Pharmacological Inhibition and Genetic Knockdown of Exchange Protein Directly Activated by cAMP 1 Reduce Pancreatic Cancer Metastasis In Vivo

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ABSTRACT

cAMP plays a critical role in regulating migration of various cancers. This role is context dependent and is determined by which of the two main cAMP sensors is at play: cAMP-dependent protein kinase or exchange protein directly activated by cAMP (EPAC). Recently, we have shown that the cAMP sensor protein EPAC1 promotes invasion/migration of pancreatic ductal adenocarcinoma (PDA) in vitro. In this study, we investigated the role of EPAC1 in invasion and metastasis of PDA in vivo, and evaluated the therapeutic potential of EPAC inhibitors as antimetastasis agents for this neoplasm. We employed an orthotopic metastatic mouse model in which the PDA cells Mia PaCa-2 were injected into the pancreas of athymic nude mice, and their local and distant spread was monitored by in vivo imaging and histologic evaluation of the number of metastatic foci in the liver. Either genetic suppression of EPAC1 or its pharmacologic inhibition with 3-(5-tert-butyl-isoxazol-3-yl)-2-[(3-chloro-phenyl)-hydrazono]-3-oxo-propionitrile, an EPAC-specific antagonist recently identified in our laboratory, decreased invasion and metastasis of the PDA cells. Mechanistically, EPAC1 promotes activation and trafficking of integrin α1, which plays an essential role in PDA migration and metastasis. Our data show that EPAC1 facilitates metastasis of PDA cells and EPAC1 might be a potential novel therapeutic target for developing antimetastasis agents for PDA.

Introduction

Pancreatic ductal adenocarcinoma (PDA) is one of the leading causes of cancer-related deaths worldwide and has a poor prognosis with ~95% mortality rate (Kamangar et al., 2006; Raimondi et al., 2009). In addition to late diagnosis, the biggest factors behind such a dismal outlook are the aggressive nature and high metastatic potential of PDA (Keleg et al., 2003). The chemoradiation therapy currently approved for the treatment of this neoplasm has minimal efficacy, and, although surgical resection when possible can be curative, recurrence with distant metastases happens in the majority of patients (Cress et al., 2006; Winter et al., 2006). One of the biggest hurdles to developing effective therapeutic strategies is our incomplete understanding of the genetic and biochemical alterations that govern the progression of this cancer from well-differentiated intraepithelial neoplasia to highly metastatic and poorly differentiated PDA (Hezel et al., 2006).

The second messenger, cAMP, plays a critical role in regulating migration of various cell types, including cancer cells. However, this role is complex and depends on tissue type as well as the cAMP sensor transducing the signal (Burdyga et al., 2013). cAMP signals are mediated by two main protein families, cAMP-dependent protein kinase and exchange protein directly activated by cAMP (EPAC) (de Rooij et al., 1998; Kawasaki et al., 1998; Kawakami et al., 1998). The latter is composed of two isoforms, EPAC1 and 2, which primarily act as guanine nucleotide exchange factors.

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that activate the small GTPases, Rap1 and Rap2 (de Rooij et al., 1998; Kawasaki et al., 1998). These small GTPases in turn mediate the vast majority of EPAC responses, which span a wide array of biologic functions, including the regulation of cell adhesion and migration in different cellular contexts (Enserink et al., 2004; Carmona et al., 2008).

It has been reported that EPAC1 is overexpressed in human PDA tissue compared with the surrounding normal pancreatic tissue (Lorenz et al., 2008). A number of studies have shown that EPAC1 plays an important role in regulating cancer migration, but this role appears to be context dependent. In melanoma, there is consensus that EPAC1 enhances metastasis, as has been shown in several in vitro and in vivo studies (Baljinnyam et al., 2010, 2011, 2014). In prostate cancer, some studies suggest that EPAC1 promotes migration and proliferation (Bailey et al., 2009; Misra and Pizzo, 2009, 2012), whereas others suggest an inhibitory role for the EPAC-selective activator 8-(4-chlorophenylthio)-2'-O-methyladenosine-3', 5'-cyclic monophosphate (007) (Grandech et al., 2009). However, it has been argued that it is actually 007's indirect activation of cAMP-dependent protein kinase, rather than activation of EPAC1, that resulted in inhibition of migration and proliferation of prostate cancer (Menon et al., 2012). In ovarian cancer, EPAC1 seems to have promigratory effects in some cell lines (Ovcar3) (Rangarajan et al., 2003) and antimigratory effects in others (ES-2) (Bastian et al., 2009). To improve the development of antimetastasis therapeutic strategies, it is of paramount importance that we determine the role of EPAC1 in the context of each cancer and elucidate the mechanism by which it boosts or attenuates migration in each case.

We have recently shown that EPAC1 enhances migration/invasion of the two pancreatic cancer cell lines AsPC-1 and Panc-1 in vitro, without affecting their proliferation (Almahariq et al., 2013), a finding that was corroborated by a latter report (Burdyga et al., 2013). In this study, we investigated EPAC1's role in the invasion and metastasis of PDA, and the potential of EPAC inhibitors as antimetastatic agents using an orthotopic metastatic PDA mouse model. We report in this work that genetic suppression and pharmacologic inhibition of EPAC1 reduce PDA metastasis. In addition, our results suggest that EPAC1 mediates Migration/invasion of PDA through regulation of integrin β1 (Itgβ1) activation and trafficking. The findings of this study have significant clinical implications, as they identify EPAC1 as a potential therapeutic target for preventing pancreatic cancer metastasis.

Materials and Methods

Cell Lines. The pancreatic ductal adenocarcinoma cell lines AsPC-1, Panc-1, and MIA PaCa-2, obtained from American Type Culture Collection (Manassas, VA), were maintained in glutamine-containing RPMI 1640 medium (Thermo Scientific HyClone, Logan, UT) supplemented with 10% fetal bovine serum (Invitrogen, Grand Island, NY) at 37°C under 5% CO₂. Stable cell lines with suppressed EPAC1 expression (Epac1-KD) or cells expressing firefly luciferase were generated using MISSION TRC (Sigma-Aldrich, St. Louis, MO) lentiviral-short hairpin RNA (shRNA) or a lentiviral-based pGL4 vector 8-(4-chlorophenylthio)-2'-O-methyladenosine-3', 5'-cyclic monophosphate, acetoxymethyl ester (007-AM), is a selective EPAC agonist (Vliek et al., 2008; Chepurny et al., 2009) (Biolog Life Science Institute, Bremen, Germany). Bisindolylmaleimide I (BIM 1), 2,6-diamino-N-{[(1-oxotridecyl)-2-piperidinyl] methyl} hexamidine (NPC 15437), and 3-[1-[3-(dimethylamino)propyl]-5-methoxy-1H-indol-3-yl)-4-(1H-indol-3-yl)-1H-pyrrrole-2,5-dione (Gö 6983) are selective protein kinase C (PKC) antagonists (Touille et al., 1991; Sullivan et al., 1992; Gschwendt et al., 1996) (Santa Cruz, Dallas, TX). The 3-(4,5-dimethyl-thiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay (Promega) was used to measure cell viability.

Antibodies were used: fluorescein isothiocyanate (FITC)-conjugated integrin β1 clone 12G10 (Abcam, Cambridge, MA) and FITC-conjugated Itgβ1 clone K-20 (Santa Cruz).

Animals. Animal experiments were conducted in 6- to 8-week-old female athymic BALB/c nu/nu mice (Charles River Laboratories, Wilmington, MA). Use of all animals was in accordance with the guidelines of the University of Texas Medical Branch Institutional Animal Care and Use Committee.

Rap1 Activation Assay. Cells were treated with 10 μM 007-AM for 10 minutes with or without pretreatment with 5 μM ESI-09 for 5 minutes. The activation of Rap1 was examined using an active Rap1 (GTP-bound) pull-down assay, as described previously (Mei et al., 2002).

Cell Viability Assay. Cells were incubated for 24 hours at 37°C under 5% CO₂ in the presence of 007-AM (10 μM), ESI-09 (5 μM), BIM I (1 μM), Gö 6983 (1 μM), or NPC 15437 (1 μM), and viability was measured by an alamarBlue cell viability assay (Life Technologies, Carlsbad, CA).

Luciferase Activity. Luciferase-transduced MIA PaCa-2 cells were transfected with Ctrl- or Epac1-specific shRNA (Epac1-KD). Cells were then lysed, and luciferase activity was determined by a Luciferase Assay System assay (Promega). Bioluminescence was measured with a LmaxII 384 microplate reader (Molecular Devices, Sunnyvale, CA).

Transwell Migration/Invasion Assay. The top chambers of 8-micron inserts (Corning, Tewksbury, MA) were coated with BD Matrigel Basement Membrane Matrix (50 μg/ml) (BD Biosciences, San Jose, CA). Cells were starved of serum for 24 hours, detached with 0.25% trypsin-EDTA, and pretreated with the EPAC agonist 007-AM (10 μM) alone or in combination with the EPAC inhibitor ESI-09 (5 μM) or PKC inhibitors BIM I (1 μM), NPC 15437 (1 μM), or Gö 6983 (1 μM) for 10 minutes in serum-free RPMI 0.2% bovine serum albumin (BSA). The bottom chamber was filled with RPMI/4% fetal bovine serum (FBS) and the same combination of drugs used for pretreatment. The cells (2 × 10⁶) were then added to the top chamber and incubated at 37°C under 5% CO₂ for 20 hours. At the end of the incubation period, cells remaining inside the insert were removed, and the rest were fixed in methanol and stained with crystal violet. The numbers of migrated cells were counted in four different fields.

Wound-Healing Assay. Cells were grown to 95–100% confluency and starved of serum for 24 hours before a scratch wound was made. The cells were treated with 007-AM (10 μM) alone or with ESI-09 (5 μM) or BIM I (1 μM) in serum-free RPMI for 30 minutes before adding 10% FBS. The cells were then incubated at 37°C under 5% CO₂. The wound was imaged at 0 and 24 hours. Healing rate was determined by calculating the percentage of wound closure according to the following equation: % wound closure = initial wound width – final wound width / initial wound width × 100. To make the results comparable across all assays, the widths of the initial wounds were all normalized to a 1-mm distance.

Surface Protein Isolation. MIA PaCa-2 cells were seeded on a fibronectin matrix (Sigma-Aldrich), starved of serum for 24 hours, and treated with 007-AM (10 μM) alone or with ESI-09 (5 μM) or BIM I (1 μM) in serum-free RPMI for 45 minutes before surface proteins
were biotinylated with EZ-link sulfo-NHS-SS-biotin (Pierce, Rockford, IL) and isolated according to the manufacturer’s instructions. Briefly, 0.25 mg/ml EZ-link sulfo-NHS-biotin was added to the cells and incubated at 4°C for 30 minutes. Biotinylation was quenched, and the cells were harvested and centrifuged at 500g for 3 minutes. Cells were solubilized with the kit’s lysis buffer containing the protease inhibitor phenylmethylsulfonyl fluoride (Sigma-Aldrich) and incubated on ice for 30 minutes. The samples were centrifuged at 10,000g for 2 minutes at 4°C, and the supernatant containing biotinylated membrane proteins was incubated with NeutraAvdin gel slurry for 60 minutes at room temperature. Then surface proteins were eluted from the column with elution buffer containing 50 mM dithiothreitol. Approximately 15 μg biotinylated protein was separated by SDS-PAGE and transferred to a polyvinylidenedifluoride membrane, and surface Itg1 was probed by Western blotting. The plasma membrane protein Na+/K+ ATPase was used as a loading control. To obtain total Itg1, cells were treated as described for surface Itg1 isolation and lysed with SDS lysis buffer (2% SDS, 10% glycerol, 60 mM Tris, pH 6.8).

Flow Cytometry Analysis of Activation and Cell Surface Expression of Itg1. MIA PaCa-2 cells were starved for 24 hours and treated with 0.25% trypsin-EDTA for 5 minutes. Cells were then treated with 007-AM (10 μM) alone or in combination with ESI-09 (5 μM) or NPC 154371 (1 μM) in serum-free RPMI/10% BSA and incubated for 15 minutes at 37°C under 5% CO2. Cells were fixed in 4% paraformaldehyde for 12 minutes, and then active Itg1 was stained with 12G10 (1:100), which only recognizes the active conformation of Itg1 (Byron et al., 2009), and total Itg1 with K20 (1:5) for 30 minutes at 4°C in phosphate-buffered saline (PBS) containing 3% BSA. The samples were analyzed by FACS (FACSCalibur, BD Biosciences). The level of active Itg1 was determined by normalizing the mean fluorescence intensity of active Itg1 staining to total Itg1 mean fluorescence intensity.

**Immunofluorescence Staining of Cell Surface Itg1.** Wild-type or Epac1-KD AsPC-1 and PANC1 cells were seeded on fibronectin-coated microscope slide cover slips and starved of serum overnight. Then cells were treated with vehicle dimethylsulfoxide or the EPAC inhibitor ESI-09 (5 μM) for ~45 minutes and fixed with paraformaldehyde (~12 minutes), followed by staining for total Itg1 with the FITC-conjugate antibody K20 (1:4) in nonpermeabilizing buffer for 2 hours at room temperature. Then nuclei were stained with the dye 4’,6’-diamidino-2-phenylindole, and cells were mounted on a microscope slide and visualized with an Olympus BX51 immunofluorescence microscope. Fluorescence intensity for Itg1 and cell nuclei was determined in four random fields for each sample using the SimplePC16 imaging software. For comparison of different samples, fluorescence intensity of Itg1 was normalized to the fluorescence intensity of the nuclei from the same imaging field.

**Orthotopic Mouse Model and In Vivo Imaging.** MIA PaCa-2 cells stably expressing firefly luciferase and transfected with nontargeting (Ctrl) or Epac1-specific shRNA-C32 (Epac1-KD) were grown to 80–90% confluency and detached with 0.25% trypsin-EDTA, washed with RPMI/10% FBS, and suspended in ice-cold PBS/Matrigel (1:1). Mice were anesthetized with isoflurane and then injected into the pancreas of athymic nude mice (Supplemental Fig. 4). Local invasion and metastasis were monitored in vivo using bioluminescence imaging, and, at the end of the experiment, metastasis was further quantified by the number of metastatic foci in the liver, which is one of the most common sites of PDA metastasis (Pai et al., 2012).

**Results**

EPAC1 Facilitates Invasion and Metastasis of MIA PaCa-2 Cells. We have previously shown that EPAC1 is overexpressed in the PDA cells AsPC-1 and Panc-1 and facilitates their invasion/migration in vitro (Almahariq et al., 2013). To further determine whether EPAC1 plays an important role in PDA metastasis in vivo, we developed an orthotopic metastatic PDA mouse model using the PDA cells MIA PaCa-2. EPAC1 is highly expressed in MIA PaCa-2 cells, and its expression was successfully suppressed by shRNA (Supplemental Fig. 1A). In contrast, EPAC2 expression is undetectable (Supplemental Fig. 1B). To verify EPAC1’s activity in these cells, we examined the impact of its activation on the level of GTP-bound Rap1 (active form). Treatment with the EPAC-specific agonist 007-AM led to a significant increase in activation of the EPAC effector Rap1, and the EPAC inhibitor ESI-09 blunted its activation (Fig. 1A). Furthermore, similar to our findings in AsPC-1 and Panc-1 cells, activation of EPAC1 with 007-AM significantly increased invasion/migration of MIA PaCa-2 cells in wound-healing and Transwell invasion/migration assays, whereas pharmacologic inhibition with ESI-09 or shRNA silencing (clone 32) of EPAC1 expression completely abolished 007-AM’s stimulatory effect (Fig. 1B, 1C).

The pharmacologic treatment had no impact on cell viability in the time frame of the employed assays (Supplemental Fig. 3). These results confirm that EPAC1 plays an important role in facilitating PDA invasion and migration in vitro and MIA PaCa-2 cells are a viable candidate for testing EPAC1’s function in PDA metastasis.

Subsequently, we transduced luciferase into Ctrl or Epac1-KD MIA PaCa-2 cells. Cells with comparable luciferase activity were then injected into the pancreas of athymic nude mice (Supplemental Fig. 4). Local invasion and metastasis were monitored in vivo using bioluminescence imaging, and, at the end of the experiment, metastasis was further quantified by the number of metastatic foci in the liver, which is one of the most common sites of PDA metastasis (Pai et al., 2012).

Suppression of EPAC1 reduced local and distant spread of MIA PaCa-2 cells (Fig. 2A) and significantly decreased metastasis to the liver (Fig. 2B, 2C).

**EPAC1 Promotes Trafficking of Itg1.** The expression, distribution, trafficking, and function of integrins are frequently altered in tumor cells in a manner that promotes cancer migration (Caswell and Norman, 2006). Itgβ1 is particularly important for invasion of PDA and plays an essential role in facilitating its metastasis (Vogelmann et al., 1999; Grzesiak et al., 2011). Several reports have implicated EPAC1 and its effector Rap1 in Itgβ1-mediated adhesion and migration of endothelial progenitor and immune cells (Lorenowicz et al., 2006; Carmona et al., 2008). Hence, we hypothesized that EPAC1 facilitates invasion/migration of PDA through an Itgβ1-related pathway. Neither activation nor pharmacologic inhibition or genetic knockdown of EPAC1 altered expression levels of Itgβ1 in MIA PaCa-2 cells (Fig. 3A). However, when cells were treated with the EPAC activator 007-AM (45 minutes), followed by biotinylation and isolation of surface proteins, the plasma membrane fraction of Itgβ1 was significantly increased, and this observed rise was completely negated by ESI-09.
treatment or knockdown of EPAC1 (Fig. 3B). In fact, inhibition or suppression of EPAC1 reduced the membrane fraction of Itgβ1 below the basal level determined for vehicle-treated parental MIA PaCa-2 cells (Fig. 3B).

Additionally, after cells were trypsinized, recovery of cell surface Itgβ1, even after only 15 minutes, was significantly enhanced by EPAC activation and attenuated by its inhibition or suppression (Fig. 3C). We also followed membrane expression...
levels of Itgβ1 by immunofluorescence in the PDA cell lines AsPC-1 and PANC-1 and found that inhibition/suppression of EPAC1 decreased the cell surface levels of Itgβ1 (Supplemental Fig. 5). Together, these results suggest that EPAC1 facilitates trafficking of Itgβ1 to the plasma membrane during invasion/migration.

**EPAC1 Promotes Trafficking of Itgβ1 through PKC.** PKC regulates integrin trafficking and has been shown to promote migration of cancer cells (Wang et al., 2011; Al-Alem et al., 2013). Of note, during the integrin trafficking cycle, PKCε mediates the movement of Itgβ1 from the perinuclear recycling compartment (PNRC) to the plasma membrane (Caswell and Norman, 2006). Numerous studies have shown that EPAC activates PKC in various cell contexts, including PKCε (Hucho et al., 2005; Borland et al., 2009; Almahariq et al., 2014). Therefore, we reasoned that EPAC1 enhances trafficking of Itgβ1 to the plasma membrane in part through PKC activation. Similarly to the impact of inhibition and suppression of EPAC1, inhibition of PKC with the PKC-specific inhibitor BIM I had no impact on total Itgβ1 expression (Fig. 4A), but completely abrogated the increase in surface Itgβ1 seen with EPAC1 activation by 007-AM (Fig. 4B). Furthermore, inhibition of PKC negated the rise in invasion/migration observed with EPAC1 activation in Transwell invasion/migration and wound-healing assays (Fig. 4C, 4D).

To confirm the specificity of the observed response to BIM I treatment, we used two other PKC-specific inhibitors (NPC 15437 and Gö 6983). These inhibitors also blocked 007-AM's stimulatory effect on invasion/migration of MIA PaCa-2 and Itgβ1 trafficking (Supplemental Fig. 6). The used PKC inhibitors had no impact on the viability of MIA PaCa-2 cells during the time frame of the essay (Supplemental Fig. 3). Together, these findings suggest that EPAC1 promotes Itgβ1 trafficking through the PKC pathway. Noticeably though, PKC inhibition (Fig. 4C, 4D) didn’t reduce invasion/migration as effectively as did the inhibition of EPAC1 (Fig. 1B, 1C). This prompted us to search for potential additional mechanisms by which EPAC1 facilitates invasion/migration of PDA cells.

**EPAC1 Promotes Activation of Itgβ1.** Integrins are usually present in an inactive conformation that has a low affinity for their ligand. A series of signaling events that involve the recruitment of various adaptor proteins to the cytosolic domain of the integrin are required for activation (inside-outside signaling) (Banno and Ginsberg, 2008). Several reports have shown that Rap1 plays a role in the integrin activation cascade (Bos, 2005; Han et al., 2006). To determine whether EPAC1 facilitates activation of Itgβ1 in PDA, we probed the activation status of Itgβ1 after EPAC1 activation or inhibition/suppression using an antibody that only recognizes the active conformation of Itgβ1 (12G10) (Byron et al., 2009). To account for the change in total surface Itgβ1 in response to altering EPAC1 activity, we normalized the fluorescence intensity of the integrin’s active conformation to the intensity of total surface integrin.

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Fig. 3. EPAC1 increases trafficking of integrin β1 to the plasma membrane. (A and B) MIA PaCa-2 cells were treated with 007-AM in the presence or absence of ESI-09, and total or plasma membrane proteins were isolated, respectively. Itgβ1 levels were probed by Western blotting, quantified by densitometry, and presented as a percentage of the indicated loading control. (C) Cells were trypsinized, and recovery of surface integrin β1 was probed by FACS. Data are presented as mean fluorescence intensity (MFI) and normalized to vehicle-treated Ctrl MIA PaCa-2 cells. **Significantly higher than vehicle-treated Ctrl cells (P < 0.01). *Significantly lower than vehicle-treated Ctrl cells (P < 0.02). Bars represent mean ± S.D. (n = 3).
Itgβ1 staining. Activation of EPAC1 by 007-AM significantly increased the fraction of active Itgβ1, and inhibition/suppression of EPAC1 lowered it below basal levels (Fig. 5). Moreover, inhibition of PKC with NPC 15437 did not affect Itgβ1 activation (Fig. 5). BIM I could not be used in FACS-based assays due to its strong and broad-spectrum autofluorescence. These results suggest that, in addition to regulating integrin trafficking, EPAC1 mediates activation of Itgβ1, and this effect is most likely PKC independent.

**Pharmacological Inhibition of EPAC Reduces PDA Metastasis.** To determine whether inhibition of EPAC1 is a potentially viable therapeutic strategy for reducing metastasis of PDA, we employed the EPAC inhibitor recently discovered by our group, ESI-09 (Almahariq et al., 2013), in the orthotopic metastatic mouse model described earlier. We have previously shown that ESI-09 has excellent pharmacological activity in vivo and is capable of protecting mice from a lethal dose of rickettsial infection and recapitulating the EPAC1 knockout phenotype (Gong et al., 2013). Treatment with ESI-09 (10 mg/kg daily starting 2 days post-tumor injection) appeared to reduce local and distant spread of MIA PaCa-2 cells and significantly decreased metastasis to the liver (Fig. 6). These results suggest that EPAC1 is a potential target for developing antimetastasis agents for treatment of PDA.

**Discussion**

Our study shows that EPAC1 promotes invasion and metastasis in PDA. Genetic suppression of EPAC1 in the PDA cell line MIA PaCa-2 inhibited their invasion/migration in vitro and local spread and metastasis to the liver in an orthotopic metastatic mouse model. Given previous reports by our group and others (Almahariq et al., 2013; Burdyga et al., 2013), showing that EPAC1 increases invasion/migration of the PDA cell lines AsPC-1 and PANC-1, it appears that, within the context of PDA, EPAC1’s promigratory role is a general one.

Mechanistically, our results show that EPAC1 promotes invasion/migration of PDA by enhancing Itgβ1 trafficking, which plays a crucial role in controlling cell mobility. During cell migration, integrins are internalized from the cell surface into endosomes and accumulate in the PNRC before being shuttled to the leading edge of the migrating cell (Caswell and Norman, 2006; Shin et al., 2012). This process is dynamic and occurs rapidly, with each step being tightly regulated by various interactions of a complex network of proteins (Shin et al., 2012). The PKC isoform PKCα is particularly important for mediating movement of Itgβ1 vesicles from the PNRC pool to the plasma membrane (Caswell and Norman, 2006). We found that the PKC-specific inhibitors BIM I and NPC 15437 negated the
stimulatory impact of the EPAC agonist 007-AM on trafficking of Itgβ1 and invasion/migration of PDA. The inhibitor NPC 15437 preferentially targets novel isoenzymes of PKC, such as PKCε (Sullivan et al., 1992). Therefore, our results suggest that it is likely through the PKCε pathway that EPAC1 promotes Itgβ1 trafficking. This is in agreement with numerous reports showing EPAC activates PKC, including PKCε, through the phospholipase C pathway (Hucho et al., 2005; Borland et al., 2009; Almahariq et al., 2014). However, we cannot rule out the involvement of other PKC isoforms, and further studies are needed to determine the specific isoenzyme targeted by EPAC1 in the context of PDA.

Interestingly, although the impact of EPAC1 inhibition/suppression on Itgβ1 trafficking was mimicked by PKC inhibition, the latter did not suppress invasion/migration of MIA PaCa-2 to the same extent as the former did, prompting us to investigate additional mechanisms underlying EPAC1’s role in PDA migration. In addition to integrin trafficking, cells control their adhesion and migration by regulating the activation status of integrins (Banno and Ginsberg, 2008). Our results show that EPAC1 facilitates the activation of Itgβ1. This is in concordance with several reports showing the EPAC effector Rap1 activates integrins, including Itgβ1, through Rap1-GTP–interacting adaptor molecule, without affecting the overall surface expression levels of these proteins (Reedquist et al., 2000; Lafuente et al., 2004). In contrast, inhibition of PKC had no impact on the activation status of Itgβ1, suggesting EPAC1’s role in this mechanism is PKC independent.

To determine the druggability of EPAC1, we used ESI-09, an EPAC-specific inhibitor recently discovered in our laboratory (Almahariq et al., 2013). This compound was able to reduce invasion and metastasis of MIA PaCa-2, suggesting that EPAC1 is a potentially viable target for antimetastasis agents. However, the in vivo role of EPAC1 in metastasis of additional PDA cell lines and/or tumors extracted from patients must be examined to better assess and confirm the therapeutic potential of this protein in PDA treatment.

The findings of our study might carry significant clinical implications, as there is a dire need for mechanism-based therapeutic strategies for pancreatic cancer, especially ones that target the Itgβ1 activation and/or trafficking pathways. Numerous studies have shown that Itgβ1 mediates the malignant phenotype of PDA and facilitates loss of epithelial integrity and oncogenic transformation in epithelial tumors (Grzesiak and Bouvet, 2006; Lee et al., 2013; Onodera et al., 2014). In fact, constitutive activation of Itgβ1 is correlated with higher grade...
canceromas (Lee et al., 2013). There are currently available some small molecules that target ITGβ1, but monobal antibodies and synthetic peptides against this integrin have shown significant clinical efficacy (Barkan and Chambers, 2011). Hence, EPAC inhibitors might provide a new approach to target ITGβ1 in cancer treatment.

Authorship Contributions
Participated in research design: Almahariq, Chao, Mei, Hellmich, Cheng.
Conducted experiments: Almahariq, Chao, Mei.
Contributed new reagents or analytic tools: Tomatedi.
Performed data analysis: Almahariq, Patrikeiev, Cheng.
Wrote or contributed to the writing of the manuscript: Almahariq, Cheng.

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Pharmacological Inhibition and Genetic Knockdown of EPAC1 Reduce Pancreatic Cancer Metastasis in vivo

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Supplementary Fig. S1. EPAC1 and EPAC2 expression in MIA PaCa-2. (A) Expression and knockdown by shRNA (clone 32) of EPAC1 in MIA PaCa-2 cells were confirmed by Western blotting. (B) EPAC2 expression in MIA PaCa-2 was probed. Mouse brain tissue was used as a positive control for EPAC2 expression.
Supplementary Fig. S2. EPAC1 knockdown decreases invasion and migration of MIA PaCa-2. (A) Cells were transfected with non-targeting shRNA (Ctrl) or an Epac1 specific shRNA (clone 28) and EPAC1 expression was probed by Western blotting. Knockdown of EPAC1 reduced invasion/migration of MIA PaCa-2 cells in invasion/migration and wound-healing assays. (B) Quantification of the number of migrating cells. (C) Quantification of wound closure presented as the distance traveled by the edge of the wound relative to the wound’s initial size. * Significantly lower than Ctrl cells (P < 0.02). Bars represent mean ± S.D. (n=3).
Supplementary Fig. S3. Effects of utilized compounds on viability of MIA PaCa-2. Cells were treated with vehicle DMSO, 10 µM 007-AM, 5 µM ESI-09, 1 µM BIM I, 1 µM NPC 15437, or 1 µM Gö 6983 for 24 hrs, and cell viability was determined by an almarBlue viability assay. RFU, relative fluorescence unit normalized to vehicle treated cells. Bars represent mean ± S.D. (n=3).
Supplementary Fig. S4. Luciferase activity in MIA PaCa-2 and MIA PaCa-2-Epac1-KD cells. MIA PaCa-2 cells were transduced with luciferase, followed by transfection with non-targeting shRNA (Ctrl) or an Epac1 specific shRNA-clone 32 (Epac1-KD). An equal number of cells ($2 \times 10^4$) was then used to examine luciferase activity in each cell line using an in vitro luciferase assay system. Bars represent mean ± S.D. (n=3)
Supplementary Fig. S5. Inhibition or suppression of EPAC1 decreases the level of surface integrin β1. Control or Epac1-KD AsPC-1 and PANC1 cells were treated with vehicle DMSO or the EPAC inhibitor ESI-09 (5 µM) for ~ 45 min and fixed with paraformaldehyde, followed by staining for total surface Itgβ1 with the FITC-conjugate antibody K20 (1:4) in non-permeabilizing buffer for 2 hr. Nuclei were stained with the dye DAPI and cells were mounted on a microscope slide. (A) Representative image showing surface Itgβ1 (green) and cell nuclei (blue) staining in AsPC-1. (B) Fluorescence intensity of Itgβ1 was normalized to that of the nuclei. * Significantly lower than vehicle-treated WT cells (P < 0.01). Bars represent mean ± S.D. (n=5).
Supplementary Fig. S6. PKC inhibition negates 007-AM’s impact on MIA PaCa-2’s migration and trafficking of integrin β1. (A) Cells were treated with 007-AM (10 µM) in the presence or absence of NPC 15437 (1µM) and invasion/migration was examined by a trans-well migration assay. (B) Cells were trypsinized, treated with 007-AM in the presence or absence of NPC 15437 in a non-permeabilizing buffer, and recovery of surface Itgβ1 was probed by FACS using the Itgβ1 antibody K20 by FACS. Mean fluorescence intensity (MFI) was normalized to vehicle-treated cells. (C) Cells were treated with 007-AM (10 µM) in the presence or absence of Gö 6983 (1µM) and invasion/migration was examined by a trans-well migration assay. ** Significantly higher than vehicle-treated WT cells (P < 0.02). * Significantly higher or lower than vehicle-treated WT cells (P < 0.05). # Significantly lower than 007-AM-treated cells (P < 0.05). Bars represent mean ± S.D. (n=3).