Periostin activates integrin α5β1 through a PI3K/AKT-dependent pathway in invasion of cholangiocarcinoma

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Abstract. Periostin (PN) is mainly produced from stromal fibroblasts in cholangiocarcinoma (CCA) and shows strong impact in cancer promotion. This work aimed to investigate the mechanism that PN uses to drive CCA invasion. It was found that ITG α 5 β 1 and α 6 β 4 showed high expression in non-tumorigenic biliary epithelial cells and in almost all CCA cell lines. PN had preferential binding to CCA cells via ITG α 5 β 1 and blocking this receptor by either neutralizing antibody or si*ITG* α 5 β 1 binding, intracellular pAKT was upregulated whereas there was no change in pERK. Moreover, PN could not activate AKT in condition of treatment with a PI3K inhibitor. These data provide evidence that PN-activated invasion of CCA cells is through the ITG α 5 β 1/PI3K/AKT pathway. Strategies aimed to inhibit this pathway may, thus, provide therapeutic benefits.

Introduction

Cholangiocarcinoma (CCA), caused by liver fluke infection, is a major public health problem in the Northeastern and Northern parts of Thailand. It is a slow growing cancer with rapid metastasis and a high mortality rate (1). The incidence of CCA has increased worldwide with the highest rate in Thailand where around 93-100 per 100,000 peoples have been diagnosed with this disease (2). The pathogenesis of CCA has been demonstrated mainly by genetic mutations of bile duct epithelial cells themselves as the result of chronic inflammation which can also create a local environment enriched with cytokines and growth factors (3). In addition to cancer cells, the role of cancer-

associated fibroblasts has been revealed in cancer pathogenesis by production of several mitogenic and pro-invasive factors into the tumor microenvironment (4). Several current reports indicate the crucial involvement of stromal fibroblasts in the progression of CCA via certain types of secreted tumorigenic substances (5-7). These substances can prime cancer cells in a paracrine mode to develop tumorigenic intracellular signaling pathways resulting in the increased cell proliferation, growth and invasion.

Almost all of the CCA stromal fibroblasts have been revealed to be α -smooth muscle actin (α -SMA)-positive cells (5,8). The fibroblasts were transformed into activated fibroblasts or myofibroblasts and were incorporated into the tumor and produced extracellular matrix proteins that led to tumor fibrosis (8). The recent work from the authors of the present group suggests that levels of α-SMA positive CCA stromal fibroblasts were correlated with poor patient survival (5). CCA fibroblasts isolated from CCA tissues have exhibited the ability to promote tumor cell proliferation and invasion (5,6). The whole gene expression analysis of CCA fibroblasts compared to normal fibroblasts has reported several differential expressed genes encoded to produce tumorigenic secreted proteins including periostin (PN) (6). The findings reported here are in support of several groups of researchers, wherein PN has been confirmed to express in CCA tissues solely from stromal fibroblasts (6,9,10). It has been proposed as a potential marker of poor prognosis in CCA patients (6). Though in vitro studies revealed the function of PN in induction of CCA cell proliferation, growth, and invasion, little is known regarding the mechanism of PN-induced CCA cell invasion that is a crucial phenomenon to activate CCA metastasis.

PN is an extracellular matrix (ECM) protein with multifunctional roles in tumorigenesis and tumor progression at each step of the transformation of normal into malignant cells and metastatic tumors (11). It has been proposed as a marker associated with cancer aggressiveness in pancreatic cancer (12,13), gastric cancer (14), breast cancer (15), thyroid carcinoma (16) and non-small cell lung cancer (17). In CCA, the impact of fibroblast-derived PN is convincing by its ability to activate cancer cell proliferation and invasion (6). To activate biological

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functions of cells, PN has been investigated for its ability to bind to integrin (ITG) receptors. In epithelial ovarian carcinoma, PN bound with ITGs avß3 and avß5 promoted cancer cell motility (18). The interference of these two ITGs by specific anti-ITG antibodies had an effect on the ability of PN to mediate cell adhesion in head and neck squamous cell carcinoma (19). For intracellular signaling, PN potently promotes metastatic growth of colon cancer by augmenting cell survival via the AKT/PKB pathway in colon cancer (20). Similarly, PN from pancreatic cancer cells activated ITGB4 and promoted invasiveness of cancer cells through the PI3K pathway (21). But in vascular smooth muscle cells, PN was demonstrated to induce cell migration through ITGs $\alpha v\beta 3$ and $\alpha v\beta 5$ and the focal adhesion kinase pathway (22). In breast cancer, PN enhanced angiogenesis, in part, from the up-regulation of the vascular endothelial growth factor receptor on endothelial cells through the ITGav_{β3}-focal adhesion kinase (FAK)-mediated signaling pathway (23). In addition, PN could induce epithelial-mesenchymal transition characteristics resulting in tumor metastasis through cross-talk between ITG $\alpha\nu\beta5$ and the epidermal growth factor receptor signaling pathways (24). It seems reasonable then to conclude that PN, in the dependent context, can activate specific ITG-mediated signal pathways and different biological responses.

Even though, it was found that the invasive property of CCA cells stimulated by PN was reduced after transient knockdown $ITG\alpha 5$ (6), little is known regarding the intracellular signaling pathway activated by PN through ITGa5-mediated cell invasion. In the present study, the ITGs expression in CCA cell lines were explored and showed abundance of ITGs $\alpha 5\beta 1$ and $\alpha 6\beta 4$. The adhesion assay revealed the propensity of CCA cells to bind PN via ITG α 5 β 1. Using cells with a low level of functional ITG α 5 β 1 by treating these cells with either si*ITG* α 5 β 1 or neutralizing anti-ITG α 5 antibody, the results showed that the PI3K/AKT-mediated, but not the ERK-mediated signaling pathway was involved in PN-stimulated CCA invasion. Since, PN is abundant in CCA tissues, understanding the role of PN in induction of cancer cell invasion may provide important information to attenuate metastasis and help identify the possible therapeutic targets.

Materials and methods

CCA cell culture. Human CCA cell lines KKU-M055, KKU-100, KKU-M139, KKU-M156, KKU-M213, KKU-M214 and KKU-OCA17 were kindly donated from Associate Professor Banchob Sripa, Khon Kaen University, Thailand. KKU-M213, KKU-M214 and KKU-OCA17 originated from well differentiated CCA tissues; KKU-M055, KKU-M139 and KKU-M156 from moderate CCA; and KKU-100 was isolated from poorly differentiated tissue (25). The non-tumorigenic immortalized bile duct epithelial cell MMNK1 was kindly provided by Professor Naoya Kobayashi, Department of Surgery, Okayama University Graduate School of Medicine and Dentistry, Japan (26). CCA cells and MMNK1 were cultured in Ham F-12 medium (Invitrogen, Carlsbad, CA, USA) supplemented with 10% fetal bovine serum (FBS), 100 U/ml penicillin, 100 μ g/ml streptomycin (Invitrogen) and anti-fungal agent. Cells were cultured in a humidified 5% CO₂ incubator at 37°C. Cells were passaged by 0.25% trypsinEDTA and those of more than 90% viability were used in further experiments.

Measurement of ITG expression pattern in CCA cell lines by real-time PCR. Total-RNA was extracted from all CCA cell lines and MMNK1 using PerfectPure RNA Cultured Cell Kit (5 Prime, Gaithersburg, MD, USA) according to the manufacturer's instructions. The cDNA was synthesized using SuperScript[™] III First-Strand Synthesis System (Invitrogen) according to the instructions. Expression levels of *ITGs* av, a5, $\alpha 6, \beta 1, \beta 3, \beta 4$ and $\beta 5$ were determined by SYBR-Green-based real-time PCR in Light Cycler® 480 II machine (Roche Applied Sciences, Indianapolis, IN, USA). The β -actin served as an internal control to adjust the amount of starting cDNA. The expression of each *ITG* was calculated by the $2^{-\Delta C_p}$ equation. In this case, $\Delta C_p = C_p(ITG) - C_p(\beta - actin)$. The sequences of genes used in this study were retrieved from PubMed (www.ncbi.nln. nih.gov) and primers were designed using Primer 3 software. A list of primers is summarized (Table I).

Flow cytometry analysis of ITG expression. For detection of the actual ITG α 5 β 1 and α 6 β 4 levels in biliary epithelial cells, cell pellets of around 1x106 were fixed in 2% formaldehyde for 15 min at room temperature. The fixed cells were incubated with 1:50 goat anti-human ITGa5 (Santa Cruz Biotechnology, Santa Cruz, CA, USA) diluted in a washing solution which was HAM/F-12 containing 2% (v/v) FBS, 1% (w/v) bovine serum albumin (BSA) and 10 mM NaN₃ for 2 h at room temperature. Cells in the washing solution were centrifuged at 400 x g for 3 times and 5 min each to remove the excess primary antibody and this was followed by staining with 1:2,000 donkey anti-goat IgG-Alexa 488 (Invitrogen) diluted in the washing solution for 1 h at room temperature with light protection. For ITG α 6 β 4 detection, 1:100 mouse anti-human ITG β 4 (Millipore, Temecula, CA, USA) diluted in washing solution was incubated with cells for 1 h at 4°C and followed by 1:100 rabbit anti-mouse IgG-FITC (Dako, Carpinteria, CA, USA) diluted in washing solution for 30 min at 4°C with light protection.

The ITG α 5 and ITG β 4 signals were determined in the FL-1 channel of a Becton Dickinson FACSort (Becton Dickinson, Franklin Lakes, NJ, USA) and data analysis was performed by CellQuest software (Becton Dickinson). The relative mean fluorescence intensity (MFI) of CCA cell lines was normalized to that of the negative control stained with secondary antibody only. Two independent experiments were performed.

Immunocytochemistry of ITG α 5 β 1 and α 6 β 4 in CCA cell lines. Immunocytochemistry was employed to localize ITG α 5 β 1 and ITG α 6 β 4 on the cell membrane. KKU-M213 (2x10⁴ cells) were cultured on sterile cover slips placed in a 24-well plate for 48 h. Cells were fixed in 4% paraformaldehyde for 15 min and blocked with 1% BSA for 30 min at room temperature. Then the cells were incubated with 1:50 goat anti-human ITG α 5 (Santa Cruz Biotechnology) for 2 h at room temperature and subsequently stained with 1:500 donkey anti-goat IgG-Alexa 488 (Invitrogen) for 1 h at room temperature with light protection.

For localization of the ITG β 4, cells were plated onto glass cover slips at a density of $4x10^4$ cells in 24-well plate. After 48 h, cells were washed twice with 1X PBS and then blocked with blocking solution (10% FBS containing 1X PBS) for

Gene	Forward primer 5'-3'	Reverse primer 5'-3'	Size (bp)	Accession no.
ITGav	TGACTGGTCTTCTACCCGC	CTCACAGATGCTCCAAACCA	121	NM_002210
ITGa5	AGTTGCATTTCCGAGTCTGG	CTCTGGGAGCACCAGATACAA	223	NM_002205
ITGa6	GGCCTTATGAAGTTGGTGGA	CTCTGGGAGCACCAGATACAA	144	NM_000210
ITG <i>β1</i>	TCCCTGAAAGTCCCAAGTGT	TTTCCTGCAGTAAGCATCCA	143	NM_033666
ITGβ3	TGGTCCTGCTCTCAGTGATG	TGAAGGTAGACGTGGCCTCT	180	NM_000212
$ITG\beta 4$	TCTCCTACCGCACACAGGA	CTTCACCTGCAGCTCTTTCC	110	NM_001005619
$ITG\beta 5$	CTCCACTCTGGGAAACCTGA	AGGACGGTCAGGTTGGACTT	188	NM_002213
β -actin	CACACTGTGCCCATCTACGA	CTCCTTAATGTCACGCACGA	162	X00351

Table I. The primers used in this study.

30 min at room temperature. Cells were further incubated with 1:500 mouse anti-human ITG β 4 (Millipore) for 2 h at room temperature and then incubated in 1:2,000 goat anti-mouse IgG-Cy3 (Jackson ImmunoResearch Laboratories Inc., West Grove, PA, USA).

The nuclei were stained with 1:1,000 Hoechst 33258 (Invitrogen) for 30 min at room temperature. The fluorescence signal was observed under the LSM 510 Meta laser scanning confocal microscope (Carl Zeiss, Jena, Germany) at the Division of Medical Molecular Biology, Office for Research and Development, Faculty of Medicine, Siriraj Hospital, Mahidol University, Bangkok, Thailand.

Neutralization of $ITG\alpha 5\beta 1$ and $ITG\beta 4$ on CCA cells. To ensure roles of ITGa5B1 and ITGB4 on PN-mediated CCA cell adhesion and invasion, neutralizing antibody specific to the ITGα5β1 heterodimer and ITGβ4 subunits was employed to block intact ITG α 5 β 1 and ITG β 4 on the cell membrane of CCA cells. KKU-M213 CCA cells were trypsinized and washed with 1X PBS two times. Cell pellets of around 1x10⁵ cells were incubated with 1:200 anti-human ITGa5_{β1} (Millipore) or 1:200 mouse anti-human ITG_{β4} (Millipore) at 37°C for 1 h. Cells in the antibody solution were centrifuged at 400 x g for 5 min to remove excess antibody. The ITG-blocked cells were then collected to explore their responses to PN-induced invasion and adhesion. The number of either adhered or invaded cells induced by PN were compared with and without antibody blocking conditions. Two independent experiments were performed.

Adhesion assay of CCA cell lines on PN-coated surface. Recombinant PN (rPN) (1 μ g) (Biovendor, Heidelberg, Germany) was coated on a 96-well plate surface at 37°C for 2 h. Cells with or without exposure to neutralizing antibodies against ITG α 5 β 1 or ITG β 4 (Millipore) were then added to each well and incubated at 37°C for 1 h. Unattached cells were removed by rinsing twice with serum-free media. The number of adherent cells was determined by an MTS assay kit (Promega, Madison, WI, USA) according to the manufacturer's instructions. The percentage of PN-induced cell adhesion was normalized to that of cells attached on 1% BSA-coated wells. Two independent experiments were performed. Invasion assay. To investigate the effect of rPN on the invasion of parental KKU-M213 cells, si*ITGa5*-treated cells, and ITG α 5 β 1-blocked cells; 2x10⁴ cells of each condition were suspended in 100 ng/ml rPN containing complete medium and cultured in the upper chamber of the MatrigelTM invasion chamber (BD Biosciences, San Jose, CA, USA) for 24 h. Invaded cells were fixed in 5% (v/v) glutaraldehyde and then hematoxylin and eosin staining was performed. The number of invaded cells was counted under an inverted microscope by two independent investigators using x100 magnification fields. The assays were done in replicates of three independent experiments. Numbers of invaded cells was compared to those without rPN treatment.

Western blot analysis of pAKT and pERK. Cells with or without ITG α 5 silencing and cells in the presence or absence of 100 μ M LY294002, a PI3K inhibitor (Calbiochem, San Diego, CA, USA) or 30 µM U0126, an ERK inhibitor (Tocris Bioscience, MO) were induced by 100 ng/ml rPN for 30 and 120 min. Then the levels of pAKT and pERK1/2 were determined using western blot analysis. Cell pellets were collected after centrifugation of cell suspensions at 400 x g for 5 min in a refrigerated centrifuge. The cell pellets were rinsed by cold 1X PBS 2 times before lysed in 1X sample buffer containing 50 mM Tris-HCl pH 6.8, 2% (w/v) SDS, 10% (v/v) glycerol, 5% (v/v) β -mercaptoethanol and 0.05% (w/v) bromophenol blue. Cell lysate was boiled for 10 min and centrifuged to remove the undissolved proteins and cell debris at 8,000 x g for 1 min. Cell extracts were then separated in 10% SDS-PAGE and transferred onto PVDF membranes (Amersham, Buckinghamshire, UK). Membranes were blocked in 5% skim milk containing TBST for 1 h at room temperature. Rabbit anti-human pAKT (Thr308) (Santa Cruz Biotechnology) at the dilution of 1:1,000 and rabbit antihuman pERK1/2 (Cell Signaling Technology Inc., Danvers, MA, USA) at the dilution of 1:2,000 were used as primary antibodies for incubation with the membrane for 1 h at room temperature. The 1:2,000 goat anti-rabbit IgG-HRP (Abcam, Cambridge, MA, USA) was used as the secondary antibody and incubated for 1 h at room temperature. The immunoreactive signals were visualized by enhanced chemiluminescense (Pierce, Rockford, IL, USA). The β -actin protein level was used as an internal control to determine the equal amount of loading proteins.

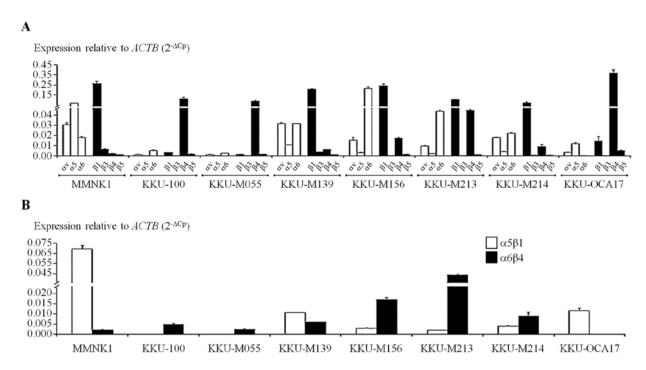


Figure 1. (A) Expression profile of ITG α - and β -subunits in immortalized bile duct epithelial cell MMNK1, KKU-K100, KKU-M055, KKU-M139, KKU-M156, KKU-M213, KKU-M214 and KKU-OCA17 CCA cell lines. (B) The possible expressions of different ITG $\alpha\beta$ heterodimers are predicted. Bars represent mean \pm SD of triplicate experiments.

Statistical analysis. The values from different independent experiments were expressed as mean \pm SD. The significance of the different data sets was determined by the Student's t-test. A P-value of ≤ 0.05 was defined as statistically significant.

Results

Integrin expression profile in CCA cell lines. ITGs that were previously reported in CCA, and in the related cancer, hepatocellular carcinoma, and those that have been revealed as PN receptors including α -subunits: αv , $\alpha 5$ and $\alpha 6$; and β -subunits: $\beta 1$, $\beta 3$, $\beta 4$ and $\beta 5$ were explored. Real-time PCR using 50 ng starting cDNA exhibited different levels of each subunit in CCA cell lines and MMNK1 cells. Among α -subunits, $\alpha 6$ was expressed at the highest level in almost of cell types except KKU-OCA17 (Fig. 1A). In addition, $\alpha 5$ -subunit showed the highest level among other α -subunits in MMNK1, but a moderate expression level was found in KKU-M139, KKU-M213 and KKU-M214. For the β -subunit, the results revealed that ITG $\beta 1$ had the highest expression level in KKU-M139, KKU-M156, KKU-M213, KKU-M214 and MMNK1, whereas KKU-K100, KKU-M055 and KKU-OCA17 had highest level of ITG $\beta 4$.

Since certain type of α -subunit ITG can be paired with a specific type of β -subunit, the predicted level of intact ITGs were presented based on the minimal level of their counterpart of either α -subunit or β -subunit. It has been shown that ITG α 5 can bind only to β 1 while ITG β 4 can only bind with α 6 (27). Moreover, ITG α v can bind to several β -subunits including β 1, β 3, β 5, β 6 and β 8. So the level of ITG β 3 may determine the maximal level of ITG α v β 3. Using the same concept, the amount of ITG β 5 would roughly present the possible maximal level of ITG α v β 5. Whereas, for ITG α 5 β 1 and ITG α 6 β 4, the expression of α 5-subunit and β 4-subunit may determine the maximal levels.

Hence, the possible maximal levels of ITGs $\alpha\nu\beta3$, $\alpha\nu\beta5$, $\alpha5\beta1$ and $\alpha6\beta4$ are shown (Fig. 1B). The expressions of both $\alpha5\beta1$ and $\alpha6\beta4$ ITGs were found in all cell types. Some cell types had ITG $\alpha5\beta1$ as the predominant expression level including MMNK1 and KKU-M139, but KKU-M156, KKU-M213 and KKU-M214 CCA cells showed predominant ITG $\alpha6\beta4$ expression. Cells with high levels of both ITG $\alpha5\beta1$ and ITG $\alpha6\beta4$ were KKU-M139, KKU-M156, KKU-M213 and KKU-M214. Interestingly, KKU-M055 showed very low levels of ITG $\alpha5\beta1$ as well as KKU-100.

Expressions of membrane ITGs $\alpha 5\beta 1$ and $\alpha 6\beta 4$ on CCA cell *lines*. To confirm the actual protein expression levels of ITG α 5 β 1 and $\alpha 6\beta 4$, FACS analysis was performed. The results revealed different amounts of these two ITGs among different types of CCA cell lines. For ITG α 5 β 1, the relative mean fluorescence intensity (MFI) showed similar levels of expression in all cell lines with the highest signal in KKU-M213 (Fig. 2A). Most of CCA cells originated from well differentiated cancers including KKU-M213 and KKU-M214 expressed high levels of ITGα6β4 similar to KKU-M139 which was derived from moderate differentiation of squamous cell types (Fig. 2B). Almost all cell lines derived from moderately differentiated types (KKU-M055 and KKU-M156) and the poorly differentiated type (KKU-100) had low expression levels of ITG α 6 β 4. In contrast, the expression of ITGa6β4 was found higher in KKU-M139, KKU-M213 and KKU-M214 than other cells, which is concordant to the results of mRNA levels. Interestingly, the highest level of ITG α 6 β 4 (relative MFI=12.4 \pm 1.6) and ITG α 5 β 1 (relative MFI=2.0 \pm 0.57) were detected in KKU-M213 (Fig. 2B).

PN mediates cell adhesion through ITGs α *5* β *1 and* α *6* β *4 receptors.* To demonstrate the impact of ITG α *5* β *1 and ITG* α *6* β *4 in*

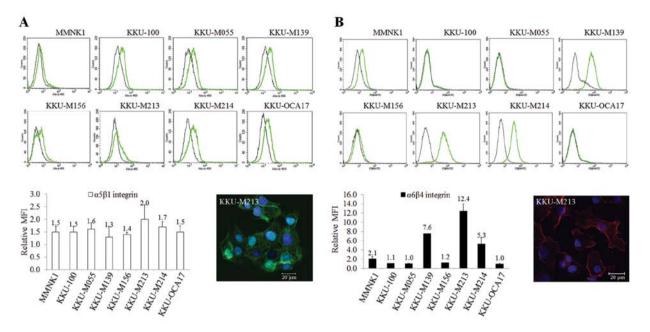


Figure 2. Flow cytometry analysis of the intact ITGs (A) α 5 β 1 and (B) α 6 β 4 CCA cell lines. Mean fluorescence intensity (MFI) of ITG α 5 β 1 shows in relative signals compared to unstained cells without adding the first antibodies presented as grey in the graph. Bar graphs represent mean ± SD of relative MFI from three independent experiments. Immunofluorescence staining indicates the membranous pattern of both ITGs on the membrane of KKU-M213 cells.

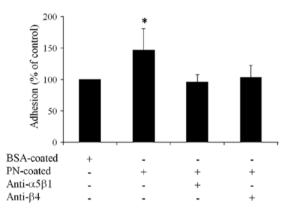


Figure 3. Adhesion assays of KKU-M213 cells in different culture coating surfaces with and without treatment with neutralizing anti-ITGs antibodies. Cells adhered onto BSA-coated plate were used as controls. Bars represent mean \pm SD of two independent experiments. An asterix represents statistical significance (P<0.05) compared to the control BSA coating surface.

PN-activated tumorigenic function of CCA cells, KKU-M213 cells having the highest level of both ITGs were used to perform the adhesion assay. PN-coated culture plates were utilized to explore the binding efficiency of the cancer cells with and without functional ITG receptors on the cell membrane after incubation with the neutralizing antibodies. The results showed that CCA cells could intrinsically bind to PN-coated surface more than to BSA-coated surfaces or negative controls with statistical significance (Fig. 3). The binding efficiency was reduced to a similar level of the negative control when cells were blocked either with the intact ITG α 5 β 1 or ITG α 6 β 4 by specific neutralizing antibodies.

PN-mediated CCA invasion via ITG\alpha5\beta1. To confirm whether cells with unavailable ITG α 5 β 1 could be induced by PN,

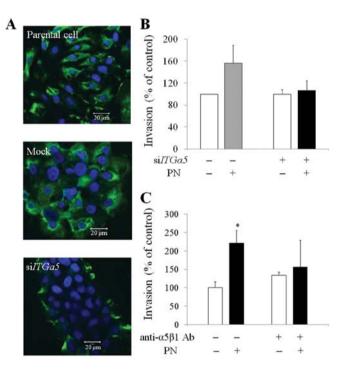


Figure 4. Invasion assay of KKU-M213 cells with and without ITG α 5 β 1. (A) Cells were treated with si*ITG* α 5 and the reduction of ITG α 5 β 1 is shown by fluorescence microscopy. (B) PN-induced invasion of si*ITG* α 5-treated cells in comparison to that in mock cells are shown. (C) Percents of invaded KKU-M213 cells when ITG α 5 β 1 is blocked with anti- α 5 β 1 antibody and untreated cells are shown in conditions with and without PN stimulation. Invasion of cells without si*ITG* α 5 or antibody treatments was used as the control. Bars represent mean \pm SD of duplicate or triplicate experiments. *P<0.05 compared to the control. The graph shows percentage of invaded cells exposed to PN compared to that of no PN treatment which was assumed to be 100%.

si $ITG\alpha 5$ -treated and anti-ITG $\alpha 5\beta 1$ antibody-treated cells were performed invasion assay with and without PN treatment. The

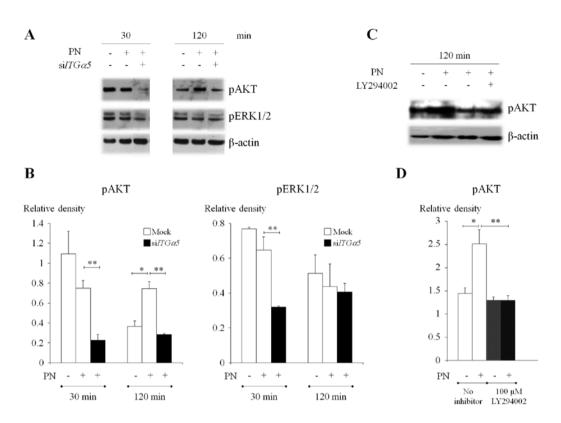


Figure 5. PN induces pAKT through PI3K pathway. (A) Western blot analysis shows level of pAKT and pERK1/2 in si*ITGa5*-treated cells and mock cells after being treated with 100 ng/ml of rPN for 30 and 120 min. The β -actin level indicates equal amounts of total protein. (B) Densitometry measurement of pAKT and pERK1/2 after normalization with the corresponding intensity of β -actin. (C) The KKU-M213 cell line was treated with rPN in the presence or absence of 100 μ M LY294002, a PI3K inhibitor, for 120 min. (D) Densitometry measurement of pAKT in cells treated with PI3K inhibitor after normalization with the corresponding intensity of β -actin. The * and ** represent P<0.05 of either pAKT or pERK in mock cells treated and untreated with rPN; and mock and si*ITGa5*-treated cells after stimulation with rPN, respectively.

intact ITGa5ß1 receptor on the membrane of CCA cells was detected by immunocytochemistry and showed that cells exposed to siITG α 5 successfully inhibited the expression of ITG α 5 β 1 on the membrane of cancer cells as compared to the intrinsic expression in parental and mock cells (Fig. 4A). The decreased PN-induced invasive capability of the $ITG\alpha 5$ -knockdown cells was revealed (Fig. 4B). Significant increases of PN-induced invasion was observed in mock cells (156±18%) as compared to cells without PN stimulation. The results revealed that ITGa5knockdown cells showed decreased PN-induced invasive capability (106±18% of invaded cells) when compared to negative controls without PN. Moreover, in cells blocking ITGa561 with neutralizing anti-ITG α 5 β 1 antibody, the results showed in a similar way that PN could not induce KKU-M213 cell invasion if there was no ITG α 5 β 1 available on the cell membrane (Fig. 4C).

PN induces pAKT through activation of PI3K pathway in CCA cells. In order to investigate the intracellular signaling pathway activated by PN via ITG α 5 β 1, cells with normal levels and ITG α 5 β 1-knockdown cells were treated with recombinant PN. The results revealed that exogenous PN could significantly induce phosphorylation of AKT in KKU-M213 parental cells at 120-min post-treatment compared to control cells without PN treatment (Fig. 5A and B). PN could not activate pAKT in cells with transient knockdown of ITG α 5, pERK1/2, however, did not change as a result of PN stimulation when compared between cells with and without ITG α 5 β 1 (Fig. 5A and B). These results

suggested that upon PN stimulation via $ITG\alpha 5\beta 1$, AKT, but not ERK, was activated.

To confirm the signaling pathway of PN-induced AKT, the pharmacological inhibitor of PI3K (LY294002) which is the upstream molecule of AKT, was applied to KKU-M213 CCA cells and the level of AKT phosphorylation was determined under the conditions with and without PN stimulation. The results showed depletion of pAKT level in responding to LY294002 treatment which confirmed the antagonist effect of this inhibitor to pAKT (Fig. 5C and D). Interestingly, PN could not induce pAKT in the PI3K-inhibited cells. Hence, in cells activated with PN, the pAKT was significantly higher in parental cells than in cells pretreated with PI3K inhibitor.

Discussion

The progression of cancer is no longer recognized as an independent aberrant event occurring only in cancer cells. Substances surrounding the cancer cells secreted from a variety of cells in the tumor microenvironment and signaling pathways induced by cancer and other cells, and cancer cell-substance interactions are thought to play crucial roles. The role of tumor-associated fibroblasts as the most abundant cell types in the tumor microenvironment and the major source of growth factors and extracellular matrix substances in tumorigenesis and tumor progression have been reported as important contributors (4,28). In the current situation drug resistance of cancer treatment is a common phenomenon

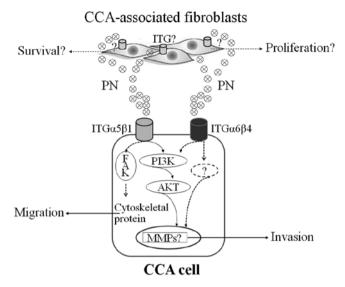


Figure 6. Proposed mechanism of PN-induced CCA migration and invasion. CCA-associated fibroblasts are the major source for PN production in CCA microenvironment. The fibroblast-derived PN acts as a ligand of ITG α 5 β 1 and α 6 β 4 receptors. Upon binding of PN and ITG α 5 β 1, the PI3K/AKT signaling pathway is activated and invasiveness property is driven in CCA cells. It is possible that PN may also activate the PI3K/AKT pathway resulting in cell proliferation and survival of CCA fibroblasts in an autocrine manner. Base on these effects, CCA fibroblasts may be used as a therapeutic target for CCA patients. The dashed lines were not investigated in this study.

in cancer patients due to the vulnerability of cancer cells to undergo genetic changes as the result of their rapid proliferation rate. Cancer fibroblasts are therefore suitable to target since they exhibit less genetic instabilities (29). Based on this concept, several studies have been performed and the obtained information has been summarized in several review articles to confirm the possibility of targeting cancer fibroblast as an additional treatment strategy in cancer patients (4,30-32).

Different cancer fibroblasts have their unique properties in the production of certain substances (6,33,34). Fibroblasts isolated from CCA tissues can produce a variety groups of tumorigenic substances which play important roles in induction of cancer cell proliferation (5). Unpublished data from the present research team has revealed migration induction when CCA cells were treated with the conditioned-medium from primary cultured-CCA fibroblasts. The gene expression profile of CCA fibroblasts has been performed by this group and the increased expression of tumor-related genes in CCA-derived fibroblasts has been reported (6). Among these genes, PN has been confirmed with high expression in the microenvironment of CCA tissues with relation to the short survival time of the patients together with tumorigenic induction in cancer cells in vitro including cell proliferation, growth, and invasion (6). Herein, the underlying mechanism of how CCA cells respond to PN-driven invasion has been explored. ITGs, as the receptors for PN (18,21,22), were explored in CCA cells. Adhesion assays indicated that ITG α 5 β 1 and ITG α 6 β 4 were the receptors for PN and the invasion assay confirmed that $ITG\alpha 5\beta 1$ was involved in PN-induced CCA cell invasion and PI3K/AKT was the signaling pathway underlying this mechanism.

Previous reports on ITG expression in CCA cells indicated that $\alpha 1$ had no expression whereas ITGs $\alpha 2$, $\alpha 3$, $\alpha 6$, $\beta 1$ and

 β 4 were expressed in almost all CCA cell lines (35-37). This is consistent with the present findings that all CCA cell lines expressed high level of ITG α 6, though the synthesis of ITGs $\alpha 2$ and $\alpha 3$ were not included in the current study. Notably, the expression of ITG α 5 is not uniformly expressed in CCA (36). The result showed the expression of ITG α 5 in some CCA cells; in particular the cells with previously reported high responses to PN-induced invasion such as KKU-M213 CCA cell lines (6). Though the expression of ITGav has mostly been reported as PN receptor, no evidence was found in CCA cells. The current results revealed that ITGav could be detected in almost all CCA cell lines but at a lower level than that of ITGs $\alpha 5$ and $\alpha 6$. These results imply that in addition to ITGs $\alpha 2$, $\alpha 3$, and α 6 previously reported, CCA cells could express α v and α 5. It is difficult to predict the level of ITG $\alpha\nu\beta$ 3 and $\alpha\nu\beta$ 5 because ITG α v can bind to several β -subunits of ITG including β 1, β 3, β 5, β 6 and β 8 (27). ITG α 5, however, can form heterodimers only with ITG β 1 (27,38). It can then be concluded that ITG α 5 β 1 may be one of the existing ITGs on the CCA cell membrane and may have an important impact in cancer progression after the stimulation by stromal PN.

For β -subunit ITGs, the present results are similar to previous reports that ITGs \beta1 and \beta4 were expressed in almost all CCA cells (35,36). ITG64 was found at higher levels in CCA cells as compared to those in MMNK1 immortalized non-tumorigenic biliary cells. Though in the previous report (35), ITGβ4 was detected in normal and proliferating biliary epithelial cells but was an inconsistent finding in CCA. This study provides conclusive evidence of the presence of ITG β 4 in CCA cells. Since ITG β 4 can bind only to ITG α 6, it is likely that the level of ITG β 4 can be roughly determined by the level of ITG α 6 β 4 presented on CCA cell membranes. In addition, this study indicated β 3 and β 5, generally form heterodimers with ITGav, as the ITGs with low expression in CCA cells. These results therefore suggested that CCA cells express high levels of ITGs $\alpha 5$, $\alpha 6$, $\beta 1$ and $\beta 4$. It is suggested that the heterodimers of ITG α 5 β 1 and α 6 β 4 may be the major ITG receptors expressed on CCA cell membrane. The flow cytometry analysis and immunofluorescence staining confirmed the presence of these two ITGs on the membrane of KKU-M213 cells. The favorable binding of PN on ITGs $\alpha 5\beta 1$ and $\alpha 6\beta 4$ indicated by the lower numbers of cells bound onto the PN-coated surface after treatment of cells with the specific neutralizing antibody. The ITGa6β4 was able to interact with PN as well as ITG α 5 β 1 at similar levels. Hence, it can be concluded that PN may influence the progressive tumor behavior of CCA cells though either ITG α 5 β 1 or ITG α 6 β 4.

PN has been demonstrated to activate cellular responses via different ITGs depending on cell type context, for example, ITG $\alpha\nu\beta\beta$ in non-small cell lung cancer (39) and ITG $\alpha\delta\beta4$ in pancreatic cancer (21). Previous work by this group reports that si*ITG* α 5-treated cells have lower PN-induced cell proliferation and invasion compared to the parental cells (6) and shows in the first report presenting the association of PN and ITG $\alpha5\beta1$ in tumor promotion of CCA cells. It is possible that the interaction of PN and ITG $\alpha5\beta1$ may be the unique phenomenon of PN in promotion of CCA. Hence, in this current report, we explored the signaling pathway starting from ITG $\alpha5\beta1$ in the induction of cancer invasion by PN. Similar to the data on the PN-activated signaling pathway reported previously (21,39), PN could activate cell invasion via the stimulation of AKT-dependent, but not ERK, pathway in CCA cells. It is well known that activation of the PI3K-AKT-dependent pathway is essential in the regulation of several different biological functions including cell survival, growth and proliferation, invasion and migration in a ligand specific manner (40-42). ITGa5_{β1} plays an important role in metastasis, invasion and poor prognosis of some cancers (43). Cancer cells with high expression of ITGa5B1 showed an increased invasiveness into 3D collagen matrices through enhancement of the contractile force (44). In addition, the role of ITG α 5 β 1 in stimulation of cell invasion has been revealed through activation of matrix metalloproteinase 2 (MMP2) in breast cancer (45). The fact that PN-ITG α 5 β 1 interaction stimulates the enhancement of cell contractile force and some MMP expressions is the possible underlying mechanisms of how PN helps tumor cells to invade and finally metastasize. Unpublished data by the present authors showed that PN induced MMP9 and MMP13 from CCA cells and activated cancer cells to migrate (data not shown) which may be the effect of PN-mediated change of cytoskeletal proteins through FAK (22). Finally, the actual downstream signaling pathway after AKT activation is of particular interest because the proper inhibitor can be proposed to apply for the attenuation of cancer progression with minimal side-effects (46).

In summary, fibroblasts are not passive bystanders in tumor environment. The current trend in cancer research is the inclusion of the cancer fibroblast as a major contributor of disease progression and suggests the inhibition of fibroblast-derived tumor-promoting factors as the first line approach. This study provides evidence for the contribution of CCA-associated fibroblast-derived PN in the activation of tumorigenic properties of CCA cells through receptor ITGs (Fig. 6). PN secreted mainly from fibroblasts binds to either ITGa5β1 or ITGa6β4 and can facilitate the invasiveness property of CCA cells. The ITG5α5β1-mediated PI3K-dependent or FAK (22) signaling pathways are activated eventually regulating several cellular responses in particular MMP production and cell migration. The obtained knowledge implies the potential of using an anti-PN antibody (47), anti-ITGa5_{β1} antibody (48), and PI3K/AKT inhibitors (49,50) to attenuate CCA progression driven by fibroblasts. Understanding the exact mechanisms responsible for fibroblast-associated cancer progression is a challenge for the future as the alternative and synergistic cancer-targeted therapy in CCA patients.

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