

MEK inhibition suppresses cell invasion and migration in ovarian cancers with activation of ERK1/2

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Abstract. The extracellular-regulated kinase (ERK) signaling pathway plays an important role in regulating the malignant potential of a cancer cell. However, the effect of ERK signaling on cancer metastasis is not clearly understood. In the present study, we examined the status of ERK activation in 88 ovarian carcinomas in order to clarify the clinicopathological and prognostic significance of phosphorylated ERK1/2 (p-ERK1/2). p-ERK1/2 expression was identified in 37 (42%) of 88 ovarian carcinomas. There was no significant correlation between p-ERK1/2 expression and any of the clinicopathological factors tested. No significant correlation between p-ERK1/2 expression and overall survival was found in patients with ovarian carcinoma treated with platinum and taxane chemotherapy ($P=0.426$). Next, to clarify the role of ERK1/2 activation in ovarian cancers, we inactivated ERK1/2 in ovarian cancer cells using the MEK inhibitor, CI-1040, which prevents ERK1/2 activation. Based on simulated wound healing and invasion chamber assays, we found that the motility and invasion of ES2 and MPSC1 cells with p-ERK1/2 were significantly reduced ($P<0.01$) after treatment with CI-1040. By contrast, CI-1040 did not have any effect on KF28 cells, which were negative for p-ERK1/2. Twist was down-regulated simultaneously with p-ERK1/2 following treatment of ES2 and MPSC1 cells with CI-1040. Immunohistochemistry of ovarian carcinoma tissue revealed that the increased expression of p-ERK1/2 significantly correlated with Twist expression ($P<0.01$). The findings in this study provide new insight into the biological role of ERK signaling in ovarian carcinomas. Additionally, our observations have an important therapeutic implication for patients with ovarian cancers that express p-ERK1/2 as these patients may potentially benefit from CI-1040 therapy.

Introduction

Ovarian cancer is the most lethal gynecological malignancy in the world (1). Its frequency has increased dramatically in the last decade. Despite the advent of newer screening tools, the majority of patients have peritoneal dissemination and distant metastasis at the time of diagnosis. With disseminated disease, treatment is often unsuccessful and the overall survival is low. Therefore, stage is one of the most important factors for determining prognosis (2). The identification of an invasion-related molecule associated with the early and rapid spread of ovarian cancer is the current focus of many investigators.

First-line chemotherapy with platinum drugs and taxanes yields a response rate of over 80%, but almost all patients relapse, and in nearly all patients, recurrent disease is incurable. Although there are well-established surgical and chemotherapeutic treatments for primary ovarian cancer, there is significant opportunity to develop drugs targeting specific molecular pathways to improve survival in recurrent disease. To do so requires a thorough understanding of the molecular pathways of ovarian carcinogenesis.

Several genetic alterations are associated with ovarian carcinogenesis, the most frequent of which are mutations in *P53*, *KRAS* and *BRAF* (3,4). Mutations of either *BRAF* or *KRAS* lead to constitutive activation (phosphorylation) of their downstream target, mitogen-activated protein kinase (MAPK), also known as extracellular signal-regulated protein kinase (ERK) (5,6). These mutations are correlated with overexpression of activated ERK1/2 in ovarian serous tumors (7). Phosphorylation of ERK1/2 activates downstream cellular targets (8,9), including a variety of cellular and nuclear proteins.

The RAS/RAF/MEK/MAPK signaling pathway plays a major role in various cellular activities including proliferation, differentiation, apoptosis, angiogenesis and migration (10-15). Suppression of MAPK activity by inhibitors, dominant-negative MEK1 mutants and anti-sense nucleotides reduces the migratory ability of certain cell types in response to extracellular growth stimulation (16-20). Given the important role of MAPK in cell function, we investigated whether the RAS/RAF/MEK/MAPK pathway is involved in the metastasis of ovarian cancer. In this study, we assessed the activation of MAPK in ovarian cancer tissues and examined the effects of a MEK inhibitor on cell motility and invasion.

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Materials and methods

Tissue samples. Formalin-fixed, paraffin-embedded tissue samples of 88 ovarian cancers, including 45 serous, 10 mucinous, 10 clear-cell and 23 endometrioid carcinomas, were used in this study. The samples were obtained from the Department of Obstetrics and Gynecology of the Shimane University Hospital. Diagnosis was based on a conventional morphological examination of H&E-stained sections, and tumors were classified according to the WHO classification. Tumor staging was performed according to the International Federation of Gynecology and Obstetrics (FIGO) classification. The clinicopathological characteristics of the patients included in this study are summarized in Table I. All patients were primarily treated with cytoreductive surgery and adjuvant platinum and taxane chemotherapy (CBDCA AUC5, 175 mg/m² paclitaxel or 70 mg/m² docetaxel). All cases received 6–12 courses of this regimen. The acquisition of tumor tissues was approved by the Shimane University Institutional Review Board. The paraffin tissue blocks were organized into tissue microarrays, which were made by removing 3-mm diameter cores of tumor from each block. Selection of the area to core was made by a gynecologic oncologist (K.N.) and pathology technician (K.I.) and was based on a review of the H&E slides.

Cell culture and cell lines. ES2 (clear-cell carcinoma) human ovarian cancer cell lines were obtained from the American Tissue Culture Center (Rockville, MD, USA). The human ovarian carcinoma cell line KF28 (serous carcinoma) was a kind gift from Dr Yoshihiro Kikuchi (Ohki Memorial Kikuchi Cancer Clinic for Women, Saitama, Japan) (10). The MPSC1 cell line was established from a low-grade serous carcinoma and was a kind gift from Dr Le-Ming Shih (Johns Hopkins Medical Institutions, Baltimore, MD, USA).

Immunohistochemistry. Expression levels of the active (phosphorylated) form of ERK1/2 and Twist were assessed by immunohistochemistry. The antibody used in this study was a rabbit polyclonal antibody that reacted exclusively with phosphorylated ERK1/2 (p-ERK1/2; Cell Signaling Technology). After antigen retrieval in a sodium citrate buffer, slides were incubated overnight at 4°C with antibodies to p-ERK1/2 (Cell Signaling Technology) and Twist (Santa Cruz Biotechnology, Santa Cruz, CA, USA) at dilutions of 1:1,000 and 1:100, respectively. This was followed by incubation with a biotinylated linker and streptavidin-horseradish peroxidase (LSAB2 System-HRP; Dako Cytomation, Carpinteria, CA, USA). The signals were visualized using ABC+ (Dako Cytomation) as the substrate-chromagen at room temperature for 10 min. Sections were counterstained with hematoxylin and mounted. The percentage of positive cells was estimated by randomly counting ~500 tumor cells from three different high-power fields (x40) within one specimen. A positive reaction was defined as discrete localization of the brown chromagen in the nucleus or cytoplasm. Slides for all samples were evaluated with a light microscope by two researchers; the researchers were blind to the clinicopathological factors. The antibody staining intensity was then analyzed in the cancer cell nuclei using the HSCORE (21). This modified HSCORE was calculated as follows: $HSCORE = \sum P_i(i)$, where i is the

intensity of staining (0, undetectable; 1, weakly positive; 2, moderately positive; 3, intensely positive) and P_i is a score based on the percentage of stained cells for each intensity, varying from 0 to 100%.

Western blot analysis. Cell lysates were prepared by dissolving cell pellets in Laemmli sample buffer (BioRad, Hercules, CA, USA) supplemented with 5% β-mercaptoethanol (Sigma, St. Louis, MO, USA). Western blotting was performed on ovarian cancer cell lines/cultures including ES2, MPSC1 and KF28. Similar amounts of total protein from each lysate were loaded and separated on 10% Tris-glycine-SDS polyacrylamide gels (Novex, San Diego, CA, USA) and electroblotted to Millipore Immobilon-P polyvinylidene difluoride membranes. The membranes were probed with an active ERK1/2 antibody (pTEpY, 1:5,000; Cell Signaling Technology) followed by a peroxidase conjugated anti-mouse or anti-rabbit immunoglobulin (1:20,000). The same membrane was probed with an antibody that reacted with total ERK1/2 (1:5,000; Cell Signaling Technology) for loading controls. Western blots were developed by chemiluminescence (Pierce, Rockford, IL, USA).

Simulated wound healing assay to assess cell motility. ES2, MPSC1 and KF28 cells were allowed to grow to confluence in 24-well plates in the presence of 5 μmol/l CI-1040 or 5 μmol/l DMSO (control). A linear wound was created by scraping the wells with an ART-1000E pipette tip (Molecular BioProduct, San Diego, CA, USA). The floating cells were removed by gentle washes in culture medium. The wound was observed 24 h later, and the number of individual cells in the wound was quantified as an average from multiple fields (at least five) at magnification x200 for each experiment.

Matrigel invasion assay. The invasion chamber assay was performed according to the manufacturer's instructions. Briefly, Matrigel-precoated transwell chambers with PET membranes containing 8-μm pores (BD Bioscience, Bedford, MA, USA) were soaked in Dulbecco's modified Eagle's medium and incubated for 60 min at 37°C. ES2, MPSC1 and KF28 cells were pre-treated with 5 μmol/l CI-1040 or DMSO (as vehicle control) for 30 min. For each well, 5x10⁴ cells in 0.5 ml of culture medium were added to the upper compartment of the transwell chambers. As a control, an equal number of uncoated BD Falcon TC companion plates were seeded with cells in parallel. After 24 h of incubation in the medium containing CI-1040 or DMSO, all non-invading cells in the upper compartment were removed using a cotton-tipped swap. Viable cells were stained with H&E and photographed (magnification x200). Cells were counted in several fields of triplicate membranes. Data were expressed as the percentage of cells that invaded through the Matrigel matrix-coated membrane relative to the cells that migrated through the control membrane.

Statistical analysis. Overall survival was calculated from the date of diagnosis to the date of death or last follow-up. There was no relationship between p-ERK1/2 expression and performance status distributions. Data were plotted as Kaplan-Meier curves, and the statistical significance was determined by the log-rank test. Data were censored when patients were lost to follow-up. Results are expressed as the mean ± SD. A value of

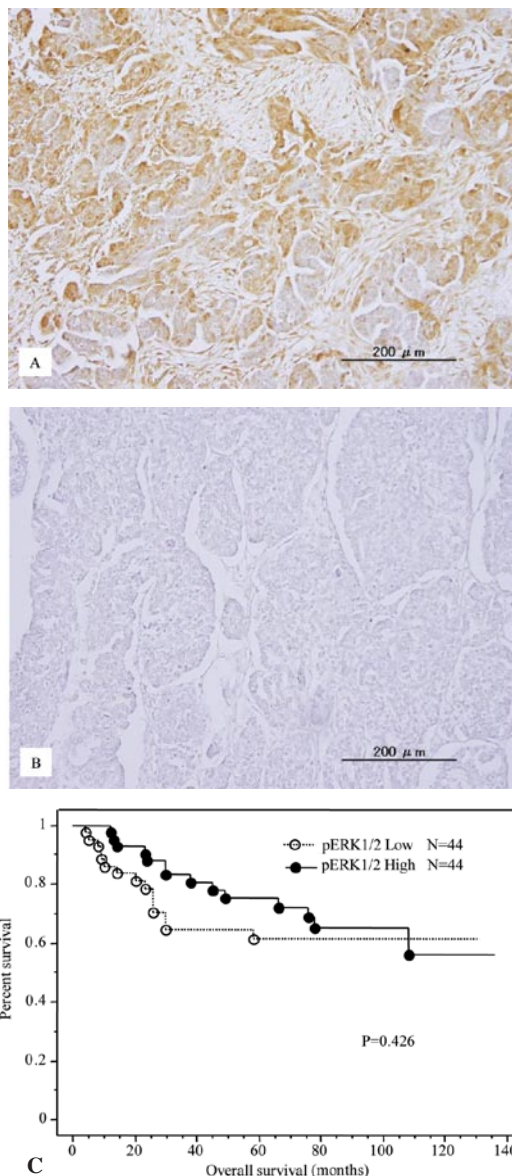


Figure 1. Immunohistochemical staining of phosphorylated extracellular-regulated kinase (p-ERK)1/2. (A) Intense immunoreactivity is present in both the nucleus and cytoplasm in this ovarian carcinoma. (B) A case showing negative staining of p-ERK1/2. (C) Kaplan-Meier survival curve in 88 patients with ovarian carcinoma according to p-ERK1/2 expression. p-ERK1/2 expression does not correlate with shorter overall survival in patients with ovarian carcinomas.

$P < 0.05$ was considered statistically significant. To generate the P-value, the Student's t-test was used. The Pearson correlation coefficient test was used to examine statistical significance in the immunohistochemical analysis values.

Results and Discussion

Relationship between p-ERK1/2 expression and clinicopathological factors. p-ERK1/2 immunoreactivity was detected in both the nuclei and cytoplasm of the tumor cells (Fig. 1). This is consistent with a previous report (22). The median HSCORE of nuclear p-ERK1/2 and the range were 18 and 0-288, respectively. Patients were stratified into one of two groups depending on the median p-ERK1/2 immunohistochemical

Table I. Association between p-ERK1/2 expression and clinicopathological factors in patients with ovarian cancer.

Factors	Patients	p-ERK1/2 immunostaining		P-value
		Low	High	
FIGO stage				
I, II	44	20	24	0.394
III, IV	44	24	20	
Grade				
G1	17	10	7	0.479
G2, G3	71	34	37	
Histology				
Serous	45	21	24	0.522
Other	43	23	20	
Age (years)				
<60	40	22	22	0.392
≥60	48	26	22	
Residual tumor				
<1 cm	47	22	25	0.522
≥1 cm	41	22	19	

HSCORE. The relationships between p-ERK1/2 expression and clinicopathological factors are shown in Table I. There was no significant correlation between p-ERK1/2 expression and the tested clinicopathological factors (Table I).

Effect of p-ERK1/2 on the prognosis of ovarian carcinomas. Next, we examined the prognostic effect of p-ERK1/2 expression. Kaplan-Meier estimates of overall survival are plotted in Fig. 1. There was no significant relationship between p-ERK1/2 expression and overall survival in patients with ovarian carcinoma ($P=0.426$). A univariate analysis demonstrated that FIGO stage III, IV ($P=0.0068$; log-rank test), patient age ≥ 60 years ($P=0.0078$; log-rank test), residual tumor ≥ 1 cm ($P<0.0001$; log-rank test) correlated with shorter overall survival.

Effects of ERK1/2 inactivation by CI-1040 lead to decreased motility and invasive capabilities in vitro. Previously, we reported the p-ERK1/2 expression in various ovarian cancer cell lines (23). Western blot analysis showed a dose-dependent effect on the expression of active ERK1/2 in MPSC1 and ES2 cells, and active ERK1/2 was not detectable 6 h after treatment of the cells with CI-1040 at a concentration of 5 μ M (Fig. 2A). To understand the phenotypic characteristics of inhibition by p-ERK1/2, we analyzed the motility and invasive abilities of MPSC1 and ES2 cells which expressed p-ERK1/2. Cell motility was investigated with a wound-healing assay. CI-1040-treated MPSC1 and ES2 cells showed 90% reduced cell motility compared to the DMSO control ($P<0.01$) (Fig. 3A and B). Following the Matrigel invasion assay, an 85 and 79% decrease in cell invasion was observed in CI-1040-treated MPSC1 and ES2 cells in comparison to the DMSO control ($P<0.05$) (Fig. 4A). These results were not observed in p-ERK1/2-negative KF28 cells following CI-1040 treatment (Fig. 4B).

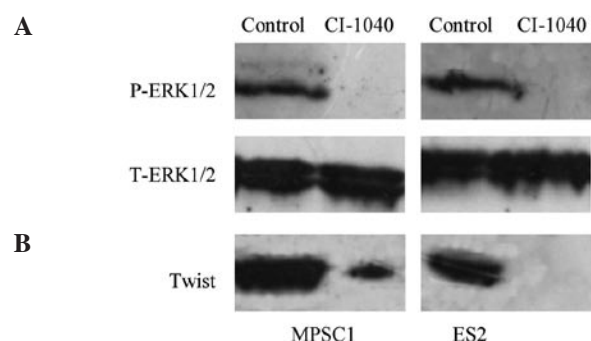


Figure 2. Western blot analysis. (A) Expression of p-ERK1/2 is undetectable in all CI-1040-treated samples. A similar amount of protein was loaded in CI-1040- and DMSO-treated (control) samples as evidenced by a similar intensity of total ERK1/2. (B) Expression of Twist is undetectable or reduced in all CI-1040-treated samples.

ERK1/2 inactivation by CI-1040 leads to decreased motility and invasion via down-regulation of Twist. Tumor progression and invasion are complex biological processes that involve the remodeling of stromal tissue by invading cells. Twist appears to play a key role in these processes. Twist expression activates dormant developmental pathways in invading tumor cells (24). Twist has recently been associated with metastasis in ovarian (25), liver (26) and breast cancer (27). Exogenous expression of Twist promotes colony formation in anchorage-independent assays (28). Our results demonstrated that MEK inhibition by CI-1040 leads to suppression of invasion and migration. Therefore, suppression of invasion-related molecules such as Twist may be one mechanism by which p-ERK1/2 suppresses metastasis. In this study, ERK1/2 inactivation by CI-1040 decreased the expression of Twist with a subsequent significant reduction in cell motility and invasiveness (Fig. 2B).

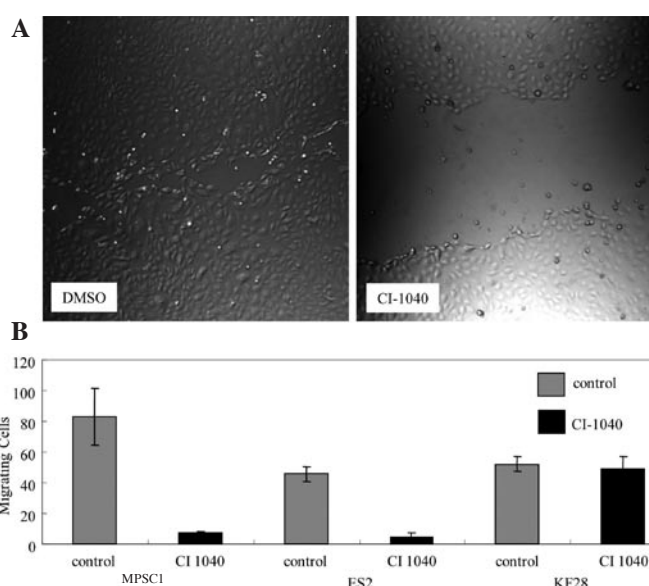


Figure 3. The effects of a MEK inhibitor on cell migration. (A) A simulated wound was created by scraping a confluent monolayer of ES2 cells. Compared to the ES2 cells treated with DMSO, there was a significant reduction in the number of CI-1040-treated ES2 cells that migrated into the wound. (B) Inactivation of p-ERK1/2 affects cell migration in ES2 and MPSC1 (p-ERK1/2-positive) but not in KF28 (p-ERK1/2-negative) cells.

Next, we analyzed p-ERK1/2 and Twist expression by immunostaining in the ovarian carcinoma samples. The Twist immunohistochemical HSCORE significantly correlated with the p-ERK immunohistochemical HSCORE ($r=0.370$, $P=0.003$; Fig. 5). Expression of Twist (Twist immunohistochemical HSCORE >0) was observed in 24% (21/88) of the analyzed tumors. Consideration of these *in vitro* findings in the context of our *in vivo* results suggests that Twist may be an

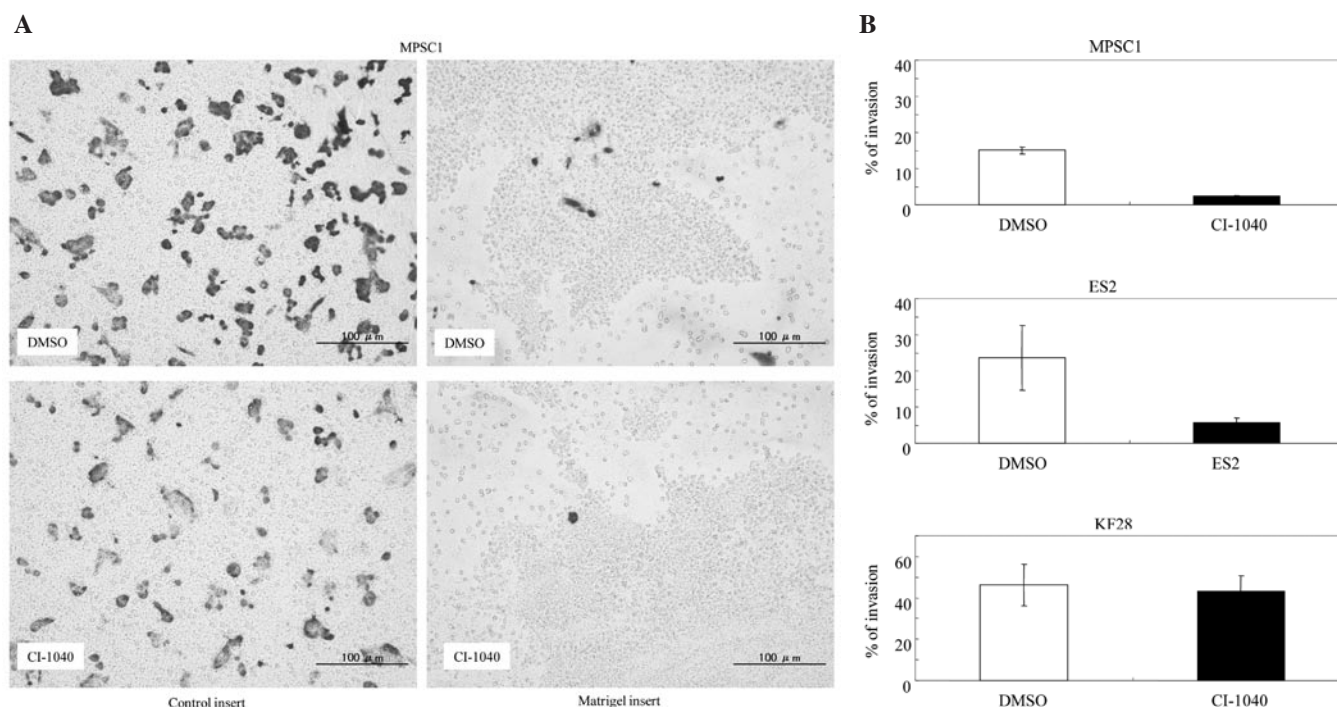


Figure 4. Matrigel invasion chamber assay. (A and B) The invasion abilities of ES2, MPSC1 and KF28 cells were evaluated by their ability to invade and penetrate through the Matrigel-coated pores on membrane inserts in transwells. DMSO-treated ES2 and MPSC1 cells show a significantly higher invasive capacity than the CI-1040-treated ES2 and MPSC1 cells. KF28 cells are not affected by CI-1040.

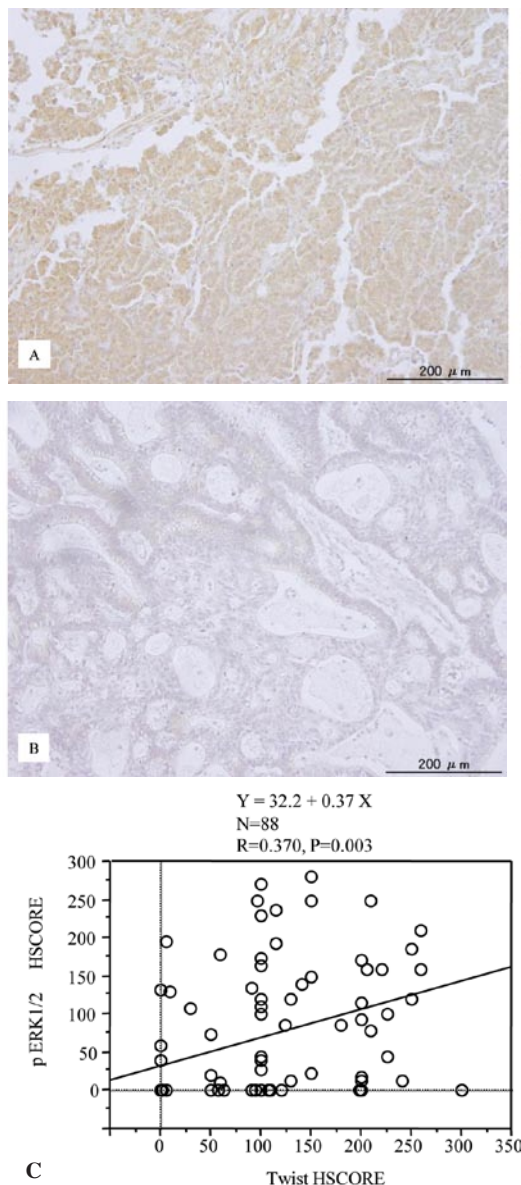


Figure 5. Twist immunoreactivity in ovarian cancer tissues. (A) Intense immunoreactivity is present in the nuclei of carcinoma cells. (B) In this sample negative staining for Twist is noted. (C) Correlation between the p-ERK1/2 immunohistochemical score and the Twist immunohistochemical score in ovarian cancer tissue. The HSCORE for p-ERK1/2 significantly correlates with the Twist immunohistochemical HSCORE ($r=0.370$, $P=0.003$).

important downstream participant in MAPK signaling. In the present model, p-ERK1/2 expression was positively correlated with the Twist expression level in the regulation of metastasis. Further studies are required to confirm whether Twist is a downstream target of ERK signaling.

The development of a highly invasive ovarian cancer phenotype requires coordinated up- or down-regulation of many signaling pathways. Identification of the active MAPK-regulated molecules in ovarian cancer is important to gain further insight into the role of the MAPK signaling pathway in the metastasis of ovarian cancer. In the present study, we observed a profound reduction in cell motility, as evidenced by wound healing and invasion assays in CI-1040-treated MPSC1 and ES2 cells with positive p-ERK1/2. The lack of effect of the MEK inhibitor on motility and invasion in KF28 ovarian

cancer cells was expected given that they do not express p-ERK1/2.

While we measured Twist expression as an effect of MAPK, other downstream effectors may also contribute to cell motility and invasion in ovarian cancer. Several studies have shown that MAPK activation leads to extracellular matrix-dependent cell spreading and migration (13,17,19). The mechanisms for this likely involve integrin activation (29), integrin-dependent adhesion or cytoskeletal organization, and phosphorylation of myosin light chain kinase, calpain or focal adhesion kinase (17-19). These activities directly or indirectly enhance cell motility. In addition to promoting cell motility and migration, constitutive activation of MAPK enhances matrix metalloproteinase activation, a key event during cellular invasion (30). The present study is consistent with previous reports showing that MAPK pathway inhibitors are potent in suppressing cell migration in tumor cells (17,31,32). Blockade of the MAPK pathway by treatment with MEK inhibitors correlates well with the inhibition of invasion of tumor cells derived from rhabdomyosarcoma, fibrosarcoma, bladder carcinoma, colon carcinoma, prostate carcinoma and breast carcinoma cell lines (33). These results suggest that the inactivation of the MAPK pathway may have an anti-metastatic effect on tumor cells.

Our results provide compelling evidence that the biological effects of the ERK signaling pathway depend on the p-ERK1/2 status. We found that ovarian carcinomas with p-ERK1/2 expression were more sensitive to cell migration and invasion inhibition by the MEK inhibitor, CI-1040. This observation suggests that ovarian carcinomas with p-ERK1/2 expression are more highly dependent on the activation of the MEK-ERK pathway for cell migration and invasion than those without p-ERK1/2 expression. Thus, inactivation of ERK1/2 results in marked inhibition of cell migration and invasion of ovarian carcinomas with p-ERK1/2 expression in comparison to only a modest effect on p-ERK1/2-negative tumors. The above observations lend strong support to the view of 'kinase addiction', in which dependence on a particular kinase pathway confers susceptibility to a kinase inhibitor (34,35).

In light of our *in vivo* and *in vitro* findings, we propose that recurrent ovarian cancer patients with p-ERK1/2 expression should be considered for MEK inhibitor (CI-1040) therapy. Additionally, patients with primary tumors that express p-ERK1/2 may also benefit from MEK inhibitor therapy in combination with conventional platinum and taxane chemotherapy. The use of a MEK inhibitor at the time of initial chemotherapy may decrease the invasive potential of any residual tumor cells, thereby extending the disease-free interval.

Thus far, the MEK inhibitor CI-1040 has fared poorly in clinical trials for breast, colon and lung cancer (36). It is possible that this may be due to non-selective use of the inhibitor. It is possible that more favorable outcomes may result when patients are stratified based on p-ERK1/2 expression status. We recommend that stratification is used in further clinical trials of MEK inhibitors in ovarian cancer patients.

In summary, we demonstrated that the phenotypic changes in cell migration and invasion in ovarian carcinomas in response to MEK inhibition depend on p-ERK1/2 status. The findings of this study provide new insight into the biological

roles of MAPK signaling in ovarian carcinomas. Additionally, our observations have an important therapeutic implication for patients with ovarian cancers expressing p-ERK1/2. Therefore, detection of p-ERK1/2 in ovarian cancers may identify patients who will benefit from CI-1040 therapy.

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