

## The Novel Quinolone CHM-1 Induces DNA Damage and Inhibits DNA Repair Gene Expressions in a Human Osterogenic Sarcoma Cell Line

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**Abstract.** 20-Fluoro-6,7-methylenedioxy-2-phenyl-4-quinolone (CHM-1) has been reported to induce cell cycle arrest and apoptosis in many types of cancer cells. However, there is no available information to show CHM-1 affecting DNA damage and expression of associated repair genes. Herein, we investigated whether or not CHM-1 induced DNA damage and affected DNA repair gene expression in U-2 OS human osterogenic sarcoma cells. The comet assay showed that incubation of U-2 OS cells with 0, 0.75, 1.5, 3 and 6  $\mu$ M of CHM-1 led to a longer DNA migration smear (comet tail). DNA gel electrophoresis showed that 3  $\mu$ M of CHM-1 for 24 and 48 h treatment induced DNA fragmentation in U-2 OS cells. Real-time PCR analysis showed that treatment with 3  $\mu$ M of CHM-1 for 24 h reduced the mRNA expression levels of ataxia telangiectasia mutated (ATM), ataxia-telangiectasia and Rad3-related (ATR), breast cancer 1, early onset (BRCA1), 14-3-3sigma (14-3-3 $\sigma$ ), DNA-dependent serine/ threonine protein kinase (DNA-PK) and O<sup>6</sup>-methylguanine-DNA

methyltransferase (MGMT) genes in a time-dependent manner. Taken together, the results indicate that CHM-1 caused DNA damage and reduced DNA repair genes in U-2 OS cells, which may be the mechanism for CHM-1-inhibited cell growth and induction of apoptosis.

Much evidence has shown that DNA damage is implicated in age-related diseases. For example, DNA single-strand breaks (SSBs) in neocortex of people with Alzheimer disease (AD) are two-fold higher than in the controls (1). Moreover, cortical neurons in AD patients contain DNA SSBs and DNA double-strand breaks (DSBs) *in situ* (2). It is also reported that DNA damage is involved in apoptosis of tumor cells (3).

Quinolone derivatives such as 2-phenyl-4-quinolones have been shown to induce cytotoxicity in many human cancer cell lines and to inhibit platelet aggregation (4-9); synthesized 2-phenyl-4-quinolone series compounds inhibited tubulin polymerization and acted as anti-mitotic agents (4-8); and synthesized 2-phenylpyrroloquinolin-4-ones inhibited the growth of hepatocellular carcinoma *in vitro* and *in vivo* (10). Recently, 20-fluoro-7-methylenedioxy-2-phenyl-4-quinolone (CHM-1) has been shown to act as an anti-invasive agent in hepatocellular carcinoma cells (11).

However, there is no available information to address whether CHM-1 induces DNA damage in U-2 OS human osterogenic sarcoma cells. Therefore, in this study, we investigated the effects of CHM-1 on DNA damage and DNA repair genes in U-2 OS cells.

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Key Words: CHM-1, DNA damage, DNA repair, gene expression, U-2 OS human osteogenic sarcoma cells.

Table I. Primers used in real-time PCR in this study. The DNA sequence was evaluated using Primer Express software.

Primer name		Primer sequence
Human <i>ATM</i>	F	TTACCTAACTGTGAGCTGTCTCCAT
	R	ACTTCCGTAAGGCATCGTAACAC
Human <i>ATM</i>	F	GGGAATCAGACTCGCTGAA
	R	CTAGTAGCATAGCTCGACCATGGA
Human <i>BRCA1</i>	F	CCAGGGAGTTGGTCTGAGTGA
	R	ACTTCCGTAAGGCATCGTAACAC
Human <i>14-3-3σ</i>	F	GCCATGGACATCAGCAAGAA
	R	GGCTGTTGGCGATCTCGTA
Human <i>DNA-PK</i>	F	CCAGCTCTCACGCTCTGATATG
	R	CAAACGCATGCCCAAAGTC
Human <i>MGMT</i>	F	CCTGGCTGAATGCCTATTTCC
	R	TGTCTGGTGAACGACTCTTGCT
Human <i>GAPDH</i>	F	ACACCCACTCCTCCACCTTT
	R	TAGCCAAATTCGTTGTCATACC

Each assay was conducted at least twice to ensure reproducibility. F, Forward; R, reverse.

### Materials and Methods

**Cell culture.** The U-2 OS human osterogenic sarcoma cell line was obtained from the Food Industry Research and Development Institute (Hsinchu, Taiwan). U-2 OS cells were plated onto 75 cm<sup>2</sup> tissue culture flasks with 90% McCoy's 5a medium (Gibco BRL, Grand Island, NY, USA). The cell medium with 2 mM L-glutamine was adjusted to contain 10% fetal bovine serum (FBS; Gibco BRL/Invitrogen, Grand Island, NY, USA), and 1% penicillin-streptomycin (100 Units/ml penicillin and 100 µg/ml streptomycin) and grown at 37°C under a humidified 5% CO<sub>2</sub> atmosphere (12).

**Flow cytometric assay for the percentage of viable human osterogenic sarcoma cells.** Approximately 2×10<sup>5</sup> cells/well of U-2 OS cells in 12-well plates were incubated with CHM-1 at final concentrations of 0, 0.75, 1.5, 3 and 6 µM, vehicle (1 µl dimethyl sulfoxide; DMSO) and 5 µM of H<sub>2</sub>O<sub>2</sub> (positive control) for 24 hours, or cells were treated with 3 µM of CHM-1 for 0, 12, 24, 36 and 48 hours. Cells in 5 ml tubes from each treatment were stained with propidium iodide (PI, 5 µg/ml) and immediately analyzed by flow cytometry (Becton-Dickinson, San Jose, CA, USA) for the percentage of viable cells as previously described (13-14).

**Comet assay for examining DNA damage in U-2 OS cells.** U-2 OS cells (2×10<sup>5</sup>/well) in 12-well plates with cell medium were incubated with CHM-1 at final concentrations of 0, 5, 10, 25 and 30 µM, vehicle (1 µl DMSO) and 5 µM of H<sub>2</sub>O<sub>2</sub> (positive control) grown at 37°C in 5% CO<sub>2</sub> and 95% air. At the end of incubation, cells were harvested for the examination of DNA damage using the comet assay as described elsewhere (13-14). Briefly, glass slides were pre-coated with 1% agarose, about 3×10<sup>4</sup> cells per gel for each treatment were centrifuged (500×g, 5 min at 4°C) and the cell pellet was then suspended in 170 µl of warm (37°C) 0.5% agarose and two 80 µl aliquots placed onto a glass slide which placed in lysis buffer (100 mM EDTA, 2.5 M NaCl, 10 mM Tris-HCl and 1% Triton X-100, adjusted to pH 10 with NaOH) for 2 hours. Then slides were washed twice with ice-cold deionised water and were transferred to

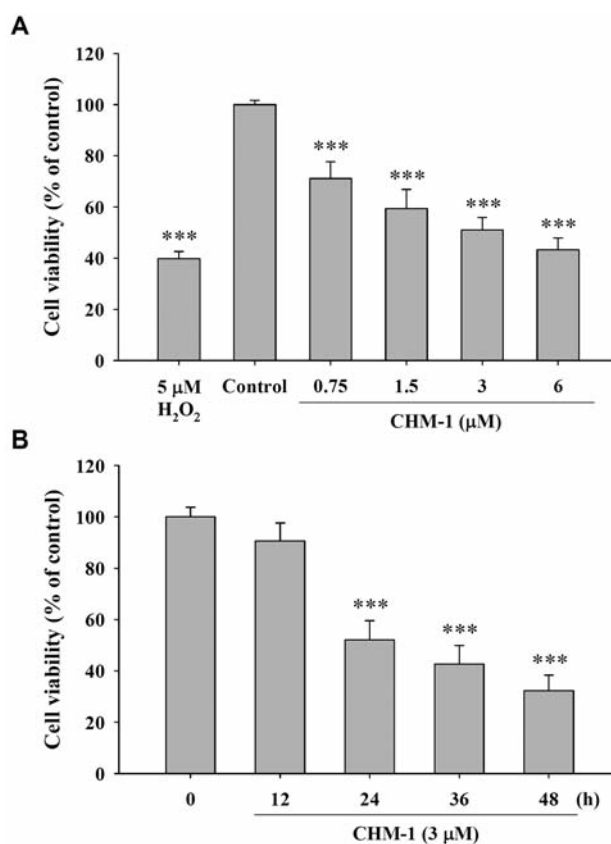


Figure 1. CHM-1 affected the percentage of viable human osterogenic sarcoma U-2 OS cells. U-2 OS cells (2×10<sup>5</sup> cells/well) were placed in 12-well plates and were incubated with CHM-1 at final concentrations of 0, 0.75, 1.5, 3 and 6 µM, vehicle (1 µl DMSO) and 5 µM of H<sub>2</sub>O<sub>2</sub> (positive control) for 24 hours (A), or cells were treated with 3 µM CHM-1 for 0, 12, 24, 36 and 48 hours (B). Cells from each treatment were stained with propidium iodide (5 µg/ml) and analyzed by flow cytometry (Becton-Dickinson, San Jose, CA, USA) as previously described. \*\*\* p<0.001 Compared to untreated control.

an electrophoresis tank which contained cold electrophoresis buffer (300 mM NaOH, 1 mM EDTA, pH 13) and incubated for 20 min to allow unwinding of the DNA. Then electrophoresis was carried out at 30 V and 300 mA for 20 min before slides were removed and flooded with neutralization buffer (0.4 M Tris-HCl, pH 7.5), and rinsed twice with deionised water. Slides were stained with 1 ml of PI (2.5 µg/ml) for 20 min (15). Comets were visualized and photographed by use of a fluorescence microscope as previously described (16-17). PI-stained DNA tails in the individual nucleus were quantified by TriTek Comet Score V 1.5 software (TriTek Corp., Sumerduck, VA, USA).

**DNA gel electrophoresis for examining DNA damage in U-2 OS cells.** U-2 OS cells (1×10<sup>6</sup>/well) in 6-well plates with cell medium were incubated with 3 µM of CHM-1 for 24 and 48 hours. At the end of incubation, cells were harvested in phosphate-buffered saline (PBS) by centrifugation and were lysed in a digestion buffer containing 0.5% sarkosyl, 0.5 mg/ml proteinase K, 50 mM Tris-HCl

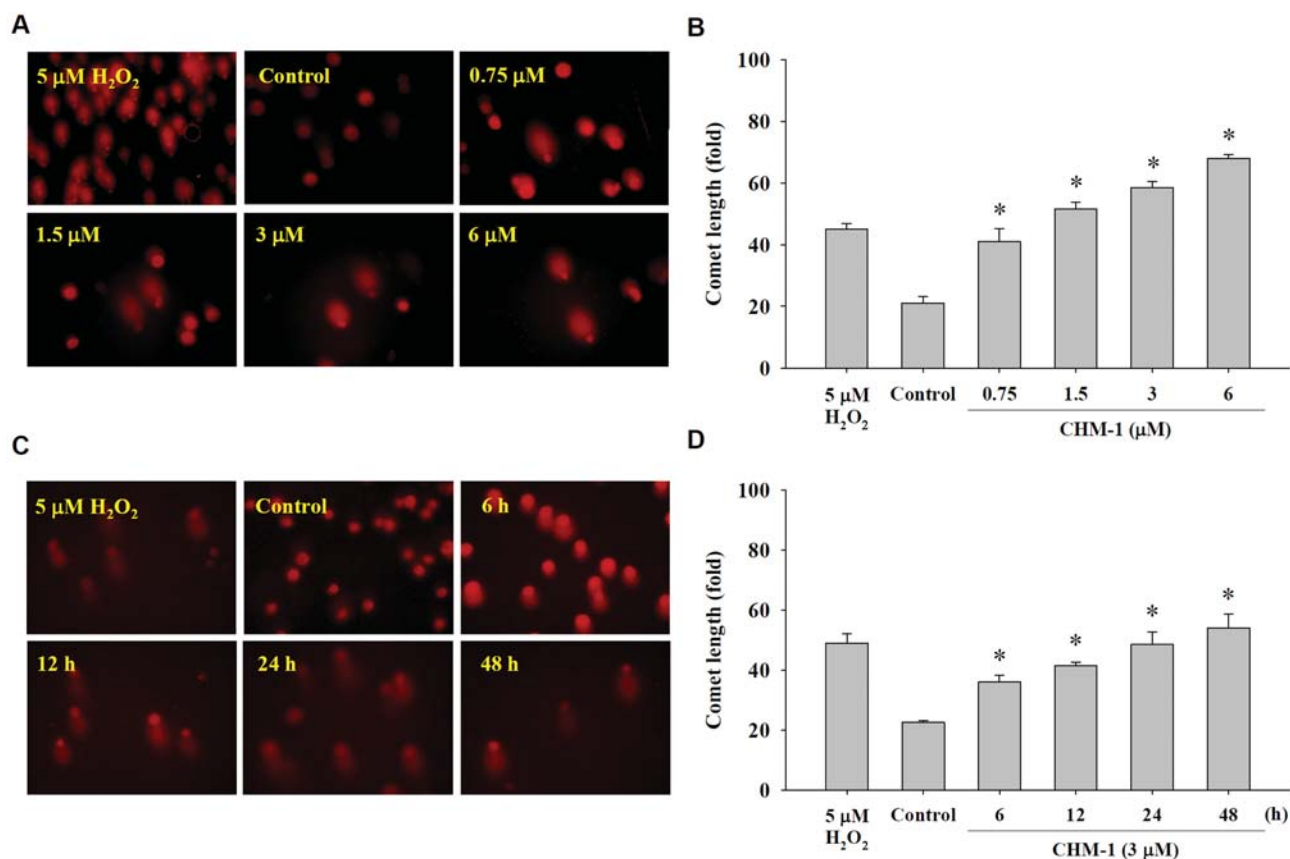


Figure 2. CHM-1-induced DNA damage in U-2 OS cells was examined by comet assay. U-2 OS cells ( $2 \times 10^5$  cells/well; 12-well plates) were incubated with different concentrations of CHM-1 for 24 hours or cells were treated with 3  $\mu$ M of CHM-1 for different time periods and DNA damage was determined by comet assay as described in the Materials and Methods. Representative images of cells are shown in panel A and C, and panel B and D show comet length (fold). \* $p < 0.05$  Compared to untreated control.

(pH 8.0) and 10 mM EDTA at 55°C overnight. Cells were then treated with 0.5  $\mu$ g/ml RNase A for 2 hours at 37°C. The genomic DNA was extracted by phenol-chloroform-isoamyl alcohol extraction (25:24:1) as previously described (14). The extracted DNA from each treatment was resuspended with 50  $\mu$ l TBE buffer. Approximately 1  $\mu$ g/ $\mu$ l (12  $\mu$ l) of DNA was loaded into each well and DNA gel electrophoresis was performed at 50 V for 90 min using 2% agarose. After ethidium bromide staining, the cells then were photographed under fluorescence light as previously described (14, 16-17).

**Real-time PCR of ATM, ATR, BRCA1, 14-3-3 $\sigma$ , DNA-PK and MGMT in U-2 OS cells.** U-2 OS cells ( $1 \times 10^6$ /well) in 6-well plates were incubated with 3  $\mu$ M of CHM-1 for 24 hours. At the end of incubation, cells were harvested in PBS by centrifugation then the total RNA from each sample was extracted by using the Qiagen RNeasy Mini Kit (Qiagen Inc., Valencia, CA, USA) as described previously (15, 18). RNA samples were reverse-transcribed for 30 min at 42°C with High Capacity cDNA Reverse Transcription Kit according to the standard protocol of the supplier (Applied Biosystems, Foster City, CA, USA). The quantitative PCR from each sample was performed as

follows: 2 min at 50°C, 10 min at 95°C, and 40 cycles of 15 s at 95°C, 1 min at 60°C using 1  $\mu$ l of the cDNA reverse-transcribed as described above, 2X SYBR Green PCR Master Mix (Applied Biosystems) and 200 nM of forward and reverse primers as shown in Table I. Each assay was run on an Applied Biosystems 7300 Real-Time PCR system in triplicates and expression fold-changes were derived using the comparative  $C_T$  method (19).

**Statistical analysis.** Student's *t*-test was used to analyze differences between exposure to CHM-1 and the untreated (control) group. All data are presented as the means  $\pm$  SD of three experiments and  $p < 0.05$  was considered significantly.

## Results

**CHM-1 reduced the percentage of viable U-2 OS cells.** The U-2 OS cells were exposed to different concentrations of CHM-1 for 24 hours or were treated with 3  $\mu$ M of CHM-1 for different time periods. The cells were collected for the determination of percentage of viable U-2 OS cells and the

results are shown in Figure 1A and B, which indicate that CHM-1 reduced the percentage of viable cells and these effects took place in a dose- and time-dependent manner.

*CHM-1-induced DNA damage in U-2 OS cells as examined by comet assay.* In earlier studies, it had been shown that CHM-1 induced cytotoxic effects on U-2 OS cells (20). In the present study, we investigated whether or not CHM-1 induced DNA damage in U-2 OS cells. The results from the comet assay are shown in Figure 2 and indicate that CHM-1 induced DNA damage in U-2 OS cells. These effects are dose dependent (Figure. 2A and B). However, the long incubation of U-2 OS cells with 3  $\mu$ M of CHM-1 led to a longer DNA migration smear (comet tail) (Figure 2C and D), indicating that CHM-1 induced DNA damage in U-2 OS cells in a time-dependent manner.

*CHM-1-induced DNA fragmentation in U-2 OS cells was determined by DNA gel electrophoresis.* DNA fragmentation is a characteristic of apoptosis (21). Here, we isolated DNA from U-2 OS cells after treatment with 3  $\mu$ M of CHM-1 for 24 and 48 hours before DNA was used for agarose gel electrophoresis. The results shown in Figure 3 indicate that CHM-1 induced DNA damage and fragmentation in U-2 OS cells, which also indicated the occurrence of apoptosis (Figure 3). The longer time (48 hours) of incubation of U-2 OS cells with CHM-1 led to more DNA damage and fragments than that of short time (24 hours) incubation. This finding suggests that CHM-1 appears to significantly induce apoptosis of U-2 OS cells.

*CHM-1 inhibited the relative expression levels of DNA damage and repair genes in U-2 OS cells as shown by real-time PCR.* For investigating whether CHM-1 affected DNA damage and repair gene expressions, U-2 OS cells were treated with 3  $\mu$ M CHM-1 for 0 and 24 hours. Total RNA was isolated from each sample and associated gene expressions were examined by real-time PCR (Figure 4). Expression levels of *ATM*, *ATR*, *BRCA1*, *14-3-3 $\sigma$* , *DNA-PK* and *MGMT* mRNA were decreased on 24 hours treatment when compared with the control group.

**Discussion**

In our primary studies, we have demonstrated that CHM-1 induced cell cycle arrest and apoptosis in many human cancer cell lines (20, 22). However, there is no available information to show CHM-1 affected DNA damage and DNA repair associated gene expression in U-2 OS cells. Herein, we used the comet assay (single-cell gel electrophoresis) to measure the levels of DNA damage from U-2 OS cells after exposure to different concentrations of CHM-1 for various time periods. It is well-documented that the comet assay is a highly sensitive technique for DNA

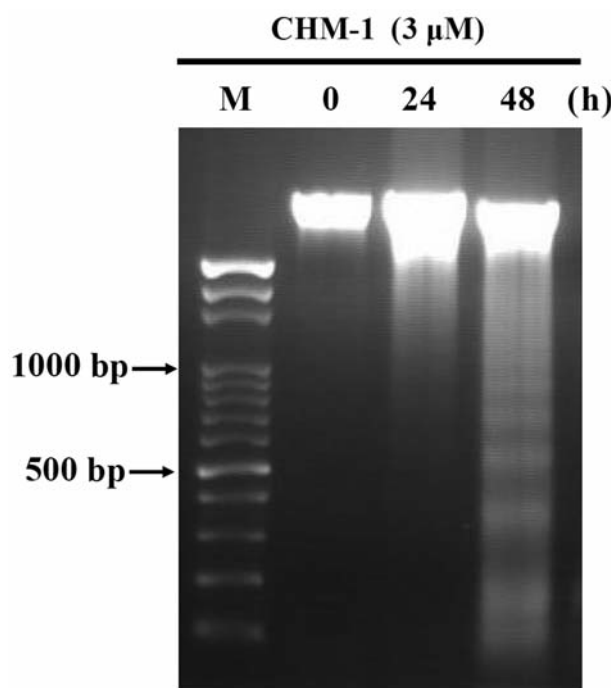


Figure 3. CHM-1-induced DNA fragmentation in U-2 OS cells was examined by DNA gel electrophoresis. U-2 OS cells were incubated with 3  $\mu$ M CHM-1 for 0, 24 and 48 hours, the cells were harvested and DNA was extracted from each treatment before DNA fragmentation was examined by DNA gel electrophoresis as described in the Materials and Methods.

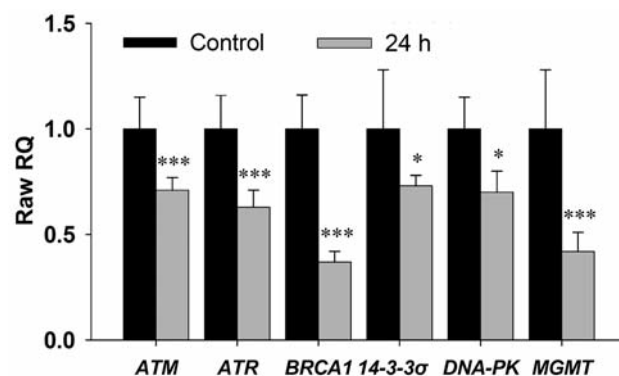


Figure 4. CHM-1-inhibited expression of DNA damage and repair genes in U-2 OS cells were examined by real-time PCR. The total RNA was extracted from the U-2 OS cells after treatment with 3  $\mu$ M of CHM-1 for 0 and 24 hours. RNA samples were reverse-transcribed cDNA and real-time PCR carried out as described in the Materials and Methods. The experiments of *ATM*, *ATR*, *BRCA-1*, *14-3-3 $\sigma$* , *DNA-PK* and *MGMT* genes related to *GAPDH* are presented. Data represent the mean $\pm$ SD of three experiments. \* $p$ <0.05 and \*\*\* $p$ <0.001 compared to untreated control.

damage examination (19, 23-25). The results showed that CHM-1 induced a significant increase in the tail moment of the comets of U-2 OS cells. H<sub>2</sub>O<sub>2</sub> was used as positive control and showed significant tail movement.

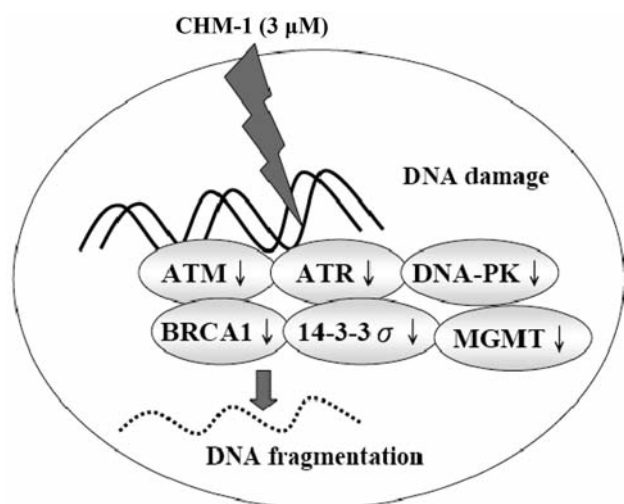


Figure 5. The possible signaling pathway for CHM-1-reduced expression of DNA damage and repair genes in a human osterogenic sarcoma cell line (U-2 OS).

These findings are in agreement with other reports which showed that CHM-1 induced DNA damage in human osterogenic sarcoma cells (20) and murine colorectal adenocarcinoma cells (22). Other reports already showed that strand-break formation during the process of excision repair may also cause DNA migration measurable in the comet assay (26-27). In cells, DNA repair can reduce DNA damage by eliminating DNA lesions. Results (Figure 3) from DNA gel electrophoresis demonstrated that CHM-1 indeed induced DNA fragmentation (apoptosis) in U-2 OS cells.

Actually, in our primary studies, we had already documented the cytotoxic effects of CHM-1 on other cells such as osterogenic sarcoma cells (20) and osterogenic sarcoma and, including U-2 OS cells (Figure. 1). Our earlier studies also showed that the reduction of cancer cell numbers by CHM-1 may be achieved through the induction of apoptosis or by antiproliferative effect (20, 22). Our results also demonstrated that CHM-1 inhibited expression of DNA repair genes such as *ATM*, *ATR*, *BRCA1*, *14-3-3σ*, *DNA-PK* and *MGMT* (Figure 4) in examined U-2 OS cells. Further studies are needed to establish the role of the interaction of CHM-1 with DNA in carcinogenesis.

In conclusion, the CHM-1 itself may induce DNA damage in U-2 OS cells *via* the inhibition of expression of DNA repair genes such as *ATM*, *ATR*, *BRCA1*, *14-3-3σ*, *DNA-PK* and *MGMT*, subsequent to DNA damage (Figure 5).

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