

Close correlation between MEK/ERK and Aurora-B signaling pathways in sustaining tumorigenic potential and radioresistance of gynecological cancer cell lines

FRANCESCO MARAMPON^{1,2*}, GIOVANNI LUCA GRAVINA^{1,2*}, VALDIMIR M. POPOV^{3,4},
LUCA SCARSELLA¹, CLAUDIO FESTUCCIA², MARIA EMILIA LA VERGHETTA², SILVIA PARENTE²,
MANUELA CERASANI², GEMMA BRUERA⁵, CORRADO FICORELLA⁵, ENRICO RICEVUTO⁵,
VINCENZO TOMBOLINI⁶, ERNESTO DI CESARE^{1,2} and BIANCA MARIA ZANI¹

¹Department of Biotechnological and Applied Clinical Sciences, University of L'Aquila; ²Department of Biotechnological and Applied Clinical Sciences, Division of Radiotherapy and Radiobiology Laboratory, San Salvatore Hospital, University of L'Aquila, I-67100 L'Aquila, Italy; ³LIPOGEN LLC, Mount Laurel, NJ 08054;

⁴Department of Cancer Biology and Medical Oncology, Kimmel Cancer Center, Thomas Jefferson University, Philadelphia, PA 19107, USA; ⁵Medical Oncology, S. Salvatore Hospital, University of L'Aquila, L'Aquila, I-67100; ⁶Department of Radiological Sciences-Oncology and Pathological Anatomy, and Department of Radiotherapy, University of Rome 'La Sapienza', I-00161 Rome, Italy

Received July 11, 2013; Accepted August 30, 2013

DOI: 10.3892/ijo.2013.2167

Abstract. Both Aurora-A and -B kinases have been implicated in tumorigenesis; and as such, they represent an attractive therapeutic target. Recent studies found that Aurora-A is a downstream target of mitogen-activated protein kinase 1/ERK2, while Aurora-B has been found to be a prognostic/predictive therapeutic target for epithelial cancer. In a wide range of human cancers, the Ras/Raf/MEK/ERK/MAP kinase pathway is enhanced and the cellular response to growth signals is known to increase. The purpose of this study was to investigate whether the MEK/ERK cascade regulates tumorigenic signaling and radioresistance via the Aurora-B-mediated pathway in a panel of gynecological cancer cell lines. Exponentially growing human endometrial (Ishikawa), cervical (HeLa), cervical (CASKI) and vulva (SiHa) cancer cells were used in culture treated with either control or MEK/ERK inhibitor or AZD1152 before and after irradiation. Western blotting, ERK1/2 siRNA transfection, growth assay in modified monolayer, Annexin V and migration/invasion assays were performed. The specific MEK/ERK inhibitor U0126 decreased the tumorigenic potential and improved the

radiation response in all cellular models. The modulation of radioresponse upon U0126 treatment positively correlated with the inhibition of phospho-ERKs and the reduction of Aurora-B kinase expression. In addition, upon U0126 treatment DNA-PKcs protein expression was found to be downregulated, indicating that the improved radiation response may be caused by decreased DNA double-strand damage repair mechanisms. The knockdown of ERK by siRNA confirmed the MEK/ERK-dependent Aurora-B kinase expression. The use of AZD1152, a selective Aurora-B inhibitor, counteracted tumorigenic potential and radioresistance phenotype by highly increasing apoptotic mechanisms in all gynecological cancer cell lines used. Evidence from our experiments show that tumorigenic potential and radiation response in gynecological cancer cells may ensue from a MEK/ERK or Aurora-B inhibition. Together with the close correlation of MEK/ERK and Aurora-B protein expression, this study underlines the potential role of a MEK/ERK/Aurora-B axis whose interruption recovers the antitumor effects of radiotherapy.

Introduction

Even though cervical, endometrial and ovarian cancers are relatively common, vulvar, vaginal, and fallopian tube cancers, and choriocarcinomas are very rare (1). Treatment of these malignancies requires a multimodal approach combining surgery, radiotherapy (RT) and/or chemotherapy. Although overall statistics have improved considerably, the outcome for patients with high-risk disease remains relatively poor, which points to a clear need for new therapeutic strategies. In situations like these, radiation therapy can be used to prevent local cancer recurrences after surgery (2-4). It has been clarified that the radiation resistance is associated with the abnormal

Correspondence to: Dr Francesco Marampon, Department of Biotechnological and Applied Clinical Sciences, Division of Radiotherapy and Radiobiology Laboratory, San Salvatore Hospital, University of L'Aquila, Via Vetoio, I-67100 L'Aquila, Italy
E-mail: f.marampon@gmail.com

*Contributed equally

Key words: U0126, gynecological cancer, Aurora-B, MEK/ERK inhibitor, radiotherapy

expression of activated oncogenes, including Ras (5,6) and Aurora-B (7). The Ras/Raf/mitogen-activated protein kinase kinase (MEK)/extracellular signal-regulated kinase (ERK) cascade regulates proliferation, differentiation, survival, motility (8-18) and when constitutively activated, mediates resistance to ionizing radiation (15). The aurora kinases are a family of oncogenic serine/threonine kinases involved in the mitotic (M) phase of the cell cycle, acting to establish the mitotic spindle, bipolar spindle formation, alignment of centrosomes on mitotic spindle, centrosome separation, cytokinesis and monitoring of the mitotic checkpoint (19-22). There are three known aurora kinases (Aurora-A, -B and -C) in human neoplastic and non-neoplastic tissues (19,22-26). Inhibition of aurora kinase activity leads to catastrophic errors of mitosis, defective cytokinesis, misaligned centrosomes, mitotic spindle malformation and apoptosis (26-30). However, the role of aurora kinases in radioresistance phenomena remains unclear. The Ras/Raf/MEK/ERKs pathway positively regulates Aurora-B expression at the transcriptional level (31). DNA double-strand break is critical in DNA lesions induced by radiation. In mammalian cells the repair of these lesions occurs by non-homologous end joining (NHEJ) requiring Ku70/Ku86 and the recruitment of the catalytic subunit of DNA-dependent PK (DNAPKcs). DNAPKcs is implicated in radioresponsiveness of embryonal rhabdomyosarcoma-uterine-derived cell lines (32-34) and human glioblastoma cell lines (35,36). DNA-PKcs are necessary for genomic stability whereas abnormal levels in cancer cell may contribute to cell proliferation, radioprotection eventually contributing to oncogenic phenotype (37-40). Herein we investigated whether MEK/ERK pathway, collaborating with Aurora-B kinase, sustains the tumorigenic potential and radioprotection in a panel of gynecological cancer cell lines.

Materials and methods

Cell culture, treatment and radiation exposure. The human Ishikawa endometrial cancer cell line was kindly provided by Professor Marcello Maggiolini (University of Calabria, Italy). The human HeLa cervical cancer cell line was kindly provided by Professor Roberto Maggio (University of L'Aquila, Italy), while the human CASKI cervical- and SiHa vulva-cancer cell line were kindly provided by Professor P.J.F. Snijders (VU Medisch Centrum, Germany). Tumor cell lines were cultured in the appropriate medium supplemented with 10% fetal calf serum (FCS). Treatments with 10 μ M MEK/ERK inhibitor U0126 (1,4-diamino-2,3-dicyano-1,4-bis[2-aminophenylthio]butadiene; Promega) or with AZD1152 (60 nM), an Aurora-B kinase inhibitor, was done as shown in the figures. Radiation was delivered at room temperature using an X-ray LINAC at the dose rate of 2.5 Gy/min. For clonogenic survival assay, exponentially growing cells, diluted to appropriate densities, were plated with complete medium in presence of U0126, AZD1152 or vehicle/control [dimethyl sulfoxide (DMSO) 0.1%] and then irradiated with graded doses (0-2-4-6 Gy). Cells were then cultured in drug-free medium for 14 days, fixed with methanol/acetic acid (10:1, v/v) and stained with crystal violet. Colonies containing >50 cells were counted. The plating efficiency (PE) was calculated as the number of colonies observed/the number of cell plated; the surviving

fraction (SF) was calculated as: colonies counted/cells seeded (PE/100).

Cell proliferation assay. Cells from adherent culture were counted using hemocytometer and tested for exclusion of trypan blue. Results represent the average of triplicate experiments, including standard error.

Western blot analysis. Proteins of whole cell lysates were assessed using the Lowry method (40) and equal amounts of proteins were separated on SDS-PAGE. The proteins were transferred to a nitrocellulose membrane (Schleicher & Schell Bioscience GmbH, Germany) by electroblotting. Immunoblottings were performed with the following antibodies directed against c-Myc, ERK1/2, phospho-ERK1/2, p27, cyclin D1, CDK2, Aurora-B, DNAPKcs and α -tubulin (B-7) (all from Santa Cruz Biotechnology, Santa Cruz, CA, USA). Peroxidase-conjugate anti-mouse or anti-rabbit IgG (Amersham-Pharmacia Biotech, UK or Santa Cruz) were used for enhanced chemiluminescence (ECL) detection.

Cell cycle and apoptosis analysis by flow cytometry. After the appropriate treatments, cells (1×10^6) were fixed for 30 min in 70% ethanol and pelleted by centrifugation (720 g; 5 min). After removal of ethanol, cells were incubated and resuspended in 1 ml of DNA staining solution (PBS containing 200 mg/ml RNase A, 20 mg/ml propidium iodide plus 0.1% Triton X-100) and left at room temperature for 60 min. Ten thousand events per sample were acquired using a FACScan flow cytometer (Becton-Dickinson, San Jose, CA, USA) and the percentage of cells in G₁/S, G₂/M and Sub-G₂/M phases of the cell cycle were determined using CellQuest software (Becton-Dickinson). Apoptosis was analyzed by using Annexin V staining (GenScript, Piscataway, NJ, USA). All cells were then measured on a FACScan flow cytometer with an argon laser at 488 nm for excitation and analyzed using CellQuest software. Apoptotic cells were detected by the percentage of Annexin V stained cells. The results were expressed as the percentage of death by apoptosis induced by a specific treatment.

Soft agar clonogenic assays. Soft agar assays were performed as previously described (34). Briefly, 2×10^3 cells were suspended in 0.3% Bacto-agar (Life Technologies) containing the appropriate medium supplemented with 10% FCS and/or 10 μ M U0126 and seeded in 60-mm cell culture plates. The plates were incubated at 37°C for 14 days. The numbers of colonies containing ≥ 6 cells were counted.

Small interfering RNA transfection. Tumor cells (1.5×10^5 cells/well) were plated in 6-well plates and grown in the appropriate medium supplemented with 10% FCS for 2 days. These cells were transfected with ERK1siRNA and ERK2 siRNA (sc-29308 and sc-44224 respectively, Santa Cruz Biotechnology). All siRNA duplexes were transfected using Oligofectamine reagent (Invitrogen) according to the manufacturer's instructions. After ERK transfection, cells were grown in the appropriate medium for 72 h.

Invasion and migration assays. Transwell membrane (Corning Costar Corp.) was used. Cancer cells were trypsinized, washed

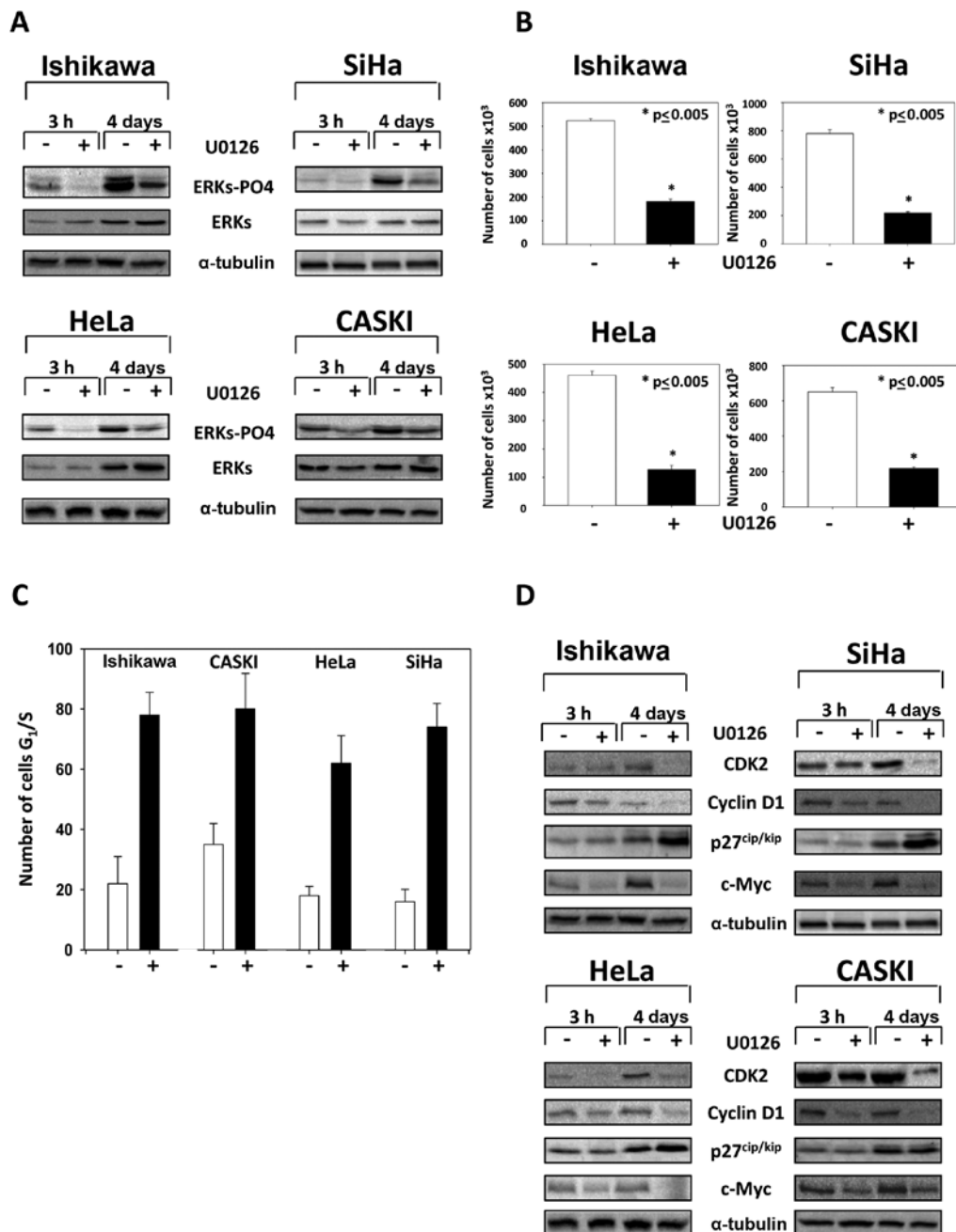


Figure 1. U0126 induces ERK inhibition and cell growth arrest in G_1/S phase of the cell cycle of Ishikawa, SiHa, HeLa and CASKI gynecological cancer cell lines. (A) Expression of protein kinases ERK1/2-P04 and ERK1/2 in untreated (-) or 3 h and 4 days U0126-treated (+) Ishikawa, SiHa, HeLa and CASKI cancer cell lines. α -tubulin was blotted as loading control. (B) Growth curve of Ishikawa, SiHa, HeLa and CASKI cancer cell lines untreated (-) or U0126-treated (+) for 4 days. Each point of the curve is the average \pm SEM of three samples. (C) Histogram showing the number of Ishikawa, SiHa, HeLa and CASKI cells in G_0/G_1 phase in untreated (-) and U0126-treated cancer cells (+) for 24 h. Similar results were obtained in three separate experiments. (D) Cell lysates from Ishikawa, SiHa, HeLa and CASKI cells untreated (-) or treated (+) with U0126 for 3 h, or 4 days were analyzed by immunoblotting with specific antibodies for indicated proteins. α -tubulin expression shows the loading of samples. Similar results were obtained in three different experiments.

and kept suspended in the appropriate medium without FCS. To the lower wells of the chambers, migration-inducing medium (with 10% FCS) was added. Upper wells were filled with serum-free medium with cells (20,000 cells per well) in the absence or in the presence of the appropriate treatments. After 8 h, filters were removed and fixed with methanol and subsequently the cells on the upper side were wiped off using a Q-tip. Filters were stained with 20% Giemsa solution. Evaluation of complete transmigration was performed under the microscope and random fields were scanned (four fields

per filter) for the presence of cells at the lower membrane side only. Invasion assays were done in a similar manner as the migration assays described above, unless the inserts were pre-coated with Matrigel (BD Biosciences).

Statistical methods. Continuous variables were summarized as the mean and standard deviation (SD) and the statistical comparisons between control and treatments were established by carrying out the ANOVA test or the t-test when appropriate. Dichotomous variables were summarized by absolute and/

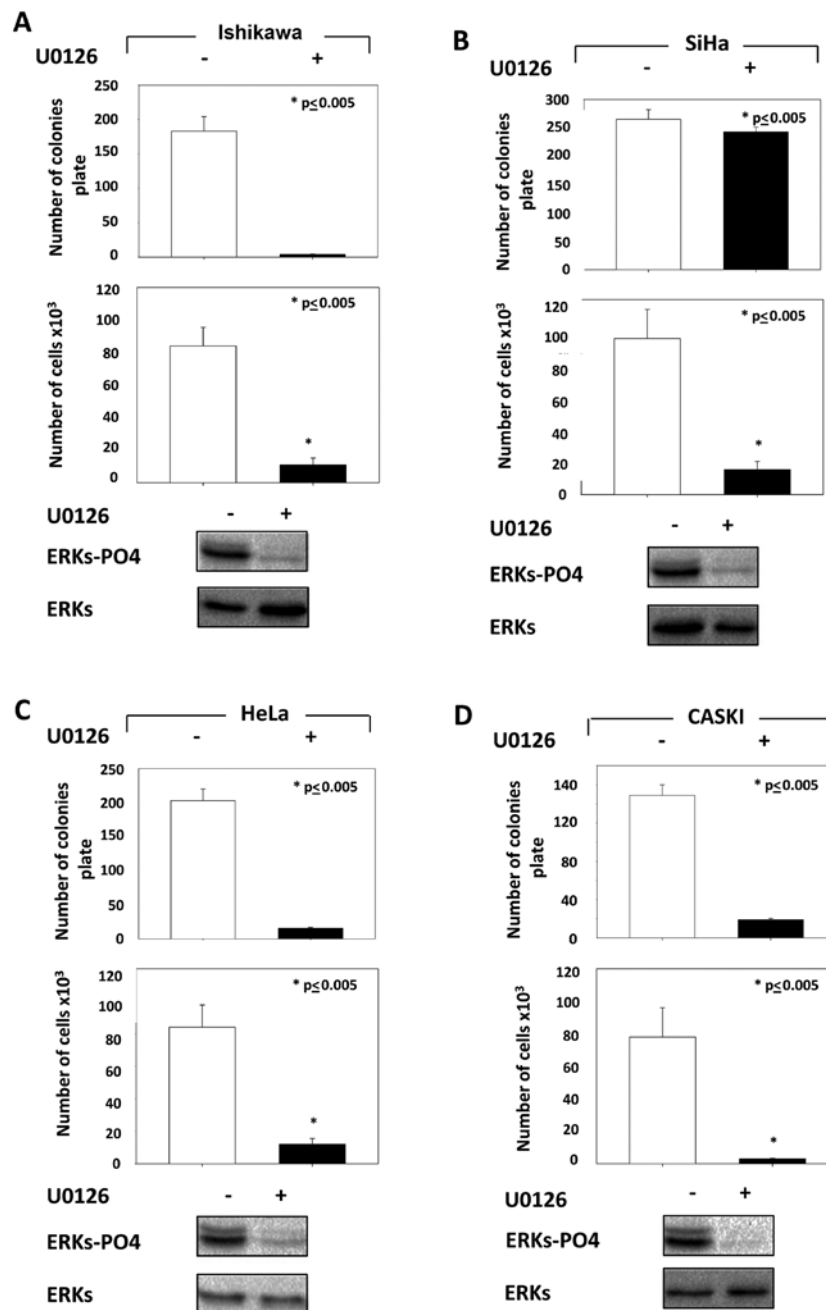


Figure 2. U0126 blocks anchorage-independent and modified monolayer growth ability of Ishikawa, SiHa, HeLa and CASKI cancer cells. (A-D, upper panel) Ishikawa, SiHa, HeLa and CASKI cells untreated (-) or treated (+) with U0126 were tested for growth in soft agar. Colonies were photographed after 14 days. (A-D, middle panel) Growth curve in modified monolayer conditions of Ishikawa, SiHa, HeLa and CASKI cells untreated (-) or treated with U0126 (+). Cells were counted after 6 days. The data shown are the mean \pm SEM of triplicates of a representative experiment. (A-D, lower panel) Expression of protein kinases ERK1/2-P04 and ERK1/2 in untreated (-) or 3 U0126-treated (+) Ishikawa, SiHa, HeLa and CASKI cancer cell lines growth in modified monolayer conditions.

or relative frequencies and statistical comparisons between control and treated groups were established by carrying out the Fisher's exact test. For multiple comparisons the level of significance was corrected according to Bonferroni correction. All tests were two-sided and were determined by Monte Carlo significance. $P < 0.05$ was considered statistically significant.

Results

Persistent ERK inhibition induces growth arrest in G_1 , inhibition of migration/invasion and promotes anchorage-independent

growth. In order to verify the effects of MEK-inhibitor, U0126, a time course experiment with or without U0126 treatment (10 μ M) was performed. Cells were treated with U0126 either for 3 h or for 4 days and subsequently processed for immunoblotting, cell count and FACS analysis. As shown in Fig. 1A and B, U0126 induced a rapid (3 h) and persistent (4 days) decrease in phospho-active ERKs (Fig. 1A) concomitant with a decrease in the proliferation rate ranging from 65.5 to 74.9% (Fig. 1B). As shown in Fig. 1C, treatment with U0126 resulted in preferential accumulation of tumor cells in the G_1 -S phase of cell cycle, with the percentage of Ishikawa, SiHa, Caski and

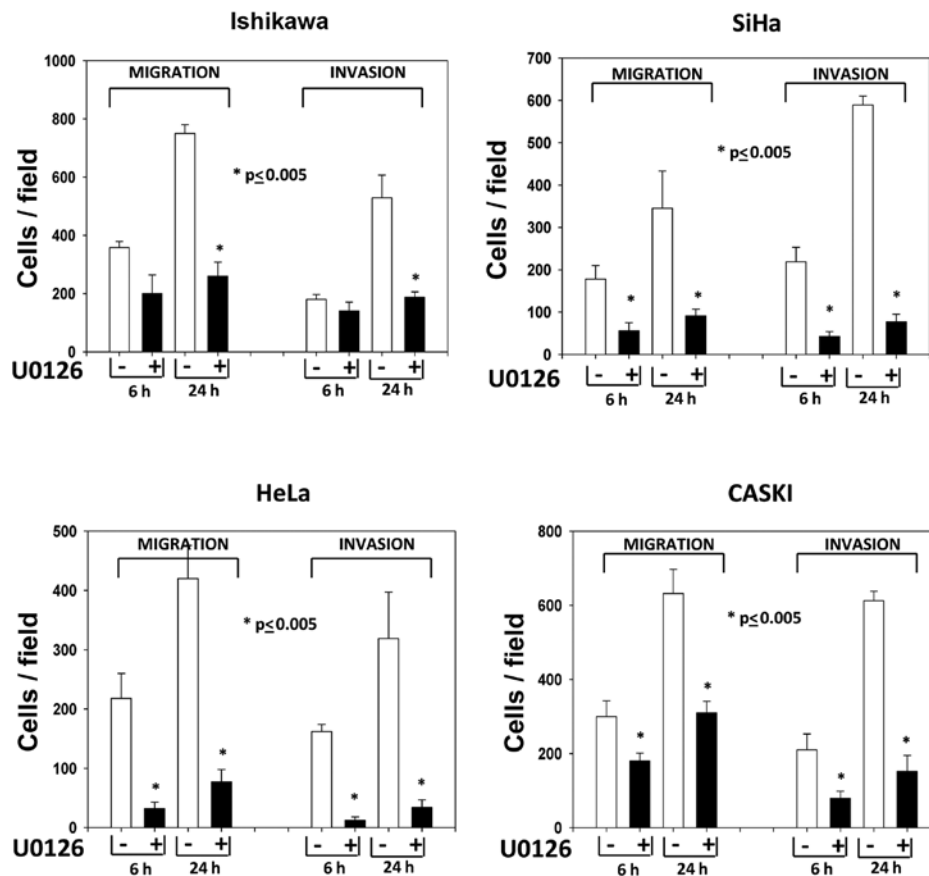


Figure 3. U0126 counteracts migration and invasion ability of Ishikawa, SiHa, HeLa and CASKI cancer cells. Transwell invasion or migration assay of Ishikawa, SiHa, HeLa and CASKI cells untreated (-) or treated (+) with U0126 for 6 or 24 h. Cells per field were counted. Similar results were obtained in three experiments.

HeLa cells in this phase of 61, 69, 74 and 77%, respectively. This phenomenon was consistent with cyclin D1 and CDK2 downregulation and p27^{Cip/Kip} upregulation (Fig. 1D). c-Myc, one of the most tumorigenic transcription factors and downstream target of ERKs, while increasing in the expression levels comparing 3 h and 4 days untreated cultured cells, was early (3 h) and persistently (≤ 4 days) inhibited by the U0126 treatment as shown in Fig. 1D.

MEK/ERK inhibition is followed by downregulation of growth signal and tumorigenic molecules. To verify whether this inhibition affects anchorage-independent growth of cancer cells, we performed soft agar and cell growth assay in modified layer. Soft agar assay showed that tumor cells cultured with vehicle only formed cell aggregates of different sizes depending on the cell line used (Fig. 2A-D, upper panel). U0126 (10 μ M) greatly affected the propensity of tumor cells to grow in modified layer (polyHEMA coated dishes) compared to controls (Fig. 2A-D, middle panel) with a decrease in the proliferation rate of 86, 84, 93 and 89% in Ishikawa, SiHa, Caski and HeLa cell lines, respectively (ANOVA test; $P < 0.01$). The decrease of phospho-active-ERK levels in tumor cells cultured on polyHEMA-coated dishes was still present after U0126 treatments (Fig. 2A-D, lower panel). Finally, U0126 (10 μ M) significantly inhibited Ishikawa, SiHa, Caski and HeLa invasion and migration both at 3 h and 4 days after treatment, with the most evident effect at the longest time (Fig. 3). All together, these

results indicate that U0126 induces growth arrest by blocking the molecular mechanism responsible of G₁/S cell cycle phase progression and reduces the signals enabling tumorigenic and metastatic potential of tumor cells.

U0126 increases the radiosensitivity of gynecological cancer cell lines by delaying DNA repair machinery and enhancing apoptotic signaling. Tumor cells were treated with U0126 (10 μ M) 24 h before the delivery of increasing doses of ionizing radiation (0-600 cGy) (Fig. 4). All cell lines were basically radioresistant at all doses tested (Fig. 4) and U0126 treatments increased the radiosensitivity with effects already evident at lowest radiation doses (Fig. 4). We further investigated whether suppressing ERKs function would influence the repair machinery of DNA double-strand breaks (DSBs) induced by irradiation. Tumor cells treated with U0126 were irradiated and the number of DNA foci representing the amount of unrepaired DSBs was counted in 4 different cell lines: i) untreated, ii) U0126-treated, iii) U0126-pretreated and iv) treated with RT (Fig. 5A-D, upper panel). For these experiments, a single dose of 4-Gy radiation was used. We have found that this dose is the most suitable to effectively distinguish the individual from combined treatments. Radiation-induced γ -H2AX foci were readily detectable at 1 h of irradiation in all treated cells (Fig. 5A-D, upper panel). Interestingly, adding U0126 together with radiation led to a substantially increased number of cells retaining these foci

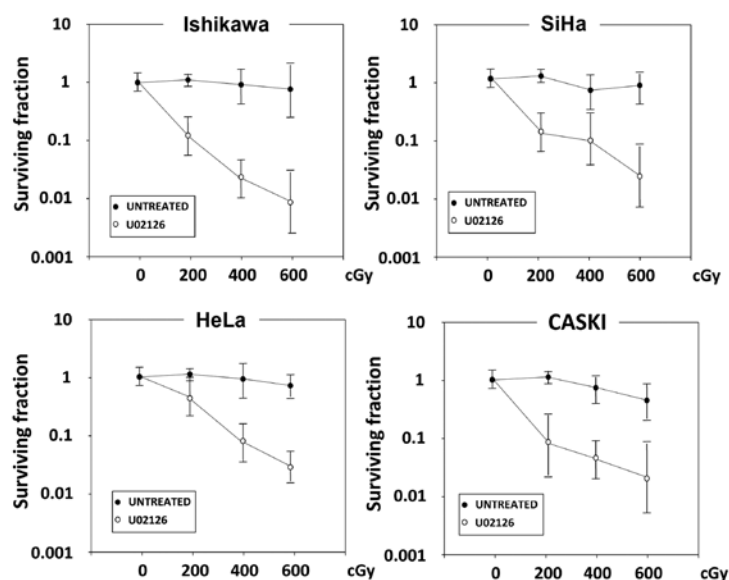


Figure 4. MEK/ERK inhibition by U0126, radiosensitizes Ishikawa, SiHa, HeLa and CASKI cells. Ishikawa, SiHa, HeLa and CASKI in the exponential phase of growth were exposed to the indicated doses of γ -radiation. Clonogenic survival was determined by counting the number of colonies containing >50 cells after 2 weeks of growth. The surviving fraction is shown in a semilogarithmic plot against radiation dose. Points, means from triplicate flasks from two to three independent experiments; bars, SE.

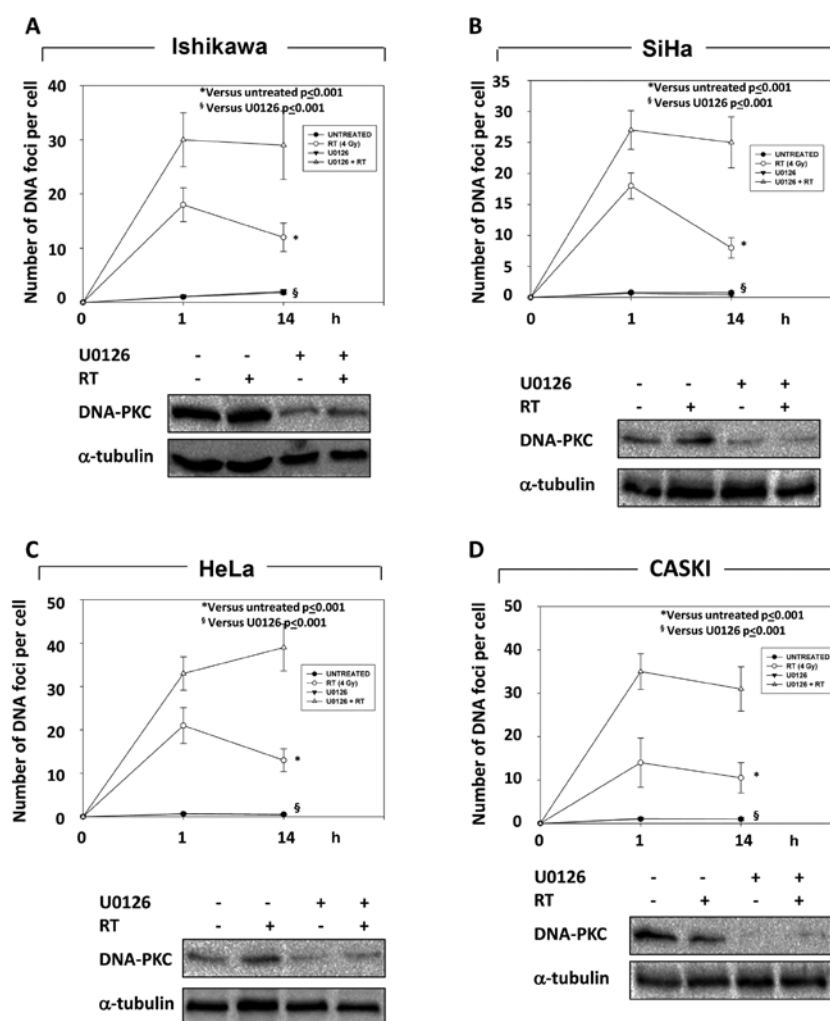


Figure 5. U0126 induces DNA damage correlated to DNA-PK protein downregulation. (A-D, upper panel) Quantification of γ -H2AX foci: DNA double-strand break profiles of Ishikawa, SiHa, HeLa and CASKI cancer cells after 24-h treatment with U0126 followed by 24 h of radiation treatment, as measured and categorized after γ -H2AX immunostaining. For each category and for each dose of radiation, the mean \pm SD of three experiments is shown. (A-D, lower panel) Expression of DNA-PK in Ishikawa, SiHa, HeLa and CASKI cancer cell lines pre-treated with U0126 and then exposed to 400 cGy of γ -radiation. α -tubulin was blotted as loading control. Similar results were obtained in three experiments.

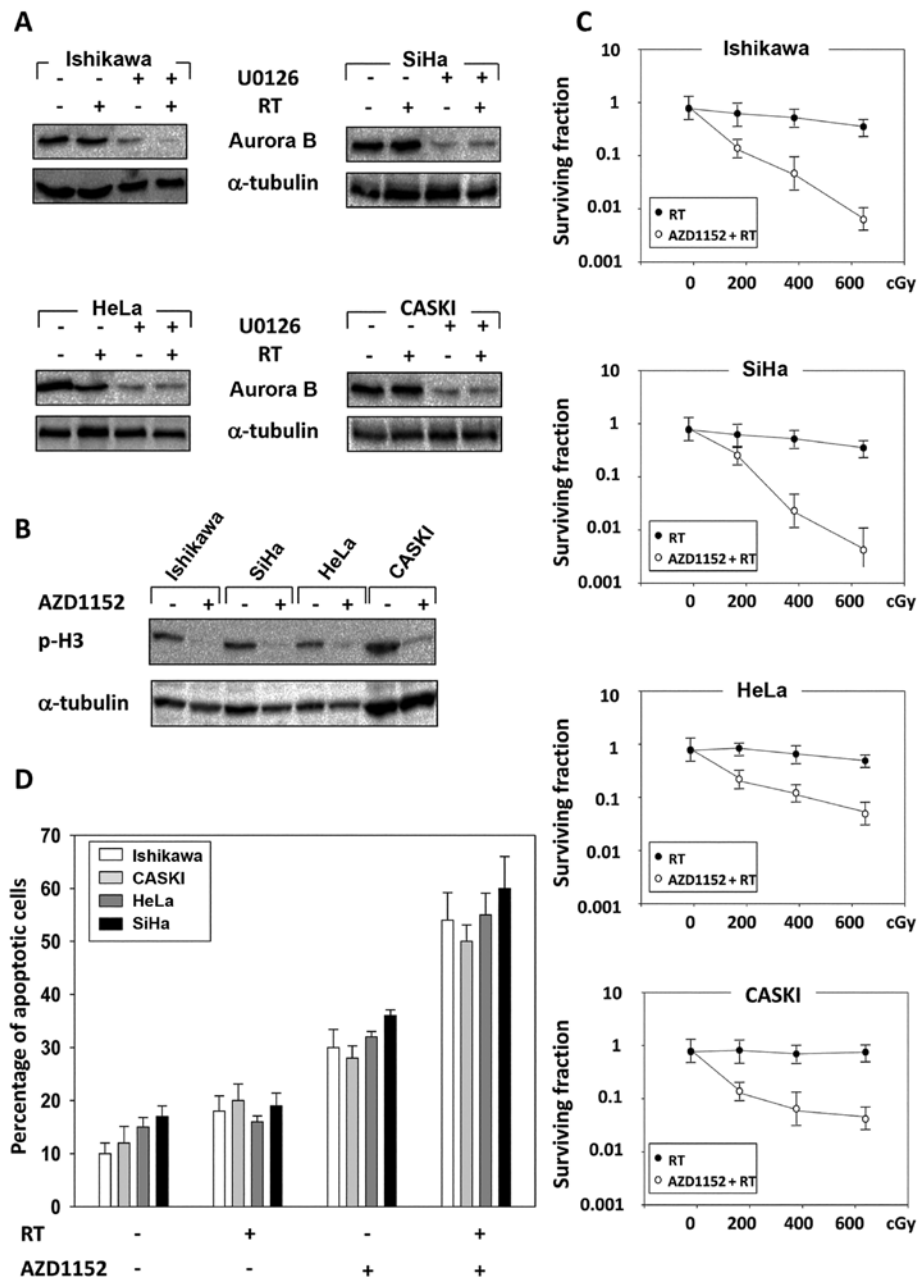
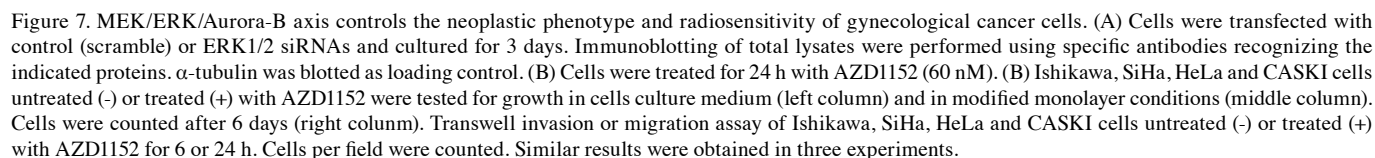


Figure 6. U0126 induces Aurora-B downregulation, inhibition of which by AZD1152 affects radiosensitivity of Ishikawa, SiHa, HeLa and CASKI cells. (A) Quantification of Aurora-B protein expression in Ishikawa, SiHa, HeLa and CASKI cancer cells lines pre-treated with U0126 and then exposed to 400 cGy of γ -radiation. α -tubulin was blotted as loading control. (B) Ishikawa, SiHa, HeLa and CASKI cancer cells after 24-h treatment with the Aurora-B inhibitor, AZD1152. The Aurora-B inhibition was assessed by quantifying the phosphorylation levels of H3. α -tubulin was blotted as loading control. (C) Ishikawa, CASKI, HeLa and SiHa in the exponential phase of growth were exposed to the indicated doses of γ -radiation in the presence or absence of AZD1152. Clonogenic survival was determined by counting the number of colonies containing >50 cells after 2 weeks of growth. The surviving fraction is shown in a semilogarithmic plot against radiation dose. Points, means from triplicate flasks from two to three independent experiments; bars, SE. (D) Annexin V staining by FACS analysis of Ishikawa, CASKI, HeLa and SiHa in the exponential phase of growth were exposed to the indicated doses of γ -radiation in the presence or absence of AZD1152. Similar results were obtained in three experiments.

for ≥ 14 h with respect to cells treated with RT alone. The difference in the percentage of cells retaining DNA foci between the two groups was significant ($P < 0.01$). Due to the fact that DNA-PKcs is usually activated in malignant tumor cells and its inactivation impairs DNA repair following ionizing irradiation we examined whether U0126 would affect DNA-PKcs expression. Of note, DNA-PKcs expression measured by western blotting at 24 h after treatments significantly decreased in all cell lines tested in the present study (Fig. 5A-D, lower panel).

Functional correlation between MEKs/ERKs and Aurora-B supports tumorigenic and radioresistant phenotype. We tested if the U0126-mediated effect on radiosensitization involved Aurora-B, known to play a central role in the control of DNA repair machinery. To this purpose cancer cells were cultured with U0126 24 h before the delivery of ionizing radiation (4 Gy) (Fig. 6) and expression levels of Aurora-B kinases was evaluated in these cell lines. As shown in Fig. 6A, MEK/ERK inhibition reduced Aurora-B expression in all cell lines used (Fig. 6A). To verify whether in these cell lines the Aurora-B



p-H3, the active phosphorylated form of histone H3 required for normal chromosomal condensation. As shown in Fig. 6B, 24 h after AZD1152 (60 nM) treatment a dramatic decrease

in p-H3 expression, consistent with inhibition of Aurora-B H3-phosphorylating activity, was observed in all cell lines. The functional inactivation of Aurora-B by AZD1152 increased the radiosensitivity of all cellular models (Fig. 6C) with a kinetic of radiosensitization similar to that observed during U0126 treatment (compare Fig. 4, with Fig. 6C). The radiosensitizing effect of AZD1152 was partially due to increased apoptotic stimulus that was greatly potentiated in association with radiotherapy (Fig. 6D) as measured by Annexin V assay. This evidence suggests a possible role of Aurora-B in the induction of pro-apoptotic stimulus upon radiation treatment.

In order to verify the existence of a direct functional link between ERK1/2 and Aurora-B, ERK1/2 RNA interference experiments were performed. Upon 72 h of treatment with ERK1/ERK2 siRNA, downregulation of total ERKs and Aurora-B protein levels was observed (Fig. 7A). Since MEK/ERK silencing caused downregulation of Aurora-B expression we further studied AZD1152 effects in cellular growth, growth in modified monolayer and in migration/invasion assays. The Aurora-B inactivation by AZ1152 significantly decreased growth rate and growth in modified monolayer (Fig. 7B, left and middle panels) as well as migration/invasion (Fig. 7B, right panels) potential of all tumor cell models. This body of evidence suggests that Aurora-B mediates tumorigenic potential of gynecological cancer cell lines controlling their invasion and metastatic potential.

Discussion

Ras/MEK/ERK signaling inhibition is considered a potential novel strategic approach for therapeutic use (10). In this study, we addressed the issue of whether MEK/ERK inhibition, by interfering with Aurora-B kinase, affects the tumorigenic phenotype and radiation response of a panel of gynecological cancer cell lines. The decrease in tumorigenic signaling induced by the MEK inhibitor, U0126, clearly demonstrated that the MEK/ERK pathway is a pre-requisite for the aberrant growth in *in vitro* and *in vivo* system of these cells. Indeed, the permanent phospho-active ERK inhibition in all tumor cell lines used was followed by growth arrest, loss of both anchorage-independent growth and migration/invasion potential. Importantly, MEK/ERK inhibition had significant effects on cell cycle distribution causing an increased accumulation of cells in G₁ phase, ranging from 65-80% depending on cell line in 24-h treatment. This anti-growth response is countersigned by a decreased levels of a number of proteins that are globally correlated with cell cycle progression (CDK2, Myc and cyclin D1) or with cell cycle inhibition (p27).

Moreover, the effects of Ras/MEKs/ERKs inhibition and the downstream target pathways on radiation response have not yet been studied in detail in gynecological tumors. With respect to intrinsic cellular radiation sensitivity, it is known that cells are more responsive to the ionizing radiation effects in G₂/M than in G₁ phase. In these lines the MEK/ERK inhibitor led a significant percent of cells (65-80%) into G₁ phase of the cell cycle within 24-h treatment, while radiation, combined with the MEK inhibitor, in just one hour gained the main effects on DNA foci. From this we can conclude that MEK inhibitor does not lower RT-mediated DNA lesions but rather helps radiosensitization as we previously reported in RMS *in vivo* and *in vitro*

models (32,34). Effects of combined MEK/ERK inhibitor and radiation are particularly evident in the persistence of DNA foci, which decays within 14 h after radiation alone. Our data point to a possible loss of DNA repair mechanisms, resulting in MEK/ERK inhibitor-induced DNA-PKcs downregulation. Our study for the first time, demonstrated a close correlation between MEK/ERK and Aurora-B in a functional axis where the loss of ERK causes Aurora-B downregulation enhancing the response to DNA damage by radiation. The idea that MEK/ERK is a therapeutic target which enhances radiosensitivity was corroborated by the MEK inhibitor-mediated loss of DNA-PKcs, a member of the DNA repair machinery.

The close correlation between MEK/ERK signaling pathway and Aurora-B is of particular significance in light of recent findings reporting that MAPKs and Aurora-B cooperate in promoting invasive growth (41-45), tumorigenic potential (46,47) and radioresistance phenomena (42-44). We successfully demonstrated that MEK/ERK kinases regulate Aurora-B expression, suggesting that MAPK pathway and Aurora-B kinase rely on a functional collaboration, which is most likely required to sustain the tumorigenic and radioresistant phenotype of the gynecological cancer cells. This evidence was supported by the findings that selective Aurora-B inhibition by AZD1152 recovered radioprotection, induced apoptotic events and elicited anti-growth responses, similarly to that seen after MEK/ERK inhibition. Aurora-B expression dependence on active ERKs is also demonstrated by the siRNA-mediated ERK silencing, which induced downregulation of Aurora-B protein expression. Collectively, our findings suggest that the modulation of MEK/ERK signaling pathway has a strong impact on Aurora-B kinase resulting in reduced tumorigenic radioresistant phenotype in all gynecological cancer cell lines.

In conclusion, results from our studies on MEK/ERK/Aurora-B axis inhibition by MEK/ERK inhibitor can be considered of high relevance in molding the new therapeutic approaches in treating tumorigenesis such as endometrial, cervical and vaginal cancers.

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