downregulation and PARP cleavage, the XAF1 and XIAP protein levels were evaluated. Expression of XAF1 protein was increased and expression of XIAP protein was decreased at 24 h following the ACF treatment and decreased expression at 6 h. Expression of XIAP was decreased at 1, 3, 10 and 24 h but increased at 6 h following ACF treatment in the MDA-MB-231 cell line. In the HS578T cells, XAF1 increased at 3, 6, 10 and 24 h but decreased at 1 h and XIAP decreased at 1, 10 and 24 h and increased at 3 and 6 h following ACF treatment (Fig. 3D). In addition, MCL-1 downregulation was also observed both in hypoxic and normal conditions (Fig. S3). The aforementioned experiments entailed that ACF treatment induces an antitumor effect by downregulating MCL-1 not HIF-1 and it may also be associated with the expression of XAF1 and XIAP in TNBC cells.

ACF induces downregulation of MCL-1 at a transcriptional-and post-translational-dependent manner. MCL-1 protein expression is regulated in multiple steps: transcription and post-translation. First, the mRNA levels of MCL-1 were evaluated using RT-qPCR. ACF decreased the levels of MCL-1 mRNA in MDA-MB-231 and HS578T cells (Fig. 4A). Secondly, it was evaluated if MCL-1 degradation occurs under CHX (10 µg/ml), an inhibitor of protein biosynthesis. ACF (10 µM) decreased MCL-1 expression even following inhibition of translation (Fig. 4B). This indicated that MCL-1 is also downregulated following translation. Next, it was evaluated whether ACF increases proteasomal degradation of MCL-1 using a proteasome inhibitor, MG-132 (10 µM). The decrease in MCL-1 by ACF was blocked using MG-132 (Fig. 4C). Interestingly, cells treated with MG-132 alone demonstrated markedly higher MCL-1 protein levels than those treated with both MG-132 and ACF. Lastly, the ubiquitination of
MCL-1 protein was evaluated. Western blot data followed by immunoprecipitation also revealed increased ubiquitination of MCL-1 protein by ACF treatment (Fig. 4D). A higher ubiquitination level in cells treated with MG-132 was also consistent with the previous experiment (Fig. 4C). However, the level of GSK3β remained unchanged when treated with ACF (Fig. S4). Collectively, these data indicated that MCL-1 downregulation by ACF occurs at transcriptional and post-translational levels.

**ACF and ABT-263 work synergistically by overcoming the resistance to ABT-263.** To determine whether ACF may be used as a treatment strategy, HS578T cells were treated with ABT-263, a BCL-2 protein inhibitor, with or without ACF. While ABT-263 or ACF alone was not so effective in inducing apoptosis, when combined with 2 µM ACF, apoptosis was induced in more than 30% of cells. Importantly, apoptosis induced by the combination of ABT-263 and ACF was five times greater than that by ABT-263 or ACF alone (Fig. 5A). As previously reported (26), ABT-263 induced MCL-1 expression in TNBC cells, and ACF treatment prevented MCL-1 induction (Fig. 5B). The combination treatment of ACF and ABT-263 revealed synergy in both MDA-MB-231 and HS578T cells (Figs. 5C and D and S3). To determine the molecular pathways involved, the expression of caspase-8 and caspase-9 was analyzed. Although the expression of active caspase-9 was detected after 24 h, caspase-8 remained unaltered when treated with ACF (Fig. S2). However, the expression of both caspases...
was not detected 6 h following ACF treatment. Interestingly, combination of ACF and ABT-263 activated both caspases suggesting the involvement of both an intrinsic and extrinsic apoptotic pathway in the combination treatment (Fig. 5E). To determine whether the synergistic effect of ACF and ABT-263 may be observed in other cancer types, GBM (U87, U343, U251) and NSCLC (A549) cells were treated with ACF and ABT-263. Immunoblotting data revealed that the ABT-263 treatment upregulated the expression of MCL-1. However, when treated with 2 µM ACF, the expression of MCL-1 was downregulated again (Fig. 6A). Addition of ACF to ABT-263 also increased apoptosis in GBM (U87, U251 and U343) and NSCLC (A549) cell lines (Fig. 6B and C) compared with ABT-263 single therapy. Analysis of CI revealed synergistic apoptosis in U251, U343 and A549 cell lines (Fig. S5). Cell viability data also suggested that the combination treatment could eliminate cancer cells in a dose-dependent manner (Fig. 6D). These data suggested that ACF could act synergistically with ABT-263 by preventing MCL-1 upregulation.

Discussion

The BCL-2 family proteins modulate the intrinsic (mitochondrial) apoptotic pathway via the balance of pro- and anti-apoptotic proteins. The BCL-2 family consists of three main subclasses depending on the location of BCL-2 homology (BH) and function: multidomain anti-apoptotic (BCL-2, MCL-1 and BCL-XL), multidomain pro-apoptotic (BAX and BAK), and BH3-only pro-apoptotic (BIM, PUMA and NOXA) (2). Anti-apoptotic BCL-2 family members regulate apoptosis by isolating the stimulants from interacting with BAX and BAK (27,28). BH3 proteins (BAD and NOXA) are considered to cause apoptosis by engaging with anti-apoptotic proteins, freeing activators to stimulate BAX and BAK (29). Furthermore, anti-apoptotic BCL-2 family proteins seldom directly inhibit BAX and BAK, both inducing cell survival directly and indirectly, by sequestering BH3-only proteins (27,28). One of the most well-known and clinically advanced BCL-2 family target therapies is ABT-737 and its clinical analog ABT-263 (navitoclax). Their molecular structure mimics BAD BH3 and binds to the BH3 binding groove of BCL-2, BCL-XL, and BCL-W (30,31). However, the potency of BCL-2/BCL-XL inhibitors has been underwhelming (32). Their major limitation is the unexpected upregulation of MCL-1. In addition, the most frequently amplified gene in inhibitions of BCL-XL and BCL-2 is MCL-1, gaining significant interest in anticancer therapeutics.

MCL-1 is overexpressed in a wide range of cancers, including both solid and hematologic malignancies such as liver, ovarian, prostate, hematologic, and breast cancers (33-38). Previous studies revealed that MCL-1 is a crucial anti-apoptotic factor in TNBC (39,40). For instance, MCL-1-knockdown TNBC cell lines revealed markedly reduced viability, while BCL-XL silencing had a modest effect (41). Furthermore, Balko et al revealed an overexpression of the MCL-1 gene in 54% of TNBC patients who underwent neoadjuvant chemotherapy (42). They suggested that MCL-1 protects TNBC cells against apoptosis induced by cytotoxic chemotherapy (37). Moreover, the overexpression of MCL-1 has been proposed as a marker of poor prognosis in TNBC (39,43). Overall, MCL-1 may be one of the prospective therapeutic targets for TNBC. Upregulation of anti-apoptotic...
BCL-2 family proteins hallmark numerous cancers and occurs via various pathways. A prototype of this mechanism is MCL-1 upregulation, which contributes to protein stabilization by genetic inactivation of the ubiquitin ligase complex protein F-box and WD repeat domain-containing 7 (FBW7) (9,44,45). Given the importance of inducing apoptosis...
Figure 6. ACF overcomes the therapeutic resistance of ABT-263. (A) U87, U343, U251 and A549 cells were treated using ABT-263 and were incubated with 0 or 2 µM ACF for 24 h, and the expression of MCL-1 was detected using western blotting. (B) U87, U343, U251 and A549 cells were treated with ACF (2 µM) and/or ABT-263 (0.5 µM) for 24 h and apoptosis was evaluated using Annexin V and 7-AAD staining followed by FACS. (C) Quantitative analysis of the percentage of cell death (Annexin V+/7-AAD-). Data are presented as the means ± standard deviation from two independent experiments. (D) U87, U343, U251 and A549 cells were treated with indicated concentrations of ACF and ABT-263 and cell viability was determined using the sulforhodamine B assay. *P<0.01 compared with non-treated and ABT alone treated cells. ACF, acriflavine; 7-AAD, 7-amino-actinomycin D; MCL-1, myeloid cell leukemia sequence 1.
in managing cancer treatment, agents inhibiting anti-apoptotic proteins have garnered interest as a therapeutic option against cancer. To overcome this MCL-1 upregulation by BCL-2 family inhibitors, trials were conducted to identify MCL-1 inhibitors in various cancers, including breast, hematologic, and lung cancers. They combined the pre-existing BCL-2 family inhibitors with novel MCL-1 inhibitors to overcome the potential resistance to BCL-2 family inhibitors and improve therapeutic outcomes (46,47).

In the present study, it was revealed that ACF could regulate MCL-1 at the transcriptional and post-translational levels in breast cancer cells. The anticancer mechanism of ACF with MCL-1 downregulation was also revealed to be effective in lung and GBM cells. The limitation of the present study on the exact mechanism of MCL-1 decrease in lung cancer and GBM cells should be explored in further studies. Our study revealed that ACF promoted cell death of TNBC cells (MDA-MB-231 and HS578T) through the intrinsic apoptotic pathway in normoxia regardless of the HIF-1 pathway. Both ubiquitin-dependent and independent degradation of MCL-1 was reported. E3 ubiquitin-ligases confer a high degree of specificity to ubiquitination by recognizing target proteins. Ubiquitination of MCL-1 is mediated by five E3 ubiquitin ligases, including MCL-1 ubiquitin ligase E3 (MULE), glycoenzyme synthase kinase 3β (GSK3β), β-transducin repeats-containing protein (β-TrCP), FBW7, and tripartite motif-containing 17 (TRIM17) (48). However, different E3 ubiquitin-ligases appear to regulate MCL-1 protein expression in different cell types exposed to different stimuli. In the present study, the exact mechanism in the post-translational regulation of MCL-1 by ACF was not defined; however, ubiquitination was detected and GSK3β was not involved. Furthermore, the combination of ACF and ABT-263 generated apoptosis synergistically. This synergistic effect may be due to their independent inhibitory mechanism. ABT-263 stimulates TNBC apoptosis by inhibiting BCL-2, BCL-XL, and BCL-W. By contrast, our study revealed that ACF hinders MCL-1 and promotes TNBC apoptosis. It was assumed that as ABT and ACF work by regulating different BCL-2 family molecules, they compensated each other to enhance the antitumor effect. Furthermore, the combination of ACF and ABT-263 generated apoptosis synergistically. This synergistic effect may be due to their independent inhibitory mechanism. ABT-263 stimulates TNBC apoptosis by inhibiting BCL-2, BCL-XL, and BCL-W. By contrast, our study revealed that ACF hinders MCL-1 and promotes TNBC apoptosis. It was assumed that as ABT and ACF work by regulating different BCL-2 family molecules, they compensated each other to enhance the antitumor effect. Furthermore, the combination of ACF and ABT-263 generated apoptosis synergistically. This synergistic effect may be due to their independent inhibitory mechanism. ABT-263 stimulates TNBC apoptosis by inhibiting BCL-2, BCL-XL, and BCL-W. By contrast, our study revealed that ACF hinders MCL-1 and promotes TNBC apoptosis. It was assumed that as ABT and ACF work by regulating different BCL-2 family molecules, they compensated each other to enhance the antitumor effect.
References


This work is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International (CC BY-NC-ND 4.0) License.