



RESEARCH ARTICLE

# Flavonoid morin inhibits proliferation and induces apoptosis of melanoma cells by regulating reactive oxygen species, Sp1 and Mcl-1

Yoon Jin Lee<sup>1</sup> · Woo Il Kim<sup>2</sup> · Soo Young Kim<sup>2</sup> · Sung Woo Cho<sup>1</sup> · Hae Seon Nam<sup>1</sup> · Sang Han Lee<sup>1</sup> · Moon Kyun Cho<sup>2</sup>

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**Abstract** Reactive oxygen species (ROS) is associated with cancer progression in different cancers, including melanoma. It also affects specificity protein (Sp1), a transcription factor. Flavonoid morin is known to inhibit growth of cancer cells, including lung cancer and breast cancer. Herein, we hypothesized that morin can inhibit cancer activities in melanoma by altering ROS generation. The aim of this study is to determine the effects of morin and its underlying mechanisms in melanoma cells. Effects of morin on cell proliferation and apoptosis were determined using standardized assays. Changes in pro-apoptotic and anti-apoptotic proteins were analyzed by western blot analysis. Cellular ROS levels and mitochondrial function were evaluated by measuring DCF-DA fluorescence and rhodamine-123 fluorescence intensities, respectively. Morin induced ROS production and apoptosis, as presented by increased proportion of cells with Annexin V-PE(+) staining and sub-G<sub>0</sub>/G<sub>1</sub> peak in cell cycle analysis. It also downregulated Sp1, Mcl-1, Bcl-2, and caspase-3 but upregulated cleaved caspase-3, Bax, and PUMA. In immunohistochemical staining, Sp1 was overexpressed in melanoma tissues compared to normal skin tissues. Collectively, our data suggest that morin can induce apoptosis of melanoma cells by regulating pro-apoptotic and anti-apoptotic proteins through ROS, and may be a potential substance for treatment of melanoma.

**Keywords** Morin · Melanoma · Proliferation · ROS · Sp1 · Mcl-1

## Introduction

Cancer is a major cause of death with a very high mortality rate in both developed and developing countries (Jemal et al. 2011). Among various types of cancers, cutaneous melanoma is one of the most common types of skin cancers with high rate of metastasis and mortality (Hofmann et al. 2000). Cutaneous melanoma arises from melanocytes thus it is deeply pigmented. It is usually primarily diagnosed by presentation of the skin lesion. Cutaneous melanoma progression is associated with oxidative stress and various cell signaling pathways (Hambright et al. 2015).

Oxidative stress interferes with normal oxidation–reduction balance. It generates reactive oxygen species (ROS) that can damage cells and cause changes in the signaling process (Choi et al. 2014). ROS are byproducts of cellular metabolism and act as important signaling molecules. They are removed through antioxidizing defense mechanisms including the Keap1/Nrf2 pathway (Nguyen et al. 2009). Increase of ROS in cancer cells under oxidative stress can activate growth signaling process and cause more cell growth to result in proliferation of cancer cells. However, due to high dependency on ROS, cancer cells react more sensitive to the increase of oxidative stress, especially in melanoma cells (Cabello et al. 2012; Lee et al. 2012; Guterres et al. 2013). Accordingly, targeting cancer cells with substances that can increase ROS beyond certain threshold can selectively kill cancer cells while having little impact on normal cells (Lau et al. 2008).

Specificity protein 1 (Sp1) is a transactivation molecule in Sp/Krüppel-like factor (KLF) group (Briggs et al. 1986).

✉ Moon Kyun Cho  
mkcho2001@hanmail.net; mkcho@schmc.ac.kr

<sup>1</sup> Molecular Cancer Research, Soonchunhyang University College of Medicine, Cheonan 31151, Republic of Korea

<sup>2</sup> Department of Dermatology, Soonchunhyang University Hospital, Seoul 04401, Republic of Korea

Sp1 can lead to activation of various genes involved in cell proliferation, cell cycles, and cell apoptosis (Karlseder et al. 1996; Black et al. 2001; Deniaud et al. 2006). Furthermore, Sp1 has been shown to be expressed at high levels in various cancers including thyroid cancer, glioma, colorectal cancer, lung cancer, pancreatic cancer, gastric cancer, and breast cancer (Chiefari et al. 2002; Wang et al. 2003; Yao et al. 2004; Wang et al. 2007; Jiang et al. 2008; Guo et al. 2010; Guan et al. 2012; Hsu et al. 2012). ROS is related to adjustment of Sp1 and its downstream proteins (Wang et al. 2008; Gandhi et al. 2012; Jin et al. 2013). Mcl-1, a protein related to Sp1 (Choi et al. 2013), is essential for the survival of various cell lineage. It is frequently amplified in human cancers, making it special among Bcl-2 family members that contribute to cell survival (Opferman et al. 2003, 2005; Dzhagalov et al. 2007; Arbour et al. 2008; Dzhagalov et al. 2008; Beroukhim et al. 2010). Bcl-2 is also one of Sp1-related proteins associated with cell survival (Duan et al. 2005).

Morin (3,5,7,2',4'-pentahydroxyflavone) is a flavonoid derived from the Moraceae family for the first time. Morin acts as anti-oxidant and neuroprotectant (Noor et al. 2012; Ola et al. 2014; Du et al. 2016). Furthermore, it has been reported that morin can control inflammatory reactions, thus inhibiting carcinogenesis and cancer progression (Kawabata et al. 1999; Brown et al. 2003). Morin exhibits anti-cancer activities by regulating cell proliferation and apoptosis in lung cancer, colorectal cancer, breast cancer, leukemia, and other human cancers (Kuo et al. 2007; Jin et al. 2014; Lee et al. 2016; Sithara et al. 2017; Yao et al. 2017). However, whether morin has an impact on the progression of melanoma and underlying mechanism is not well known. In the present study, we investigated cellular responses, including cellular ROS levels, mitochondrial function, apoptosis, and Sp1 expression, in melanoma cells treated with morin.

## Materials and methods

### Tissue sample collection and preparation

Melanoma specimens were obtained from patients who had undergone surgery in the Department of Plastic and Reconstructive Surgery between December 2015 and November 2018. The institutional review board of Soonchunhyang University Seoul Hospital reviewed and approved this research protocol involving the use of tissue samples. Six melanoma tissues were obtained from three male and three female patients and were diagnosed by pathologist. Normal skin tissues were collected from the backs of six women who had breast reconstruction with a latissimus dorsi flap. For immunohistochemical analysis,

archival formalin-fixed, paraffin-embedded tissues were used.

### Cell culture

The human melanoma cell lines, G361 and SK-MEL-2, and human fibroblast cell line, KEL FIB, were purchased from the American Type Culture Collection (Manassas, VA, USA). These cells were cultured at 37 °C in a submerged monolayer in DMEM supplemented with 5% fetal calf serum, 100 U of penicillin/mL, and 100 µg of streptomycin/mL in a humidified air with 5% CO<sub>2</sub>.

### Cell viability assay

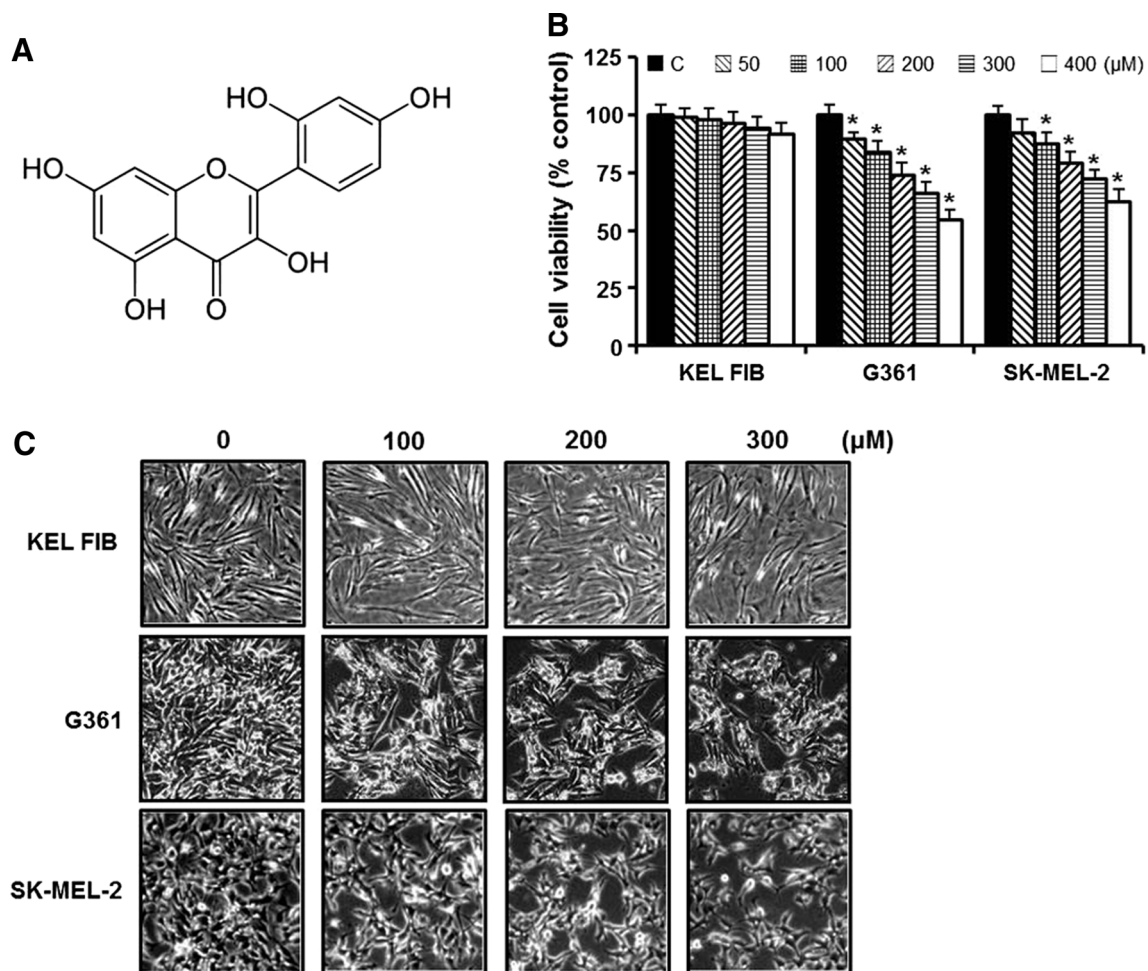
The cells were seeded into a 96-well microtiter plate at density of  $5 \times 10^3$  cells/well and then treated with 0, 50, 100, 200, 300, or 400 µM morin (BP0959; Biopurify Phytochemicals Ltd., Chengdu, China) for 24, 48, and 72 h, of which chemical structure is shown in Fig. 1a. After incubation, cells were reacted with tetrazolium dye (MTT, Sigma, St. Louis, MO, USA, final concentration: 0.1 mg/mL) for 4 h and mixed with 200 µL of dimethyl sulfoxide. Absorbance value at wavelength of 540 nm was measured with a GloMax-Multi Microplate Multimode Reader (Promega, Madison, WI, USA). The percentage of viable cells was determined by comparing absorbance values with those of vehicle-treated control cells (100%).

### Annexin V-PE binding assay

Apoptotic-cell distribution was determined with a Muse™ Annexin V & Dead Cell kit (MCH100105; Merck Millipore Co., Darmstadt, Germany) according to the manufacturer's protocol. The kit includes a fluorescent-dye phycoerythrin (PE) conjugated to Annexin V to detect phosphatidylserine on the external membrane of apoptotic cells and 7-AAD (7- amino-actinomycin D) as a dead cell marker. Briefly, trypsinized cells were harvested by centrifugation at 500×g for 7 min at 4 °C, mixed with Muse™ Annexin V and Dead Cell reagent, and analyzed using a Muse Cell Analyzer (Merck KgaA, Darmstadt, Germany).

### Measurement of intracellular ROS levels

Intracellular ROS levels were evaluated by measuring DCF-DA (Sigma) fluorescence intensity. Briefly, cells were stained with 10 µM DCF-DA for 30 min at 37 °C in the dark. Following incubation, cells were washed twice with 1x PBS, trypsinized, resuspended in 1x PBS, and immediately analyzed with a MACSQuant Analyzer flow cytometer and MACSQuantify™ software (Version 2.5;



**Fig. 1** Effects of flavonoid morin in KEL FIB, G-361, and SK-MEL-2 cells **a** Chemical structure of morin [2-(2,4-dihydroxyphenyl)-3,5,7-trihydroxy-4H-1-benzopyran-4-one]. **b** The percentage of cell viability was measured by MTT assay. Cells were treated with various concentration (0, 50, 100, 200, 300, 400  $\mu\text{M}$ ) of morin for 48 h. **c** Phase contrast images of KEL FIB, G361 and SK-MEL-2 treated with morin (0, 100, 200, 300  $\mu\text{M}$ ). The quantitative data were shown as mean  $\pm$  SD for three independent experiments. \* $P < 0.05$  compared to untreated controls

MiltenyiBiotec GmbH, BergischGladbach, Germany). DCF fluorescence was detected using a 530 nm bandpass filter. Each measurement was based on the mean fluorescence intensity of  $1 \times 10^4$  cells.

#### Mitochondrial membrane potential (MMP) disruption assay

Cells were seeded into six-well culture plates at density of  $5 \times 10^4$  cells/well and cultured for 24 h. Cells were then incubated with morin alone or in combination with *N*-acetylcysteine (NAC, Sigma) for another 48 h. These cells were then trypsinized, harvested by centrifugation at  $500 \times g$  for 7 min at  $4^\circ\text{C}$ , washed twice with  $1 \times \text{PBS}$ , and stained with serum-free DMEM medium containing Rhodamine 123 (Sigma, 30 nM) at  $37^\circ\text{C}$  for 30 min.

Fluorescence intensity was measured and analyzed using a MACSQuant analyzer flow cytometer and MACSQuantify<sup>TM</sup> software.

#### Cell cycle assay analysis

Percentages of cells in  $G_1$ , S, and  $G_2/M$  phases were measured by quantifying DNA content in PI-stained cells. Briefly, trypsinized cells ( $\sim 10^6$  cell/mL) were pelleted by centrifugation at  $500 \times g$  for 7 min at  $4^\circ\text{C}$ , fixed in 70% ice-cold ethanol overnight at  $-20^\circ\text{C}$ , and incubated with Muse<sup>TM</sup> Cell Cycle reagent (Merck Millipore, Billerica, MA, USA). Data from  $1 \times 10^4$  single-cell events were collected with a MACSQuant Analyzer flow cytometer and analyzed using MACSQuantify<sup>TM</sup> software (Version 2.5; MiltenyiBiotec GmbH, BergischGladbach, Germany).

## Western blot analysis

Total cell lysates were extracted with a  $1 \times$  RIPA buffer. Cell lysates containing 30  $\mu$ g of proteins were resolved on NuPAGE 4–12% bis-tris polyacrylamide gel and then electro-transferred onto a PVDF membrane. The membrane was probed with 1:500 diluted primary antibodies in casein blocking buffer at 4 °C for 24 h followed by incubation with 1:5000 dilution of secondary antibody coupled to horseradish peroxidase in casein blocking buffer at room temperature for 2 h. Signals were detected using an ECL detection kit. Membranes were then stripped and re-blotted with anti- $\beta$ -actin antibody (A2228; Sigma, St. Louis, MO, USA) as a loading control. The following primary antibodies were used: specific antibodies against Sp1 or Caspase-3 were purchased from Santa Cruz Biotechnology (Dallas, TX, USA), and specific antibodies against Mcl-1, Bcl-2, Cleaved Caspase-3, Bax, and PUMA were purchased from Cell Signaling Technology (Danvers, MA, USA). Secondary antibodies used were HRP-conjugated anti-rabbit IgG, anti-goat IgG, and anti-mouse IgG, and these were purchased from Santa Cruz Biotechnology (Dallas, TX, USA).

## Small interfering RNA (siRNA) transfection

RNA interference assay was performed using an Sp1-targeting siRNA (HSS110105, HSS110106, HSS186048; Invitrogen, Carlsbad, CA, USA). Briefly, cells were seeded onto six-well and 96-well plates, and then transfected at 40% confluency with the Sp1-targeting siRNA duplex or the Stealth RNAi negative control duplex (12935-200; Invitrogen, Carlsbad, CA, USA) using lipofectamine RNAi MAX (Invitrogen, Carlsbad, CA, USA). They were processed for western blotting.

## Immunohistochemical analysis

For immunohistochemical (IHC) analysis, four micron paraffin sections were prepared and stained according to the avidin–biotin staining technique. Briefly, slides were deparaffinized and cleared in xylene, rehydrated with ethanol, and placed in citrate buffer (0.01 M, pH 6.0) solution, with subsequent heating using ultrashort wave for 15 min. The sections were incubated overnight at 4 °C with anti-Sp1 antibody (1:500), followed by incubation in biotin-labelled secondary antibody for 30 min and then streptavidin-peroxidase for another 30 min. The staining was visualized using 3,3'-diaminobenzidine tetrahydrochloride. Negative controls were stained at the same time and in the same way, substituting the primary antibody with PBS. Assessment of IHC was scored as 0, 1 + (mild), 2 + (moderate), and 3 + (strong)

## Results

### Effects of morin treatment on fibroblast and melanoma cells

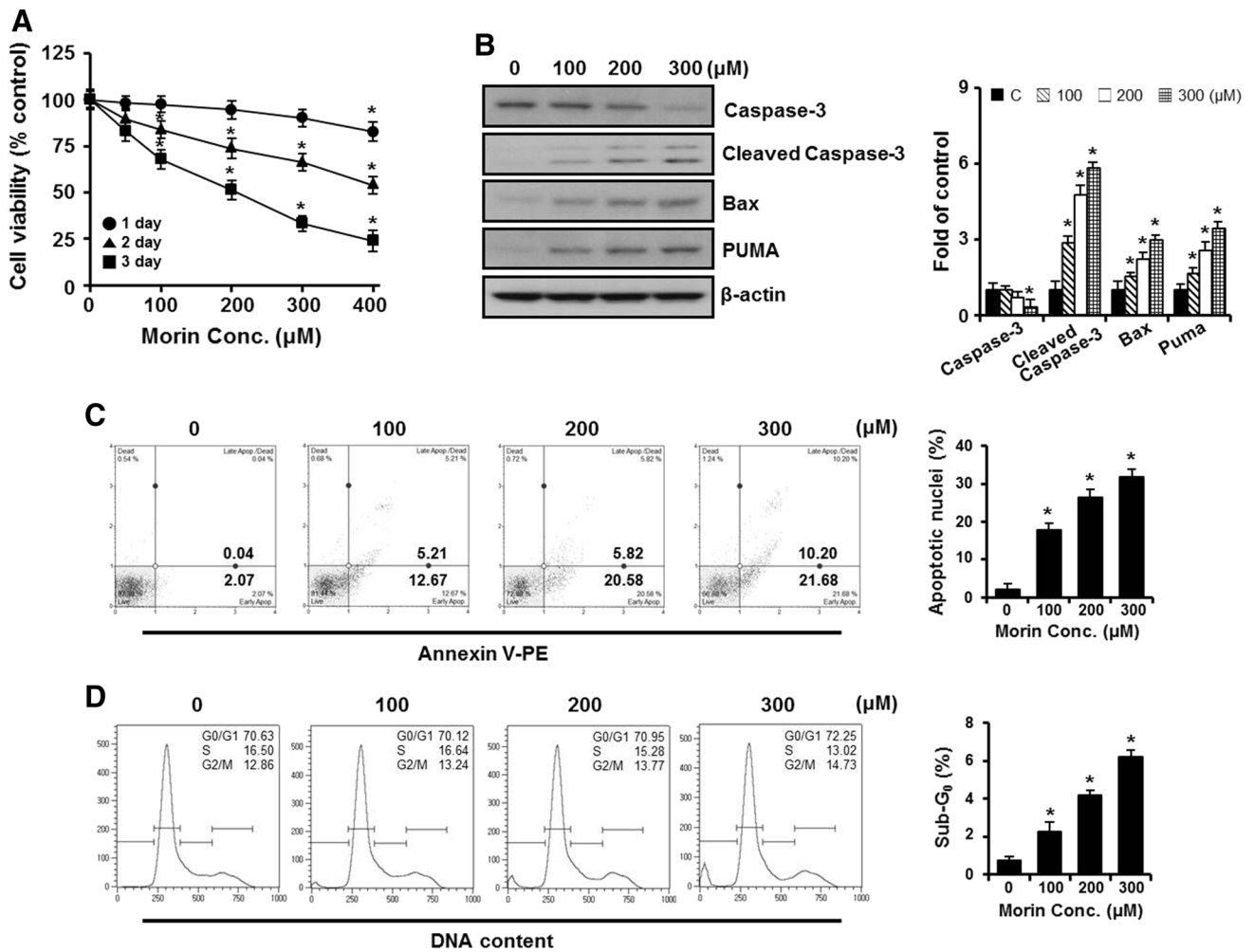
To determine whether morin shows cancer cell-specific cytotoxicity in melanoma cells, both cell viability and morphology was analyzed by MTT assay and phase contrast images, respectively, in KEL FIB cells and melanoma cells, including G361 and SK-MEL-2 cells. As shown in Fig. 1b, the viability of G361 and SK-MEL-2 cells decreased dose-dependently in response to different concentrations (50, 100, 200, 300 and 400  $\mu$ M) of morin, while the viability of KEL FIB cells didn't change significantly. Furthermore, at post 48 h treatment with 0, 100, 200, and 300  $\mu$ M of morin, the morphology of melanoma cells changed dramatically and some cells appeared to be detached compared to KEL FIB cells (Fig. 1c). These results suggest that morin may have preferential cytotoxicity on melanoma cells.

### Apoptotic effects of morin treatment on G361 cells

We investigated whether effects of morin treatment on cell viability were associated with apoptosis. As shown in Fig. 2a, cell viability of G361 cells was decreased by treatment of morin in time and dose-dependent manner. Then, we experimented to figure out changes of apoptosis-associated proteins by western blotting. The results showed that levels of cleaved caspase-3, Bax, and PUMA were increased while level of caspase-3 was decreased, as the concentration of morin was increased (Fig. 2b). The proportion of cells showing early and late phases of apoptosis was increased as concentration of morin increased (Fig. 2c). Cell cycle analysis demonstrated that the amount of sub- $g_0/g_1$  phase cells meaning apoptosis was increased with rising concentrations (100, 200, and 300  $\mu$ M) of morin (fig. 2d). With the increasing morin concentrations, the percentage of cells in  $G_2/M$  phase was increased, meaning a delay in  $G_2$  to M phase transition, and those in  $G_1$  and S phases were decreased compared to untreated controls. These findings suggest that morin may inhibit cell growth and promote apoptosis of G361 cells.

### Effects of morin treatment on oxidative stress of G361 cells

To determine whether apoptotic effects of morin treatment was correlated with oxidative stress, cells were treated with morin and intracellular ROS levels were analyzed by flow cytometry using DCF-DA. In the representative histogram, treatment with 0, 100, 200, and 300  $\mu$ M of morin for 48 h increased the production of ROS to 2.20, 15.57, 25.54, and



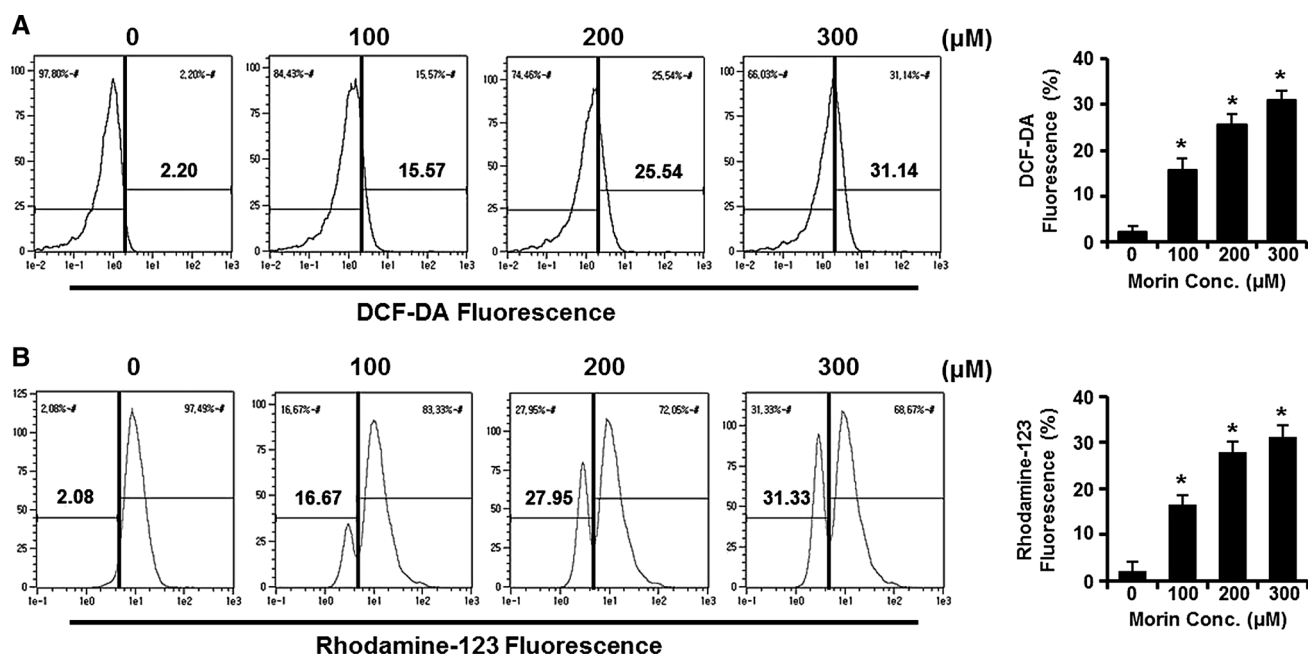
**Fig. 2** Apoptotic effects of morin on G361 cells. Cells were treated with various concentration (0, 100, 200, 300, 400 µM) of morin for 48 h. **a** The percentage of cell viability was measured by MTT assay. **b** The levels of apoptosis-associated proteins were measured after morin treatment by western blotting. β-actin was used as a loading control. **c** The percentage of apoptotic cells after Annexin V-PE binding was analyzed using a Muse cell analyzer. **d** Cell distribution in G<sub>0</sub>/G<sub>1</sub>, S, and G<sub>2</sub>/M phases was analyzed using flow cytometry after staining with propidium iodide (20 µg/mL). The quantitative data were shown as mean ± SD for three independent experiments. \*P < 0.05 compared to untreated controls

31.14%, respectively (Fig. 3a). Mitochondrial transmembrane potential ( $\Delta\Psi_m$ ) was also measured by flow cytometry using fluorescent dye rhodamine-123. As shown in Fig. 3b, morin caused a dose-dependent decrease in the number of cells with intact  $\Delta\Psi_m$ . The percentage of cells with  $\Delta\Psi_m$  loss increased 2.08, 16.67, 27.95, and 31.33% when morin concentration used for treatment was 0, 100, 200, and 300 µM, respectively. These findings suggest that the stability of  $\Delta\Psi_m$  is significantly impaired by morin due to increased ROS levels.

**Apoptotic effects of reactive oxygen species following morin treatment on G361 cells**

To determine whether apoptosis was caused by ROS in G361 cells, cells were treated with morin and ROS

scavenger N-acetylcysteine (NAC), alone or in combination. As shown in Fig. 4a, western blot analysis demonstrated that morin treatment decreased expression levels of Sp1, Mcl-1, Bcl-2, and caspase-3 but increased expression levels of cleaved caspase-3, Bax, and PUMA in G361 cells. However, morin-induced changes of protein expression were effectively reversed in the presence of NAC. In addition, pretreatment of cells with NAC resulted in significant improvement of cell viability (Fig. 4b). Changes of cell morphology were also reduced based on phase contrast images (Fig. 4c). NAC pretreatment also decreased the proportion of cells undergoing apoptosis based on Annexin V-PE staining results (Fig. 4d) and the sub-G<sub>0</sub>/G<sub>1</sub> peak (Fig. 4e) compared to morin treatment alone. These results suggest that morin would lead to apoptosis of G361 cells



**Fig. 3** Quantification of ROS formation and loss of mitochondrial membrane potential by Rhodamine 123 staining. G361 cells were treated with various concentrations of morin (0, 100, 200, 300 μM) for 48 h. **a** After incubation, cells were stained with DCF-DA and analyzed using flow cytometer. Data analyzing fluorescence intensity from triplicate measurements. **b** After incubation, cells were stained with Rhodamine 123 analyzed using flow cytometer. Data analyzing fluorescence intensity from triplicate measurements. \* $P < 0.05$  compared to untreated controls

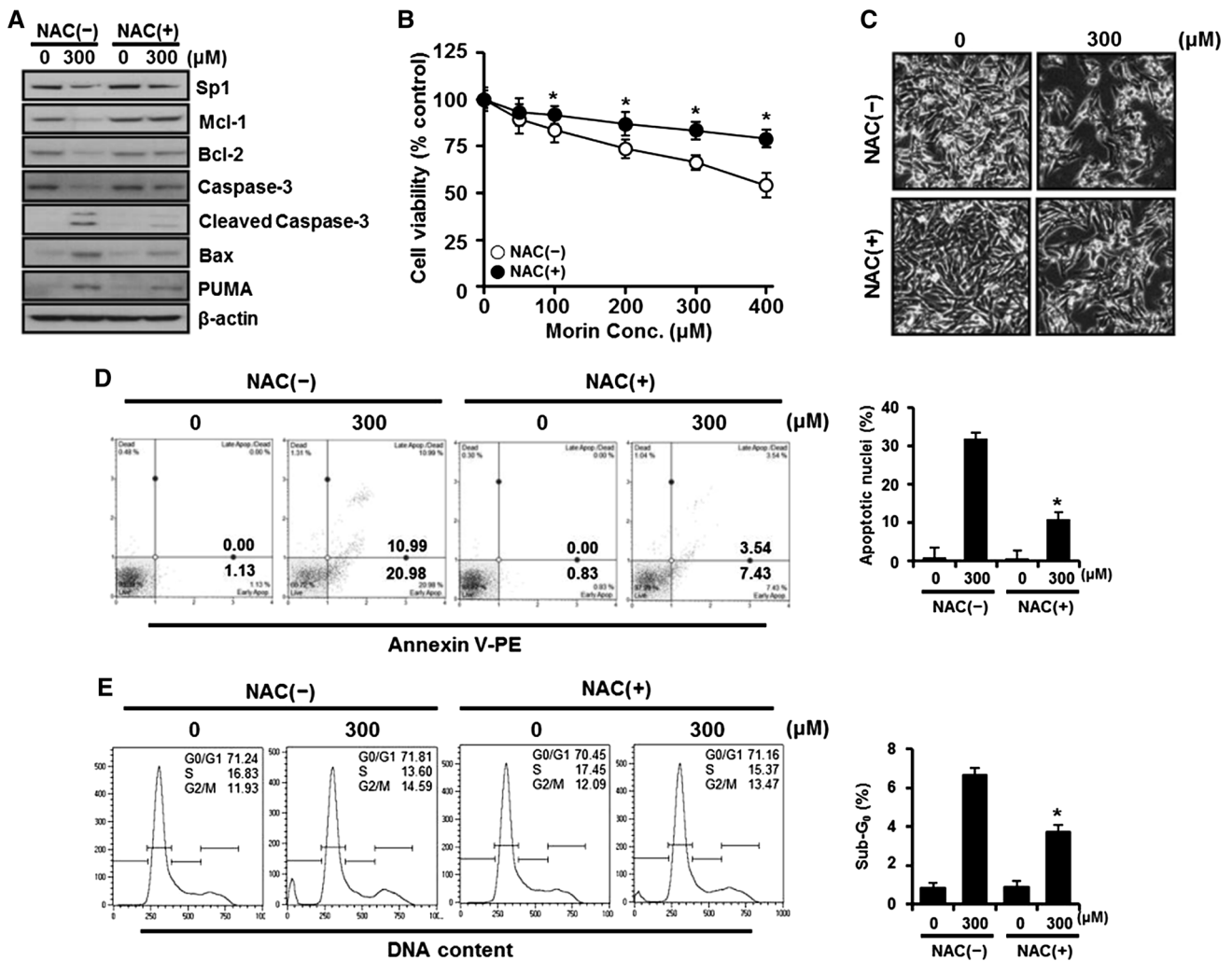
through ROS by regulating Sp1, Sp1-related proteins, and pro-apoptotic proteins.

#### Pro-oxidant effects of morin on G361 cells

To confirm that morin could act as a pro-oxidant in G361 cells, we used NAC to reduce intracellular ROS levels. After cells were treated with morin and NAC, intracellular ROS levels were analyzed by flow cytometry with DCF-DA. As shown in the representative histogram (Fig. 5a), treatment with morin at final concentration of 300 μM for 48 h produced ROS to approximately 30.93%. However, the accumulation of ROS was significantly attenuated by NAC during morin treatment to approximately 17.44%. The  $\Delta\Psi_m$  was also measured by flow cytometry with fluorescent dye rhodamine-123. As shown in Fig. 5b, the percentage of cells with  $\Delta\Psi_m$  loss increased from 1.90 to 34.37% when morin concentration was increased from 0 to 300 μM. However, pretreatment with NAC reduced it to approximately 1.76% and 14.34%, respectively. These findings suggest that ROS scavenging could prevent mitochondrial damage by protecting substantial loss of  $\Delta\Psi_m$  in morin-treated G361 cells, while the stability of  $\Delta\Psi_m$  was highly damaged by morin.

#### Effects of morin as a Sp1 inhibitor on melanoma

Suppression of Sp1 is known to activate apoptotic signals for cell death. Here, we examined the effect of morin as a Sp1 inhibitor on melanoma cells. Western blot analysis showed that levels of Sp1, Mcl-1, and Bcl-2 proteins were decreased in G361 and SK-MEL-2 cells while levels of those proteins were almost unchanged in KEL FIB cells, as the concentration of morin was increased (Fig. 6a). Next, the effects of morin on Sp1, Mcl-1, and Bcl-2 expression have been determined by western blotting following Sp1 silencing in G361 and SK-MEL-2 cells. Expressions of Sp1, Mcl-1 and Bcl-2 proteins were downregulated by treating Sp1-targeting siRNA in both cell types; however, these findings were recovered by NAC pretreatment (Fig. 6b). To investigate the association between Sp1 and cell viability, we examined whether Sp1 knockdown affects sensitivity to morin treatment. When cells transfected with Sp1-targeting siRNA were treated with morin, the viability of G361 and SK-MEL-2 cells was reduced to approximately 18.7% and 16.1%, respectively, in morin-untreated cells, and approximately 18% and 14.9%, respectively, in morin-treated cells, compared with that of respective control siRNA-treated cells (Fig. 6c). Next, we examined whether a decrease in the cell viability following Sp1 knockdown is associated with ROS. As shown in Fig. 6d, the increased ROS levels by morin were further

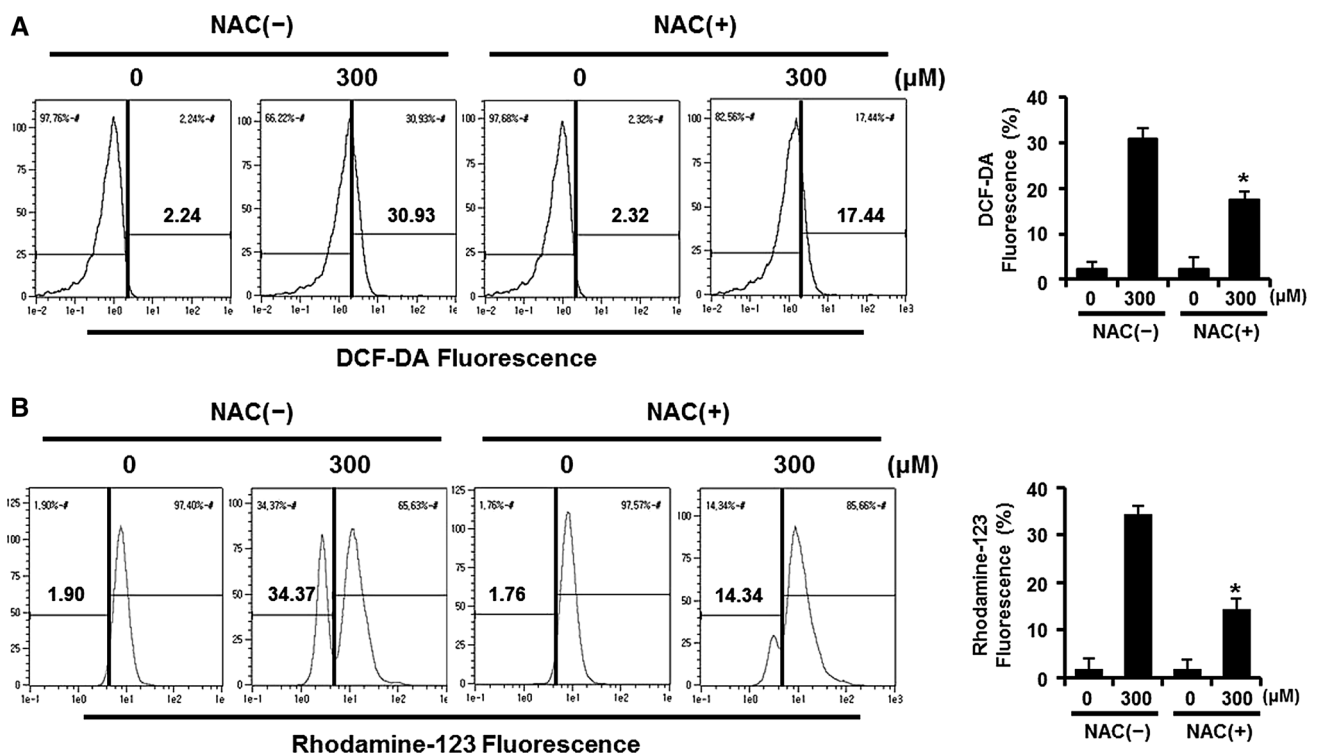


**Fig. 4** Apoptotic effects of reactive oxygen species following morin treatment on G361 cells. Cells were pretreated with 5 mM NAC for 1 h prior to exposure to 0 or 300 µM concentrations of morin for another 48 h. **a** The levels of Sp1, anti-apoptotic and pro-apoptotic proteins in total cell lysates were measured by western blotting. **b** The percentage of viable cells was measured by MTT assay. **c** Phase contrast images of cells treated with morin. **d** The percentage of apoptotic cells after Annexin V-PE binding was analyzed using a Muse cell analyzer. **e** Cell distribution in G<sub>0</sub>/G<sub>1</sub>, S, and G<sub>2</sub>/M phases was analyzed using flow cytometry after staining with propidium iodide (20 µg/mL). The quantitative data were shown as mean ± SD for three independent experiments. \**P* < 0.05 compared to untreated controls

elevated by Sp1 knockdown in both G361 and SK-MEL-2. To confirm overexpression of Sp1 in melanoma tissues, Sp1 proteins was examined by immunohistochemistry in the paraffin sections of both six human melanoma tissues and six normal skin tissues. As shown in Fig. 6e, Sp1 was overexpressed significantly in melanoma tissues compared to normal skin tissues. Four melanoma tissues showed 3 + (strong) positive staining of Sp1 while two melanoma tissues demonstrated 2 + (moderate) positive staining of Sp1. However, all of normal skin tissues showed 1 + (mild) positive staining of Sp1. Taken together, these findings demonstrate that morin may induce apoptosis through downregulation of antiapoptotic Mcl-1 and Bcl-2, which is regulated, at least in part, by the ROS-linked suppression of Sp1 expression in melanoma cells.

### Discussion

Oxidative stress is induced by imbalance between oxidative substances and antioxidizing defense mechanisms. Oxidative stress is shown to be related to various skin diseases including skin cancer (Klaunig et al. 1995; Marrett et al. 2003). Melanoma, one of the most serious types of skin cancers, is related to oxidative stress (Hambright et al. 2015). It has been reported that oxidative stress was increased in melanoma cells through various mechanisms (Sander et al. 2003). For example, an increase in superoxide anion and a decrease in hydrogen peroxide have been found in melanoma cells. This environment makes the cell easier for oxidation. In addition, the signaling pathway that strengthens the aggressiveness of melanoma cells



**Fig. 5** Pro-oxidant effects of morin on G361 cells. Cells were pretreated with 5 mM NAC for 1 h prior to exposure to 0 or 300  $\mu\text{M}$  concentrations of morin for another 48 h. **a** The levels of cellular ROS were measured using flow cytometry after staining with DCF-DA (10  $\mu\text{M}$ ). A shift of DCF fluorescence to the right indicates an increase in ROS. **b** The levels of  $\Delta\Psi\text{m}$  were measured using a flow cytometry after staining with rhodamine 123 (30 nM). \* $P < 0.05$  compared to untreated controls

characterized by high proliferation speeds and drug resistance is activated (Pervaiz and Clement 2007). However, there is an intriguing report that while increased ROS can contribute to carcinogenesis by causing DNA damage, ROS levels beyond a threshold can cause cell aging and cell death, therefore functioning as an anti-tumor substance (Reuter et al. 2010). Hence, substances which induce ROS can also increase oxidative stress in cancer cells, leading to activation of apoptotic signals.

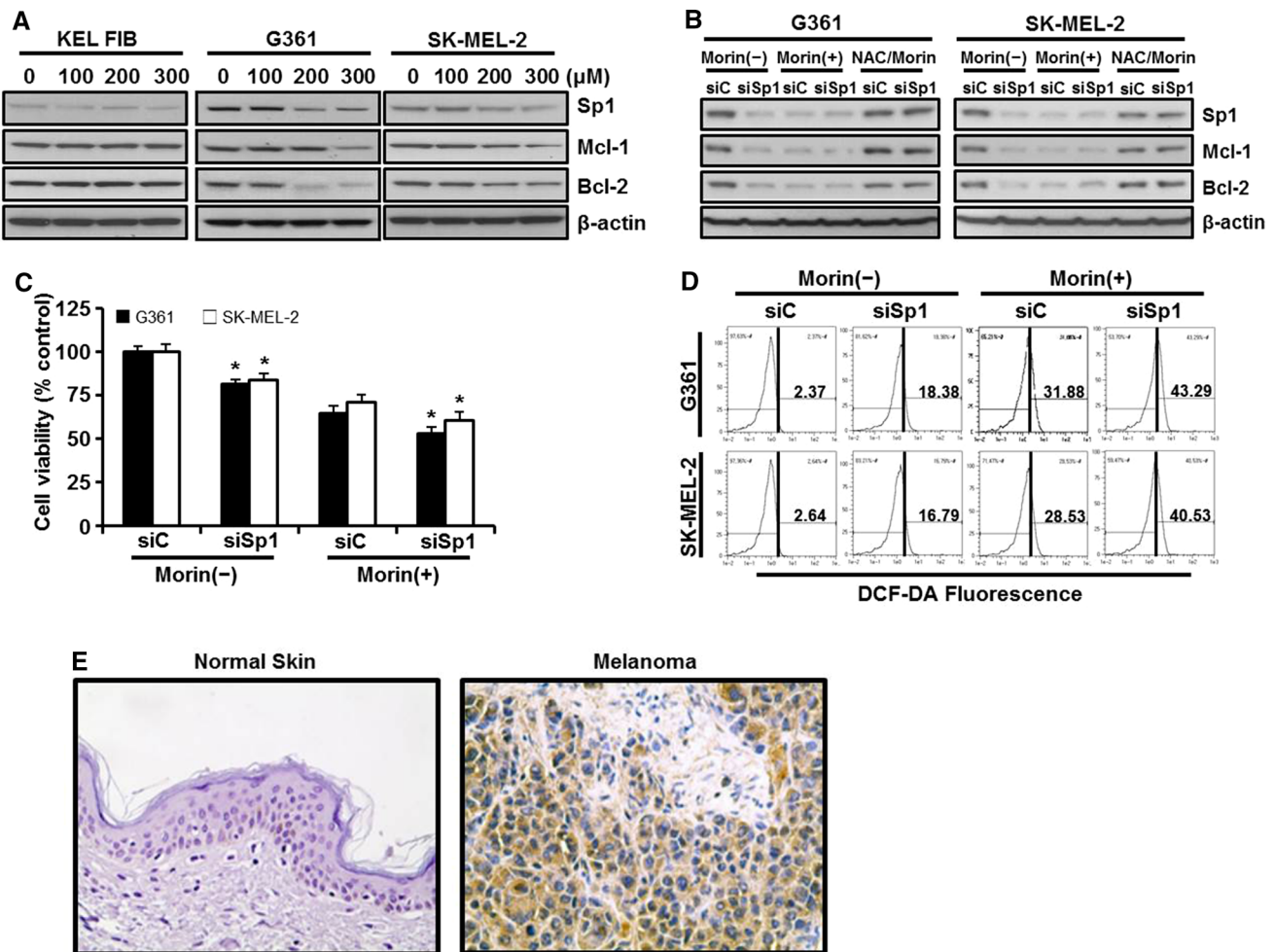
Morin is originally produced from the Moraceae family such as mulberry figs and other Chinese herbs. It has also been discovered as a yellow pigment from almond hulls and old fustic (Lee et al. 2008). Morin is a flavonoid substance having structure of [2-(2, 4-dihydroxyphenyl)-3,5,7 trihydroxy-4H-1-benzopyran-4-one]. It is widely used in herbal medicines for its biological activities such as anti-inflammation, antioxidizing, and anti-cancer effects (Kim et al. 1993; Lee et al. 1993; Sivaramakrishnan and Niranjali Devaraj 2009; Jin et al. 2014). It has been revealed that morin induced apoptosis in various human cancer cells including lung cancer, colorectal cancer, breast cancer, and leukemia (Kuo et al. 2007; Jin et al. 2014; Lee et al. 2016; Sithara et al. 2017; Yao et al. 2017).

Due to limited information, it was very difficult to determine quantitative levels of the cellular morin that can

be achieved in vitro conditions versus in vivo situation. However, recent studies have shown that cancer-specific apoptosis occurred after treating high concentration of morin in breast cancer cells (Jin et al. 2014), whereas any significant changes in cell viability did not be observed in normal lung fibroblasts, even though morin was treated up to 500  $\mu\text{M}$  (Lee et al. 2017). Consistently, our data in melanoma cells and fibroblasts support the previous results, indicating that morin has a selective cytotoxic effect on malignant cells. However, large well-designed clinical studies are required to further elucidate the effect of morin.

Our results suggest that apoptosis signaling activated by morin treatment may be associated with increased ROS levels and mitochondrial damage in human melanoma G361 cells. Although there has been no report to confirm relationship between ROS generation and anti-cancer effects of morin in melanoma cells, some papers reported that morin causes cell death by increasing cellular ROS levels in human leukemia cells and colorectal cancer cells (Kuo et al. 2007; Hyun et al. 2015; Sithara et al. 2017). Here, our data underline an importance of Sp1 as an ROS-regulating molecule. Sp1 related to cell proliferation and cell survival contributes to cancer formation (Prathyusha et al. 2017). This rationale is supported by our findings that Sp1 was more expressed in melanoma tissues than in





**Fig. 6** Expression of Sp1 in melanoma cells and tissues. **a** The levels of Sp1, Mcl-1, and Bcl-2 proteins were measured following morin treatment by western blotting in KEL FIB, G361, and SK-MEL-2 cells. Strongly positive staining in melanoma tissue (IHC stain, × 400). **b** G361 and SK-MEL-2 were transfected with 10 nM of Sp1-targeting siRNA (siSp1) or Stealth RNAi control (siC) for 24 h, after which cells were with either morin (300 μM) or NAC (5 mM) for 48 h. The levels of Sp1, Mcl-1, and Bcl-2 proteins were measured by western blotting. **c** The percentage of viable cells following treatment of either Sp1-targeting siRNA or morin was measured by MTT assay. **d** The levels of cellular ROS were measured in melanoma cells treated with either Sp1-targeting siRNA or morin by using flow cytometry after staining with DCF-DA (10 μM). **e** Representative immunohistochemical staining for Sp1 protein expression in paraffin-embedded normal skin and melanoma tissue. Weakly positive staining in normal skin (IHC stain, × 400). The quantitative data were shown as mean ± SD for three independent experiments. \* $P < 0.05$  compared to untreated controls

normal skin tissues, and that Sp1 knockdown decreased the viability of G361 and SK-MEL-2 cells, suggesting that overexpression of Sp1 may contribute to melanoma formation. So far, no in vivo studies have been performed to confirm expression of Sp1 protein in melanoma tissues. Furthermore, our findings that morin-induced cytotoxicity was accompanied by the down-regulation of antiapoptotic proteins, including Sp1, Mcl-1 and Bcl-2, and the up-regulation of pro-apoptotic proteins, including cleaved caspase-3, Bax and PUMA, in G361 cells might be at least in part explained by ROS-induced pro-oxidant effects. This possibility is supported by pretreatment with antioxidant NAC, which significantly recovered a series of morin-induced changes such as cell viability, the levels of apoptosis

promoting and apoptosis-inhibiting proteins, cellular ROS levels, and  $\Delta\Psi_m$ . Mcl-1, a protein that inhibits apoptosis, has promoter rich in GC that can directly bind to Sp1 (Choi et al. 2013). Bcl-2, a well-known protein that can inhibit apoptosis, also has promoter that can bind to Sp1 (Duan et al. 2005). In this study, the reduced viability of melanoma cells and the downregulation of Mcl-1 and Bcl-2 expression following Sp1 knockdown suggest that Sp1 may contribute to cell apoptosis and act as a transcriptional regulator of Mcl-1 and Bcl-2. Considering that ROS is a trigger factor for intrinsic apoptosis through interaction with mitochondrial permeability transition complex proteins (Tsumimoto and Shimizu 2007; Bajpai and Nagaraju 2017), dysregulation of ROS by morin treatment in G361

cells can promote mitochondrial apoptosis by downregulating Sp1 and Sp1-related proteins such as Mcl-1 and Bcl-2. Conversely, Sp1-linked molecular pathway may be essential for cell survival in melanoma.

Interestingly, our data demonstrated increased ROS production following downregulation of Sp1 in both G361 and SK-MEL-2 cells. These results suggest that Sp1 may protect cells from oxidative stress. In consistent with our results, there are some reports that Sp1 could attenuate ROS generation to increase cell survival, while downregulation of Sp1 could increase intracellular ROS levels. The upregulation of Sp1 increased Zinc finger protein 179 (Znf179) through the transcriptional activation, which protected cells from ROS-induced damage via upregulation of antioxidant proteins, including peroxiredoxin III (Prx3) and superoxide dismutase 2 (SOD2) (Chuang et al. 2017). Moreover, Sp1 is also known as a transcription activator of SOD1 representing a protection against NO toxicity via scavenging the radical superoxide (Baldelli et al. 2008).

Taken together, we observed that morin-induced prooxidant environment led to a series of cancer cell-specific apoptotic process in melanoma cells, as evidenced by increased proportion of cells with Annexin V-PE(+) staining and sub-G<sub>0</sub>/G<sub>1</sub> peak in cell cycle analysis, the downregulation of Sp1, Mcl-1, and Bcl-2, and upregulation of cleaved caspase-3, Bax, and PUMA. Despite the fact that future research, including large well-designed clinical studies, is warranted, morin may be a potential candidate substance in treatment of melanoma by activating apoptosis pathway through ROS-mediated damage and Sp1 suppression in vitro.

#### Compliance with ethical standards

**Conflict of interest** The authors declare no conflict of interest.

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