

Apurinic endonuclease 1 promotes the cisplatin resistance of lung cancer cells by inducing Parkin-mediated mitophagy

ZHENG LI, YONGQIANG WANG, LINBO WU, YALU DONG, JING ZHANG,
FAN CHEN, WEI XIE, JIANGUO HUANG and NING LU

Cancer Center, Xinjiang Military Area General Hospital, Urumchi, Xinjiang Uygur Autonomous Region 830000, P.R. China

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Abstract. Platinum-containing doublet chemotherapy is the cornerstone of lung cancer treatment; however, cisplatin resistance is a major obstacle in the treatment of lung cancer. However, the mechanism underlying this resistance has not been fully elucidated. Previous studies have shown that serum apurinic/apyrimidinic endonuclease 1 (APE1) levels in patients with NSCLC are inversely associated with progression-free survival after platinum-containing doublet chemotherapy, and can serve as a biomarker for predicting disease prognosis and treatment efficacy. The present study was designed to investigate the role played by APE1 in the resistance of lung cancer to cisplatin. The levels of mitochondrial apurinic endonuclease 1 (m-APE1) and total APE1 (t-APE1) protein in a cisplatin-resistant A549 cell line (A549/DDP) and cisplatin-sensitive A549 cells were analyzed by western blotting. Mitochondrial membrane potential was detected by using the JC-1 staining method. The cisplatin-resistance of APE1-overexpressing A549 cells and APE1-silenced A549/DDP cells was assessed by cell apoptosis and colony formation assays. The results revealed that cisplatin-resistant A549 cells contained high levels of APE1, and exhibited elevated levels of autophagy. The levels of m-APE1 and t-APE1 protein were increased in the A549/DDP cells when compared with these levels in the A549 cells. Overexpression of APE1 and Mia40 enhanced the cisplatin resistance and autophagy of the A549 cells. APE1 knockdown restored the cisplatin sensitivity and reduced the levels of LC3II and Parkin in the A549/DDP cells, but promoted the release of cytochrome *c*. Furthermore, Parkin silencing or treatment with 3-methyladenine (3-MA, an autophagy inhibitor) promoted the apoptosis of APE1-overexpressing A549 cells, indicating that Parkin-mediated mitophagy plays an important role in the APE1-induced cisplatin resistance

of A549 cells. In conclusion, APE1 promotes the cisplatin resistance of lung cancer cells by inducing Parkin-mediated mitophagy.

Introduction

Lung cancer is one of the most common causes of cancer-related morbidity and mortality, and approximately 1.5 million new cases of lung cancer are diagnosed annually worldwide. Furthermore, lung cancer accounts for 13% of all newly diagnosed cancers worldwide (1,2). Approximately 85% of lung tumors are non-small cell lung cancers (NSCLCs), and adenocarcinoma is the most prevalent cancer subtype (3). The high incidences of lung cancer recurrence and metastasis are the two main causes of the failure to successfully treat NSCLC. Standard therapies for patients with locally advanced or metastatic lung cancer consist of radiotherapy and chemotherapy. However, most patients fail to achieve long-term survival with such treatments as they become resistant to chemotherapy (4). Chemotherapy resistance plays an important role in the recurrence and metastasis of NSCLC.

Cisplatin, approved by the FDA in 1978, is one of most effective chemotherapy drugs commonly used in the treatment of various types of solid tumors, including testicular, head and neck, ovarian, esophageal, cervical, and non-small cell lung cancer tumors (5,6). Cisplatin binds to DNA in cellular nuclei and mitochondria to form cisplatin-DNA adducts, which exert cytotoxic effects by blocking the replication and transcription of DNA in cancer cells (7). As cisplatin-based chemotherapy is commonly used to treat patients with advanced NSCLC, cisplatin resistance leads to the failure of this type of chemotherapy (8). Therefore, it is critical that we gain a better understanding of the mechanisms of acquired cisplatin resistance, and develop safe methods for reversing drug resistance. Apurinic/apyrimidinic endonuclease 1 (APE1) is a key, multi-functional DNA repair enzyme that plays important roles in repairing DNA damage, controlling protein reduction/oxidation, and modulating the activity of transcription factors (9,10). The mitochondrial intermembrane space assembly (MIA) pathway and Mia40 protein play pivotal roles in trafficking APE1 into mitochondria via the formation of disulfide bonds (11). APE1 provides most of the total AP endonuclease activity found in human

Correspondence to: Dr Ning Lu, Cancer Center, Xinjiang Military Area General Hospital, 359 Friendly North Road, Shayibake, Urumchi, Xinjiang Uygur Autonomous Region 830000, P.R. China
E-mail: luning407@sina.com

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cells, and is essential for proper functioning of the base excision repair (BER) pathway. APE1 also helps to regulate gene transcription, and participates in many important pathways, (e.g. the NF- κ B, AP-1, Myb, HIF-1 α and p53 pathways) via redox-dependent mechanisms (10,12). Importantly, APE1 accumulates in the cell nuclei and cytoplasm of various tumors, including NSCLC tumors. Abnormal APE1 expression in cancer patients is associated with a poor prognosis, tumor cell invasion, metastasis and angiogenesis, as well as resistance to radiotherapy and chemotherapy (13-15). Previous research has shown that serum APE1 levels in patients with NSCLC are inversely associated with progression-free survival after platinum-containing doublet chemotherapy, and can serve as a biomarker for predicting disease prognosis and treatment efficacy (16). However, the role played by APE1 in the resistance of lung cancer to cisplatin remains unclear.

To explore the role of APE1 in cisplatin resistance, the levels of APE1 and Mia40 expression were assessed in a cisplatin-resistant A549 cell line. Moreover, the cellular behaviors of cisplatin-resistant A549 cells were examined after altering their levels of APE1 and Mia40 expression, and a subsequent Parkin-mediated autophagy analysis was also performed. We believe that our results provide new insights into the cisplatin resistance of NSCLC cells, and will facilitate the development of new clinical therapies.

Materials and methods

Cell culture and establishment of a cisplatin-resistant subline. A549 lung cancer cells were obtained from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China) and cultured in RPMI-1640 medium (Gibco; Thermo Fisher Scientific, Inc.) supplemented with 10% heat-inactivated fetal bovine serum (FBS; HyClone; GE Healthcare), 100 U/ml penicillin, and 100 mg/ml streptomycin (Gibco; Thermo Fisher Scientific, Inc.) at 37°C in a humidified atmosphere containing 5% CO₂. A cisplatin-resistant A549 cell line (A549/DDP) was derived from A549 cells by incubating A549 cells with progressively increasing concentrations of cisplatin. After each treatment, the surviving cells were re-expanded and conventionally propagated for 4 generations in cisplatin-free medium. The relative levels of cisplatin resistance were determined using the clonogenic assay.

CCK-8 assay. Viability of the treated cells was determined by using a CCK-8 assay kit (Dojindo, Japan) according to the manufacturer's instructions. Briefly, 5,000 cells in 5 or 100 μ l of medium were seeded into each well of a 96-well culture plate. Next, 10 μ l of CCK-8 reagent was added to each well and incubated for 2 h at 37°C. After incubation, the absorbance of each well at 450 nm was measured with a microplate reader (Thermo Fisher, Finland).

Colony formation assay. Cells were seeded into the wells of a 12-well plate (1 \times 10³ cells per well) and cultured for 10-14 days. After culture, the cells were fixed with 20% methanol and stained with 0.1% crystal violet. Representative microscopic fields of the stained cells were photographed, and the numbers of colonies were counted.

Quantitative real-time polymerase chain reaction (qPCR). TRIzol reagent (Invitrogen; Thermo Fisher Scientific, Inc.) was used to extract the total RNA from cells according to the manufacturer's instructions; after which, a reverse transcription reagent kit (Takara, Tokyo, Japan) was used to reverse transcribe the total RNA into cDNA. The cDNA was detected according to the protocol used for SYBR Green Master Mix (Thermo Fisher Scientific, Inc.). qPCR analysis for relative expression level of mRNA was determined according to the procedures as follows: 95°C for 2 min, followed by 40 cycles of 94°C for 20 sec, 58°C for 20 sec, 72°C for 20 sec, and finally extension at 72°C for 4 min. The Agilent Stratagene Mx3000P Sequence Detection system (Agilent Technologies) was used to perform qPCR analysis. The primers used for qPCR were synthesized by Invitrogen/Thermo Fisher Scientific, Inc. and their sequences were as follows: APE1 forward, 5'-CCAGCC CTGTATGAGGACC-3' and reverse, 5'-GGAGCTGACCAG TATTGATGAGA-3'; Mia40 forward, 5'-ATGACCCCAACG ATCCATACG-3' and reverse, 5'-GGGGATAGAGGTCTG GGTATTT-3'; GAPDH forward, 5'-TGTGGGCATCAATGG ATTGG-3' and reverse, 5'-ACACCATGTATTCCGGGT CAAT-3'. GAPDH was used as an internal reference gene. The relative levels of gene expression were calculated using the 2^{- $\Delta\Delta C_q$} method (17).

Cell apoptosis assay. The A549 cells (5 \times 10⁵) were seeded in 6-well plates (Corning Inc.) and cultured for 24 h, then the A549 cells were transfected as described below. Forty-eight hours later, the treated cells were collected, washed, fixed, and permeabilized; after which, they were stained using Annexin V-FITC/PI (KeyGen) according to the manufacturer's instructions. After 15 min of staining, the number of apoptotic cells was determined by flow cytometry (BD Biosciences).

Mitochondrial membrane potential. The mitochondrial membrane potential was detected by using the JC-1 staining method. Briefly, the A549 cells (5 \times 10⁵) were seeded in 6-well plates (Corning Inc.) and cultured for 24 h. Then the A549 cells were transfected as described below. Forty-eight hours later, the cells were harvested, washed with PBS, and then incubated for 30 min with 5 μ M JC-1 (Cell Signaling Technology). After incubation, the cells were analyzed by flow cytometry (BD Biosciences).

Isolation of subcellular fractionation. The cytosolic fractions from the cultured A549 cells were isolated using a biochemical fractionation method using Mitochondria Isolation Kit for Cultured Cells (Pierce; Thermo Fisher Scientific, Inc.) according to the manufacturer's instructions. Briefly, the cells after the indicated treatment were lysed in lysis buffer A [20 mmol/l Tris (pH 7.4), 150 mmol/l NaCl, 2 mmol/l EDTA, and 1 % Triton X-100 with protease and phosphatase inhibitors] for 20 min at 4°C, and mitochondria were extracted in a Dounce homogenizer in mitochondrial buffer. The supernatant was further centrifuged to pellet the mitochondria, and the resulting supernatant was stored as the cytosolic fraction. Subcellular fractionation and western blot analysis were used to detect the cytochrome *c* content in the cytosol and mitochondria.

Western blot analysis. Cells (2×10^6) were collected and lysed in RIPA buffer. A BCA Protein Assay Kit (Pierce Biotechnology; Thermo Fisher Scientific, Inc.) was used to measure the protein concentrations in the lysates. Next, 50 μ g samples of total protein were separated by 12% SDS-PAGE, and the separated protein bands were transferred onto PVDF membranes (EMD Millipore). The membranes were first incubated with primary antibodies against APE1 (Abcam, cat. no. ab137708; dilution 1:1,000), Mia40 (Abcam; cat. no. ab87033, dilution 1:1,000), GAPDH (Abcam; ab8245, dilution 1:5,000), COX4 (Abcam; cat. no. ab33985, dilution 1:1,000), LC3 (Abcam; cat. no. ab48394, dilution 1:1,000), cytochrome *c* (Abcam; cat. no. ab133504, dilution 1:1,000), and Parkin (Abcam; cat. no. ab77924, dilution 1:1,000), followed by incubation with an HRP-conjugated goat anti-rabbit antibody (Wuhan Boster Biological Technology, Ltd.; cat. no. BA1054, dilution 1:20,000) or HRP-conjugated goat anti-mouse antibody (Wuhan Boster Biological Technology, Ltd.; cat. no. BA1051, dilution 1:20,000). Immunostaining of the protein bands was detected by enhanced chemiluminescence (ECL) reaction (Kibbutz Beit Haemek), and staining intensity was analyzed with an Alpha Innotech Flour Chem-FC2 imaging system (Alpha Innotech).

Cell transfection. To force overexpression of APE1 and Mia40 in cells, pcDNA 3.1 vectors (Genechem) containing APE1 or Mia40 plasmids were co-transfected into the A549 cells using Lipofectamine 2000 transfection reagent (Invitrogen; Thermo Fisher Scientific, Inc.). An empty vector served as a negative control. Specific small interfering RNAs (siRNAs) purchased from RiboBio, Inc. were used to knock down APE1 and Parkin expression in the cells. The siRNAs used were si-APE1 (5'-UACUCCAGUCGUACCAGACCU dTdT-3'), si-Parkin (5'-AUUUCUUGACCUUUUCUCCAC dTdT-3'), and a scrambled control siRNA (5'-CCAUGAGGU CAUGGUCUGdTdT-3'). Lipofectamine 2000 transfection reagent was used for all siRNA transfections. After 48 h of transfection with plasmid or siRNA, the transfected A549 cells were used for subsequent experiments.

Immunofluorescence and confocal microscopy. Briefly, treated cells were fixed in 4% paraformaldehyde and then permeabilized with 0.5% Triton X-100 for 15 min. Next, the cells were washed with PBS, blocked with 5% BSA in PBS, and then incubated with the primary antibody (anti-APE1, dilution 1:500), overnight at 4°C, followed by incubation with a secondary antibody that was conjugated with Alexa Fluorescence 568 (Invitrogen, Thermo Fisher Scientific, Inc.; cat. no. A-11011; dilution 1:1,000). DAPI was used for nuclear staining (Sigma-Aldrich, Merck KGaA; cat. no. D9542, dilution 1:5,000). The stained cells were visualized by confocal fluorescence microscopy.

Statistical analysis. All statistical analyses were performed using GraphPad Prism 5.0 (GraphPad Software, Inc.) and SPSS version 16.0 software (SPSS, Inc.). Results are presented as the mean \pm SEM of values obtained from at least three independent experiments. Statistical significance was determined by one-way analysis of variance (ANOVA) followed by Dunnett's test. A P-value <0.05 was considered statistically significant.

Results

Cisplatin-resistant A549 cells exhibit high levels of APE1 and autophagy. First, we established a cisplatin-resistant A549 cell line (A549/DDP) by incubating A549 cells with progressively higher concentrations of cisplatin. The A549 cells were assessed for viability after they had been treated with 0, 1, 2, 3, 4, 5 and 6 μ mol/l cisplatin, respectively (Fig. 1A). A549/DDP cells were assessed for viability after treatment with 0, 10, 20, 30, 40, 50, and 60 μ mol/l cisplatin, respectively. The A549/DDP cell line was significantly more resistant to cisplatin, as the IC₅₀ values for the A549 and A549/DDP cells were 4.10 and 25.91 μ mol/l, respectively. Next, we examined the levels of mitochondrial APE1 (m-APE1) and total APE1 (t-APE1) in both cell lines. We found that both the m-APE1 and t-APE1 protein levels were increased in the A549/DDP cells when compared with those in the A549 cells (Fig. 1B and C). Immunofluorescence assays confirmed the increased t-APE1 protein levels in A549/DDP cells (Fig. 1E), suggesting that APE1 might be transported into the mitochondria. As the MIA pathway is responsible for APE1 trafficking into the mitochondria, and Mia40 is directly involved in APE1's mitochondrial translocation, we analyzed the levels of Mia40 protein. We found that the levels of Mia40 protein were also increased in the A549/DDP cells when compared with those in the A549 cells (Fig. 1C). Furthermore, our findings suggest that the increased m-APE1 protein levels in A549/DDP cells are mediated by Mia40.

Accumulating evidence suggests that the autophagy process in cancer cells is involved in their resistance to chemotherapy (18-20). As shown in Fig. 1D and F, the levels of LC3II were increased in the A549/DDP cells. This finding suggests that cisplatin-resistant A549 cells have higher levels of autophagy.

APE1 and Mia40 overexpression enhances the cisplatin resistance of A549 cells. To investigate the roles of APE1 and Mia40 in A549 cells, we performed transfection studies that forced the overexpression of APE1 and Mia40 in A549 cells. The transfection efficiencies are shown in Fig. 2A and Fig. S1A. Interestingly, we found that overexpression of Mia40 increased the levels of mitochondrial APE1 (m-APE1) protein in the A549 cells (Fig. 2B). Moreover, the levels of t-APE1 and Mia40 protein in the A549 cells were increased after the cells were transfected with APE1 and Mia40 plasmids, respectively (Fig. 2C). These results indicate that Mia40 promotes the translocation of APE1 into mitochondria. Immunofluorescence assays confirmed the increase in APE1 expression (Fig. 2D). We also assessed the viability of A549 cells overexpressing APE1 and Mia40, and found that overexpression of both APE1 and Mia40 enhanced the cisplatin resistance of the A549 cells (Fig. 2E). Colony-forming assays showed that overexpression of both APE1 and Mia40 enhanced the colony formation efficiency of the A549 cells (Fig. 2F). In addition, APE1 and Mia40 overexpression decreased the apoptosis rate of the A549 cells after treatment with cisplatin (Fig. 2H and J). As APE1 and Mia40 interact with each other via disulfide bond formation, and Mia40 levels directly affect the translocation of APE1 into mitochondria, where it helps to maintain the integrity of mitochondrial DNA (11), it is possible that the overexpression

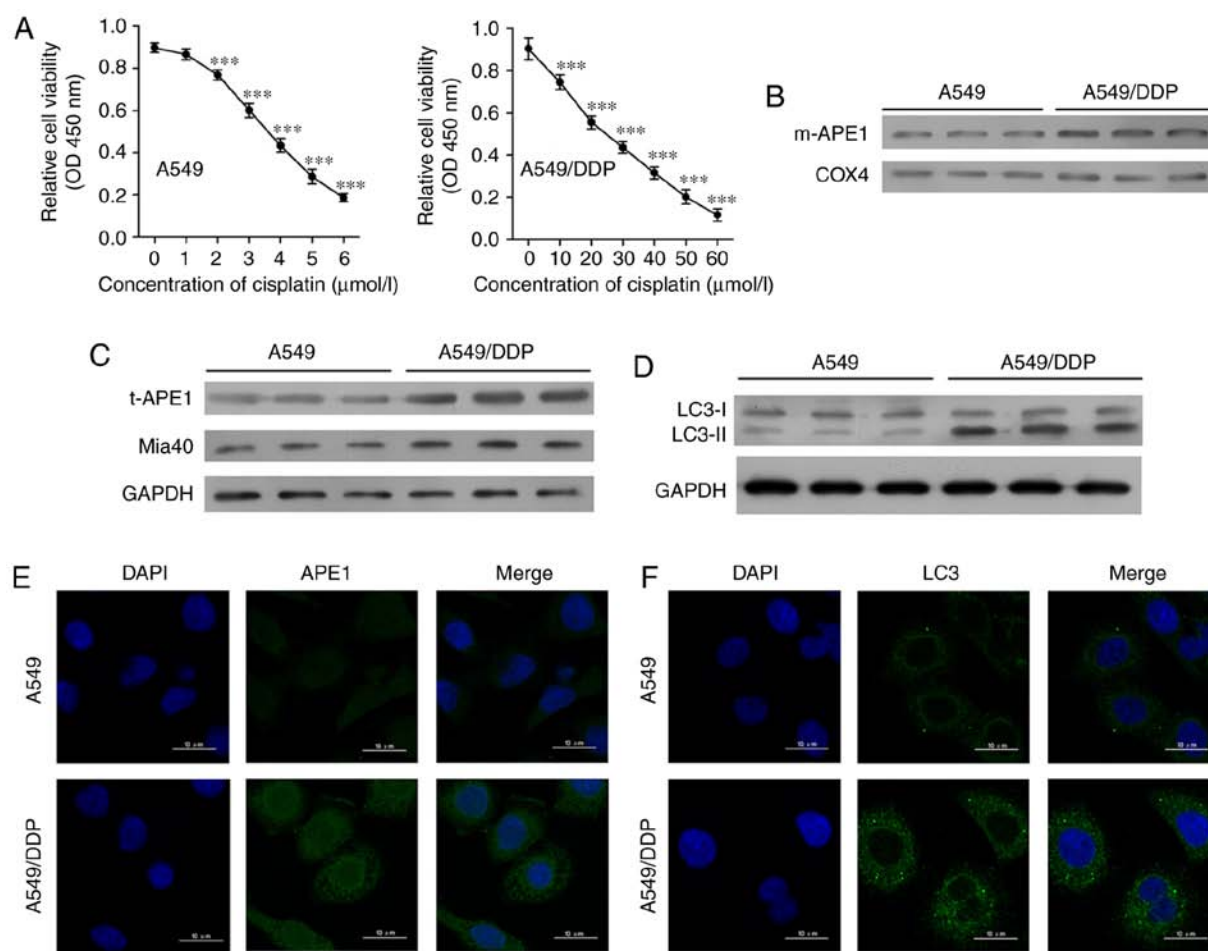


Figure 1. Cisplatin-resistant A549 cells exhibit high levels of APE1 and autophagy. (A) A549 cells were treated with 0, 1, 2, 3, 4, 5 and 6 μmol/l cisplatin for 24 h, and the A549/DDP cells were treated with 0, 10, 20, 30, 40, 50 and 60 μmol/l cisplatin for 24 h. Cell viability was assessed by the CCK-8 assay. (B) Western blot analysis was performed to analyze the levels of mitochondrial APE1 (m-APE1) protein in A549/DDP and A549 cells. COX4 was used as loading control for the mitochondrial APE1 protein. (C) Western blot analysis of the total APE1 (t-APE1) and Mia40 protein levels in A549/DDP and A549 cells. GAPDH was used as a loading control. (D) Western blot analysis of the LC3 protein levels in A549/DDP and A549 cells. (E and F) Immunofluorescence assays were performed to assess the total APE1 and total LC3 levels in A549 and A549/DDP cells. Data represent results obtained from three independent experiments (mean ± SEM of triplicate samples). ***P<0.001, vs. 0 μmol/l cisplatin-treated A549 or A549/DDP cells. APE1, apurinic/apyrimidinic endonuclease 1; LC3, microtubule-associated protein 1A/1B-light chain 3.

of Mia40 enhances cisplatin resistance by promoting the translocation of APE1 into mitochondria. We next used flow cytometry to assess the mitochondrial membrane potential of transfected A549 cells. As shown in Fig. 2G, the overexpression of both APE1 and Mia40 enhanced the mitochondrial membrane potential. Furthermore, overexpression of APE1 and Mia40 also promoted autophagy in A549 cells by upregulating LC3-II expression, and decreasing the levels of cytochrome *c* protein in the cytosolic fraction (Fig. 2I). It is possible that APE1 induced Parkin-mediated mitophagy, because as shown in Fig. 2I, overexpression of APE1 increased Parkin expression, and overexpression of Mia40 further increased the levels of Parkin protein in the A549 cells.

APE1 knockdown promotes the cisplatin sensitivity of A549/DDP cells. To further investigate the role of APE1 in A549/DDP cells, we performed transfections to knock down APE1 expression those cells. The transfection efficiency is shown in Fig. 3A. Both the m-APE1 and total APE1 (t-APE1) protein levels in A549/DDP cells were significantly decreased by APE1 knockdown (Fig. 3B and C). The decrease in t-APE1

protein was confirmed by an immunofluorescence assay (Fig. 3D). APE1 knockdown increased the cisplatin sensitivity of A549/DDP cells (Fig. 3E). In addition, APE1 knockdown reduced the colony formation efficiency and promoted the apoptosis of A549/DDP cells (Fig. 3F, G and J). The mitochondrial membrane potential of A549/DDP cells was significantly downregulated after APE1 knockdown (Fig. 3H). With regards to cell mitophagy, APE1 knockdown reduced the levels of LC3-II and Parkin in the A549/DDP cells; however, it significantly upregulated the levels of cytochrome *c* protein in the cytosolic fraction (Fig. 3I). Collectively, APE1 was shown to play an important role in the cisplatin resistance of A549 cells, and regulate Parkin-mediated mitophagy.

Parkin-mediated mitophagy plays an important role in the APE1-induced cisplatin resistance of A549 cells. Given that APE1 overexpression and knockdown were shown to regulate Parkin-mediated mitophagy, we next investigated whether Parkin-mediated mitophagy is involved in cisplatin resistance. Here, the study was divided into three groups as follows: a) APE1 overexpression group;

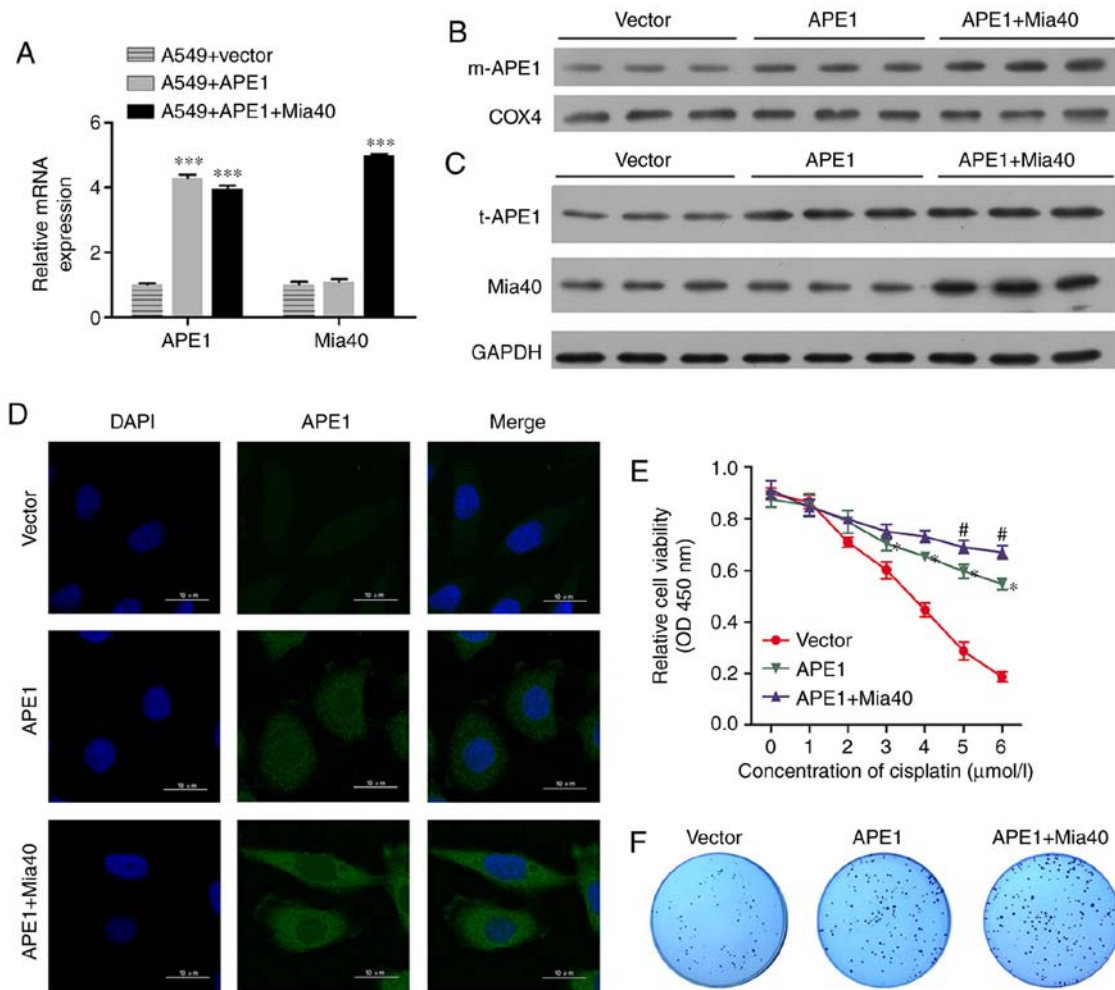


Figure 2. APE1 and Mia40 overexpression increases the cisplatin resistance of A549 cells. A549 cells were transfected with vectors containing an APE1 overexpression plasmid and Mia40 overexpression plasmid as indicated. (A) Transfection efficiency was assessed by qPCR. (B) Western blot analysis of the mitochondrial APE1 (m-APE1) protein levels in A549 cells after transfection. COX4 was used as loading control for mitochondrial APE1. (C) Western blot analysis was used to analyze the total APE1 (t-APE1) and Mia40 protein levels in A549 cells after transfection. GAPDH was used as a loading control. (D) Immunofluorescence assays were performed to assess the total APE1 levels in A549 cells after transfection. (E) The transfected A549 cells were treated with 0, 1, 2, 3, 4, 5, and 6 $\mu\text{mol/l}$ cisplatin for 24 h, and their viability was assessed by the CCK-8 assay. (F) Colony-formation assays were performed to analyze the colony formation efficiency of the transfected A549 cells. * $P < 0.05$, *** $P < 0.001$, vs. the vector group. # $P < 0.05$, vs. the APE1 group. APE1, apurinic/apyrimidinic endonuclease 1; LC3, microtubule-associated protein 1A/1B-light chain 3.

b) APE1 overexpression+Parkin siRNA group; c) APE1 overexpression+3-MA group. 3-MA (5 mM) (Sigma-Aldrich; Merck KGaA, M9298) was used as an inhibitor of autophagy according to the product introduction. In A549 cells, Parkin was significantly knocked down by si-Parkin transfection (Fig. S1B). As shown in Fig. 4A, Parkin knockdown and 3-MA treatment significantly decreased the levels of autophagy in the APE1-overexpressing A549 cells, and also increased the levels of cytochrome *c* protein in these cells (Fig. 4A). With regards to cisplatin resistance, both Parkin knockdown and 3-MA treatment significantly increased the cisplatin sensitivity of the APE1-overexpressing A549 cells (Fig. 4B). Moreover, Parkin knockdown and 3-MA treatment reduced the colony formation efficiency and significantly promoted the apoptosis of the APE1-overexpressing A549 cells (Fig. 4C-E). We also assessed the mitochondrial membrane potential of these cells, and found that their mitochondrial membrane potential was reduced by Parkin knockdown and 3-MA treatment (Fig. 4F). In brief,

Parkin-mediated mitophagy was shown to play an important role in the APE1-induced cisplatin resistance of A549 cells.

Discussion

Autophagy is a selective autophagic process that provides metabolites needed for biosynthesis and energy production in response to stressful conditions (21). Autophagy induced by metabolic and therapeutic stress can play a dual role (22). For its pro-death role, excessive autophagy in cancer cells induces 'autophagic cell death' or 'type II programmed cell death.' For its pro-survival role, autophagy eliminates damaged organelles and recycles cellular degradation products (23,24).

Mitochondrial-specific autophagy (mitophagy) is an important mitochondrial quality control system that selectively degrades excess or damaged mitochondria via an autophagic process. Similarly, mitophagy plays a dual role in cancer cell drug resistance that depends on the specific micro-environmental condition and cell type (25). Mitophagy can promote

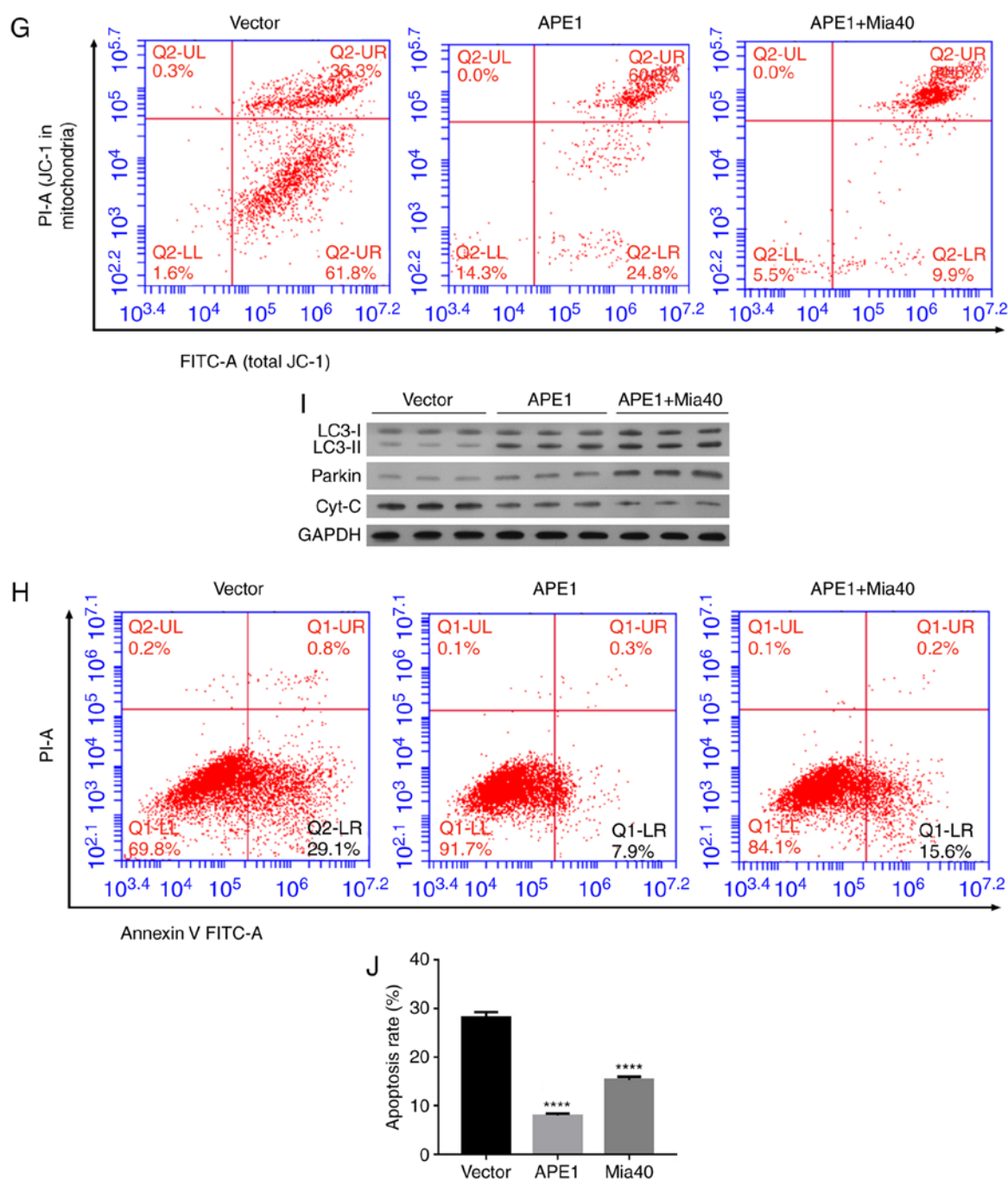


Figure 2. Continued. (G) Flow cytometry was used to analyze the cellular distribution of JC-1 after transfection. (H) Transfected A549 cells were treated with cisplatin, and their rate of apoptosis was analyzed by flow cytometry. (I) The transfected A549 cells were subjected to fractionation to obtain the cytosolic fraction. Western blot analysis was performed to analyze the levels of LC3, total cytochrome *c*, and Parkin in the transfected A549 cells. (J) The cell apoptosis rate is shown in a histogram. GAPDH was used as loading control. Data represent results obtained from three independent experiments (mean \pm SEM of triplicate samples). **** $P < 0.0001$ vs. the vector group. APE1, apurinic/aprimidinic endonuclease 1; LC3, microtubule-associated protein 1A/1B-light chain 3.

the survival of various types of cancer cells and tumors exposed to cytotoxic stress by degrading damaged mitochondria and reducing the levels of reactive oxygen species in mitochondria (26,27). Paradoxically, excessively damaged mitochondria may induce cell metabolic disorders and force cancer cells to undergo 'autophagic cell death.' While chemotherapeutic agents such as cisplatin induce oxidative stress and mitochondrial dysfunction in cancer cells, mitophagy selectively degrades excessive or damaged mitochondria via

autophagy (28). Therefore, the progress of mitophagy helps to mediate drug resistance in cancer cells. In our study, the cisplatin-resistant A549 cell line displayed an increased level of mitophagy when compared with the classical A549 cell line, and inhibition of autophagy by 3-MA treatment or Parkin knockdown reduced the cisplatin resistance of A549 cells. Interestingly, we found accumulations of APE1 throughout cisplatin-resistant A549 cells, and in the mitochondria of those cells. In addition, our findings demonstrated for the first time

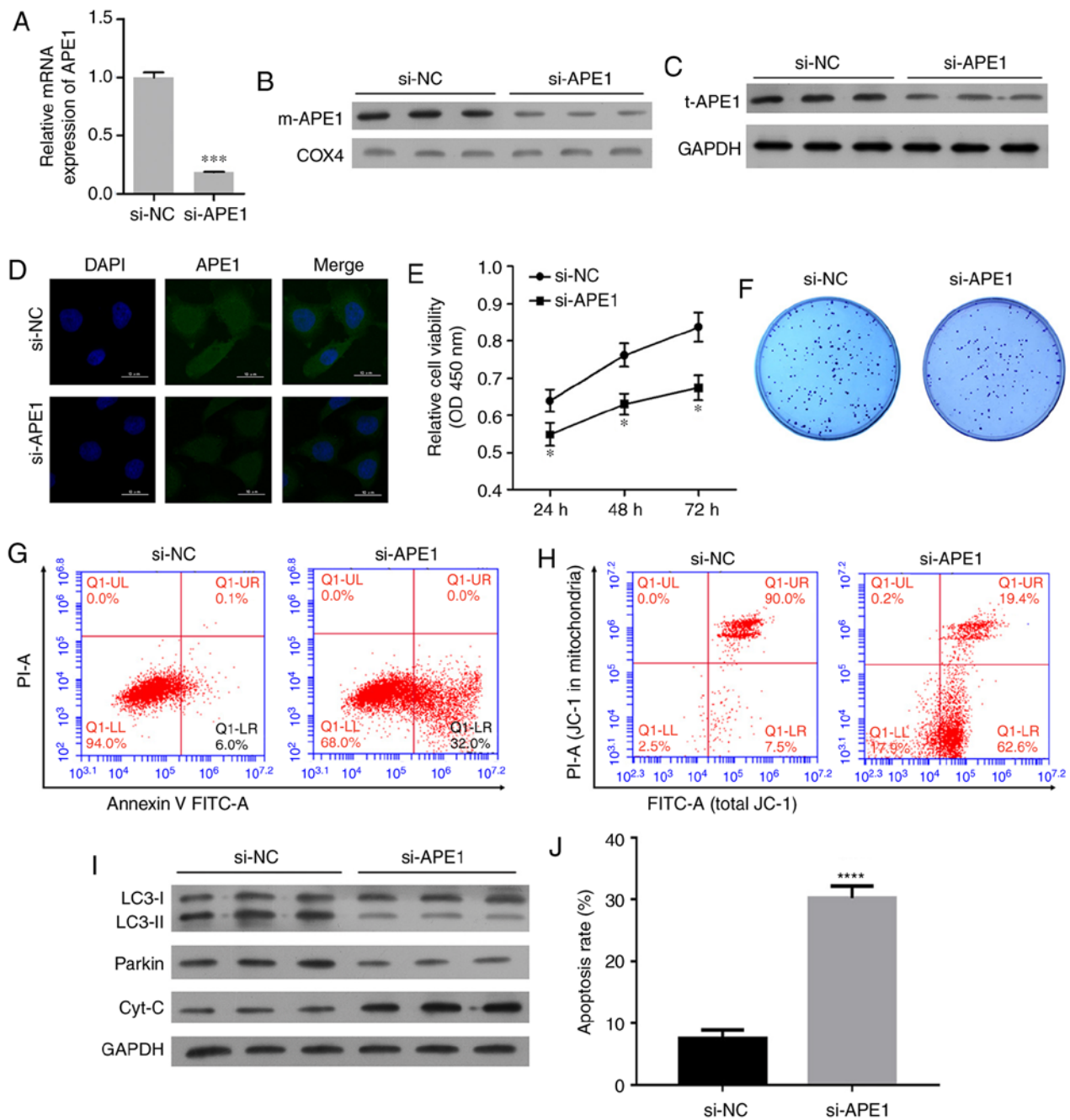


Figure 3. APE1 knockdown increases the cisplatin sensitivity of A549/DDP cells. A549/DDP cells were transfected with small interfering RNA (si-APE1 or si-NC). (A) Transfection efficiency was assessed by qPCR. (B) The levels of mitochondrial APE1 (m-APE1) in the transfected A549/DDP cells were analyzed by western blotting. COX4 was used as loading control for mitochondrial APE1. (C) The total APE1 (t-APE1) protein levels in transfected A549/DDP cells were analyzed by western blotting. GAPDH was used as loading control. (D) Immunofluorescence assays were performed to assess the total APE1 levels in A549/DDP cells after transfection. (E) Transfected A549/DDP cells were treated with 15 μ mol/l cisplatin for 24 h, and their viability was assessed by the CCK-8 assay. (F) Colony formation assays were performed to analyze the colony formation efficiency of the transfected A549/DDP cells. (G) The transfected A549/DDP cells were treated with cisplatin, and their rate of apoptosis was analyzed by flow cytometry. (H) Flow cytometry was used to analyze the distribution of JC-1 after transfection. (I) The transfected A549/DDP cells were subjected to fractionation to obtain the cytosolic fraction. Western blot analysis was performed to analyze the levels of LC3, total cytochrome c, and Parkin. (J) The cell apoptosis rate is shown in a histogram. GAPDH was used as loading control. Data represent the results obtained from three independent experiments (mean \pm SEM of triplicate samples). * P <0.05, *** P <0.001 and **** P <0.0001 vs. the si-NC group. APE1, apurinic/apyrimidinic endonuclease 1; LC3, microtubule-associated protein 1A/1B-light chain 3.

that APE1 can induce Parkin-mediated mitophagy in A549 cells (Fig. 5).

Parkin, an E3 ubiquitin ligase, selectively degrades damaged mitochondria via mitophagy in mammalian cells (29,30). For Parkin-mediated mitophagy, Parkin requires PINK1, a Ser/Thr kinase, to assist in sensing a mitochondria's

functional status (31-33). The efficiency of mitophagy depends on the levels of ubiquitin and deubiquitinases present in mammalian cells (33-35). Defects of Parkin are often associated with various pathologies, including cancer. A previous study showed that Parkin deficiency promoted tumorigenesis in a mouse model. However, Parkin is often

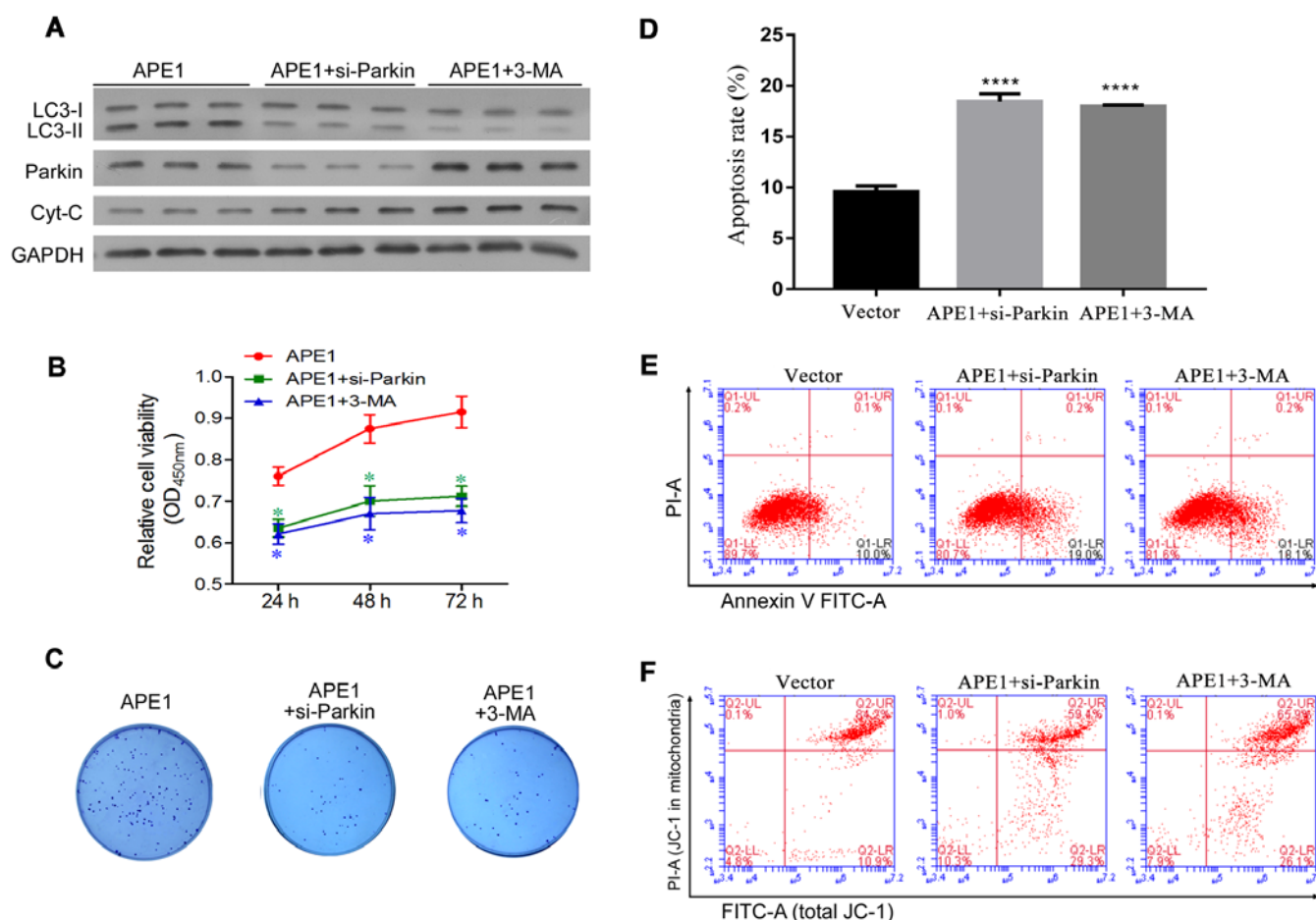


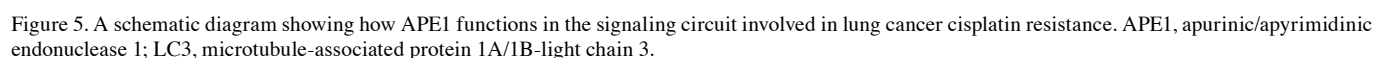
Figure 4. Parkin-mediated mitophagy plays an important role in the APE1-induced cisplatin resistance of A549 cells. A549 cells were first transfected with an APE1 overexpression plasmid. Next, the APE1-overexpressing A549 cells were transfected with small interfering RNA (si-Parkin or si-NC) or treated with 3-MA. (A) A549 cells after the relevant transfections (APE1, si-Parkin or si-NC) and their indicated treatment were subjected to fractionation to obtain the cytosolic fraction. Western blot analysis was performed to analyze the levels of LC3, total cytochrome *c*, and Parkin in the cytosolic fraction. GAPDH was used as loading control. (B) Following the indicated transfection and treatment, the A549 cells were co-cultured with 3 μ mol/l cisplatin for 24 h, and their viability was assessed by the CCK-8 assay. (C) Colony formation assays were performed to analyze the colony formation efficiency of the transfected A549 cells after they had received their indicated treatment. (D) The cell apoptosis rates are shown in a histogram. (E) The apoptosis rates of the transfected A549 cells treated or not with 3-MA were analyzed by flow cytometry. (F) The distribution of JC-1 in the transfected A549 cells was analyzed by flow cytometry. * $P < 0.05$ and **** $P < 0.0001$ vs. the APE1 group. 3-MA, 3-methyladenine (an autophagy inhibitor); APE1, apurinic/apyrimidinic endonuclease 1; LC3, microtubule-associated protein 1A/1B-light chain 3.

inactivated in many tumors, and its role remains unclear. Parkin was reported to promote resistance to apoptosis independent of mitophagy (36). However, Villa *et al* (25) reported that E3 ubiquitin ligase (ARIH1/HHARI) protects against chemotherapy-induced cell death by Parkin-mediated mitophagy (25). Our study showed that the levels of Parkin in cisplatin-resistant A549 cells were increased, and that Parkin knockdown increased the cisplatin sensitivity of APE1-overexpressing A549 cells. Thus, Parkin-mediated mitophagy in A549 cells promoted cisplatin resistance of these cells. Our results are in accordance with the conclusion previously reached by Villa *et al* (25).

As we found that Parkin-mediated mitophagy in A549 cells promotes the cisplatin resistance of these cells, strategies for inhibiting mitophagy should receive increased attention as methods for overcoming cisplatin resistance and improving the efficacy of cancer therapy. However, the functional outcomes produced by mitophagy-induced cell death or survival can depend on the type of cancer treatment

administered, and the results achieved can be confusing. In breast cancer, inhibition of mitophagy by liensinine markedly promoted the sensitivity of the cancer cells to classical chemotherapeutic agents, including doxorubicin, paclitaxel, vincristine, and cisplatin (37). Moreover, treatment with liensinine significantly inhibited late-stage mitophagy and reduced the viability of breast cancer cells, but increased their rate of apoptosis. An *in vivo* assay showed that inhibition of mitophagy inhibited tumor growth in the MDA-MB-231 xenograft model (37). Our study demonstrated that inhibition of mitophagy by Parkin knockdown and 3-MA treatment significantly enhanced the cisplatin sensitivity of tumor cells. Our findings provide a potential therapeutic strategy for treating lung cancer.

Our study has some limitations that should be mentioned. First, due to limited research funds, only one cell line was used in the study, and this may weaken the reliability of our findings. Thus, it is important to verify our findings in other NSCLC cell lines before reaching a final conclusion. Second, although we



In conclusion, our study revealed a novel mechanism by which APE1 promotes cisplatin resistance in lung cancer cells. In the cisplatin-resistant cell line, APE1 was highly expressed and was translocated into the mitochondria via its interaction with Mia40. This mitochondrial translocation of APE1 increased the mitochondrial membrane potential, decreased the levels of cytochrome *c* protein, and induced Parkin-mediated mitophagy, resulting in the cisplatin resistance of lung cancer cells.

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Not applicable.

Competing interests

The authors declare that they have no competing interests.

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