Overexpression of p21-activated kinase 1 promotes endometrial cancer progression

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Abstract. Endometrial cancer (EC) is the most common gynecologic malignancy, but the molecular events involved in the development and progression of EC remain unclear. P21-activated kinase 1 (Pak1) plays important roles in cell motility and survival. This study investigated the clinical significance of Pak1 expression and its functional roles in EC. The expression of Pak1 in clinical samples and EC cell lines was evaluated. The effects of Pak1 on EC cell functions were determined by either overexpressing it via plasmid transfection or depleting its expression using short hairpin RNA (shRNA) in human EC cell lines. Pak1 was overexpressed in clinical samples of EC compared with normal endometrium. High Pak1 expression in EC was positively correlated with lymph node metastasis, advanced disease stage and poor histological differentiation. Pak1 overexpression was also observed in multiple human EC cell lines. In EC cell lines, Pak1 overexpression promoted cell proliferation, migration, invasion and anchorage-independent growth in vitro. Conversely, shRNA-mediated stable knockdown of Pak1 reduced cell proliferation, migration, invasion and anchorage-independent growth. In addition, ectopic Pak1 overexpression protected EC cells from apoptosis, along with decreased caspase-3 activation. These results suggest that Pak1 plays important roles at multiple stages of EC progression.

Introduction

Endometrial carcinoma is the most common malignancy of the female genital tract, with an estimated 46,470 new cases and 8,120 deaths in the United States in 2011 (1). In other countries, such as China, its incidence is on the rise (2). For young women who wish to retain their fertility, only 50% show a complete response to hormonal treatment (usually progestins), while 25% suffer a relapse after a temporary response (3). Patients with metastatic and refractory cancer are at a significantly higher risk of morbidity and mortality, with a median survival of 8-16 months (4). The poor prognosis is attributed to the significant failure rate of traditional adjuvant therapy, including radiotherapy, chemotherapy and hormonal therapy, following tumor debulking (5). To improve the outcome of patients with endometrial carcinoma, it is important to investigate molecular pathways that are critical to the development of the disease, and to identify novel targets for therapy.

The p21-activated kinases (Paks) are a family of non-receptor serine/threonine kinases that integrate various signaling pathways that are vital to normal cell survival and function (6). Based on their sequence and functional similarities, six mammalian Paks are subdivided into two groups: group I (pak1-3) and group II (pak4-6) (7). Pak1 was originally identified as a downstream effector of the Rho family small GTPases Cdc42 and Rac1, which play a fundamental role in cytoskeleton reorganization and motility (8,9). Subsequent studies have revealed that Pak1 is also involved in regulating cell growth (10), apoptosis (11), adhesion (12) and angiogenesis (13), all of which are important for tumorigenesis and metastasis.

Pak1 is widely expressed in a variety of normal tissues, such as brain, muscle and spleen (14). Previous studies indicated that Pak1 expression is significantly increased in breast, colorectal and ovarian cancer (15-17). Functional studies have also implicated Pak1 in cell transformation and tumor cell invasion (14). For instance, Pak1 has been shown to promote non-small cell lung cancer (NSCLC) cell motility and invasiveness by phosphorylation of CRK-II (c-Crk) (18). In addition, Pak1 was able to promote mammary epithelial hyperplasia through phosphorylation and transactivation of estrogen receptor-α (ERα) (10), which is also involved in the pathogenesis of endometrial cancer (EC) (19). However, the expression pattern and the function of Pak1 in EC remain unknown.
In this study, we sought to determine whether Pak1 is involved in the pathogenesis of EC. The expression of Pak1 in EC cell lines and tissues was assessed and correlated with clinicopathological parameters. We also investigated the effect of Pak1 on EC cell proliferation, migration, invasion, anchorage-independent growth and survival. Our results indicate that Pak1 is a multifunctional regulator of the progression of EC.

Materials and methods

Clinical samples. A total of 42 paraffin embedded EC tissues (including 10 metastasis-positive lymph nodes) were retrieved from 32 patients with endometrial carcinoma at the Department of Pathology at the International Peace Maternity and Child Health Hospital, affiliated to Shanghai Jiao Tong University, China, from December 2009 to November 2011. The diagnosis of each sample was established by two pathologists according to the criteria of the International Federation of Gynecology and Obstetrics (FIGO). These patients ranged in age from 44 to 78 years (mean, 59 years). None had hormone therapy, radiotherapy, or chemotherapy prior to surgery.

Twenty-three normal endometrial samples were obtained from patients who underwent a hysterectomy to treat other diseases, such as myoma or adenomyosis. These patients ranged in age from 27 to 64 years (mean, 46 years). Ten endometrial hyperplasia tissues (5 without atypical cases and 5 with atypical cases) were collected from patients who had undergone hysteroscopic examination due to irregular bleeding. The collection of clinical samples was approved by the Ethics Committee of the Medical Faculty of Shanghai Jiao Tong University, China. All patients provided written informed consent.

Immunohistochemistry analysis. The immunohistochemical staining procedure was as follows: formalin-fixed sections were deparaffinized with xylene and hydrated in alcohol; for antigen retrieval, slides were immersed in citrate buffer (pH 6.0) and boiled for 20 min. These sections were then incubated in 0.3% H₂O₂ in methanol for 30 min to inactivate endogenous peroxidase activity. Non-specific reaction was blocked by incubation with 5% bovine serum albumin for 10 min. Sections were incubated with biotinylated secondary antibody of the same species was used as a negative control. Subsequently, slides were incubated with biotinylated secondary antibody (Mrbiotech, Emeryville, CA, USA). Primary antibodies included: Pak1 forward: 5'-AGTTTCAAGAGATTAG GATGATGA-3', reverse: 5'-AATACACAGGCTTTGTA TGAG-3', and β-actin forward: 5'-CAGCCATGTCGTTTG CTATCCAGG-3', reverse: 5'-AGGTCGACGCGAGATG GCATG-3'. Relative transcript expression levels were calculated using the 2^(-ΔΔCt) method using β-actin as internal reference. All data were obtained in triplicate in three independent experiments.

Western blot analysis. Cells were harvested with ProteoJET Mammalian Cell Lysis Reagent (MBI Fermentas, Burlington, ON, Canada) with a protease inhibitor cocktail (Roche Diagnostics, Basel, Switzerland). A total of 60 µg protein was separated by SDS-PAGE, and transferred to a polyvinylidene fluoride (PVDF) membrane. Membranes were incubated with primary antibodies. Signal was detected using BeyoECL Plus (Beyotime, Shanghai, China). Primary antibodies used included: Pak1 forward: 5'-AGTTTCAAGAGATTAG GATGATGA-3', reverse: 5'-AATACACAGGCTTTGTA TGAG-3'; and β-actin forward: 5'-CAGCCATGTCGTTTG CTATCCAGG-3', reverse: 5'-AGGTCGACGCGAGATG GCATG-3'. Relative transcript expression levels were calculated using the 2^(-ΔΔCt) method using β-actin as internal reference. All data were obtained in triplicate in three independent experiments.

Real-time qRT-PCR. Total RNA was extracted from cells using TRizol reagent (Invitrogen). cDNA was reverse-transcribed from total RNA using Prime Script RT reagent kit (Takara Inc., Otsu, Japan). Real-time PCR was performed using SYBR Premix Ex Taq (Takara Inc.) and analyzed with an ABI Prism 7000 Sequence Detection System. The oligonucleotide primers used included: Pak1 forward: 5'-AGTTTCAAGAGATTAG GATGATGA-3', reverse: 5'-AATACACAGGCTTTGTA TGAG-3'; and β-actin forward: 5'-CAGCCATGTCGTTTG CTATCCAGG-3', reverse: 5'-AGGTCGACGCGAGATG GCATG-3'. Relative transcript expression levels were calculated using the 2^(-ΔΔCt) method using β-actin as internal reference. All data were obtained in triplicate in three independent experiments.

Cell culture and isolation of human endometrial epithelial cells. Four EC cell lines were purchased from ATCC (American Type Culture Collection, Rockville, MD, USA). All cell lines were cultured in DMEM/F12 (1:1) (Gibco, Auckland, New Zealand) and supplemented with 10% fetal bovine serum (FBS) (HyClone, Logan, UT, USA) in a 37°C, 5% CO₂ incubator.

Primary human endometrial epithelial cells were cultured as described by Zhang et al (21) and Osteen et al (22). Briefly, the normal endometrial tissue was minced into small pieces and digested with 0.1% trypsin-EDTA for 30 min at 37°C. Using serial filtration, tissue debris was separated with 150-µm aperture sieves. Epithelial glands were retained on 40-µm aperture sieves and backwashed with PBS. The separated endometrial cells were cultured in DMEM/F12 (1:1) supplemented with 10% FBS. All specimens were confirmed as normal by histological analysis.

In vitro migration and invasion assays. Cell migration and invasion assays were performed using culture medium-treated 6.5-mm transwell chambers with 8.0 µm polycarbonate membranes (Corning LifeScience, Corning, NY, USA).

Proliferation assays. Cells (2x10⁴ cells/well) were plated in 96-well plates. Cell number was measured every 24 h by MTT assay following the manufacturer’s instructions (Beyotime). The OD value of each well was detected at 570 nm. Medium was changed every other day. Each experiment was repeated in triplicate.

Plasmid and transfection. To stably express Pak1 in AN3CA and Ishikawa cells, the cells were transfected with Pak1 expressing vector or the empty control vector pEGFP-N1 (Clontech Laboratories, Palo Alto, CA, USA) using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA), and then selected with G418 (800 µg/ml; Sigma Chemical, St. Louis, MO, USA). To stably silence Pak1, the cells were transfected with a set of shRNA constructs (GeneCopoeia, Germantown, MD, USA) against human Pak1 or the scrambled control vector psiHIV-U6 (GeneCopoeia), and then selected with puromycin (0.5 µg/ml; Sigma Chemical).
Matrigel (BD Biosciences, San Jose, CA, USA) coated transwell membrane was used for invasion assay. Cells were seeded in serum-free medium on the upper compartment of each transwell chamber at a density between $2 \times 10^5$-$1 \times 10^6$ cells/ml (200 µl/chamber). The lower chamber was filled with culture medium containing 10% FBS. After 24 h of incubation, non-migrated or non-invaded cells were removed from the upper chamber. The invaded or migrated cells on the lower side of inserts were fixed and stained with 5% crystal violet. Attached cells were lysed with 10% ethyllic acid, and absorbance (proportional to the number of cells) was measured at 590 nm.

Soft agar colony assays. Cells were seeded in 0.3% top agar in growth medium over a layer of 0.6% agar in a 6-well plate at a density of $1 \times 10^4$ cells/ml. After 3 weeks of incubation, colonies of $>50$ cells were produced. Only colonies with $>50$ cells were counted and photographed with an inverted microscope. All assays were performed at least three times in triplicate.

Hoechst staining. Cells were seeded in 24-well plates and treated with TNF-α (20 ng/ml; Peprotech, Rocky Hill, NJ, USA). At the indicated time point, cells were washed in PBS and fixed in 4% (wt/vol) paraformaldehyde for 16 h at 4°C. Cells were washed in PBS and permeabilized with 0.2% Triton X-100 for 20 min at room temperature. Permeabilized cells were then treated with Hoechst 33258 (10 mg/ml) for 30 min at room temperature. Photomicrographs of cells were captured under fluorescence at a magnification of x400.

Flow cytometry. Annexin V-TRITC apoptosis detection kit (KeyGen Biotech, Shanghai, China) was used to identify the percentage of apoptosis, following the manufacturer's instructions. Briefly, after stimulation, both floating and attached cells were collected by low speed centrifugation. After washing in PBS, cells were stained with Annexin V-TRITC and 7-AAD. Following incubation for 30 min at room temperature, the cells were analyzed by flow cytometry (Becton Dickinson FACScan, Immunocytochemistry Systems, San Jose, CA, USA). Tests were performed in triplicate.

Caspase activity assay. The enzymatic activity of caspase-3 was measured according to the manufacturer's instructions (Caspase-3 Activity Assay kit, Beyotime). After 0-12 h of TNF-α exposure, $2 \times 10^6$ cells were collected and suspended in cell lysis buffer. Cell lysates were mixed with reaction buffer at 37°C for 1 h. An increase in absorbance at 405 nm was used to quantify the activation of caspase-3 activity. Caspase activity was expressed as percentage of enzyme activity compared to control. Control groups received 0.1% DMSO. All the experiments were carried out in triplicate.

Statistical analysis. All tests were carried out with SPSS 16.0 (Microsoft, Redmond, WA, USA) or Prism (GraphPad,
San Diego, CA, USA). Each experiment was performed at least three times. Where applicable, data are shown as the means ± SD. The two-tailed Student's t-test or Mann-Whitney U test were used for comparison of data between two groups. Differences were considered statistically significant at P<0.05.

Results

Pak1 is overexpressed in endometrial cancer tissues and cell lines. Immunohistochemistry staining showed that Pak1 protein was predominantly localized to the cytoplasm of endometrial epithelial cells. There was only weak or no staining in normal endometrium, whereas moderate to strong Pak1 immunostaining was found in endometrial hyperplasia and endometrial carcinoma tissues. The most intense staining was found in lymph node metastases (Fig. 1A).

To account for both stain intensity and the uniformity of stain, a composite histoscore (percentage epithelium stained x stain intensity) was calculated. Analyzed by the Kruskal-Wallis rank test, cytoplasmic Pak1 expression (Pak1 composite histoscore) progressively increased in the epithelial cells of endometrial hyperplasia, invasive carcinomas, and lymph node metastases relative to the normal endometrium (P<0.001; Table I). We next explored the correlation of Pak1 expression levels with clinicopathological parameters in endometrial carcinomas. Significantly higher cytoplasmic Pak1 expression was found in carcinomas of advanced stage (stages III and IV) and poor histological differentiation (grade 3) (all P<0.05; Table I). Additionally, increased Pak1 expression was significantly associated with depth of myometrial invasion, cervix involvement and vascular space invasion (all P<0.05; Table I). However, there was no significant difference between endometrioid and non-endometrioid (serous and clear cell histological subtypes) endometrial carcinomas (P=0.051; Table I).

<table>
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<th>Characteristics</th>
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<td>Late (III-IV)</td>
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Significance of difference (P-value) between categories was analyzed by *Kruskal-Wallis rank test and ^Mann-Whitney U test, respectively. Those with significant P-values are underlined. A total of 10 specimens from eight patients.
Moreover, using real-time qRT-PCR and immunoblotting, we detected higher levels of Pak1 mRNA (Fig. 1B) and protein (Fig. 1C) in EC cell lines compared with primary cultured normal endometrial epithelial cells. The elevated Pak1 expression in endometrial carcinoma determined above suggests that overexpression of Pak1 might contribute to the malignant progression of human EC.

**Pak1 promotes endometrial cancer cell proliferation.** Immuno-histochemical staining showed that Pak1 expression was associated with the progression of EC. To investigate the potential role of Pak1 in EC cell proliferation, AN3CA and Ishikawa cells were stably transfected with Pak1 expressing vector or empty control vector (Fig. 2A). Proliferation assays revealed that stable overexpression of Pak1 significantly induced AN3CA and Ishikawa cells compared with cells transfected with control vector (P<0.001, P<0.001). Conversely, shRNA-mediated knockdown of Pak1 in AN3CA and Ishikawa cells resulted in significantly decreased cell migration and invasion. Cells transfected with shRNA expression vector showed >50% defective migration and invasion compared with cells transfected with scrambled control vector (Fig. 3C and D) (P<0.001, P<0.001). These results indicate that Pak1 is an important participant in EC cell invasion and metastasis.

**Pak1 promotes endometrial cancer cell anchorage-independent growth.** We next investigated whether Pak1 is required in anchorage-independent growth, a hallmark of oncogenic transformation (23). The soft agar assays were performed to test the transformation of EC cells (24). Pak1 expressing vector and control vector transfected EC cells were plated in soft agar and scored for colony formation after 3 weeks. As shown in Fig. 4A, there was a significant difference in the number and size of colonies between vector-transfected control cells and Pak1 overexpression AN3CA and Ishikawa cells (P<0.001, P<0.01). By contrast, stable knockdown of Pak1 almost completely abolished colony formation of EC cells in soft agar (Fig. 4B). These findings indicate that Pak1 promoted EC cell anchorage-independent growth.
Pak1 protects endometrial cancer cells from apoptosis via inhibition of caspase-3 activation. Since overexpression of Pak1 results in increased growth of EC cells, we sought to determine whether Pak1 also plays a protective role against apoptosis in EC cells. To elucidate this possibility, vector control cells and Pak1 overexpressing EC cells were treated with TNF-α. As shown in Fig. 5A, at 2 h, more control cells had condensed nuclei compared to Pak1 overexpressing cells, as determined by Hoechst staining. At 12 h, flow cytometry using Annexin V and 7-AAD showed that a higher percentage of apoptotic cells were observed in control cells compared with Pak1 overexpressing cells (Fig. 5B and C). To further elucidate the mechanism by which Pak1 acts, we tested the activation of caspase-3, a well known downstream target of the TNF-α induced death pathway (25). We found that overexpression of Pak1 markedly abolished TNF-α induced caspase-3 activation (Fig. 5D). Our results indicate that, at least in response to TNF-α, Pak1 plays an essential role in protecting cells from apoptosis via inhibition of caspase-3 activation.

Discussion

Pak1 is upregulated and activated in several human tumor types (6). However, the expression of Pak1 in EC tissue and its relationship with pathological parameters remains unknown. In the present study, significantly increased Pak1 protein expression in clinical EC samples and cell lines was detected relative to normal endometrial tissues and primary cultured normal endometrial epithelial cells. Furthermore, Pak1 protein expression is additionally increased in lymph node metastases. These data indicate that Pak1 expression is specifically increased in the most advanced lesions of EC. We found a significantly positive correlation between Pak1 expression with advanced disease stage and poor histological differentiation, which further suggests that Pak1 expression is increased with EC progression. Collectively, these data indicate that overexpression of Pak1 might contribute to malignant progression of human EC.

Pak1 has been shown to be involved in the rapid proliferation of cancer cells (17,26). Pak1 promotes proliferation
Figure 4. Pak1 promotes EC cell anchorage-independent growth. (A) Stable EC cell lines expressing Pak1 were assayed for anchorage-independent growth in soft agar. AN3CA and Ishikawa cells that had been stably transfected with the empty vector were used as controls. Colony formation (≥50 cells) was assessed using a colony counter. (B) Knockdown of Pak1 led to a significant decrease in the colony-forming ability of EC cells. Representative images (left) were captured with an inverted phase contrast microscope (magnification, x200). Columns (right), mean number of colonies from three independent experiments, each in triplicate; bars, SD; **P<0.01, ***P<0.001.

Figure 5. Pak1 prevents caspase-3 activation and protects cells from TNF-α-induced apoptosis. (A) Nuclear condensation after 2 h of TNF-α (20 ng/ml) treatment was detected by fluorescence microscopy analysis of control and stable Pak1-expressing EC cells stained with Hoechst 33258. Photographed representative numbers of apoptotic cells are shown. (B) FACS analysis for Annexin V and 7-AAD staining in control and stable Pak1-expressing EC cells. (C) Control and stable Pak1-expressing EC cells were treated with 20 ng/ml TNF-α, and cells were harvested at the indicated number of hours. (D) Caspase-3 activity was expressed as percentage of enzyme activity compared to 0 h; **P<0.01, ***P<0.001. All the experiments were carried out in triplicate.
through the NF-xB-dependent pathway that regulates cyclin D1 transcription in breast cancer (17). Knockdown of Pak1 in colorectal cancer cells decreased proliferation and delayed the G1/S cell-cycle transition (26). In this study, we were able to confirm effect of Pak1 on proliferation from two aspects: Pak1 ectopic expression significantly enhanced proliferation whereas shRNA-mediated knockdown of Pak1 efficiently impeded proliferation in EC cells. Pak1 is a common point of convergence of growth factor signaling (27), and also directly interacts with estrogen receptor (ER) in ER-positive breast cancer (28). It has been proposed that Pak1 may constitute an important point of cross-talk between growth factors and the ER (29). Cross-talk between ER and growth factor signaling has emerged as a critical factor in endocrine resistance in EC (30). If this is the case, the levels and activation of Pak1 could potentially affect the action of anti-estrogen therapies. In experimental breast cancer model systems, overexpression of Pak1 correlates with the development of ligand-independent stimulation of ERα signaling, which is associated with sensitivity to tamoxifen (a selective anti-estrogen, which has anti-estrogenic effects in the breast but weak estrogenic effects in the endometrium) (28,31).

Therapies that target Pak1 may therefore represent a strategy to increase the hormonal response in EC.

We further investigated the effect of Pak1 on enhancing cell migration and invasion of EC cell lines. Tumor cell migration and invasion requires coordinated reorganization of the actin cytoskeleton (32). Pak1 is thought to regulate actin reorganization through several reported substrates (6). For example, Pak1 phosphorylates LIM-kinase at threonine 508 within kinase’s activation loop, and increases the phosphorylation and activation of cofilin, thus regulating actin depolymerization (9). Inhibition of Pak1 abolishes filamentous actin (F-actin) flow in the lamella, displaces myosin IIA from the cell edge, and decreases focal adhesion turnover (33). Pak1-knockdown cells have significantly impaired migration and invasion (16,34,35). In the present study, we also showed that downregulation of Pak1 expression reduced migration and invasion of EC cells.

Anchorage-independent growth is an important hallmark of oncogenic transformation (23). The expression of kinase-active Pak1 mutants significantly stimulates anchorage-independent growth of breast cancer cells in soft agar in a preferential mitogen-activated protein kinase (MAPK)-sensitive manner (36). Pak1 also phosphorylates dynein light chain 1 (DLC1) on serine 88, and promotes breast cancer cell anchorage-independent growth and tumor formation (37). In addition, Pak1 simultaneously activates MAPK and MET signaling, that permit human mammary epithelial cells to form anchorage-independent colonies (38). Herein, we showed that knockdown of Pak1 reduces, and overexpression enhances, the colony-forming ability of EC cells. These results demonstrate the ability of Pak1 expression to stimulate the growth of EC cells in an anchorage-independent manner, and indicate that Pak1 may play an essential role in oncogenic transformation.

The efficacy of cancer treatments depends not only on the cellular damage they cause but also on the cell’s ability to respond to these damages by inducing the apoptotic response (5). Pak1 has been shown to be involved in the regulation of several important pro-apoptotic pathways (6). For instance, Pak1 has been reported to directly phosphorylate and inactivate the pro-apoptotic functions of B-cell lymphoma 2 (BCL2) antagonist of cell death (BAD), which results in cell survival (39). In our study, Pak1 was found to protect EC cells from apoptosis induced by TNF-α via inhibition of caspase-3 activation. Alteration of this signal transduction pathway leading to apoptosis has been reported in drug-resistant cells (5). For instance, reduced caspase-3 activity is associated with cisplatin resistance in human ovarian cancer (40). In human bladder carcinoma, activation of AKT increases paclitaxel resistance by increasing Bad phosphorylation, leading to decreased release of mitochondrial cytochrome c and caspase-3-mediated apoptosis (41). In EC, caspase-3 expression is associated with patient survival (42). We found that expression of Pak1 leads to an inhibition of caspase-3 activation, which might be associated with cisplatin and/or paclitaxel resistance in EC. A recent study showed that dual inhibition of Pak1 and inhibitor of apoptosis proteins (IAPs) efficiently increased effector caspase activation and apoptosis of NSCLC cells (11). Thus, inhibition of Pak1 to enhance chemosensitivity of EC may provide a promising therapeutic option to putative targeted therapy. However, whether Pak1 is involved in protecting cancer cells from apoptosis induced by chemotherapy remains to be determined.

In summary, to the best of our knowledge, we showed for the first time that overexpression of Pak1 plays important roles at multiple stages of EC progression. Our results showed that Pak1 protein is overexpressed in endometrial carcinoma, particularly in the most advanced lesions. We also found that Pak1 affects EC cell proliferation, migration, invasion and anchorage-independent growth in vitro. In addition, Pak1 regulates cell apoptosis through inhibition of caspase-3 activation. Our study demonstrated that Pak1 is a multifunctional regulator of EC, and supports further characterization of Pak1 as a therapeutic target.

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