



Roscovitine inhibits glycogen synthase kinase 3 beta signaling and exerts apoptotic effect with an increase in reactive oxygen species generation in neuroblastoma cells

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Abstract

Roscovitine (ROSC) is a selective cyclin-dependent kinase (CDK) inhibitor against CDK2, 7 and 9. ROSC's anti-proliferative and anti-cancer activities have been well-documented in both *in vivo* and *in vitro* studies against several cancer types. Glycogen synthase kinase 3 (GSK3) is a serine/threonine protein kinase that has a role in the regulation of glycogen synthase. It also has a role in multiple cellular processes and disease conditions. A member of the GSK3 family, GSK3 β , has been implicated in many human malignancies including neuroblastoma. The specific inhibition of GSK3 β reduced neuroendocrine markers and suppressed neuroblastoma (NB) cell growth. NB is a malign pediatric disease with diverse types of tumors and high heterogeneity. Lately, GSK3 β targeted therapy models are being investigated for NB therapy. The action of ROSC on GSK3 β , however, is not fully understood. In this study, we showed that ROSC exerts anti-proliferative and apoptotic activity in SK-N-AS neuroblastoma cells by increasing reactive oxygen species (ROS) generation, which can be prevented by N-acetyl-cysteine administration. ROSC treatment inhibited GSK3 β signaling by promoting Ser9 inhibitory phosphorylation. ROSC at low doses can be a drug candidate to modulate GSK3 β signaling in NB cells.

Keywords: Neuroblastoma, roscovitine, cell cycle, Apoptosis, GSK3 β

1. Introduction

As an extracranial solid tumor, NB is the second most frequent cancer in children, indicated as the neural crest-derived malignancy of the peripheral nervous system. NB might manifest anywhere along the sympathetic nervous system; however, most tumors are in the adrenal gland medulla and the abdomen (1–3). NB attributes disproportionately, significant morbidity and mortality can be seen among children, or sometimes they can develop into a benign ganglioneuroma, and even a spontaneous and complete regression can be detected (4,5). Therefore, NB is remarkable because of its heterogeneity. Disseminated metastases are present in nearly 75% of cases in children over the age of 1. Stage IV usually represents an aggressive tumor resistant to chemotherapy and is generally untreatable. The molecular mechanisms lying behind the NB are usually somatic genetic alterations, which lead to overexpression of oncogenes and inactivation of tumor suppressor genes. For instance, 20% to 25% of NB cases occur due to the amplification of MYCN proto-oncogene (6). Lately, anaplastic lymphoma kinase (ALK) was also identified as an oncogene related to NB diagnosis (7). Besides, deletions in

chromosomes 1p and 11q have also been detected and constitute a poor prognosis in patients (8,9). Recently, altered GSK3 (glycogen synthase kinase) expression was observed in NB patients due to its role in MYCN regulation (10). Studies indicated that GSK inhibitors can induce cell death in cancer cells via apoptosis (11–13). Duffy et al. showed that GSK3 inhibitor-treated NB cells exhibited altered multiple signaling pathways that contributed to the loss of cell viability (10). In addition, a GSK3 inhibitor, lithium chloride, successfully induced cancer cell death without causing harmful effects to epithelial cells (14,15).

Glycogen synthase kinase 3 β (GSK3 β) is a critical serine/threonine kinase in glycogen metabolism (16). There are two isoforms in mammals, GSK3 α and β , weighing 51 and 47 kDa, respectively. In addition, GSK3 β is also highly expressed in the central nervous system and stabilizes microtubule-associated proteins via phosphorylation (17). For this reason, GSK3 β deprivation causes fatal consequences in the embryonic stage. GSK3 β activity is tightly regulated in cells, especially from the Serin 9 and Serin 21 residues at the N-

terminus of the protein. Upon the inhibitory phosphorylation from these residues, the release of axin and β -catenin from the complex occurs, leading to the translocation of β -catenin to the nucleus. β -catenin, a transcription factor, migrates to the nucleus and coordinates the expression of essential factors in neuron survival, such as c-Jun (18). In addition, the Wnt signaling pathway inhibits GSK3 β activity by causing phosphorylation of the Serine (Ser) 9 residue (19). With this regulation, GSK3 β is involved in many cell signaling pathways, such as cell adhesion, division, transcription mechanism, and tau phosphorylation. Therefore, GSK3 β is associated with many pathological conditions as a molecular target. In addition to the phosphorylation of β -catenin in the sub-signaling pathways, it plays a decisive role in cell survival-death fate with its effects on c-Jun and interacting with IRS-1 (20). GSK3 β is an essential factor for neuronal cell survival. However, excessive GSK3 β activity can cause cell death for many cell types and neurons (21).

Among the 538 human kinases, cyclin-dependent kinases (CDKs) were examined in detail since they have significant roles in cellular processes such as cell division and cell death (22,23). The cell cycle process is a remarkably conserved phenomenon in eukaryotes, and it is highly regulated to provide proper cell division. As said before, CDKs have critical functions as cell cycle regulators working with other associated cyclins to arrange the cell cycle progression. CDK1, CDK2, CDK4, and CDK6 are essential for DNA replication, mitotic progression, and growth regulatory signal responses; however, transcriptional regulation of CDK7, CDK8, and CDK9 also plays a role (24). Abnormal activation of CDKs leads to abnormal cell cycle progression and tumorigenesis. Therefore, pharmacological inhibitors of CDKs are attracting attention as potential anti-cancer agents, encouraging scientists to optimize and characterize, revealing a new strategy for cancer therapy (23).

Roscovitin (ROSC) is a CDK inhibitor, a purine derivative that inhibits CDK1, CDK2, CDK5, CDK7, and CDK9. However, it is an inhibitor with weak binding to CDK4, CDK6, and CDK8 (25). It shows an affinity for the ATP binding cleft of the kinase in competition with ATP. This binding in the catalytic region occurs by direct co-crystallization with CDK2 (26). It is a synthetic inhibitor studied in clinical studies as a candidate drug for some oncogenic indications (27). ROSC's anti-tumor activity in various cancer types was demonstrated on multiple cell lines (28). The therapeutic efficacy of ROSC against neurodegenerative diseases, cardiovascular disorders, viral infections, and parasitic protozoa was investigated at the preclinical level (29). In addition, ROSC is used for macrophage function and killing multi-drug-resistant bacteria (30). Although there is a wide variety of protein kinase inhibitors, the search for their optimization and characterization continues and offers promising results. In addition, ROSC exerts a synergistic effect with anti-cancer agents such as doxorubicin, taxol, 5-fluorouracil, vinblastine,

alemtuzumab, paclitaxel, trastuzumab, cisplatin, radiation, irinotecan, etoposide, and tamoxifen (31). Phase I and II trials of ROSC were carried out in various cancer treatments and tested in monotherapy and combination therapies (32). ROSC treatment induced cell cycle arrest and apoptosis in many cancer types. This process affects many pathways, including RAS-MAPK, NF- κ B, p53, estrogen receptor, and JAK-STAT signaling pathways (31). In addition to *in vitro* experiments, *in vivo* analyses were carried out, and the effects of ROSC in mono and combined therapies were investigated in various mouse and rat models (33). Studies have shown that Tau protein has serine and threonine phosphorylation sites targeted by protein kinases such as GSK3 β and CDK5 (34). ROSC, a CDK inhibitor, is also seen as a potential GSK3 β inhibitor. Inhibition of CDK5 by ROSC has neuroprotective effects in the modulation of Tau / GSK3 β and ERK / PPAR γ / CREB signals in neuronal damage and cognitive dysfunction (35). In addition, the ROSC effect in GSK3 β signaling was detected in Human T-cell leukemia virus type I (HTLV-I) cells (36). It is also known that the therapeutic agent ROSC shows GSK3 β activity in combined applications in the treatment process of Alzheimer's disease (AD) (37). In light of the above literature data, this study aims to understand the anti-cancer properties of ROSC as a CDK and potential GSK3 β inhibitor.

2. Materials and methods

2.1. Cell Culture

The NB cell line SK-N-AS (CRL-2137) was obtained from the American Type Culture Collection (ATCC). Cells were grown in DMEM (Dulbecco's Modified Eagle Medium) medium containing 10% fetal bovine serum, 10 U/ml penicillin/streptomycin antibiotic supplement, and 0.1 mM non-essential amino acid and incubated 5% CO₂ incubator at 37 °C (Heracell 150; Thermo Electron Corporation, Waltham, MA, USA).

2.2. MTT Assay

Cells were seeded at 1×10^4 density per well in 96-well plates and exposed to 1 and 10 μ M ROSC for 24 h. 10 μ l 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide dye (5 mg/ml) (Roche, Indianapolis, USA) was added to each well and cells were kept at 37°C for 4 h. The resulting formazan crystals were solubilized in 200 μ l dimethyl sulfoxide (DMSO). The density of the solubilized formazan was read at 570 nm spectrophotometrically (Bio-Rad, Hercules, CA, USA) (38).

2.3. Trypan Blue Dye Exclusion Assay

Cells were seeded at 1×10^5 density in 6-well plates (TPP, Zollstrasse, Switzerland) and treated with ROSC dose-dependently within 96h. After trypsinization (Trypsin EDTA (0.25%), Gibco, USA), and centrifugation, cells were exposed to 0.4% (w/v) Trypan Blue (Gibco -Life Technologies, USA) and cell culture media at a 1:1 ratio. 10 μ l of cells were counted by a dual-chamber 0.1 mm deep Neubauer improved hemocytometer (39).

2.4. Clonogenic Assay

Cells were seeded at 1×10^4 density into 6-well plates and treated with increasing concentrations of ROSC for 24 h. Then, the media was removed, and cells were washed with 1X PBS fixed with methanol:acetic acid (3:1) for 5 min. Following the removal of the fixing agent, cells were stained with 0.5% crystal violet in methanol for 30 min, washed with distilled water, and the images were taken under light microscopy (40).

2.5. Soft agar colony formation assay

A mixture containing a 1:1 ratio of DMEM (20% FBS and 2% penicillin/streptomycin) and 1% agar was prepared for soft agar analysis. The resulting mixture was distributed as 1 ml per well on 6-well plates and incubated until solidification. Later, 2.5×10^5 cells (untreated or treated samples) were added to each well with 0.5 % agar and were applied at a 1:1 ratio with 2X DMEM medium. Samples were incubated at 37 °C for 15 days and visualized under the light microscope (Olympus, IX70) (41).

2.6. Fluorescence Microscopy

2.6.1. Propidium iodide staining

SK-N-AS cells (1×10^5) were seeded into 12 well plates and treated with ROSC (0-30 μ M) for 24 h. Following a dose-dependent ROSC treatment, the cells were washed once with 1X PBS and stained with propidium iodide (PI) (50 mg/ml stock concentration in 1X PBS) fluorescent probe and were incubated for 10 min in the dark. ROSC-induced cell death was detected with fluorescence microscopy (Olympus, Tokyo, Japan) (42).

2.6.2. 3,3'-Dihexyloxycarbocyanine iodide (DiOC6) staining

SK-N-AS cells were seeded at a density of 1×10^5 /well into 12 well plates. Following exposure of cells to ROSC (0-30 μ M), they were washed once with 1X PBS and then stained with 4 nM 3,3'-dihexyloxycarbocyanine iodide (DiOC6) (40 nM stock concentration in DMSO; Calbiochem, La Jolla, CA, USA) fluorescent probe. Mitochondrial membrane potential (MMP) disruption was visualized by fluorescence microscopy (Zeiss, AX10) (excitation/emission: 482/ 504 nm) (43).

2.6.3. 4',6-diamidino-2-phenylindole (DAPI) staining

The cells were seeded in 12-well plates at 1×10^5 cells/well density and treated with purvalanol (0-30 μ M) for 24 h. After treatment, the cells were washed once with 1X PBS. The cells were stained with 1 μ l/ml 4',6-diamidino-2-phenylindole (DAPI) (1 mg/ml stock concentration in 1X PBS) fluorescent probe and were incubated for 10 min in the dark. Dose-dependent ROSC-induced nuclear DNA fragmentation was visualized using fluorescence microscopy (Zeiss, AX10) (excitation/emission: 350/ 465 nm) (44).

2.6.4. 2',7'-dichlorofluorescein-diacetate (DCFH-DA) staining

Fluorometric analysis of DCFH-DA staining

After drug administration in cell culture, cells were stained with DCFH-DA (1 μ M final concentration) and incubated at 37

°C for 10 min. Cells were washed with 1X PBS and analyzed with the Accuri C6 flow cytometer (BD Biosciences, Oxford, UK). Data analysis was performed with the BD Accuri C6 software (BD Biosciences) (excitation/emission: 495/529 nm) (45).

2.7. Protein isolation and immunoblotting

SK-N-AS cells were seeded in 60 mm² Petri dishes at a 5×10^5 density. Following the seeding overnight, they were exposed to increasing concentrations of ROSC for 24h. Later, cells were collected twice by scraping using 1X PBS and precipitated by centrifugation at 13,200 rpm for 1 min. After removing the supernatant, M-PER lysis buffer was added to the cells (Thermo Fischer Scientific). The samples were incubated for 20 min. at RT in a shaker and later were centrifuged at 13,200 rpm for 20 min at +4 °C. The supernatants were kept as total protein isolates at -80°C. PhosSTOP (Sigma Aldrich, Schnellendorf, Germany) was used during phosphorylated protein isolation. NE-PER Nuclear and Cytoplasmic Extraction protocol (Thermo Fisher Scientific) was followed for the nuclear and cytoplasmic protein isolation. The Bradford method was used for protein quantification, and bovine serum albumin (BSA) was used for the standard curve graphs. For the immunoblotting, isolated proteins were mixed with 4X Laemmli loading buffer, kept for 5 min at 95 °C, and 30 μ g protein samples were loaded into 12% acrylamide/bis-acrylamide gels. Following the running step, samples were transferred to polyvinyl fluoride (PVDF) membranes activated by methanol. Later, membranes were blocked using 5% skim milk prepared in 1X TBS containing 0.1% Tween 20 (TBS-T) at RT for 1h. After the blocking step, membranes were incubated at +4 °C with the selected primary antibodies at a ratio of 1:1000 (Cell Signaling Technology). Following washing steps with 1X TBS-T, membranes were incubated with secondary antibodies overnight. The secondary antibody application was followed by the washing step and subjected to a chemiluminescence solution. The signals from HRP-conjugated secondary antibodies were detected by the Chemidoc MP Imaging system (Bio-Rad Laboratories, Hercules, CA). All results were repeated at least three times, and the representative blots were given (39).

2.8. Flow cytometric analysis of apoptosis

2×10^5 cells/well were seeded in 6 well plates and treated with increased concentrations of ROSC for 24 h. Later, cells were collected following trypsinization and the cell pellet were collected and re-suspended in 300 μ l of binding buffer with RNase (100 μ g/ml), 3 μ l FITC-conjugated annexin V, and 3 μ l propidium iodide (PI, BD Biosciences, San Jose, CA, USA) obtained with FITC Annexin V Apoptosis Detection Kit I (BD Biosciences, Bedford, MA).

Samples were kept for 15 min at 37 °C in the dark. Later, the obtained samples were analyzed by BD Accuri C6 software (BD Biosciences) (46).

2.9. Cell cycle analysis

Cells were seeded in six-well plates with a density of 5×10^5 cells/well; following 24 h 1 and 10 μM of ROSC treatment, drug-containing media were discarded, and the cells were collected by trypsinization. Following a centrifuge step at 13,200 rpm for 2 min, the pellet was kept, and cells in the pellet were fixed in 70% ethanol. Cell suspensions were exposed to RNaseA and propidium iodide for staining. The cell cycle distribution was then analyzed by flow cytometry using BD Accuri C6 software (BD Bioscience) (46).

2.10. Statistics

All the experiments were statistically analyzed by GraphPad Prism 9 software (<http://www.graphpad.com/>). Error bars in the graphs were generated using \pm standard deviation (SD) values. A statistical significance test was utilized by using ANOVA Bonferroni's multiple comparisons test. $*p < 0.05$ was taken as a level of significance. Results were repeated at least three times. The immunoblotting results shown are representative of three separate experiments.

3. Results

3.1. Roscovitine treatment decreased cell viability and survival of SK-N-AS cells

To check the effects of ROSC on the cell viability of the SK-N-AS cells, an MTT assay was performed. ROSC was delivered to cells with increasing doses (0-50 μM). The results from three separate experiments were analyzed and shown in Fig. 1A. 1 and 10 μM ROSC treatment caused a significant reduction in viability, especially in 10 μM concentration. The trypan blue dye exclusion results are shown in a time-dependent manner in Fig. 1B. SK-N-AS cells were treated with 1 and 10 μM ROSC in a time-dependent manner for 96h, and the cells were counted with a hemocytometer. The numbers of viable cells were given in a graph. The number of cells treated with 10 μM ROSC was significantly reduced compared to control untreated cells or treated with 1 μM ROSC. Later, we investigated the apoptotic potential of ROSC on SK-N-AS cells. Following the exposure of cells to 1 and 10 μM ROSC for 24 h, bright DAPI-stained nuclei were observed dose-dependently under a fluorescent microscope, suggesting the DNA condensation might increase because of the drug treatments. We also observed that the PI stained cells significantly increased, especially after 10 μM ROSC treatment, indicating that dead cells accumulate following 24h at higher concentrations (Fig. 1C). Finally, diminished fluorescence was observed in cells with disrupted mitochondrial membrane potential loss, which was detected with DiOC₆ staining following cell exposure to increasing ROSC concentrations in SK-N-AS cells.

3.2. ROSC treatment decreased the colony formation potential of SK-N-AS cells

We also investigated the ability of ROSC to diminish the colony formation potential of SK-N-AS NB cells. To better visualize the ROSC effect, we seeded an increasing number of cells, 5×10^3 and 7.5×10^2 , respectively, in a 6-well plate and

exposed to 1 and 10 μM ROSC for 24h. Fig. 2A showed a remarkable difference in the colonies formed, especially after 10 μM ROSC doses. The effect of ROSC on colony formation was also determined with a soft agar assay. The diameters of the colonies were significantly reduced or completely disappeared after 10 μM ROSC treatment. A similar effect was not observed after the exposure of cells to 1 μM ROSC.

3.3. ROSC triggered apoptotic cell death in SK-N-AS cells

Flow cytometric analyses were performed to demonstrate the efficacy of ROSC in triggering apoptosis and to evaluate the cell cycle distribution following 1 and 10 μM ROSC treatment in SK-N-AS cells. When the quadrants were analyzed, it was clear that as the doses of the administered drug increased, the early and late apoptotic population percentages elevated (Fig. 3A). In addition, we also determined that 1 μM ROSC treatment caused a subG1 population percentage increase. However, the number of viable cells was low rather than a dose-dependent increment in subG1 population percentage, and the cycle distribution after 10 μM ROSC exposure did not change (Fig. 3B). Since the annexinV-PI staining and the increase in subG1 populations were observed in SK-N-AS cells due to ROSC treatment, we investigated the expression of pro- and anti-apoptotic proteins by immunoblotting. For this purpose, we evaluated Puma, Bid, Bax, and pro-caspase 9, caspase 3 and 7, Bcl-2, Bcl-xL, and PARP expressions. We observed no significant expression change in Puma after 1 mM ROSC treatment; however, a 10 μM dose was able to diminish it. The sharp decrease in Bid levels as a response to ROSC treatment indicated that the extrinsic apoptosis pathway might also be affected (Fig. 3C). While Bax protein levels elevated after 1 μM of the drug, the expression level was controlled after 10 μM drug exposure to SK-N-AS cells (Fig. 3C).

On the other hand, anti-apoptotic Bcl-2 and Bcl-xL protein expressions were downregulated time-dependently, which demonstrates that mitochondria-dependent apoptosis was induced after ROSC treatments (Fig. 3C). When we checked the expression profiles of caspase proteins, we observed that pro-caspase9 significantly decreased after drug exposure dose-dependently, and the active forms of caspase 3 and 7 were elevated. As a direct proof of apoptosis, the cleavage of PARP protein was observed following 10 μM ROSC treatment in SK-N-AS cells.

3.4. Reactive oxygen species played a role in ROSC-induced apoptosis in SK-N-AS cells

To understand whether ROS levels were affected by ROSC treatment in SK-N-AS cells and whether the elevated levels of ROSC are related to cell viability loss, we performed fluorescence staining and flow cytometry using DCFH-DA. First, flow cytometry determined a significant increase in ROS levels in both 1 and 10 μM ROSC-treated cells (Fig. 4A). The increase was calculated as 2-fold and 2.5-fold for 1 and 10 μM ROSC treatments compared to control samples in SK-N-AS cells (Fig. 4A). Later, we used NAC as a ROS scavenger to ensure that ROS levels affect cell viability after ROSC

treatments. First, we determined whether DCFH-DA levels changed after NAC combined treatment with ROSC. As shown in Fig. 4B, NAC presence diminished the ROSC-induced ROS production by 3-fold (Fig. 4B). NAC itself did not cause ROS

production. Later, we also observed that decreased relative cell viability, approximately 50% by 10 μM ROSC treatment, was completely restored with the combined treatment of NAC and 10 μM ROSC (Fig. 4C).

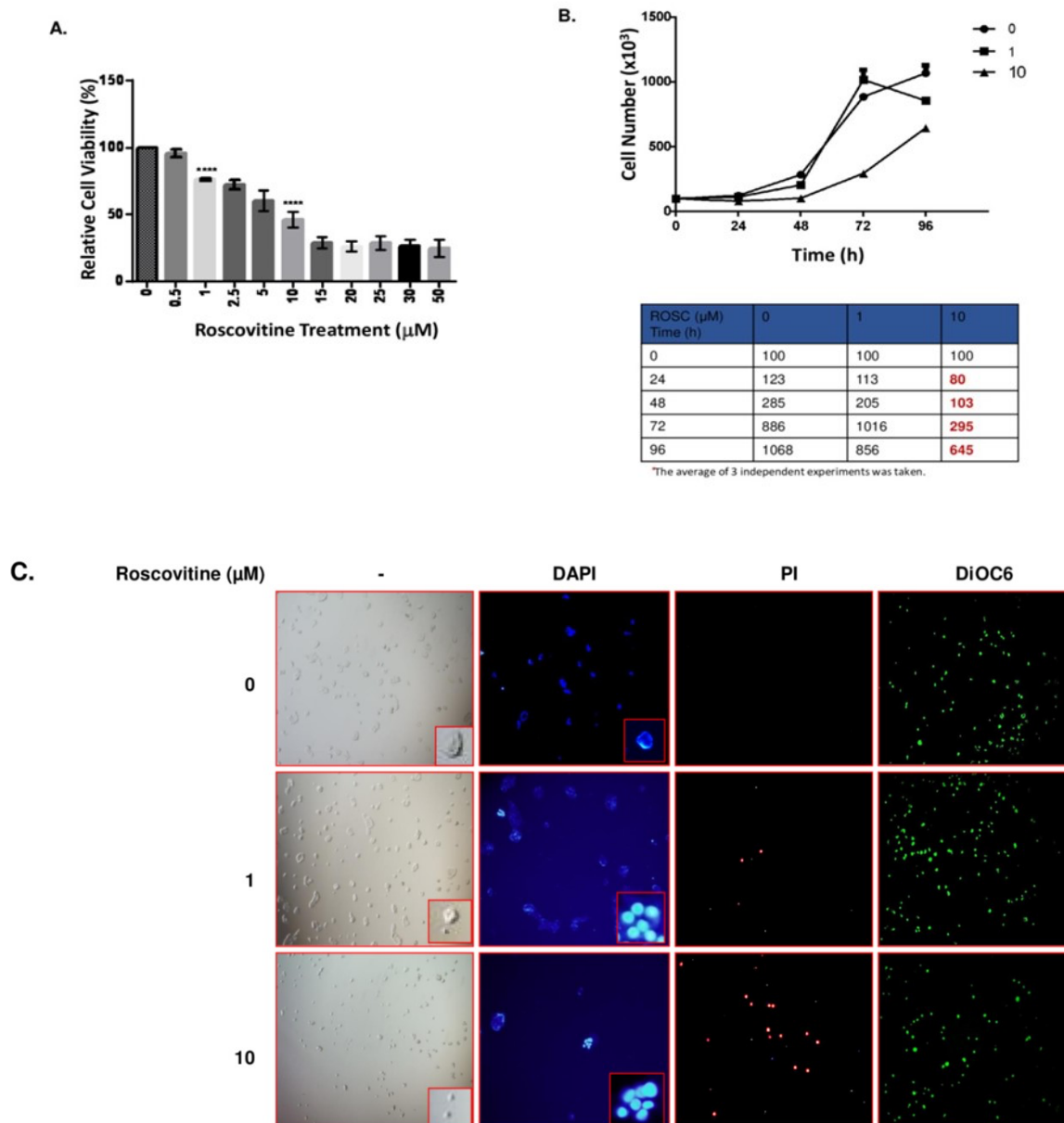


Fig. 1. The effects of ROSC administration on SK-N-AS cell viability. **A.** Cells were incubated with different concentrations of ROSC (0-50 μM) for 24 h and then analyzed with MTT assay. Results presented as the average of 3 independent experiments with 4 replicates. **B.** 0, 1, and 10 μM ROSC were applied to SK-N-AS cells in a time-dependent manner (0-96h) and counted with a hemocytometer. **C.** SK-N-AS cells were exposed to 1 μM and 10 μM ROSC for 24 h. Later, cells were stained with DiOC6, propidium iodide (PI), and DAPI and examined under the fluorescence microscope.

3.5. ROSC inhibited GSK-3 β signaling and prevented the nuclear translocation of β -catenin

Since ROSC-induced apoptosis was clarified in the SK-N-AS cell line, western blotting analysis was performed to determine whether ROSC affected GSK-3 β signaling. Total protein isolation was performed from SK-N-AS cells treated with 1 and 10 μM ROSC with the control group, and 25 μg protein

sample per well was separated in 12% SDS-PAGE gels. After the immunoblotting protocols, the membranes were visualized with the help of chemiluminescence to detect the protein bands. To enlighten GSK3 β signaling, we detected GSK3 β , pGSK3 β from the Ser9 domain, β -catenin. Histone H3 and β -actin were used as loading controls. Cells treated with 1 and 10 μM ROSC exhibited a decreased GSK3 β expression dose-dependently.

On the other hand, the inhibitory phosphorylation of GSK3 β from the Ser9 domain was increased dose-dependently as well compared to control samples, suggesting that GSK3 β signaling was disrupted in response to ROSC treatment in SK-N-AS cells (Fig. 5A). Accordingly, the downstream target of GSK3 β signaling and a vital transcription factor with the cancer progression and metastasis-inducing roles, β -catenin expression, was greatly diminished significantly after 10 μ M

ROSC treatment. Further, we investigated the nuclear translocation of β -catenin in response to ROSC treatment. Our results indicated that 10 μ M of ROSC treatment significantly reduced the expression of β -catenin in the nucleus, suggesting ROSC prevented nuclear translocation (Fig. 5B).

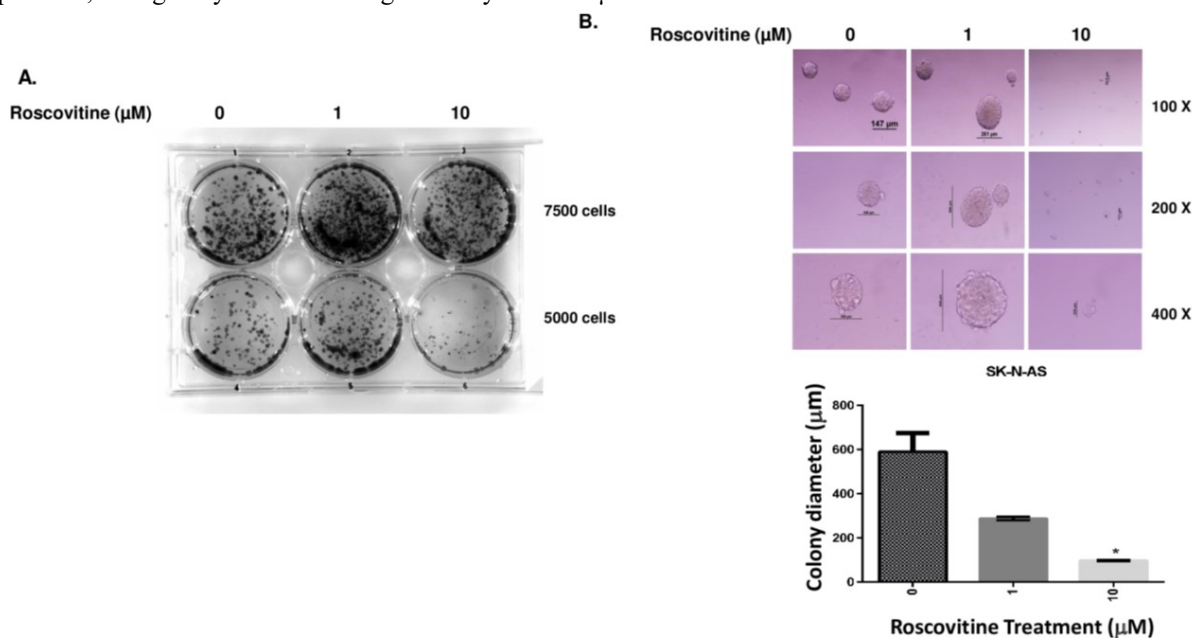


Fig. 2. The effect of ROSC treatment on colony formation potential of SK-N-AS cells. **A.** Cells were seeded at a concentration of either 5×10^3 or 7.5×10^3 / well and treated with increased concentrations of ROSC for 24h. A colony formation assay was performed, and cells were stained with crystal violet. The experiment was repeated at least three times, and the representative image was given. **B.** Neuroblastoma cells treated with 1 and 10 μ M ROSC for 24 h were analyzed with soft agar colony formation assay, and the diameter of each colony was examined under a light microscope.

4. Discussion

NB, one of the most common solid tumor types of infancy, originates from the malignancy of neural crest cells that affect young children at the diagnosed age of 17 months. The heterogeneity of the tumor-initiating cells provokes therapy resistance, consequences of disseminated metastasis, and poor survival. The current treatment strategies include surgery, chemotherapy, and/ or radiation therapy for NB patients. However, almost half of the high-risk patients might face low survival lengths due to tumor recurrence due to failures of conventional therapy strategies (47). Therefore, the identification of novel treatment approaches is urgently required based on identifying key drivers for the arising of NB cells and the molecular pathways behind them. For instance, previous studies have indicated that anaplastic lymphoma kinase (ALK) is the second key driver for NB development, and its inhibition has been the furthest therapy approach in clinical applications (48).

Several studies have analyzed that GSK3 β is a serine-threonine kinase that has a critical role in human cancers, either tumor promoter or suppressor manner (49). GSK3- β plays a

role in the initiation and progression of NB via correlated key signaling pathways: Wnt, MAPK, PI3K/AKT/mTOR, and p53 (49–51). Also, the GSK3 β expression is highly correlated with the MYCN expression, which is the crucial oncogenic driver of NB. Moreover, the intersection of GSK3- β with ALK is the second crucial oncogenic driver for NB (52). Similarly, the other tau kinases harboring the Src kinase family, mitogen-activated kinases (MAPK), and cyclin-dependent kinase 5 (CDK5) have been seen as a target for inhibition of tau phosphorylation (37,53). Given this knowledge, GSK3- β inhibition has been considered a potential therapeutic target for NB progression. Several studies indicated that inhibition of GSK3- β resulted in decreasing proliferation potential of NB cells (47,54). The anti-proliferative effect of LY2090314 has been demonstrated, one type of a GSK-3 inhibitor in NB cell lines harboring SK-N-AS, SH-SY-5Y, and NGP (47). Moreover, the dual inhibition of GSK3- β and CDK5 has also been studied for developing promising neuroprotective strategies in both *in vitro* and *in vivo* studies (55). To date, two GSK3 beta inhibitors, Tideglusib and LY2090314, have been proposed to be well-tolerated drug candidates for the clinic (56).

ROSC is one type of CDK inhibitor (CDK1,2,5,7 and 9) and blocks the cell cycle at G1/ S or G2/M stages based on time-, dose- and cell type-dependent manner (31,57,58). For instance, Kolodziejcki and colleagues showed that ROSC induces cell cycle arrest at the G2/M phase in A172 and G28 glioblastoma cell lines (59). The study indicated that ROSC has therapeutic potential in various types of cancer, both preclinical and clinical stages. Several studies have also

detected the synergistic effect of ROSC and other anti-cancer agents (60). For instance, Pandey and colleagues showed in their 2019 study that a combination of ROSC and temozolomide restricts the growth of glioma cells via enhancing caspase-mediated apoptosis and autophagy mechanism (61).

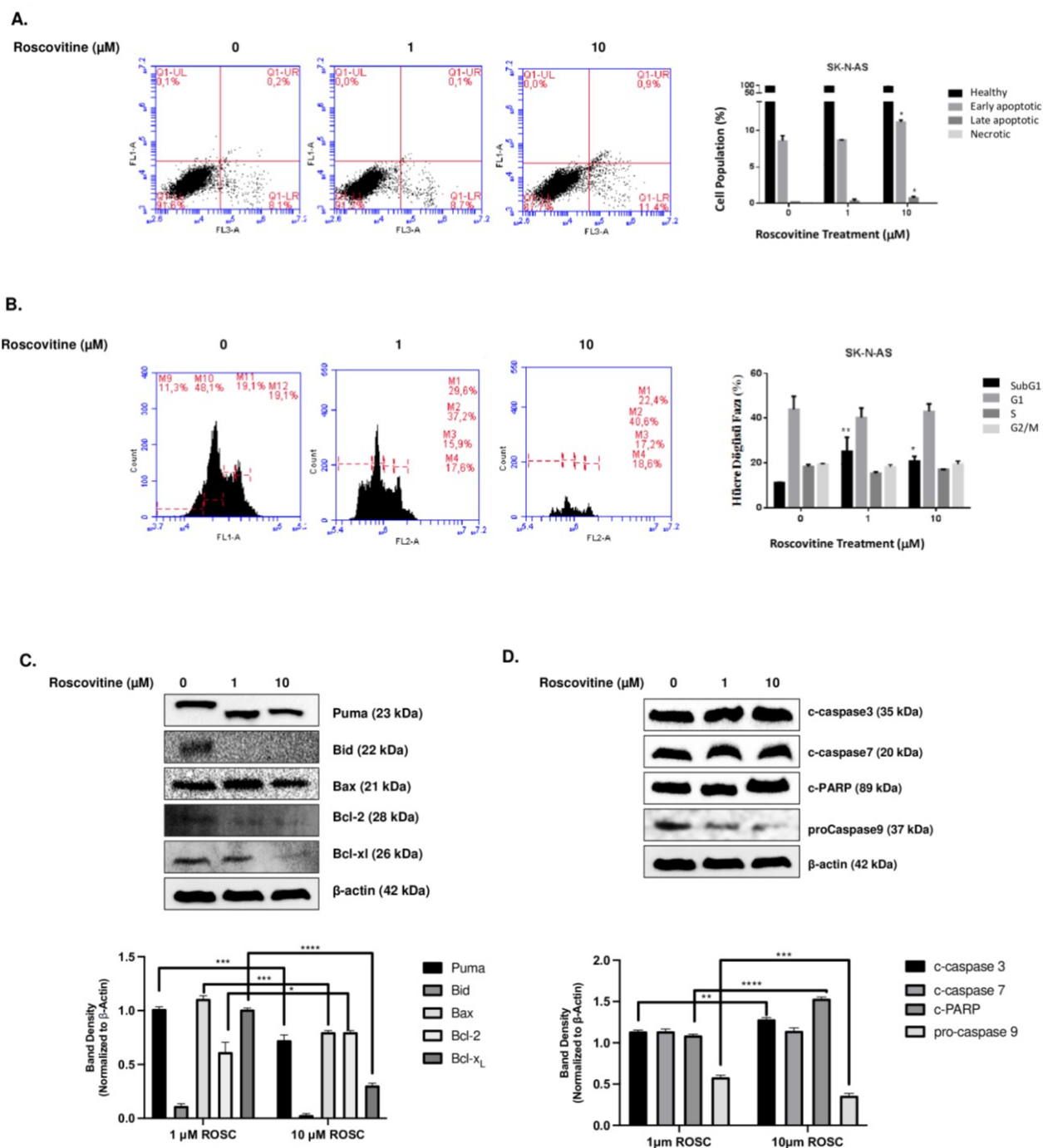


Fig. 3. ROSC induced apoptosis in a caspase9-dependent way and downregulated pro-apoptotic Bcl-2 family members in SK-N-AS cells. **A.** Annexin V-PI staining was performed to determine apoptotic cell populations. Annexin V-FITC and PI fluorescence were reported on the x-axis and y-axis, respectively. Numbers presented in the four quadrants represent the percentage of viable (lower left), necrotic (upper left), early apoptotic (lower right), and late apoptotic (upper right) cells in SK-N-AS cells. **B.** Cells were treated with 1 and 10 μM of ROSC, and the flow cytometry analysis for cell cycle phase distribution was performed following PI staining in SK-N-AS cells. **C. and D.** Increased concentrations of ROSC treatment were performed in the SK-N-AS cell line, and total protein isolation was performed. A total of 25 μg of protein was separated

with 12% SDS-PAGE. After the immunoblotting procedures, the bands of the relevant proteins were determined by the chemiluminescence method. β -actin was used as a loading control.

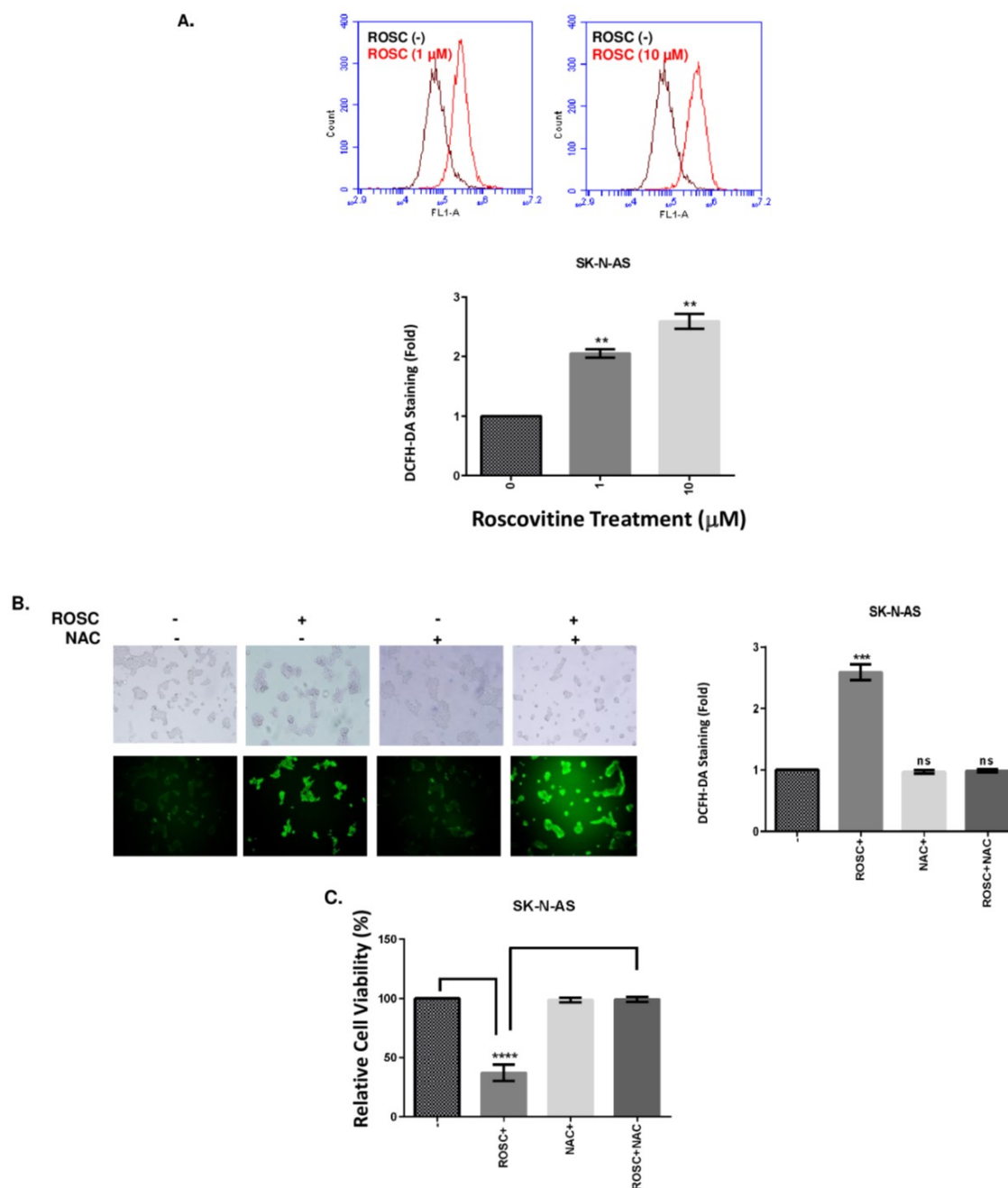


Fig. 4. Demonstration of the involvement of ROS in ROSC-induced cell death in SK-N-AS cells. **A.** The flow cytometric analysis of DCFH-DA staining was analyzed in the presence and absence of ROSC in the SK-N-AS cell line. **B.** NAC effect on ROS generation after dose-dependent treatment of ROSC was examined after DCFH-DA staining in SK-N-AS cells. **C.** The analysis of DCFH-DA staining after combined treatment of NAC and ROSC was evaluated with flow cytometry, and the results of three independent experiments were given as a graph.

Within the scope of this study, while investigating the role of ROSC to induce apoptosis in NB cells, this study also aimed to reveal downstream targets after ROSC administration. In this regard, firstly, we aimed to put forward the effect of ROSC to diminish the cell viability of NB cells. Our results showed that ROSC significantly inhibited the proliferation of human SK-N-AS cells in a time- and dose-dependent manner. The

IC₅₀ values of ROSC change between 15-100 μ M in the literature on various types of cancer (31,57,59). For instance, 20 μ M ROSC doses have been selected for HeLa and HCT116 cancer cell lines (62,63).

Moreover, Goodyear and colleagues have used 10 μ g/ml of ROSC for the MDA-MB-231 cell line (64). The cytostatic

effect of 25 μ M ROSC has been indicated in SH-SY-5Y cells, one of the NB cell lines used in the literature (65). Based on the previous data, the sub-toxic concentration range of ROSC is 5-20 μ M for brain cell lines (61). In our study, we observed that 20% to 50% decreased cell viability of the SK-N-AS cells after administration of 1 μ M and 10 μ M ROSC, respectively. We further analyzed the effect on ROSC (1 μ M and 10 μ M) in long-term colony formation assay. Our results indicated that 10 μ M ROSC significantly diminished the colony formation

potential of the SK-N-AS cells as the parallel colony formation result of the U87 and C6 rat glioma cell line published by Pandey and colleagues (61). A significant decrease was detected in the colony diameters with the application of 1 and 10 μ M ROSC. These results confirmed the anti-proliferative effect of ROSCs within cytotoxic doses (1 and 10 μ M) in SK-N-AS cell lines for subsequent experiments.

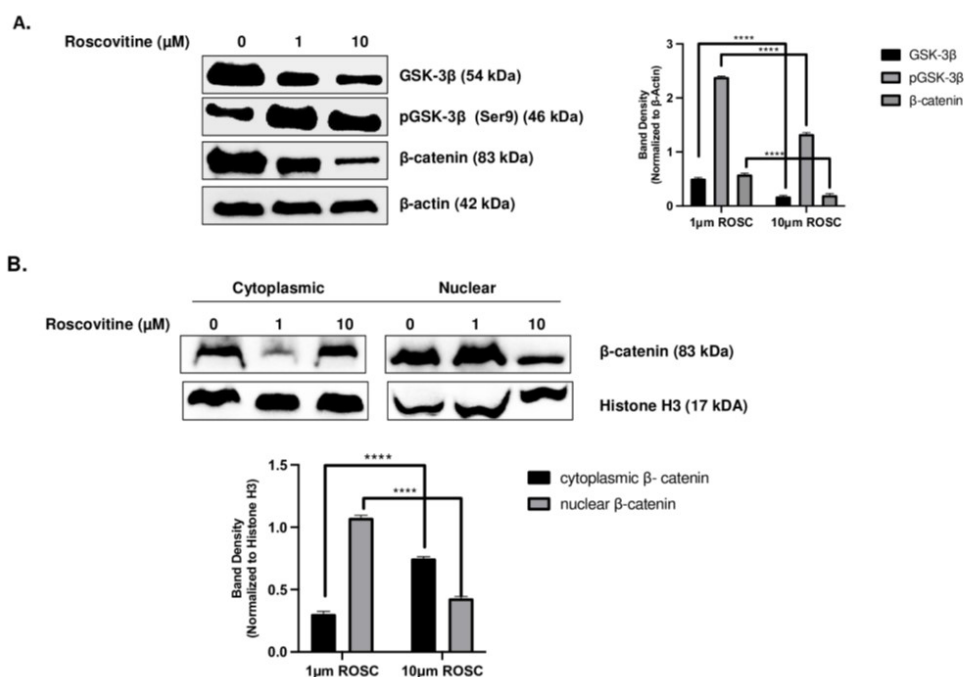


Fig. 5. Demonstration of the efficacy of ROSC administration on GSK3 β signaling. Total protein isolation was performed after increasing concentrations of ROSC in SK-N-AS cells for 24 h. A 25- μ g protein sample was prepared for each well and subjected to 12% SDS-PAGE separation. After the immunoblotting procedures, the bands of relevant proteins were visualized by the chemiluminescence activity of the secondary antibodies. β -actin was used as a loading control. Histone H3 was used as a nuclear fraction loading control.

Several investigations have proposed anti-proliferative and anti-apoptotic properties of ROSC as both solid and hematological malignancies (66). The study of Ozfiliz-Kilbas et al. revealed that ROSC triggers apoptosis and autophagy rely on unfolded protein response in HeLa cervical cancer cell lines (63). The other aim of our study was to enlighten the cell death mechanism type as a result of ROSC administration in NB cells. Similar to our results, a previous study showed that ROSC exhibited a dose-dependent G2/M and sub-G1 fraction increase in the A172 cell line (59). Studies investigating the effect of ROSC on cell cycle phase distribution in colorectal cancer cells have revealed that the drug can block the cell cycle in every stage by causing apoptotic induction (67).

Similarly, our flow cytometer results of annexin V-PI double staining indicated increasing sub-G1 populations after the ROSC treatment, thus inducing cell death. Apoptotic biomarkers were also analyzed to confirm whether loss of cell viability induced by ROSC resulted from apoptotic cell death.

The Bcl-2 family members, either pro- or anti-apoptotic, have critical roles in intrinsic and extrinsic apoptosis pathways. Previous studies indicated that ROSC administration resulted in the expression changes of these family members (65,68). For instance, in ROSC-treated B-cell chronic lymphocytic leukemia cell lines, Mcl-1 expression, a member of the anti-apoptotic family, was downregulated, whereas Bax, a pro-apoptotic protein, was upregulated (69). The high expression of Bcl-2 proteins has been indicated in NB tissues, and it has been suggested that the decrease in Bcl-2 expression should result from a therapeutic intervention for NB (70,71). For this purpose, we investigated the expression changes of apoptotic markers by immunoblotting. 1 and 10 μ M ROSC treatments significantly reduced Bcl-2 and Bcl-xL. Whereas pro-apoptotic marker expressions, Bax and Puma, were increased.

On the other hand, Bid truncation was also observed. Truncated Bid is a pro-apoptotic protein that can be induced via the activation of caspase, a member of the extrinsic

apoptotic pathway. ROSC and extrinsic apoptotic pathway correlation have been investigated in previous studies. For example, in the study of Kim et al. ROSC treatment promotes TRAIL (tumor necrosis factor-related apoptosis-inducing ligand) induced apoptosis in glioma cells, U87MG, and T98 (72). Our data showing the involvement of caspases and PARP cleavage indicated the caspase-dependent apoptosis following ROSC administration to SK-NAS cells. In the literature, ROSC-induced mitochondria-mediated and caspase-dependent apoptosis have been investigated (61,62). In the Arisan et al. study, ROSC-induced apoptotic cell death was caspase-dependent in the MCF-7 cell line (62). The increased expression of cleaved caspase families, PARP, and decreased expression of pro-caspases have been detected during the induction of apoptosis in many studies (61,62). These results demonstrated ROSC-induced mitochondria-dependent apoptosis in the SK-N-AS cell line.

Based on the existing knowledge in literature, elevated levels of ROS have been detected in cancer cells, and it is induced in various types of programmed cell death (73). In the next step of our study, we elucidated whether ROSC affected ROS generation through apoptotic induction in the SK-N-AS cell line. The relationship between ROS production and CDK5 activation, a ROSC target, has been analyzed in the HEK293 cell line in the Sandoval et al. (2018) study. Their research evaluated that the direct activation of CDK5 increased ROS production in the HEK293 cell line (74). According to our data, ROS generation increased significantly following each concentration of ROSC. Compared to ROSC alone, the combination of ROSC with NAC, a well-known ROSC scavenger, significantly prevented cell viability loss in NB cells. These results suggested that ROSC is an agent that led to the formation of ROS in SK-N-AS cell lines. Moreover, our data also put forward that NAC reversed the cytotoxic effect of ROSC by preventing intercellular ROS production when combined. According to the literature, the protective effect of NAC from oxidative stress has been elucidated in SH-SY-5Y and HepG2 cell lines (75) (76).

GSK3 β is the one key signaling mediator in various cellular processes such as DNA repair, cell cycle, cell proliferation, cell signaling, and metabolic pathways. Therefore, many studies suggested that GSK3 β plays a crucial role in cancer therapy. It is also involved in multiple signaling pathways, as mentioned above. According to previous findings, GSK3 β has either a tumor-promoting or suppressive role, depending on the cell type (77). Kotliovara et al. indicated that the inhibition of GSK3 β leads to glioma cell death via induction of apoptosis. In another study by Dickey A. et al., GSK3 β inhibition promoted the induction of apoptosis in the Neuro-2A cell line (78). In our study, when cells were exposed to 10 μ M ROSC, the expression of GSK3 β was decreased with a subsequent increase in the expression of the phosphorylated form of GSK3 β from Ser9. This result suggested that ROSC might diminish GSK3 β activity. The inhibition of GSK3 β has been

shown to cause β -catenin, a downstream target of GSK3 β , stabilization and accumulation in the cytosol. Therefore, its nuclear translocation and gene expression regulating role is prevented [(51). Aberrant nuclear accumulation of GSK3 β has been previously analyzed as a hallmark of cancer in various malignant tumors (56). The reduction of β -catenin nuclear translocation via Wnt signaling inhibition has been an accepted marker of cell survival of many neuronal and cancer cell lines (75,79). Therefore, the further step of our study was to investigate the expression changes of β -catenin in both cytosolic and nuclear localization after the ROSC treatment. Also, we revealed that β -catenin levels in SK-N-AS cells were downregulated progressively in the cytoplasm and increased in the nucleus after 1 μ M ROSC treatment.

On the contrary, when the cells were exposed to 10 μ M ROSC, β -catenin expression was lowered from the cytoplasm to the nucleus. As a result, it is suggested that ROSC promotes apoptosis by inhibiting GSK-3 β and β -catenin in SK-N-AS cell lines. The inhibitors of β -catenin signaling that can cause β -catenin downregulation, promoting ubiquitination or nuclear translocation blockage, have been accepted as anti-cancer drug candidates with therapeutic efficacy (80). ROSC, as well, has also been investigated for the effect of β -catenin expression and translocation. Studies showed that ROSC is a good candidate drug acting on β -catenin signaling in several cancer cells including prostate and cervical cancers (81,82). However, the exact relation has not been bridged in different cancer types, including NB.

Consequently, our results suggested that ROSC decreases NB cell viability and shrinks the NB cell colony diameters while inducing apoptosis. In addition, ROSC also elevated reactive oxygen generation (ROS) production, which can be prevented by N-acetyl-cysteine (NAC). ROSC also inhibited GSK3 β signaling, which was proved by inhibiting the nuclear translocation of β -catenin as the downstream target transcription factor of GSK3 β signaling after 10 μ M ROSC treatment. These results proved that ROSC administration triggers the intrinsic apoptotic cell death by triggering ROS production, and it also affected GSK3 β signaling and prevented β -catenin translocation to the nucleus in SK-N-AS cell lines. A further step of our study requires detailed data regarding the CDK5 and GSK3 β -related signaling pathways of ROSC to develop a new therapeutic modality for NB.

Conflict of interest

No conflict of interest was declared by the authors.

Ethical statement

This work does not involve any animal or human subjects. The authors are aware of the details of the research that are presented in the current manuscript and gave their consent to the publication.

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Authors' contributions

Concept: P.O.Y., E.D.A., Design: P.O.Y., Data Collection or Processing: Z.D., E.K., N.D., B.G., A.K.G., Analysis or Interpretation: P.O.Y., E.D.A., Literature Search: Z.D., E.K., N.D., B.G., A.K.G., P.O.Y., Writing: Z.D., E.K., N.D., A.K.G., E.D.A., P.O.Y.

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