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Interleukin-8 increases VEGF and neuropilin expression and stimulates ERK activation in human pancreatic cancer

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Abstract

Interleukin-8 (IL-8) is associated with tumorigenesis by promoting angiogenesis and metastasis. Although upregulation of IL-8 is indicated in many cancers, its function in pancreatic cancer has not been well characterized. In this study we examined the expression of IL-8 on pancreatic cancer cells and clinical tissue specimens, and investigated the effect of exogenous IL-8 on gene expression, and signaling in human pancreatic cancer cells. We found that pancreatic cancer cells expressed higher amount of IL-8 mRNA than normal human pancreatic ductal epithelium (HPDE) cells. IL-8 mRNA was also substantially overexpressed in 11 of 14 (79%) clinical pancreatic-adenocarcinoma samples compared with that in their surrounding normal tissues. Exogenous IL-8 upregulated the expression of VEGF₁₆₅, and NRP-2 in BxPC-3 cells, one of human pancreatic cancer cell lines. IL-8 expression was inducible by hypoxia mimicking reagent cobalt chloride. In addition, IL-8 activated ERK1/2 signaling pathway in BxPC-3 cells. Our studies suggest that IL-8 might be a malignant factor in human pancreatic cancer by induction of VEGF and NRP-2 expression and ERK activation. Targeting IL-8 along with other anti-angiogenesis therapy could be an effective treatment for this malignancy.

Keywords

IL-8; VEGF; cytokine; ERK; pancreatic cancer

Introduction

Interleukin-8 is an inflammatory cytokine upregulated in both cancer and chronic inflammatory diseases of the pancreas 1. It is linked to pancreatic cancer tumorigenesis primarily through its regulation of angiogenesis and metastasis 2. Human umbilical vein endothelial cells (HUVECs) proliferation and angiogenesis were both increased when co-cultured with pancreatic cancer cells, or cultured with exogenous interleukin-8 (IL-8). The increase of cell proliferation and angiogenesis of HUVEC can be blocked by IL-8 neutralizing antibodies 3. IL-8 and vascular endothelial growth factor (VEGF) expressions were found to be upregulated in a highly metastatic pancreatic cancer cell line Colo357L3.6pl compared with the parental cell Colo357FG, likely mediated through a mitogen activated protein kinase (MAPK) pathway 4. IL-8 transfection also made nonmetastatic human melanoma cells metastatic 5.

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Previous studies have shown that IL-8 is overexpressed in most human pancreatic cancer tissues and established cell lines². Higher IL-8 levels in pancreatic cancer patient serum were associated with significant weight loss, but not with performance status⁶. Additionally, the expression levels of IL-8 appear to correlate with their tumorigenic and metastatic potential in an orthotopic xenograft model^{2, 7–10}. Furthermore, treatment with exogenous IL-8 increased the invasiveness of human pancreatic cancer cell line Panc-1 cells in a matrigel assay, while blocking IL-8 inhibited the growth of another human pancreatic cancer cell line Capan-1 cells^{11, 12}. Blocking IL-8 in pancreatic cancer cell Hup-T4 cells decreased their growth and their ability to attach to endothelial cells, suggesting that IL-8 is an autocrine mitogenic factor important for metastasis¹³.

The expression of IL-8 can be induced by many stimuli including lipopolysaccharide, phorbol 12-myristate 13-acetate (PMA), IL-1, and tumor necrosis factor (TNF). Several stress factors, such as hypoxia, acidosis, nitric oxide (NO), and cell density, also significantly influenced the production of IL-8 in human pancreatic cancer cells⁷. IL-8 expression is shown to be regulated by nitric oxide (NO) secreted from endothelial cells. Cells modified to produce physiologic levels of NO showed increased IL-8 production compared to low NO producing cells, and the nitric oxide synthase (NOS) inhibitor NMA blocked this effect¹⁴. IL-8 has also been shown to be involved in cancer hypoxia pathway, in which IL-8 expression is regulated by hypoxia inducible factor-1 (HIF-1), NF- κ B, and KRAS^{10, 15}.

In this study we investigated the expression of IL-8 in a number of pancreatic cancer cell lines, culture medium, and surgical tissue samples. We also studied the effects of IL-8 in regulating other key molecules in pancreatic cancer growth and angiogenesis, and the signaling pathway induced by IL-8 in pancreatic cancer.

Materials and Methods

Chemicals and reagents

Human IL-8 and cobalt chloride (CoCl₂) were purchased from Sigma (St. Louis, MO). EGM-2 Bullet kit, trypsin-EDTA and trypsin neutralization solution were purchased from Clonetics (Walkersville, MD). The Ambion “RNAqueous-4PCR” kit and DNA removing kit were obtained from Ambion (Austin, Texas). The iQ SYBR Green supermix, Bioplex phosphoprotein and cytokine kits were purchased from Bio-Rad (Hercules, CA).

Cell culture and human tissue specimens

Human pancreatic cancer cell lines (Panc-1, MIA PaCa-2, BxPC-3, Capan-2, HPAF-II, and Panc03.27) were obtained from the American Type Culture Collection (ATCC, Rockville, MD). The human pancreatic ductal epithelium (HPDE) cells were provided as a generous gift from Dr. Ming-Sound Tsao from the University of Toronto, Canada^{16, 17}. Panc-1 cells were cultured in DMEM with 10% fetal bovine serum (FBS) at 37°C with 5% CO₂. MIA PaCa-2 cells were cultured in DMEM with 10% FBS and 2.5% horse serum at 37°C with 5% CO₂. BxPC-3 and Panc03.27 cells were cultured in RPMI 1640 medium with 10% FBS at 37°C with 5% CO₂. HPAF-II cells were cultured in Eagle’s MEM with 10% FBS at 37°C with 5% CO₂. Capan-2 cells were cultured in McCoy’s medium with 10% FBS at 37°C with 5% CO₂. HPDE cells were cultured in keratinocyte serum-free (KSF) medium supplied with 5 ng/mL EGF and 50 μ g/mL bovine pituitary extract (Invitrogen, Carlsbad, CA). Human pancreatic adenocarcinoma specimens were collected from patients who underwent surgery according to an approved human protocol (H-16215) at the Baylor College of Medicine (Houston, TX).

Real time RT PCR

Total RNA was extracted from six pancreatic cancer cell lines (Panc-1, MIA PaCa-2, BxPC-3, Capan-2, HPAF-II, and Panc03.27 cells), HPDE cells, and clinical specimens using an Ambion “RNAqueous-4PCR” kit following the manufacture’s instruction (Austin, Texas) as described previously¹⁸. Specific primers for IL-8 were designed with the Beacon Designer 5.1 software (PREMIER Biosoft International, Palo Alto, CA). The primer sequences for human IL-8 gene are: Sense 5’GGCACAACTTTTCAGAGACAGCAG3’; and anti-sense 5’GTTTCTTCCTGGCTCTTGTCCTAG3’.

The mRNA for IL-8 was analyzed by real time RT PCR using iCycler system (Bio-Rad, Hercules, CA). PCR reaction included the following components: 100 nM each primer, diluted cDNA templates and iQ SYBR Green supermix, and running for 40 cycles at 95°C for 20 sec and 60°C for 1 min. Each cDNA sample was run as triplicates and the corresponding no-reverse transcriptase (RT) mRNA sample was included as a negative control. The β -actin primer was included in every plate to avoid sample variations. The mRNA level of each sample for each gene was normalized to that of the β -actin mRNA. The relative mRNA level was presented as unit values of $2^{[Ct(\beta\text{-actin}) - Ct(\text{gene of interest})]}$ ¹⁸. For IL-8 and hypoxia treatment, subconfluent pancreatic cancer cells were treated with IL-8 (100 ng/mL) or different concentrations of CoCl_2 for 24 h. Total RNA was extracted and the mRNA levels for VEGF₁₆₅, VEGF receptors neuropilin-1 (NRP-1), neuropilin-2 (NRP-2), and IL-8 were analyzed by real time RT PCR as above. The primer sequences for human VEGF₁₆₅ gene are: Sense 5’CCAGCAGAAAGAGGAAAGAGGTAG3’; and anti-sense 5’CCCCAAAAGCAGGTCACTCAC3’. The primer sequences for human NRP-1 gene are: Sense 5’ AAGGTTTCTCAGCAAACCTACAGTG 3’; and anti-sense 5’ GGGGAAGAAGCTGTGATCTGGTC 3’. The primer sequences for human (NRP-2 gene are : Sense 5’ GATTCTGGGATGGGGACAGTGA 3’; and anti-sense 5’ GGTGAACCTTGATGTAGAGCATGGA 3’¹⁸.

Bio-Plex phosphoprotein assay

Pancreatic cancer cells at $1.5 \times 10^5/\text{mL}$ were treated with IL-8 (100 ng/mL) for 0, 5, 15, 30, or 60 min. Protein lysates were prepared using cell lysis kit (Bio-Rad) on samples collected at each indicated time point. The presence of p-ERK1/2 was detected by Bio-Plex phosphoprotein assay kit (Bio-Rad) and the Phosphoprotein Testing Reagent kit (Bio-Rad) according to the manufacturer’s protocol as described previously¹⁹. Briefly, 50 μL of cell lysate (100–450 μg protein/mL) was plated in the 96-well filter plate coated with anti-p-ERK1/2 antibody and incubated overnight on a platform shaker at 300 rpm at room temperature. After incubation with detection antibodies, data from the reaction was then acquired and analyzed using the Bio-Plex suspension array system (Luminex 100 system) from Bio-Rad Laboratories. The total proteins for ERK were tested using the Bio-plex total protein assay kit (Bio-Rad).

Bioplex Cytokine Assay

Panc-1, MIA PaCa-2, BxPC-3, and HPDE cells were treated with IL-8 (100 ng/mL) for 24 h, and the supernatant was collected. Cytokine concentrations were determined using the Bioplex multiplex Human Cytokine Assay kit (Bio-Rad) and the Cytokine Reagent kit (Bio-Rad) according to the manufacturer’s protocol and was described previously^{19, 20}. Briefly, 50 μL of culture supernatants or cytokines standards were plated in a 96 well filter plate coated with a multiplex of antibodies against the above mentioned cytokines and incubated overnight on a platform shaker at 300 rpm at room temperature. Data from the reaction was then acquired and analyzed using the Bio-Plex suspension array system (Luminex 100 system) from Bio-Rad Laboratories as described above.

Statistical analysis

Data from real-time PCR, and bioplex assay were expressed as mean $\pm$ SEM. Significant differences were determined by paired Student's *t*-test (two tails).

Results

IL-8 was overexpressed in human pancreatic cancer cell lines and tissues

We examined IL-8 mRNA expression in six human pancreatic cancer cell lines and HPDE cells. IL-8 was differentially increased in five cell lines (MIA PaCa-2, BxPC-3, Capan-2, HPAF-II, and Panc03.27) compared with that in HPDE cells (Fig. 1A). Bio-plex cytokine assay was used to determine the amount of IL-8 protein in the culture media from Panc-1, MIA PaCa-2, and BxPC-3 cell lines. The culture media from the HPDE cell line was included as a control. As shown in Fig. 1B, the IL-8 protein levels in MIA PaCa-2 and BxPC-3 cells were increased by 900%, and 304%, respectively, as compared to that in HPDE cells, which is consistent with the real time PCR results.

We also examined the IL-8 expression in 14 pairs of human pancreatic cancer tissues with the surrounding normal tissues. IL-8 mRNA was substantially overexpressed in 11 of 14 (79%) clinical pancreatic-adenocarcinoma samples compared with that in their surrounding normal tissues (Fig. 1C). Overall fold increase of IL-8 mRNA expression in pancreatic cancer tissue was 2.6 times that in the surrounding normal tissues. Therefore, human pancreatic cancer is associated with increased expression of IL-8.

IL-8 upregulated the expression of key molecules for angiogenesis and cell proliferation

To investigate the effect of IL-8 on the expression of key molecules controlling angiogenesis and cell proliferation in pancreatic cancer, we examined the mRNA levels of VEGF₁₆₅, NRP-1, and NRP-2 in BxPC-3 cells upon treatment with 100 ng/mL of IL-8. VEGF₁₆₅ is a critical molecule for promoting angiogenesis, and in pancreatic cancer, the neuropilins (NRP-1 and NRP-2) may act as alternate receptors for VEGF₁₆₅¹⁸. As shown in Fig. 2, VEGF₁₆₅ and NRP-2 mRNAs were significantly increased by IL-8 treatment in BxPC-3 cells. Therefore, IL-8 may be involved in upregulation of VEGF and NRPs in pancreatic cancer.

IL-8 was induced under hypoxia conditions in pancreatic cancer

To determine if IL-8 is involved in response to hypoxia in pancreatic cancer, and if there is correlation between IL-8 and VEGF under hypoxia conditions, we treated MIA PaCa-2 cells with different doses of cobalt chloride (CoCl₂), the hypoxia mimicking reagent, and found that IL-8 mRNA level was increased by hypoxia in a dose-dependent manner. At 200 μ M of CoCl₂, IL-8 mRNA was increased by 203% (Fig. 3). Similarly, VEGF₁₆₅ mRNA level was also increased by CoCl₂ in a dose-dependent manner as described before²¹, which indicates that IL-8 is inducible by hypoxia, and the levels of IL-8 and VEGF might correlate under hypoxia conditions in pancreatic cancer cells.

IL-8 activates ERK signaling pathway

To examine the signal molecules involved in the IL-8 pathway in pancreatic cancer, we investigated the phosphorylation of ERK in pancreatic cancer cells treated by IL-8. BxPC-3 cells were stimulated with 100 ng/mL of IL-8 for 0, 5, 15, 30, or 60 minutes. The percentage of phosphorylated ERK versus the total ERK proteins was then measured using a Bioplex phosphoprotein assay. As shown in Fig. 4, the addition of exogenous IL-8 dramatically increased the phosphorylation of ERK in BxPC-3 cells, which peaked at 15 minutes before dropping off towards baseline levels at 60 minute.

Discussion

These data demonstrated that IL-8 is overexpressed in human pancreatic cancer cell lines and clinical specimens. IL-8 protein was also elevated in the cultured medium of human pancreatic cancer cells compared to HPDE cells. Exogenous IL-8 caused the upregulation of VEGF₁₆₅, and NRP-2 in pancreatic cancer cells. IL-8 mRNA was found to be increased under hypoxia condition. Furthermore, we have found that IL-8 activates the MAPK pathway, with significant increased phosphorylation of ERK observed in pancreatic cancer cells treated by IL-8. Our findings reinforce the hypothesis that IL-8 might serve as a prognostic factor for pancreatic cancer, and is an important target for potential therapeutic intervention.

Previous studies have shown that IL-8 was overexpressed in pancreatic cancer, caused increased MMP-2 activity and played an important role in the invasiveness of human pancreatic cancer [11, 12, 22]. The immunohistochemical analysis of 40 surgically resected human pancreatic cancer demonstrated that positive staining for IL-8 was 50%, and positive staining for both IL-8 and IL-8 receptors was 40% [11]. In contrast, immunoreactivity for IL-8 was significantly suppressed in normal pancreatic tissues. Our data is consistent with the literature, showing that IL-8 protein levels in MIA PaCa-2 and BxPC-3 cells were increased by 900%, and 304%, respectively, as compared with that in HPDE cells. IL-8 mRNA was also found to be substantially overexpressed in 11 of 14 (79%) clinical pancreatic-adenocarcinoma samples compared with that in their surrounding normal tissues with an overall 2.6-fold increase. Therefore, human pancreatic cancer is associated with increased expression of IL-8. IL-8 expression has also been found in various human cancers, including brain tumors, breast cancer, gastric cancer, lung cancer, and prostate cancer [2, 23–27], indicating a critical role of IL-8 in cancer pathogenesis.

We also showed that IL-8 increased the expression of VEGF₁₆₅ and NRP-2. As indicated in our previous study, VEGF₁₆₅ is an important angiogenic factor, while NRPs serves as receptors for VEGF in pancreatic cancer cells [18]. It has been shown that IL-8 promotes the metastasis of pancreatic cancer through increased angiogenesis, particularly involving VEGF [3, 4, 10, 13, 28, 29]. Our data further implicate IL-8 as a pro-angiogenic cytokine that helps the spread of distant metastasis by inducing neovascularization, and possibly promoting the survival of the tumor mass in general by maintaining a rich capillary network to accommodate the heavy nutrient requirements of this aggressive cancer. Other studies have reported a direct effect on tumor growth, in which decreased IL-8 level by IL-8 antisense oligonucleotide caused decrease in the growth of a pancreatic cancer cell FG cells in mice after intrapancreatic implantation, which correlated with decreased tumor angiogenesis [7]. IL-8 is also a powerful inflammatory cytokine associated with chronic diseases of the pancreas [1]. This study, along with others, strongly indicates that IL-8 plays a critical role in angiogenesis and pancreatic cancer progression by influencing several aspects of creating an environment conducive to the development and spread of pancreatic adenocarcinoma. Blocking IL-8 and IL-8 receptor CXCR2 significantly inhibited angiogenesis [3, 28].

Both IL-8 and VEGF are important components in cellular response to hypoxia, a common event in cancer, including human melanoma, colon cancer and pancreatic cancer [2]. Mizukami et al found that besides the preserved expression of VEGF, IL-8 was induced by hypoxia in a colon cancer cell line DLD-1^{HIF-kd}, in which the hypoxia inducible factor-1 (HIF-1) was knocked out, but not in HIF-1 wild type DLD-1^{HIF-wt} cells. This induction was mediated by activation of NF-κB and KRAS. A neutralizing antibody to IL-8 significantly inhibited angiogenesis and tumor growth in DLD-1^{HIF-kd} but not DLD-1^{HIF-wt} xenografts, indicating the functional significance of IL-8 in the anti-angiogenesis and hypoxia therapy of solid tumor. Strategies that inhibit hypoxia pathway which is mostly mediated by HIF-1 may be more effective when IL-8 is simultaneously targeted [15]. In our study, we found that IL-8 was

upregulated in a dose-dependent manner in response to hypoxia mimicking reagent CoCl_2 in pancreatic cancer cells. Our previous study also showed that both VEGF_{165} and another pro-inflammatory cytokine IL-6 were inducible by hypoxia²¹. The correlation of IL-8 and VEGF under hypoxia conditions warrants further investigation. The interaction between IL-8, VEGF, HIF-1, and maybe other cytokines such as IL-6 are critical in response to hypoxia in solid tumors, and may become targets for combinational cancer therapy.

It has been suggested that the mitogen activated protein kinase (MAPK) pathway is involved in IL-8 and VEGF activation⁴. We have also examined possible cellular mechanisms of IL-8 in pancreatic cancer cells. We found that ERK was activated after treatment with exogenous IL-8 in BxPC-3 cells. The MAPK pathway plays important role for cell growth and survival, and its activation has been shown to be associated with tumorigenesis^{20, 30, 31}. Previous studies have shown that constitutive activation of several signaling pathways in tumor cells is associated with increased IL-8 expression^{8, 10}. Several molecules have been reported to stimulate IL-8 expression in tumor cell lines through MAPK cascades such as IL-1 α , which upregulates the secretion of IL-8 in human pancreatic cancer cells via activation of ERK-1/2, p38 MAPK, AP-1, and NF- κ B^{3, 32}. It has also been shown that activation of Src and its downstream pathways ERK-1/2 and p38 MAPKs played important roles in upregulating IL-8 secretion in pancreatic cancer cells³³. Inflammatory cytokines and chemokines including IL-8 can also activate ERK pathway possibly through protein kinase C (PKC). IL-8 binds to two different receptors on cell surface, CXCR1 and CXCR2. We found that in six pancreatic cancer cells we examined (Panc-1, MIA PaCa-2, BxPC-3, Capan-2, HPAF-II, and Panc03.27), CXCR1 was highly expressed in five of them (except for in MIA PaCa-2 cells) compared with HPDE cells, but CXCR2 expression was low in all six pancreatic cancer cell lines. A few other studies have also shown various expressions of IL-8 receptors in different pancreatic cancer cell lines such as Capan-1 and SUI-2 cells^{12, 34}. Upon IL-8 binding, CXCR1 and CXCR2 are phosphorylated, and have been shown to mediate MAPK activation³⁵. This autocrine loop, in which IL-8 exerts a positive feedback for ERK/MAPK activation might be important for pancreatic cancer progression. Therefore, our data suggest that IL-8 act as a direct growth and survival factor on pancreatic cancer cells. This complements well our picture of IL-8 as a multifaceted regulator of gene expression that can regulate multiple pathways including angiogenesis, metastasis, and response to hypoxia in pancreatic cancer. Further investigation is needed to elute more clearly the cellular mechanisms employed by IL-8.

In conclusion, we have found that IL-8, a multifunctional inflammatory cytokine, is over-expressed in both pancreatic cancer cell lines and in malignant tissue samples from pancreatic cancer patients. We have also demonstrated IL-8 was inducible under hypoxia condition and was able to upregulate key molecules known to promote angiogenesis and metastasis possibly through MAPK pathway. Our study indicates that this cytokine remains an important focus in pancreatic cancer and suggests the need for further research to investigate the role and mechanism of IL-8, and for therapeutic intervention with potent combinational treatment which includes both anti-angiogenesis and suppressing pro-oncogenic cytokines.

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A.

B.

Fig. 1.

Expression of IL-8 mRNA in human pancreatic cancer cells and tissues. (A). The IL-8 mRNA expression in six human pancreatic cancer cells, and control HPDE cells were determined by realtime RT PCR. (B). The IL-8 protein expression in three human pancreatic cancer cells, and control HPDE cells were determined by Bio-plex cytokine assay. (C). The IL-8 mRNA expression in 14 pairs of human pancreatic cancer specimens and surrounding normal tissues were determined by realtime RT PCR. All data shown are the mean $\pm$ SEM of three separate experiments. * $p < 0.05$ as compared to control HPDE cells.

Fig. 2.

Effect of IL-8 on the mRNA levels of VEGF₁₆₅ and NRPs in pancreatic cancer cells. Serum-starved BxPC-3 cells were treated with 100 ng/ml of IL-8 for 24 h. Relative mRNA levels of VEGF₁₆₅ and NRPs were presented as $2^{[Ct(\beta \text{ actin}) - Ct(\text{gene})]}$ (n=10). *p < 0.05 as compared to IL-8-untreated controls.

Fig. 3.

Effect of CoCl₂ on IL-8 expression in pancreatic cancer cells. MIA PaCa-2 cells were treated with 10, 50, 100, 150, and 200 μM of CoCl₂ for 24 h. Relative mRNA levels of IL-8 was determined (n=10). *p < 0.05 and **p < 0.01 as compared to CoCl₂-untreated controls.

Fig. 4. IL-8 induces phosphorylation of ERK. BxPC-3 cells were treated with 100ng/ml IL-8 for 5, 15, 30, or 60 min and tested for the presence of phosphorylated and total ERK by Bio-Plex phosphoprotein assay. The values plotted show the ratio of phosphoprotein to total protein. Data are expressed as mean $\pm$ SEM of triplicate values from two separate experiments. **P<0.001 as compared to untreated control (0 min).