

# The homologous recombination protein RAD51 is a promising therapeutic target for cervical carcinoma

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**Abstract.** RAD51 is one of the pivotal enzymes for DNA double-strand break (DSB) repair by the homologous recombination (HR) pathway, which implies it as a promising and novel target for cancer therapy. Recent findings have indicated RAD51 protein is overexpressed in a variety of tumors. The high-expression of RAD51 is related to poor prognosis. RAD51 is involved in the repair of DNA damage and the generation of genetic diversity by an evolutionarily conserved mechanism. However, the exact mechanism of RAD51 in the progression of cervical cancer remains unclear. RI-1 is a small molecule that inhibits the central recombination protein RAD51. In this study, we found that RAD51 was highly expressed in invasive squamous cervical cancer (SCC). The administration of RI-1 inhibited cell growth *in vitro* and reduced growth of tumor xenografts *in vivo* with cervical cancer cells (HeLa and SiHa). Further investigation suggested that RAD51 protein significantly promoted the cell cycle transition from the G0/G1 to S phase. In addition, the inhibition of RAD51 reduced the level of the cell cycle related protein cyclin D1, but increased the levels of p21 mRNA and protein. As a DNA DSB repair enzyme, the expression of RAD51 in tumor cells possibly affects their sensitivity to anti-cancer agents. Additionally, in experiments using cisplatin and ionizing radiation, RI-1 treated cervical cancer cells, HeLa and SiHa, were sensitized to a greater extent than the untreated control. Thus, HR inhibition of RAD51 may provide yet another mechanism of therapeutic target for the chemosensitization and radiosensitization of cervical cancer with RI-1. Collectively, our data demonstrated for the first time that inhibition of RAD51 suppressed the cervical cancer cell proliferation and the growth of cervical cancer xenografts by attenuating cell cycle transition, which could be a functional link between *RAD51* and cyclin D1 and p21.

## Introduction

Worldwide, cervical cancer is the second most common malignancy after breast cancer, and remains a leading cause of cancer-related death among females in developing countries. There were an estimated 527,600 new cervical cancer cases and 265,700 deaths worldwide in 2012 (1). Currently, surgery, platinum based chemotherapy and radiotherapy play important roles for cervical cancer treatment. Nevertheless, acquired resistance to platinum and radiation is considered a main element for tumor relapse and metastasis. Therefore, to develop more effective therapeutic strategies and in-depth research into the molecular and biologic mechanisms of oncogenesis for cervical cancer is critical. Although the development of cervical cancer is intimately associated with high-risk human papillomavirus (HPV) infection (2), not all patients infected with HPV ultimately develop cervical cancer. Thus, various molecular dysfunction, including the activation of oncogenes and inactivation of suppressor genes, are also essential for cervical cancer development (3,4).

The *RAD51* gene is homologous to the *E. coli* RecA and yeast *RAD51* genes, which are involved in the repair of DNA double-strand breaks and also play important roles in recombination repair and various SOS responses to DNA damage by  $\gamma$ -irradiation and alkylating reagents (5). *RAD51* plays a role in several cellular processes, including genomic integrity, cell cycle regulation, apoptosis and tumor formation. RAD51 is overexpressed in a variety types of tumors, including cervical cancer (6), non-small cell lung cancer (7), breast cancer (8), ovarian cancers (9), pancreatic cancer (10), melanoma and glioblastoma (11). The overexpression of RAD51 causes improper and hyper-recombination, namely contributing to genomic instability and genetic diversity, which might drive regular cells towards neoplastic transformation or further contribute to cancer progression and metastasis (12,13). Additionally, in Fanconi anemia-like patients, phenotype-derived mutation in *RAD51* plays a role in protection of DNA during the course of ICL repair, which is independent of RAD51's ability to maintain genomic integrity (14). Accumulating evidence has indicated that *RAD51* not only is involved in the progression of carcinogenesis, but also plays a part in resistance to anti-cancer treatments (16,17). The RAD51 protein is a specific HR-related target for cancer therapy. Treatment of leukemia, prostate cancer, pancreatic adenocarcinoma, lung carcinoma

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and glioma cells with imatinib can decrease RAD51 expression and sensitize them to experimental chemotherapy and radiotherapy *in vitro* and *in vivo* (17,18).

The molecular mechanisms of *RAD51* in cervical carcinoma are largely unclear. In the present study, we demonstrated that RAD51 was overexpressed in cervical carcinoma and examined the effects of the RAD51 inhibitor RI-1 on the proliferation, cell cycle and sensitivity of cervical cancer cells. Budke *et al* revealed that RI-1 made RAD51 inactive by directly binding covalently to human RAD51 protein at cysteine 319. Accordingly, RI-1 inhibited the formation of subnuclear RAD51 foci in cells in response to DNA damage, without affecting formation of the replication protein A focus (15). Our study suggested that RI-1 largely inhibits the growth of cervical cancer cells *in vitro* and *in vivo* by arresting the cell cycle. Moreover, RI-1 decreased resistance to platinum and ionizing radiation. Therefore, in addition to classical function in HR, RAD51 can also regulate the progression of the cell cycle, and thus RI-1 might have promising therapeutic effects against cervical carcinoma.

## Materials and methods

**Cell lines and cell culture.** Human cervical cancer cell lines (CaSki, C33A, HeLa and SiHa) were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). C33A, HeLa and SiHa cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM; Sigma-Aldrich, St. Louis, MO, USA), and CaSki was cultured in RPMI-1640 (Sigma-Aldrich), supplemented with 10% heat-inactivated fetal bovine serum (FBS; Invitrogen, Carlsbad, CA, USA). All cell lines were maintained at 37°C in a humidified 5% CO<sub>2</sub> incubator.

**Tissue specimens.** The human specimens (n=107) were collected from patients at the Second Affiliated Hospital of Xi'an Jiaotong University from 2010 to 2015. Of the 107 samples 43 were normal cervical (NC) tissues and 64 were squamous cervical cancer (SCC) tissues. All of the procedures followed approved medical ethics practices. None of the patients had received chemotherapy, immunotherapy or radiotherapy before specimen collection. The histological classifications and clinical staging were done in accordance with the International Federation of Gynecology and Obstetrics classification system.

**Immunohistochemistry and RAD51 inhibitor.** Formalin-fixed, paraffin-embedded tissue sections were analyzed in an immunohistochemical study. After being placed in a 60°C incubator overnight, four micrometer-thick sections were deparaffinized with xylene and rehydrated in a series of ethanols. Then the sections were incubated in heat-induced epitope retrieval in citric acid buffer (pH 7.0) for 2 min. Endogenous peroxidase was blocked at room temperature using 3% hydrogen peroxide in methanol for 10 min. Then slides were incubated with mouse monoclonal anti-RAD51 (1:100; Bioscience, Boston, MA, USA) at 4°C overnight, followed with incubation of goat anti-mouse immunoglobulin at room temperature for 30 min. The slides were counterstained with hematoxylin (7). RAD51 immunostaining was

presented by a semi-quantitative immunoreactivity score (IRS), which (negative 0-3, weak 4-7, strong 8-12) was evaluated by multiplying the values for staining intensity (scored as 0, no staining; 1, light brown; 2, brown; 3, dark brown) and the values for percentage of positive cells (scored as 0, <10%; 1, 10-25%; 2, 25-50%; 3, 50-75%; 4, >75%) in each sample. An overall score of ≤3 was defined as negative and a score of >3 as positive. All specimens were evaluated by two pathologists in a blinded manner.

The RAD51 inhibitor RI-1 was purchased from Selleck Chemicals (Houston, TX, USA), according to the manufacturer's instructions. RI-1 was dissolved in DMSO at a stock concentration of 1 mM.

**Western blot analysis.** Exponentially growing cells were harvested and resuspended in PBS pH 7.4 (Gibco). Then the cells were lysed by lysis buffer mixed with protease inhibitor cocktail (Roche, Mannheim, Germany). The lysate was mixed with Laemmli sample buffer containing 5% 2-mercaptoethanol and boiled for 5 min. Equal amounts of protein extracts were separated on a 10% SDS/PAGE, blotted onto an activated polyvinylidene difluoride membrane (Millipore, Billerica, MA, USA) and blocked in TBST with 5% dried milk. Membranes were probed with anti-RAD51 (1:1000, Bioscience), anti-GAPDH (1:1000, Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-cyclin D1 (1:500, Santa Cruz Biotechnology), anti-p21 (1:500, Santa Cruz Biotechnology) at 4°C overnight. Then, they were probed with goat anti-mouse secondary antibodies (Thermo Fisher Scientific, Grand Island, NY, USA) for 1 h.

**Cell growth and cell viability assays.** Cells were seeded in triplicates at a density of 5x10<sup>4</sup> cells with 2 ml of media into 35-mm tissue culture dishes for 7 days. The numbers of cells were manually counted after harvesting using a hemocytometer under light microscopy every two days. Cell viability assays were performed by applying 3-(4,5-dimethylthiazol-yl)-2,5-diphenyl tetrazolium bromide (MTT; Sigma-Aldrich) dye to cells that were seeded in 96-well plates with 800 cells in each well, as described in a standard protocol. Then, the number of live cells was determined by the absorbance at 490 nm (Bio-Rad, Hercules, CA, USA).

**Tumor xenograft experiment.** To assess the tumorigenicity *in vivo*, cells (1x10<sup>7</sup>) in the exponential growth phase were collected and were suspended in 200- $\mu$ l phosphate-buffered saline and then injected subcutaneously into the posterior side of 4- to 6-week-old female BALB/c-nude mice purchased from Shanghai SLAC Laboratory Animal Co., Ltd. (Shanghai, China). Xenograft tumor volume (V) was calculated using the length (a) and width (b) by  $V=ab^2/2$ . To study the function of the RAD51 inhibitor RI-1, 5 mg/kg in 200- $\mu$ l phosphate-buffered saline was injected intraperitoneally every other day into BALB/c nude mice (n=6) when the tumor volume reached 120 mm<sup>3</sup>. In this analysis, the negative control group (n=6) received saline. The animal experimental protocols were evaluated and approved by the Animal Care and Use Committee of the Medical College of Xi'an Jiaotong University. The mice were sacrificed and tumors were dissected and weighed at the end of the experiment.

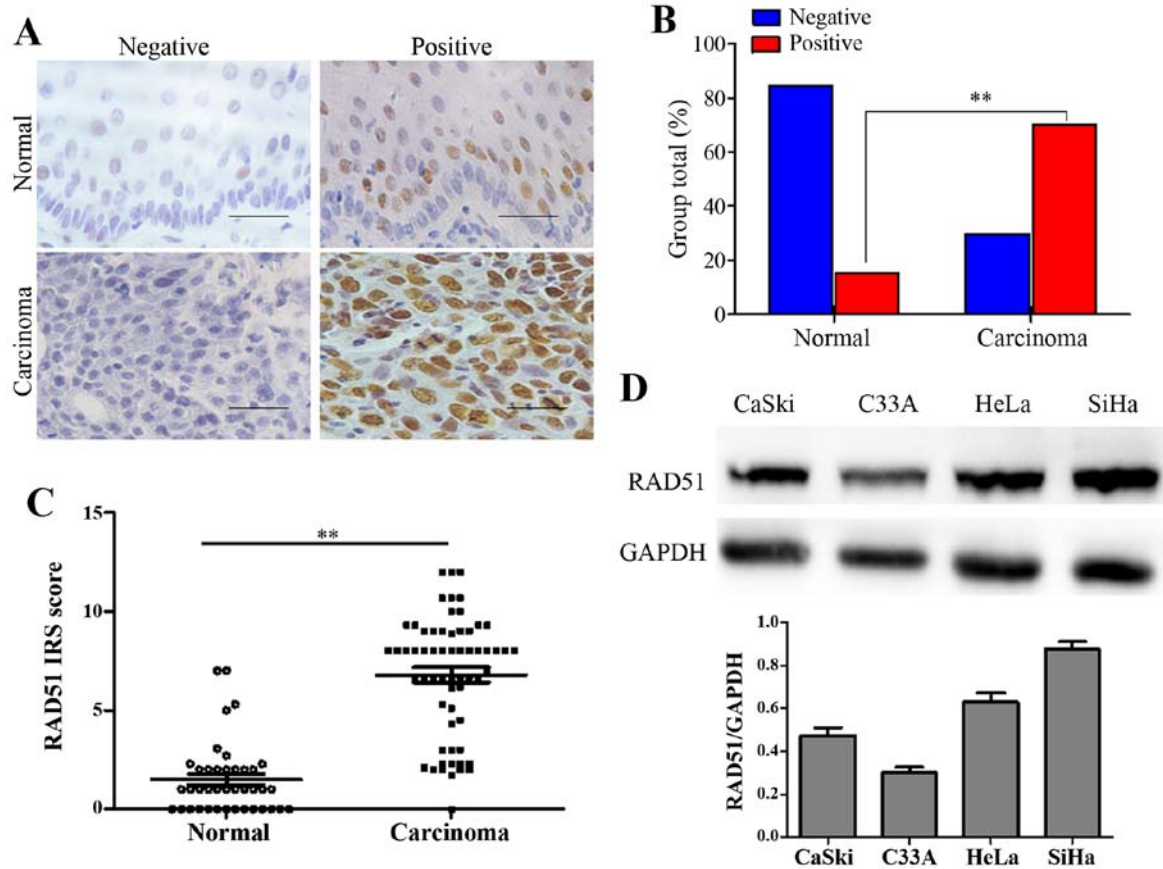


Figure 1. RAD51 expression in normal cervix and cervical carcinoma tissues. (A) Immunohistochemistry results showing RAD51 expression in normal cervix and cervical carcinoma tissues; scale bar, 50  $\mu$ m. (B) RAD51 staining is classified into negative and positive, and the percentage of tissues in each group is shown. (C) The IHC scores of RAD51 staining in the normal cervix and carcinoma tissues are shown. (D) The expression of RAD51 in Caski, C33-A, HeLa and SiHa cells was measured by western blotting, the relative expression of RAD51 was calculated based on western blot analyses, GAPDH was used as an internal control. \*P<0.05, \*\*P<0.01.

Table I. Rad51 expression in tissue specimens.

Specimens	Total	Rad51 staining		P-value
		Negative, no. (%)	Positive, no. (%)	
Normal	43	37 (86.05)	6 (13.95)	
Carcinoma	64	17 (26.56)	47 (73.44)	<0.01

**Flow cytometry analysis.** Cell cycle analysis was performed using flow cytometry (FACScan; Becton Dickinson, Franklin Lakes, NJ, USA) according to the manufacturer's instructions. Cells were harvested and fixed in 70% ice-cold ethanol overnight at 4°C. Thirty minutes before FACS analysis, the samples were washed with PBS, treated with 1 mg/ml RNase A and then stained with 20  $\mu$ g/ml propidium iodide (Sigma-Aldrich). Cell cycle distribution was analyzed using a FACSCalibur flow cytometer with Mod-Fit LT software.

**Quantitative real-time PCR.** Total RNA was extracted from exponentially growing cells with the TRIzol reagent (Invitrogen). Total cDNA was used as a template for PCR amplification. Quantitative real-time PCR was performed

using the IQ5 Real-time PCR Detection System (Bio-Rad) in triplicates and the following primers: GAPDH (GCACCGT CAAGGCTGAGAAC and TGGTGAAGACGCCAGTGG); Cyclin D1 (AAACAGATCATCCGCAAACAC and GTT GGGGCTCCTCAGGTTTC) and p21 (GCAGACCAGCATG ACAGATTTC and CGGATTAGGGCTTCCTCTTG). The protocol was 95°C for 30 sec, 40 cycles of 95°C for 5 sec and 60°C for 30 sec, and then a dissociation stage. The results were analyzed via the  $\Delta\Delta$ Ct method using GAPDH as the house-keeping gene.

**Cell survival assay.** Cells were treated with cisplatin (Sigma-Aldrich; 0-32 nmol/l) for 24 h. In other experiments, the cells were irradiated with doses between 0 and 18 Gy under aerobic conditions, at room temperature, using a <sup>137</sup>Cs unit at a dose rate of 2.5 Gy/min.

**Statistical analysis.** All of the statistical analyses were performed using GraphPad Prism 5.0 software (GraphPad Software, La Jolla, CA, USA). All data are expressed as mean  $\pm$  standard deviation (SD). Student's t-test was used for comparisons between two groups and one-way or two-way analysis of variance (ANOVA) test was used to analyze statistical differences between groups under different conditions. P<0.05 was considered to indicate a statistically significant difference.

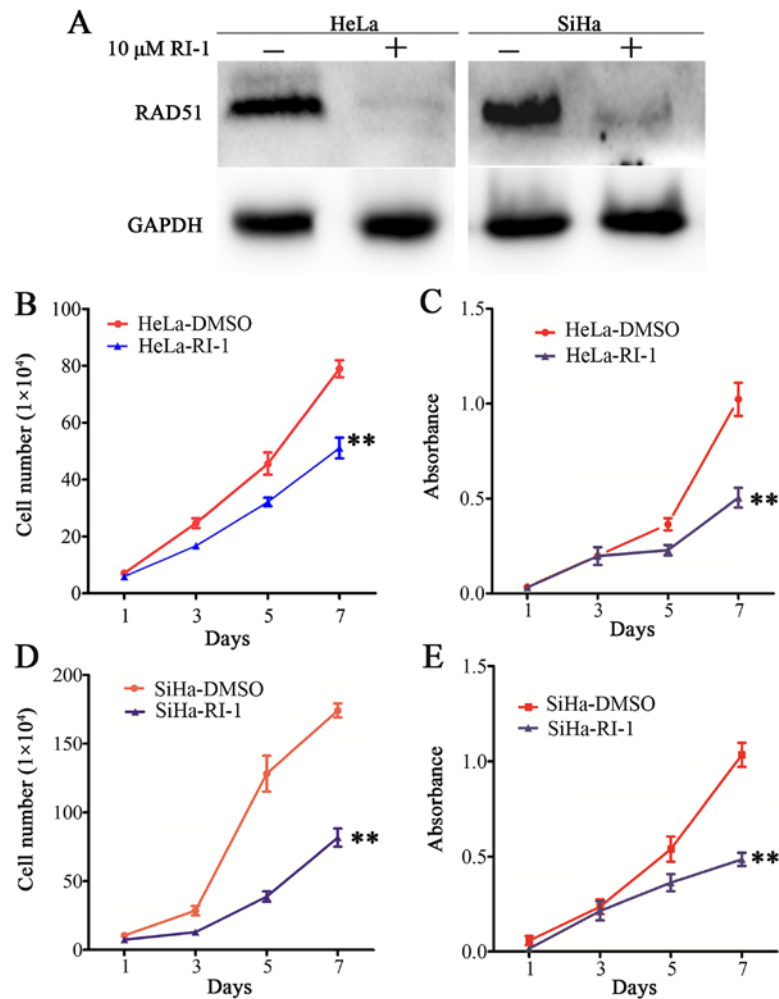


Figure 2. The RAD51 inhibitor RI-1 suppresses the proliferation of cervical cancer cells *in vitro*. (A) HeLa and SiHa cells were treated with either DMSO or 10  $\mu$ M RAD51 inhibitor RI-1 for 24 h, and the RAD51 protein was measured by western blotting. Cell growth and MTT assays were performed using RI-1- or DMSO-treated HeLa cells (B and C) and RI-1- or DMSO-treated SiHa cells (D and E). The values are presented from independent experiments performed in triplicates as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ .

## Results

**RAD51 expression in normal cervix and squamous cervical carcinoma.** To explore the RAD51 expression levels and its association with normal or neoplastic cervical tissues, we first conducted the immunohistochemistry assay using paraffin-embedded normal cervix and squamous cervical cancer tissues. RAD51 staining was observed in the nuclei of positive cells in different cervical tissues (Fig. 1A). The number of specimens with positive RAD51 staining gradually increased from 13.95% (6/43) in the normal cervical tissues to 73.44% (47/64) in the cervical cancer tissues (Table I and Fig. 1B,  $P < 0.01$ ). IHC score results revealed that the immunoreactivity score (IRS) of RAD51 staining was 1.6 for the normal cervical tissues and 6.8 for the cervical cancer tissues ( $P < 0.01$ , Fig. 1C).

Next, using western blot assay, we found that RAD51 showed different expression levels in CaSki, C33-A, HeLa and SiHa cells (Fig. 1D). In particular, higher levels of the RAD51 protein appeared in the HeLa and SiHa cells.

**The RAD51 inhibitor RI-1 suppresses the proliferation of cervical cancer cells *in vitro*.** Having shown that RAD51

expression was increased in cervical cancer progression, the role of RAD51 in cervical cancer was functionally evaluated. We conducted experiments using the RAD51 inhibitor RI-1. We found that RAD51 was almost eliminated using 10  $\mu$ M RI-1 in HeLa and SiHa cells that expressed higher levels of RAD51 protein by western blotting (Fig. 2A). Next, we tested whether the inhibition of RAD51 activity would suppress cervical cancer cell proliferation. Cell growth curve and MTT assays showed that compared to controls, RI-1 significantly reduced cell growth and viability in HeLa (Fig. 2B and C,  $P < 0.01$ ) and SiHa (Fig. 2D and E,  $P < 0.01$ ). These results demonstrated that RI-1 suppressed the proliferation of cervical cancer cells *in vitro*.

**The RAD51 inhibitor RI-1 suppresses the growth of cervical cancer xenografts *in vivo*.** To investigate whether RI-1 has similar inhibitory effect on tumorigenicity *in vivo*, female athymic nude mice (6 mice per group) were injected subcutaneously with HeLa and SiHa cells. When tumor formation reached 120 mm<sup>3</sup>, the volume of tumors was measured every three days, and we intervened with an intraperitoneal injection of RI-1 (5 mg/kg in 200  $\mu$ l PBS) or normal saline. The net

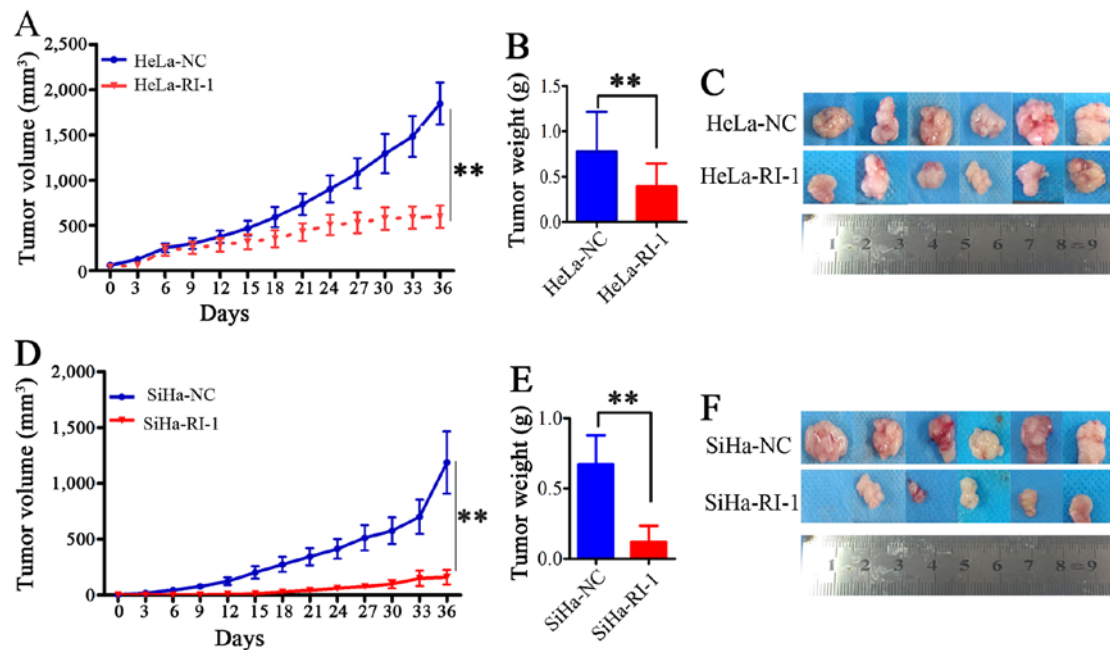


Figure 3. The RAD51 inhibitor RI-1 suppresses the growth of cervical cancer xenografts *in vivo*. (A-C) Tumor growth curves and tumor weights are shown for RI-1-treated HeLa and control cells. The NC group was treated with saline (n=6) and the RI-1 (n=6) group was treated with RI-1 for 36 days (intraperitoneal injection every other day). (D-F) Tumor growth curves and tumor weights are shown for RI-1-treated SiHa and control cells. The NC group was treated with saline (n=6) and the RI-1 (n=6) group with RI-1 for 36 days (intraperitoneal injection every other day). The values presented are of independent experiments performed in triplicates as the mean  $\pm$  SD. \*P<0.05, \*\*P<0.01.

weights of the sacrificed mice were recorded upon termination of the experiment. As shown in Fig. 3A, compared to the negative controls, the RI-1-treated mice exhibited significantly inhibited tumor growth in HeLa cells (P<0.01). The average volume and weight of xenografts of the RI-1-treated groups were markedly reduced (Fig. 3A-C, P<0.01). The results were similar in SiHa cells (Fig. 3D-F, P<0.01). These results indicated that RI-1 suppressed tumor formation of cervical cancer cells *in vivo*.

*RI-1 suppresses the proliferation of cervical cancer cells by attenuating the cell cycle transition from G0/G1 to S phase through acting on the expression of cell cycle associated protein cyclin D1 and p21.* To investigate the mechanism of RI-1-mediated suppression of cervical cancer cell proliferation, cell cycle analyses of RI-1-HeLa and SiHa and control cells was performed by fluorescence-activated cell sorting (FACS). The percentage of cells in S phase significantly decreased from 47.25% for HeLa-DMSO cells to 31.56% for HeLa-RI-1 cells (Fig. 4A and B, P<0.05). A similar effect was observed in SiHa-RI-1 cells, which had 20.34% in the S phase compared to 33.12% of control cells (Fig. 4C and D, P<0.05). This result showed that the RAD51 inhibitor RI-1 may arrest the cell cycle transition at the G1/S phase and further suppress the proliferation of cervical cancer cells.

Further experiments showed that the mRNA level of the cell cycle associated protein cyclin D1 was reduced while that of p21 increased after RI-1 treatment in HeLa and SiHa cells (Fig. 4E, P<0.05). The administration of RI-1 for 48 h resulted in a downregulation of cyclin D1 and upregulation of p21 by western blotting (Fig. 4F, P<0.01). Collectively, these results suggested that inhibition of RAD51 blocked the cell cycle transition possibly through interaction with cyclin D1 and p21.

*Inhibition of RAD51 enhances the sensitivity of cervical cancer cells to chemotherapeutic drugs and irradiation.* Given the decreased expression and function of RAD51 in cervical cancer cell lines after using RI-1, we investigated whether RI-1 could lead to an increase in tumor cell sensitivity to cisplatin and radiation. We observed that cervical cancer cells, HeLa and SiHa, when treated with RI-1 had significantly reduced survival compared to cells treated with DMSO in response to cisplatin (Fig. 5A and B). In response to radiotherapy, HeLa and SiHa cells treated with RI-1 had also significantly reduced survival compared to cells treated with DMSO (Fig. 5C and D). Thus, we concluded that RI-1 can sensitize cervical cancer cells to chemotherapeutic drugs and radiation.

## Discussion

Elevated expression of RAD51 is associated with tumor aggressiveness and is known to confer treatment resistance in a variety of tumors, including those in ovarian cancer (19), breast cancer (16), lung tumors (20), pancreatic adenocarcinoma (13) and malignant gliomas (21). Furthermore, downregulation of RAD51 protein by RAD51 antisense oligonucleotides, RNA interference (22), aptamers (23) or small molecule inhibitors against RAD51 could be used to sensitize tumors to chemotherapy or radiation. Recent studies have identified that impaired replication and intra-S mediated CHK1 signaling by RAD51 led to higher genomic instability and thus drove tumorigenesis (24). The 3'-untranslated region of RAD51 was directly bound by tumor-suppressing miR-34a and thus regulation of homologous recombination and double-strand break repair was inhibited in NSCLC cells (20).

In our study, RAD51 expression was found to be significantly elevated in cervical cancer tissues compared to normal

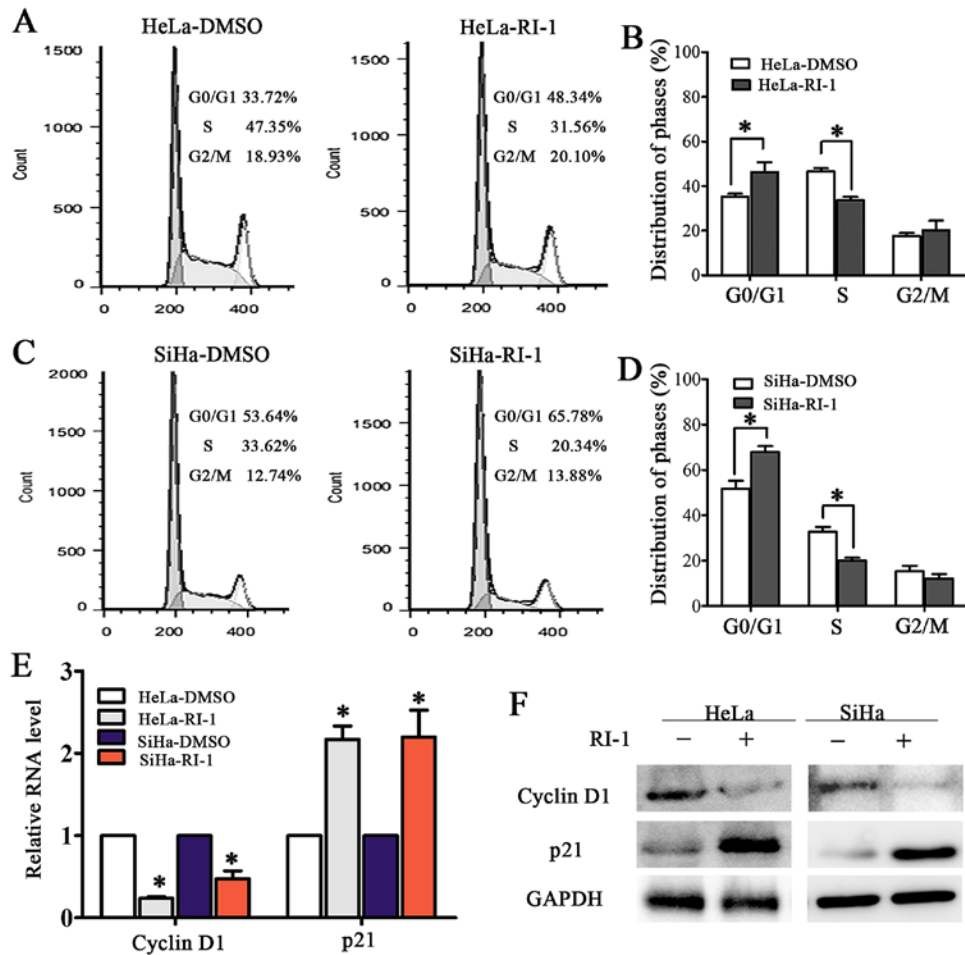


Figure 4. The RAD51 inhibitor RI-1 suppresses the proliferation of cervical cancer cells by attenuating the cell cycle transition from G0/G1 to S phase through its effect on the expression of cell cycle associated protein cyclin D1 and p21. Cell cycle progression was analyzed by FACS, and the results of quantitative analysis of the cell cycle distribution are shown for RI-1-treated HeLa cells and control cells (A and B) and RI-1-treated SiHa cells and control cells (C and D). (E) The relative mRNA levels of cyclin D1 and p21 in RI-1-treated HeLa and SiHa cells as determined by quantitative real-time PCR. (F) The expression of cyclin D1 and p21 in RI-1-treated HeLa and SiHa cells was determined by western blotting. The values presented are from independent experiments performed in triplicates as the mean  $\pm$  SD. \*P<0.05, \*\*P<0.01.

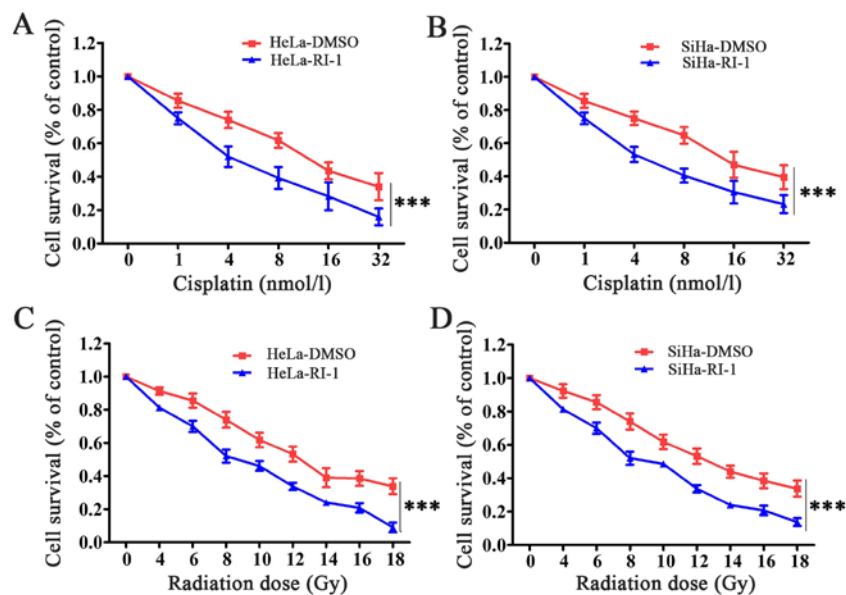


Figure 5. The RAD51 inhibitor RI-1 enhances the sensitivity of cervical cancer cells to chemotherapeutic drugs and irradiation. (A) RI-1-treated HeLa and (B) SiHa cells were treated with different concentrations of cisplatin for 24 h and subjected to MTT assay. (C) RI-1-treated HeLa and (D) SiHa cells were irradiated with the indicated doses for 24 h and evaluated with MTT assay. The mean survival for three independent experiments are shown. The values presented are from independent experiments performed in triplicates as the mean  $\pm$  SD. \*P<0.05, \*\*P<0.01. \*\*\*P<0.001.

cervical tissues (Fig. 1), consistent with results from previous studies of non-small cell lung cancer (7), breast carcinoma (8), prostate cancer (25), chronic myeloid leukaemia (CML), melanoma and glioblastoma (11). Subsequently, RI-1 was applied to further explore the function of RAD51 in cervical carcinogenesis. RI-1 could directly and specifically disrupt HsRAD51, and inhibit sub-nuclear accumulation of HsRAD51 protein at sites of DNA damage, thereby, this inhibitory activity sensitizes tumor cells to cross-linking chemotherapy (15). We further performed western blotting to evaluate the inhibitory effect of RI-1 on RAD51. A previous study indicated that the combination of minocycline and MMC in NSCLC can synergistically inhibit cell proliferation and reduce cell viability *in vitro*, while overexpression of RAD51 expression can restore cell viability upon minocycline and MMC cotreatment (26). In our study, the results showed that inhibition of RAD51 significantly suppressed the proliferation of cervical cancer cells *in vitro* and tumor growth *in vivo* (Figs. 2 and 3). In a previous study, p21 was investigated because it was well-known as a major mediator of G1 arrest and an inhibitor of G1 Cdks after DNA damage (27). Thus, the RAD51 foci-positive cells could also be arrested during the G1 phase. This would be consistent with the observation that G0/G1 arrest upon serum starvation induced by overexpressed of RAD51 protein. However, endogenous Rad51 foci were not observed in G1 phase cells. Overexpression of RAD51 could increase its function, which improved DNA recombination in S/G2 phase (28). Our cell cycle analyses suggested that RI-1 prevented the transition from G0/G1 to S phase in cervical cancer cells. In addition, RI-1 treatment decreased cyclin D1 and increased p21 mRNA and protein levels (Fig. 4). Nevertheless, the particular mechanism between RAD51 and cyclin D1 and p21 should be further investigated. Recent studies have revealed that the enhanced RAD51 protein level in tumor cells is associated with high DNA repair capacity, elevated recombination rates and increased resistance against radiotherapy and chemotherapy (10). Our study showed that downregulation of RAD51 protein could indeed sensitize cervical cancer cells to chemotherapy and radiation (Fig. 5).

In summary, our findings demonstrate that inhibition of RAD51 attenuates cell proliferation and tumor formation in cervical cancer cells by arresting the cell cycle via cyclin D1 and p21. Therefore, RAD51 might be a promising therapeutic strategy for the treatment of cervical cancer.

### Acknowledgements

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