



### **Original Article**

# The Histone Lysine-specific Demethylase 1 Inhibitor, SP2509 Exerts Cytotoxic Effects against Renal Cancer Cells through Downregulation of Bcl-2 and Mcl-1

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Lysine-specific histone demethylase 1 (LSD1), also known as KDM1A, can remove the methyl group from lysine 4 and 9 at histone H3, which regulates transcriptional suppression and activation. Recently, high expression of LSD1 in tumors has been shown to be involved in cancer cell proliferation, metastasis, and poor prognosis. We found that SP2509, a potent and reversible inhibitor of LSD1, induced apoptosis in human renal carcinoma (Caki and ACHN) and glioma (U87MG) cells. Pharmacological inhibition and siRNA-mediated silencing of LSD1 expression effectively downregulated anti-apoptotic proteins such as Bcl-2 and Mcl-1. Ectopic expression of these proteins markedly attenuated SP2509-induced apoptosis. At a mechanistic level, we found that inhibition of LSD1 downregulated Bcl-2 at a transcriptional level. Interestingly, protein expression of Mcl-1 was modulated at a post-translation level. Our results reveal that LSD1 could induce apoptotic cell death in renal carcinoma cells through downregulation of Bcl-2 and Mcl-1.

Key Words KDM1A, SP2509, Bcl-2, Mcl-1, Apoptosis

### INTRODUCTION

Histone methylation has been known as a key event for transcriptional regulation of genes, and many researchers have considered that histone methylation is an irreversible epigenetic marker [1]. However, Shi et al. [2] have reported that histone methylation is a reversible process through the discovery of the nuclear amine oxidase homolog LSD1 with histone demethylating activity. Histone demethylases are classified into 2 classes depending on the catalytic mechanisms; the flavin adenine dinucleotide-dependent amine oxidases (LSD1 and LSD2) and Jumonji domain containing proteins [3,4]. Specially, LSD1 binds to the promoter regions, demethylases lysine 4 and 9 at histone H3 (H3K4me1/2 and H3K9me1/2), and acts as a repressor or an activator of transcription. Demethylation of H3K4me1/2 or H3K9me1/2 by LSD1 induces gene transcription suppression or activation via inhibition of mono- and dimethyl group(s), respectively [5,6].

Pathologically, expression of LSD1 is elevated in many cancer cells including neuroblastoma [7], leukemia [8], lung [9], colon [10], and breast cancer [11], which is associated

with cancer cell proliferation, metastasis and poor prognosis [12,13]. Therefore, dysregulation of LSD1 can potentiate cancer treatment through inhibition of cancer cell proliferation and metastasis. Kumar et al. [14] have reported that LSD1/2 is highly expressed in renal cell carcinoma with clinical correlation as prognostic markers. Recently, multiple reversible LSD1 inhibitors have been developed. Among them, SP2509 is a non-competitive and potent inhibitor of LSD1 with anti-proliferative and anti-cancerous activities [15]. SP2509 inhibits proliferation through induction of ER stress response, as well as induces apoptosis in Ewing sarcoma cells [16,17]. Co-treatment with SP2509 and panobinostat (pan-HDAC inhibitor) was synergistically cytotoxic against acute myeloid leukemia [16]. Although the anti-cancer effects of SP2509 have been reported in several cancer cells, its molecular mechanism is still not clear.

In this study, we have found that SP2509 induces apoptosis by downregulation of Bcl-2 and Mcl-1 in human renal cancer cells.

Received June 18, 2020, Revised June 23, 2020, Accepted June 23, 2020

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#### **MATERIALS AND METHODS**

#### **Cell cultures and materials**

Human renal carcinoma cells (Caki and ACHN), human glioma carcinoma cells (U87MG), and human normal TCMK-1 cells were obtained from the American Type Culture Collection (ATCC, Manassas, VA, USA). Human renal mesangial cells were purchased from Lonza (Basel, Switzerland). All cells were cultured in appropriate medium containing 10% FBS (Welgene, Gyeongsan, Korea), 1% penicillin-streptomycin and 100 μg/mL gentamycin (Thermo Fisher Scientific, Waltham, MA, USA). R&D System (Minneapolis, MN, USA) supplied the recombinant zVAD-fmk. MG132 and lactacystin were purchased from Calbiochem (San Diego, CA, USA) and Biomol Research Laboratories (Plymouth Meeting, PA, USA), respectively. Cycloheximide (CHX) and β-actin (A5441) were purchased from Sigma-Aldrich (St. Louis, MO, USA). The PCR primers were purchased from Macrogen (Seoul, Korea). Anti-PARP (#9542S), anti-DR5 (#2764S), anti-cleaved caspase-3 (#9661S), and anti-Bcl-xL (#2764S) antibodies were supplied by Cell Signaling Technology (Beverly, MA, USA). Anti-c-FLIP (ALX-804-961-0100) and ani-USP9X (H00008239-M01) antibodies were obtained from Enzo Life Sciences (San Diego, CA, USA) and Abnova (Taipei City, Taiwan), respectively. Anti-Bim (AB17003), anti-Bax (554104), anti-caspase-3 (610322), and anti-XIAP (610762) antibodies were products of BD Biosciences (San Jose, CA, USA). Anti-clAP2 (SC-7944), anti-Mcl-1 (SC-12756), anti-Bcl-2 (SC-783), anti-LSD1 (SC-271720), and anti-β-TrCP (SC-33123) antibodies were provided from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

### Flow cytometry analysis and Western blotting

For apoptosis analysis, cells were fixed with 100% ethanol for 2 hours at 4°C, resuspended in 250  $\mu L$  of 1.12% sodium citrate buffer (pH 8.4) containing RNase (12.5  $\mu g/mL)$ , and incubated for 30 minutes at 37°C. After that, cellular DNA was stained with 250  $\mu L$  of propidium iodide (50  $\mu g/mL)$ . The stained cells were analyzed by flow cytometer (BD Biosciences). To examine protein expression, cells were lysed in RIPA lysis buffer and centrifuged at 12,000  $\times g$  at 4°C for 15 minutes. The proteins were separated by SDS-PAGE electrophoresis and transferred to an Immobilon-P membrane (GE Healthcare Life Science, Pittsburgh, PO, USA). The specific membranes were detected using an Immobilon Western Chemiluminescent HRP Substrate (EMD Millipore, Darmstadt, Germany).

### DNA fragmentation and 4', 6'-diamidino-2phenylindole (DAPI) staining

To detect apoptosis, DNA fragmentation was performed using the Cell Death Detection ELISAPLUS kit (Boehringer Mannheim, Indianapolis, IN, USA). For DAPI staining, the cells were fixed with 1% paraformaldehyde, washed with PBS, and stained with 300 nM DAPI solution (Roche, Mannheim, Germany) for 5 minutes. The nucleus condensation was tested by fluorescence microscopy (Carl Zeiss, Jena, Germany).

### **DEVDase activity assay**

Caki cells were treated with the indicated concentrations of SP2509 for 24 hours, harvested and incubated with reaction buffer containing substrate (acetyl-Asp-Glu-Val-Asp p-nitroan-ilide

### Reverse transcription-PCR and quantitative real-time PCR (qPCR)

Total cellular RNA was exacted using the Trizol® reagent (Life Technologies, Gaithersburg, MD, USA). cDNA was obtained using M-MLV reverse transcriptase (Gibco-BRL, Gaithersburg, MD, USA) [17]. The following primers were used for the amplification of human Bcl-2, Mcl-1, and actin: Bcl-2 (forward) 5'-GGT GAA CTG GGG GAG GAT TGT-3' and (reverse) 5'-CTT CAG AGA CAG CCA GGA GAA-3'; Mcl-1 (forward) 5'-GCG ACT GGC AAA GCT TGG CCT CAA-3' and (reverse) TT ACA GCT TGG ATC CCA ACT GC-3'; and actin (forward) 5'-GGC ATC GTC ACC AAC TGG GAC-3' and (reverse) 5'-CGA TTT CCC GCT CGG CCG TGG-3'. For PCR, we used Blend Taq DNA polymerase (Toyobo, Osaka, Japan) with primers targeting Bcl-2, Mcl-1 and actin: Bcl-2 (forward) 5'-GGT GAA CTG GGG GAG GAT TGT-3' and (reverse) 5'-CTT CAG AGA CAG CCA GGA GAA-3'; Mcl-1 (forward) 5'-ATG CTT CGG AAA CTG GAC AT-3' and (reverse) 5'-TCC TGA TGC CAC CTT CTA GG-3'; and actin (forward) 5'-CTA CAA TGA GCT GCG TGT G-3' and (reverse) 5'-TGG GGT GTT GAA GGT CTC-3'. The amplified products were separated by electrophoresis on a 2% agarose gel and detected under ultraviolet light. For qPCR, SYBR Fast qPCR Mix (Takara Bio Inc., Shiga, Japan) was used, and reactions were performed on Thermal Cycler Dice® Real Time System III (Takara Bio Inc.) [18].

#### **Transfection**

For knockdown of LSD1 using siRNA (Santa Cruz Biotechnology), cells were transfected using Lipofectamine RNAi-MAX Reagent (Invitrogen, Calshad, CA, USA). For constructing stable cell lines, Caki cells were transfected in a stable manner using Lipofectamine 2000 (Invitrogen) with the pcDNA3.1(+)/Bcl-2 and pcDNA(3.1+)/Mcl-1. After incubation for 48 hours, cells were replaced with fresh medium and selected by the G418 (700  $\mu g/mL$ ). To measure luciferase activity, Bcl-2/-751 and Bcl-2/-1281 promoter-constructs were transfected into the Caki cells using Lipofectamine 2000 (Invitrogen). After transfection, cells were treated with 2  $\mu$ M SP2509 for 24 hours, and lysates were incubated with luciferase substrates. Aliquots of the supernatant were used for the luciferase assay according to the manufacturer's instructions (Promega, Madison, WI, USA).

### Statistical analysis

The data were analyzed using a one-way ANOVA and post-hoc comparisons (Student-Newman-Keuls) using the Statistical Package for Social Sciences ver. 22.0 software (IBM Corp., Armonk, NY, USA).

### **RESULTS**

### The LSD1 inhibitor SP2509 induces apoptosis in human renal Caki cells

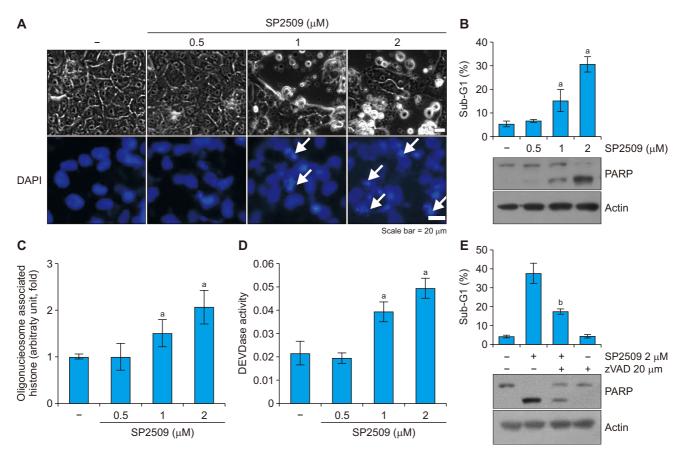
LSD1 is highly expressed in multiple cancer cells. We investigated whether the LSD1 inhibitor, SP2509 could induce apoptosis in renal carcinoma Caki cells. SP2509 induced apoptosis-related morphological changes, such as nuclear chromatin condensation, and dose-dependently increased sub-G1 population, PARP cleavage and cytoplasmic histone-associated DNA fragments (Fig. 1A-1C). We investigated the involvement of caspases in SP2509-induced apoptosis. SP2509 increased caspase-3 (DEVDase) activity in a

dose-dependent manner (Fig. 1D). Moreover, a pan-caspase inhibitor z-VAD blocked SP2509-induced increases in sub-G1 population and PARP cleavage (Fig. 1E). Therefore, these results indicate that the LSD1 inhibitor SP2509 induces caspase-dependent cancer cell death.

## Inhibition of LSD1 decreases expression levels of anti-apoptotic Bcl-2 family proteins

To investigate the molecular mechanism responsible for SP2509-induced apoptosis, we analyzed the expression levels of apoptosis-related proteins. SP2509 induced down-regulation of Bcl-2 and Mcl-1, whereas other proteins did not change (Fig. 2A). Next, we checked mRNA levels of Bcl-2 and Mcl-1 in SP2509 treated cells. As shown in Figure 2B, SP2509 inhibited Bcl-2 mRNA expression, whereas the Mcl-1 mRNA level did not change (Fig. 2B).

To further investigate whether inhibition of LSD1 is associated with modulation of Bcl-2 and Mcl-1 expression, we used LSD1 siRNA. Knockdown of LSD1 decreased Bcl-2 and Mcl-



**Figure 1. LSD1 inhibitor SP2509 induces apoptosis.** (A) Caki cells were treated with SP2509 (0.5-2 μM) for 24 hours. The cell morphology and nuclear condensation were examined by using an interference light microscope. White arrows showed nuclear chromatin condensation. (B) The sub-G1 population and protein expression were analyzed by flow cytometry and Western blotting. (C) The fragmentation of the nuclei was determined by using a DNA fragmentation assay kit. (D) Caspase activity was determined by using the caspase DEVDase assay kit. (E) Caki cells were treated with SP2509 (2 μM) in the presence or absence of a pan-caspase inhibitor, z-VAD-fmk (20 μM) for 24 hours. The sub-G1 population and protein expression were analyzed by flow cytometry and Western blotting. The values in graphs represent the mean  $\pm$  SD of three independent samples. DAPI, 4', 6'-diamidino-2-phenylindole.  $^aP$  < 0.01 compared to the treatment of SP2509.

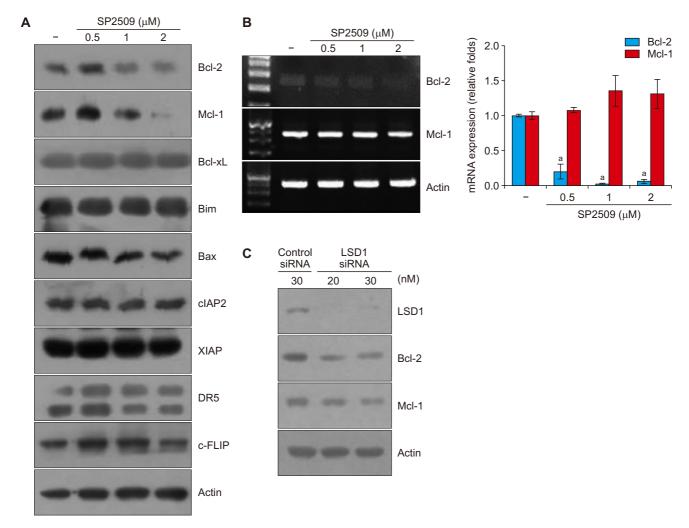


Figure 2. Inhibition of LSD1 induces downregulation of Bcl-2 and Mcl-1 expression. (A) Caki cells were treated with SP2509 (0.5-2 μM) for 24 hours. The protein expression levels of anti-apoptotic (Bcl-2, Mcl-1, and Bcl-xL), pro-apoptotic (Bim and Bax) Bcl-2 family, IAP family (cIAP2 and XIAP), DR5, c-FLIP, and actin were examined by Western blotting. (B) The mRNA levels of Bcl-2, Mcl-1, and actin were determined by reverse transcription PCR and real-time quantitative PCR. (C) Caki cells were transfected with LSD1 siRNA for 24 hours and the protein levels were measured by Western blotting. The graph represents the mean ± SD from independent experiments. <sup>a</sup>P < 0.01 compared with the control.

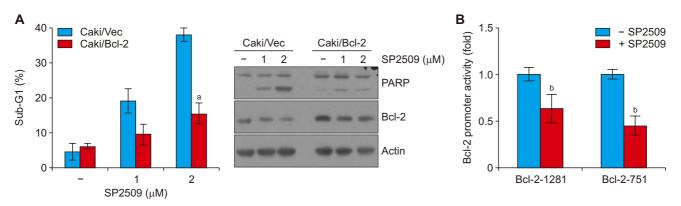


Figure 3. SP2509-induced Bcl-2 downregulation contributes to induction of apoptosis. (A) Vector cells (Caki/Vec) and Bcl-2 overexpressing cells (Caki/Bcl-2) were treated with SP2509 (1-2  $\mu$ M) for 24 hours, the protein expression and sub-G1 were examined by Western blotting and flow cytometry. (B) Caki cells were transiently transfected with a plasmid harboring the luciferase under the control of the Bcl-2/-751 and Bcl-2/-1281 promoter and then treated with 2  $\mu$ M SP2509 for 24 hours. The luciferase activity was analyzed. The graphs represent the mean  $\pm$  SD from independent experiments.  $^{a}P$  < 0.01 compared with the control in Caki/Vector.  $^{b}P$  < 0.01 compared with the control.

1 protein expression similar to the effect of LSD1 inhibitor (Fig. 2C). Therefore, these results suggest that inhibition of LSD1 induces downregulation of Bcl-2 and Mcl-1 anti-apoptotic proteins.

### SP2509 induces downregulation of Bcl-2 in human renal Caki cells

To confirm the functional importance of the Bcl-2 reduction in SP2509-induced apoptosis, we analyzed apoptosis using

overexpression of Bcl-2 stable cell lines. Ectopic expression of Bcl-2 inhibited SP2509-induced enhancement of sub-G1 population and PARP cleavage (Fig. 3A). We found that Bcl-2 mRNA expression was downregulated by SP2509 treatment (Fig. 2B). Therefore, we investigated the transcriptional regulation of Bcl-2, and found that SP2509 decreased Bcl-2 promoter activity (Fig. 3B). These data indicate that SP2509-induced downregulation Bcl-2 protein is modulated by transcriptional regulation.

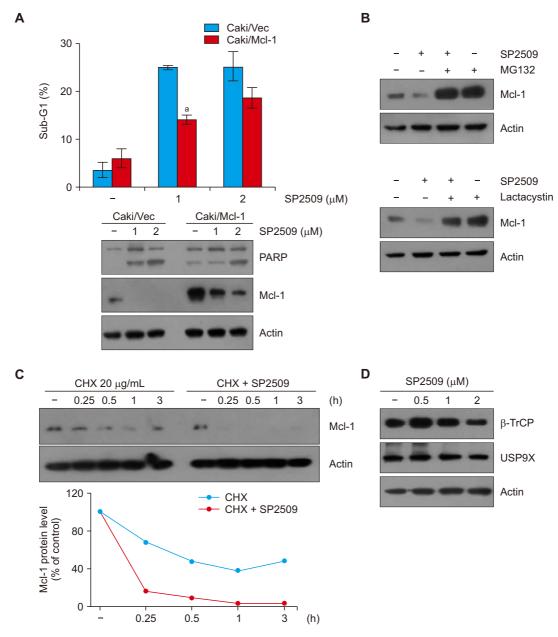


Figure 4. LSD1 inhibitor SP2509 downregulates McI-1 expression in Caki cells. (A, B) Vector cells (Caki/Vec) and McI-1-overexpressing cells (Caki/McI-1) were treated with SP2509 (2  $\mu$ M) for 24 hours. (B) Caki cells were pretreated with MG132 (1  $\mu$ M) and lactacystin (2.5  $\mu$ M) for 30 minutes, then treated with SP2509 (2  $\mu$ M) for 24 hours. (C) Caki cells were treated with or without SP2509 (2  $\mu$ M) in the presence of cycloheximide (CHX) (20  $\mu$ g/mL) for the indicated time periods. (D) Caki cells were treated with SP2509 (0.5-2  $\mu$ M). The protein expression levels were examined by Western blotting. (A) The graph represents the mean  $\pm$  SD from independent experiments.  $^{a}P$  < 0.01 compared with the SP2509-treated Caki/Vector.

### SP2509 inhibits McI-1 protein in human renal Caki cells

Next, we examined the involvement of Mcl-1 downregulation in SP2509-induced apoptosis. When we treated Mcl-1-over-expressing cells with SP2509, apoptotic population was less than that of vector transfected cells (Fig. 4A). Because SP2509 did not alter Mcl-1 mRNA expression, we explored the degradation of Mcl-1 through post-translational regulation. Two proteasome inhibitors, MG132 and lactacystin, reversed SP2509-induced Mcl-1 downregulation (Fig. 4B). Moreover, when we examined Mcl-1 protein stability using CHX, an inhibitor of protein biosynthesis, CHX plus SP2509 significantly degraded Mcl-1 protein compared to CHX alone (Fig. 4C). Mcl-1 protein stability is modulated via the ubiquitin-proteasome system. Therefore, we tested the alteration of E3 ligase and deubiquitinase capable of regulating Mcl-1 in

SP2509-treated cells. However, SP2509 did not increase E3 ligase  $\beta$ -TrCP and decrease USP9X expression (Fig. 4D).

### SP2509 selectively induces apoptosis in human renal cancer vs. normal cells

We next examined the effect of SP2509 on apoptosis in other cancer (renal carcinoma and glioma) and normal cells. SP2509 treatment effectively induced apoptosis and down-regulation of Bcl-2 and Mcl-1 expression in renal carcinoma ACHN and U87MG glioma cells (Fig. 5A). In contrast, normal cells (MC and TCMK-1) did not induce morphological change and sub-G1 population by SP2509 (Fig. 5B). Therefore, these data suggest that SP2509 may selectively induce apoptosis in cancer cells.

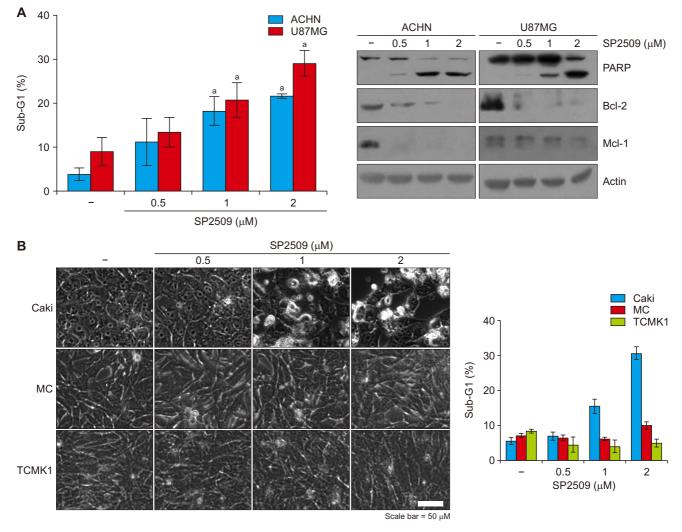


Figure 5. The effect of treatment of SP2509 in various cell lines. (A) Cancer cells (ACHN and U87MG) and normal cells (mesangial cell [MC] and TCMK1) were treated with SP2509 (0.5-2  $\mu$ M) for 24 hours. The sub-G1 cell population was examined by flow cytometry and the protein expression was measured by Western blotting. (B) Cell morphology was determined by interference light microscope. The graphs represent the mean  $\pm$  SD from independent experiments.  $^{a}P$  < 0.01 compared with control.

#### DISCUSSION

Histone methylation and demethylation imbalances are closely related to tumorigenesis and development [19]. SP2509 is known as a LSD1 inhibitor which exhibits anti-cancer effects. However, anti-cancer effect and molecular mechanisms of SP2509 have not been investigated in renal carcinoma cells. Our results reveal that pharmacological inhibition of LSD1 (SP2509) enhances apoptosis in cancer cells but not in normal cells. SP2509 downregulated Bcl-2 expression at the transcription and Mcl-1 expression at the posttranscriptional levels

To date, many reversible LSD1 inhibitors have been discovered and their activites were tested in cancer cells. Previously, CBB1007 was found to cell viability in pluripotent and ovarian cancer cells [20,21]. In case of 2-PCPA, it induces autophagic flux and cell cycle arrest in human osteosarcoma cells [22]. We first found that knockdown or pharmacologic inhibition of LSD1 downregulates Bcl-2 and Mcl-1 protein expression (Fig. 2A and 2C). SP2509 effectively decreased the protein expression of Bcl-2, and overexpression of Bcl-2 inhibited SP2509-induced cell death (Fig. 2A and 3A). Because SP2509 inhibited Bcl-2 mRNA levels and Bcl-2 promoter activity, suppression of transcription was involved in Bcl-2 downregulation by SP2509 (Fig. 3B). It has been reported that Bcl-2 transcription is regulated by diverse transcription factors, such as p53, NF-κB, c-Myc, and WT1 [23]. p53 is a tumor suppressor and negatively regulates Bcl-2 transcription [24,25]. However, SP2509 did not increase p53 expression (data not shown) in our system, so we rule out the possibility of p53-mediated Bcl-2 regulation. As mentioned above, NFκB can increase Bcl-2 expression at the transcriptional level, whereas c-Myc decreases Bcl-2 transcription [26,27]. In addition, Wilms' tumor suppressor gene WT1 positively regulates transcriptional level of Bcl-2 by suppressing promoter activity [28]. Therefore, further studies are required to investigate the molecular mechanism responsible for SP2509-mediated Bcl-2 suppression.

Our results indicate that SP2509-mediated downregulation of Mcl-1 is associated with the ubiquitin-proteasome pathway. Proteasome inhibitors reversed SP2509-mediated Mcl-1 downregulation (Fig. 4B). Ding et al. [29] demonstrated that E3 ligase β-TrCP is involved in Mcl-1 degradation, resulting in tumor suppression and sensitization to anti-cancer drugs. Schwickart et al. [30] have also reported that deubiquitinase USP9X stabilizes Mcl-1 expression. Based on these, we checked the alteration of E3 ligase and deubiquitinase capable of regulating Mcl-1 in SP2509-treated cells. However, SP2509 did not change  $\beta$ -TrCP and USP9X expression (Fig. 4D). Mule/ARF-BP1 and FBXW7 induced Mcl-1 ubiquitination and degradation [31]. In addition, FBXW7 cooperates with  $\beta$ -TrCP which results in the induction of GSK3 $\beta$ -mediated Mcl-1 degradation [32]. Therefore, it'll be of interest to elucidate involvement of these E3 ligases in SP2509-induced degradation of Mcl-1.

Taken together, our results show that downregulation of Bcl-2 and Mcl-1 play a critical role in SP2509-induced apoptosis in renal carcinoma Caki cells, but not in normal cells.

### **ACKNOWLEDGMENTS**

This work was supported by an NRF grant funded by the Korea Government (MSIP) (2014R1A5A2010008 and NRF-2019R1A2C2005921).

### **CONFLICTS OF INTEREST**

No potential conflicts of interest were disclosed.

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