Dendritic cells fused with different pancreatic carcinoma cells induce different T-cell responses

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Background: It is unclear whether there are any differences in the induction of cytotoxic T lymphocytes (CTL) and CD4+CD25high regulatory T-cells (Tregs) among dendritic cells (DCs) fused with different pancreatic carcinomas. The aim of this study was to compare the ability to induce cytotoxicity by human DCs fused with different human pancreatic carcinoma cell lines and to elucidate the causes of variable cytotoxicity among cell lines.

Methods: Monocyte-derived DCs, which were generated from peripheral blood mononuclear cells (PBMCs), were fused with carcinoma cells such as Panc-1, KP-1NL, QGP-1, and KP-3L. The induction of CTL and Tregs, and cytokine profile of PBMCs stimulated by fused DCs were evaluated.

Results: The cytotoxicity against tumor targets induced by PBMCs cocultured with DCs fused with QGP-1 (DC/QGP-1) was very low, even though PBMCs cocultured with DCs fused with other cell lines induced significant cytotoxicity against the respective tumor target. The factors causing this low cytotoxicity were subsequently investigated. DC/QGP-1 induced a significant expansion of Tregs in cocultured PBMCs compared with DC/KP-3L. The level of interleukin-10 secreted in the supernatants of PBMCs cocultured with DC/QGP-1 was increased significantly compared with that in DC/KP-3L. Downregulation of major histocompatibility complex class I expression and increased secretion of vascular endothelial growth factor were observed with QGP-1, as well as in the other cell lines.

Conclusion: The present study demonstrated that the cytotoxicity induced by DCs fused with pancreatic cancer cell lines was different between each cell line, and that the reduced cytotoxicity of DC/QGP-1 might be related to the increased secretion of interleukin-10 and the extensive induction of Tregs.

Keywords: dendritic cell, cytotoxic T lymphocyte, interleukin-10, pancreatic cancer, regulatory T cell

Introduction
Pancreatic and biliary cancers are relatively resistant to chemotherapy and radiation and may therefore provide an opportunity for testing the potential of immunotherapy. Clinical trials of dendritic cell (DC) vaccination for advanced pancreatic cancer patients have revealed that this therapy can stimulate an antitumoral T cell response and prolong the survival of refractory pancreatic cancer patients, indicating a promising treatment modality.1–5 A critical issue in optimizing DC vaccines is the choice of tumor-associated antigen for DC loading. Some tumor antigens such as mucin 1 and α-galactosylceramide have been identified as potential targets for the immunotherapy of pancreatic carcinoma.6–10 Some reports indicated the efficacy of combined therapy with DC vaccination and chemotherapeutic agents of gemcitabine.11–13 However, immunity
against a single antigen may be ineffective in tumors with heterogeneous cell populations and may carry a risk of inducing tumor antigen escape variants. Although a recent study has demonstrated that the generation of cytotoxic T lymphocytes (CTL) against three or more tumor antigens correlated with clinical response, for many tumors, no or only few antigenic epitopes have been identified. Several recent studies have demonstrated the efficacy of the induction of antitumor immunity through the generation of fused DCs with tumor cells. In this strategy, different tumor-associated antigens, including those that are as yet unidentified, are processed endogenously and presented by major histocompatibility complex (MHC) class I pathway in the context of costimulatory signals.

The mechanisms that mediate immune tolerance to cancer are not well understood, but recent findings have revealed them to be multifactorial, including downregulation of MHC class I molecules, loss of tumor antigens, defective death receptor signaling, and generation of immunosuppressive cells such as regulatory T cells (Tregs). Major populations of these cells are CD4+CD25high Tregs that specifically express the forkhead transcription factor, forkhead/winged helix transcription factor gene (FOXP3) and CD4+ type I Tregs that secrete high levels of interleukin-10 (IL-10). Tregs secrete the immunoregulatory cytokines, IL-10 and transforming growth factor (TGF)-β1, and inhibit T-cell proliferation. Tregs are significantly increased in a population of peripheral blood lymphocytes and tumor-infiltrating lymphocytes in some epithelial cancers. Prostaglandin E2 (PGE2) induces FOXP3 gene expression and Treg function in CD4+ T cells.

Very few reports have evaluated the efficacy of DC vaccination using pancreatic cancer cell lines, being almost limited to Panc02. However, antitumor immunity based on DCs has not yet been compared among various pancreatic carcinoma cell lines, and it remains unclear whether there are any differences in induction of CTL and Tregs among DCs fused with different pancreatic carcinoma cells. We selected four representative human pancreatic cancer cell lines to compare the ability to induce cytotoxicity by human DCs fused with different human pancreatic carcinoma cell lines and to elucidate the causes of variable cytotoxicity among pancreatic carcinoma cell lines.

**Materials and methods**

**Reagents**

Recombinant human granulocyte macrophage colony-stimulating factor (GM-CSF) and IL-4 were purchased from Primimmune KK (Osaka, Japan). IL-2, lipopolysaccharide (LPS) from *Escherichia coli*, and fluorochrome-labeled PKH-26 and PKH-67 were purchased from Sigma-Aldrich (St Louis, MO, USA).

**Cell culture of pancreatic carcinoma cell lines**

A human pancreatic carcinoma cell line, Panc-1 (undifferentiated carcinoma), was obtained from the Cell Resource Center for Biomedical Research, Tohoku University (Miyagi, Japan). The human pancreatic carcinoma cell lines, KP-1NL (adenocarcinoma), QGP-1 (carcinoma of islet cell), and KP-3L (adenosquamous carcinoma) were purchased from Health Science Research Resources Bank (Osaka, Japan). Cells were maintained in 75 cm2 cell culture flask with RPMI 1640 (Life Technologies, Carlsbad, CA, USA) supplemented with 10% fetal calf serum (FCS) (Life Technologies), 100 units/mL penicillin, 100 µg/mL streptomycin, and 50 µg/mL gentamycin, and were grown at 37°C with 5% CO2 in a humidified atmosphere.

**Preparation of DCs from peripheral blood**

Monocyte-derived DCs were generated as described previously. Briefly, peripheral blood mononuclear cells (PBMCs) were isolated from peripheral blood from four healthy volunteers by Ficoll-Hypaque gradient centrifugation and were subsequently suspended in 6-well culture plates for 1 hour at 37°C. The nonadherent cells were removed. The adherent cells were harvested and cultured in RPMI 1640 supplemented with 10% FCS, 50 µM 2-mercaptoethanol, 1 µM pyruvate, 50 µg/mL gentamycin, 25 µg/mL GM-CSF, and 10 µg/mL IL-4. On day 6, DCs were harvested from the nonadherent and loosely adherent cells.

**Preparation of DCs fused with carcinoma cells**

Fused DCs were generated as described previously. Briefly, DCs were harvested at day 6. DCs were mixed with carcinoma cells treated with 10 mg/mL mitomycin C (Sigma-Aldrich) for 20 minutes at a 3:1 ratio, and were incubated in serum-free RPMI 1640 containing 50% polyethylene glycol (Sigma-Aldrich) for 7 minutes. After slowly diluting with serum-free RPMI 1640, the cells were washed and resuspended in RPMI.
1640 supplemented with 10% FCS, 25 µg/mL GM-CSF, and 10 µg/mL IL-4. DCs cocultured with carcinoma cells at the same ratio in RPMI 1640 supplemented with FCS, GM-CSF, and IL-4 were used as “norfused DCs.” The next day, fused DCs and nonfused DCs were induced to mature with 1 µg/mL LPS for 24 hours. To determine fusion efficiency, DCs were labeled with PKH26 red fluorescent dye, and tumor cells were labeled with PKH67 green fluorescent dye prior to cell fusion. Individual cells or fused DCs were analyzed by flow cytometry and fluorescence microscopy. As a control, DCs generated from PBMCs and stimulated with LPS were examined, which were used as “DCs alone.”

Flow cytometry analysis

Cells were washed with 0.01 M phosphate-buffered saline, pH 7.4 and incubated with antibodies directed against MHC class I (W6/32; mouse IgG2α; Dako, Glostrup, Denmark), MHC class II (HLA-DR) (mouse IgG1; Nichirei, Tokyo, Japan), CD40 (LOB-11; mouse IgG1; Santa Cruz Biotechnology, Santa Cruz, CA, USA), CD80 (MAB104; mouse IgG1; Immunotech, Marseille, France), and CD86 (BU63; mouse IgG1, Ylem, Roma, Italy) for 30 minutes on ice. After washing with phosphate-buffered saline, the cells were incubated with fluorescent isothiocyanate (FITC)-conjugated goat anti-mouse IgG (Beckman Coulter, Marseille, France) for 30 minutes on ice. In the case of FITC-conjugated antibody against ligand (Beckman Coulter), cells were incubated for 30 minutes on ice. The appropriate respective mouse isotype controls (Dako) were employed. Samples were washed, fixed in 4% paraformaldehyde, and analyzed by FACS-Calibur and CellQuest software (BD, Franklin Lakes, NJ, USA).

Cytotoxicity assay

Fused DCs, nonfused DCs, and DCs alone were cocultured with autologous nonadherent PBMCs in a 1:20 ratio, in the presence of 25 units/mL IL-2. Fifty percent of the medium was replaced on days 3 and 5, and the cultures were restimulated by freshly isolated DCs with culture medium containing 25 units/mL IL-2 on day 7. On day 14, the stimulated PBMCs were harvested and washed three times in serum-free medium and applied as effectors at various effector/target ratios. Pancreatic carcinoma cells as target cells (1 × 10^6) were cocultured with effector cells for 4 hours at 37°C. To measure lactate dehydrogenase (LDH) release from targets, the supernatants were harvested and cytotoxicity was examined using the LDH Cytotoxicity Detection Kit (Takara Bio, Tokyo, Japan) as described previously. Briefly, 100 µL of LDH substrate was added to the 96-well flat-bottomed plate and incubated for 30 minutes at room temperature. The optical absorbance of red formazan was determined spectrophotometrically at a wavelength of 490 nm. Spontaneous release of LDH by effectors or targets was assessed by incubation of the effectors or targets alone, respectively. Maximum release of LDH was determined by incubation of the targets in 1% Triton X-100. The percentage of specific cytotoxicity was calculated according to the following formula:

\[ \text{% cytotoxicity} = 100 \times \left( \frac{\text{Experimental} - \text{Effector spontaneous} - \text{Target spontaneous}}{\text{Target spontaneous}} \right) \]

Quantification of cytokine secretion

The supernatants of the cocultures were harvested 48 hours after the second stimulation with fused DCs, and the concentrations of human cytokines, interferon-γ, IL-10 (Endogen, Rockford, IL, USA), TGF-β1, and PGE₂ (R&D Systems, Minneapolis, MN, USA) were quantified by enzyme-linked immunosorbent assay (ELISA). In other experiments, the supernatants of the media were harvested from 48-hour-cultured respective pancreatic carcinoma cells, DCs, and fused DCs at a density of 2 × 10^5/mL in 6-well plate, and the concentrations of human cytokines, vascular endothelial growth factor (VEGF) (Endogen), IL-10, TGF-β1, and PGE₂ were quantified by ELISA.

Reverse transcription-polymerase chain reaction (RT-PCR) for IL-21

RNA was extracted from pancreatic carcinoma cells and PBMCs using RNeasy Mini Kit (Qiagen, Hilden, Germany) and was reverse-transcribed by QIAGEN OneStep RT-PCR Kit (Qiagen) according to the manufacturer’s instructions. Amplification was performed in a total volume of 25 µL for 35 cycles of 30 seconds at 94°C, 30 seconds at 54°C, and 30 seconds at 72°C. PCR products were resolved by electrophoresis on 2% agarose gels (Funakoshi, Tokyo, Japan), the bands were visualized by ethidium bromide (Bio-Rad Laboratories, Hercules, CA, USA) staining, and photographs of the gels were taken. The positive control for the expression
of IL-21 mRNA was RNA from normal PBMCs treated in anti-CD3-coated plates for 24 hours.

Primers used to detect IL-21 mRNA (GenBank accession number: BC066258) were 5′-ACAGACTA ACATGCCCTTCA-3′ for the forward primer, and 5′-TCTTCACTTCCGTGTTCT-3′ for the reverse primer, which produced PCR products of 134 bps. IL-21 primers were synthesized by Nihon Gene Research Laboratories (Miyagi, Japan). Primers used to detect β-actin mRNA were 5′-GATCAGCAAGCAGGATATG-3′ for the forward primer, and 5′-GGCCATTCTCCTTAGAGA-3′ for the reverse primer, which produced PCR products of 390 bps.

**Analysis of CD4+CD25high T cells in cocultured PBMCs**

PBMCs cocultured with fused DCs or DCs alone for 14 days were washed and incubated with a FITC-conjugated antibody against CD4 (13B8.2; mouse IgG1; Beckman Coulter) and a phycoerythrin-conjugated antibody against CD25 (B1.49.4; mouse IgG2a; Immunotech) for 30 minutes on ice. Mouse isotype controls were employed. Cells were washed again and analyzed by flow cytometry using a live gate set around 2000 cells. Positive cells were counted in at least 50 fields under a light microscope (×200).

**Expression of FOXP3 and IL-10 in cocultured PBMCs**

Cocultured PBMCs were immunostained for FOXP3 to recognize CD4+CD25high Tregs. Briefly, PBMCs cocultured with fused DCs or DCs alone for 14 days were washed, and cytospins from PBMCs were air-dried and fixed for 10 minutes in acetone. Endogenous peroxidase activity was blocked with methanol containing 0.3% hydrogen peroxide. Monoclonal antibody anti-FOXP3 (ab20034; mouse IgG1; Abcam, Cambridge, England) was incubated at 4°C overnight on the cytospin slides. The labeled streptavidin-biotin-peroxidase method (Ultratech HRP Streptavidin-biotin Universal Detection System; Dako) was used, and a positive reaction was detected as a brown color with 3,3′-diaminobenzidine (Dojin Chemicals, Kumamoto, Japan). The slides were counterstained with 1% methyl green (Muto Pure Chemicals, Tokyo, Japan). Positive cells were counted at least 50 fields under a light microscope (×200) by two observers independently. As a control, FOXP3+ T cells in PBMCs before coculture were analyzed, respectively. Next, phycoerythrin-conjugated antibody against IL-10 (JES3-19F1; rat IgG2a; PharMingen, San Diego, CA, USA) was incubated at 4°C overnight on the cytospin slides. The number of positive cells was counted in at least 50 fields under a fluorescence microscopy (×200).

**Statistical analysis**

Data were expressed as the mean ± standard deviation. Statistical significance was determined by the Student’s t-test for paired data using StatView-software (Abacus Concepts, Berkeley, CA, USA). Differences were considered statistically significant for P < 0.05.

**Results**

**Characterization of DCs fused with pancreatic carcinoma cells**

Fusion efficiency of DCs labeled with PKH26 and carcinoma cells labeled with PKH67 was confirmed by fluorescence microscopy. The population of fused DCs was 43.9% ± 4.85% of total cells by flow cytometry. The expression of MHC class II and costimulatory molecules on DCs was then analyzed by flow cytometry. Unstimulated (immature) nonfused DCs strongly expressed MHC class II (HLA-DR) and CD40, and low levels of CD80 and CD86 (Figure 1). Nonfused DCs stimulated by LPS (mature DCs) strongly expressed MHC class II and costimulatory molecules such as CD80, CD86, and CD40. The immunophenotype of fused DCs (Panc-1 [Figure 1], KP-1NL, KP-3L, and QGP-1 [data not shown]) was similar to that of mature DCs.

**Induction of cytotoxicity against pancreatic carcinoma cell lines**

To assess the induction of antitumor immune response by fused DCs against the pancreatic carcinoma cell lines, fused DCs were cocultured with autologous PBMCs. As a control, PBMCs were also cocultured with nonfused DCs or DCs alone. PBMCs cocultured with DCs fused with Panc-1 (DC/Panc-1), KP-1NL (DC/KP-1NL), or KP-3L (DC/KP-3L) induced significant cytotoxicity against tumor targets compared with those cocultured with DCs alone (P < 0.05; Figure 2A). By contrast, PBMCs cocultured with DCs fused with QGP-1 (DC/QGP-1) induced only a low level of cytotoxicity and there was no significant difference between fused DCs and controls (nonfused and DCs alone). Furthermore, when the cytotoxicity of PBMCs cocultured with fused DCs was compared among pancreatic carcinoma cell lines, the level of cytotoxicity in DC/QGP-1 was significantly lower compared with that of other carcinoma cell lines (P < 0.05; Figure 2B).
It was investigated whether or not the four carcinoma cell lines produced immunoregulatory cytokines, such as IL-10, TGF-β₁, VEGF, and PGE₂ by ELISA. IL-10 and TGF-β₁ were secreted at very low levels from the four carcinoma cell lines (Figure 3A). A greater amount of VEGF, which suppresses the differentiation and maturation of DCs, was secreted from Panc-1 and QGP-1 compared with KP-3L (P < 0.05). PGE₂ was secreted from KP-3L at a higher level than from the other cell lines. IL-21 mRNA, that likely suppressed the maturation of DCs and failed to induce antigen-specific T cell proliferation, was not expressed in any tumor cells by RT-PCR (Figure 3B). Moreover, to confirm the maturation of fused DCs, the expression of MHC class II and costimulatory molecules, such as CD80, CD86, and CD40, on DC/QGP-1 or DC/KP-3L was investigated by flow cytometry. The expression of these molecules was increased, consistent with the immunophenotype of mature DCs, and no suppression of the maturation of fused DCs was observed.

**Secretion of immunoregulatory cytokines from pancreatic carcinoma cell lines**

It was investigated whether or not the four carcinoma cell lines produced immunoregulatory cytokines, such as IL-10, TGF-β₁, VEGF, and PGE₂ by ELISA. IL-10 and TGF-β₁ were secreted at very low levels from the four carcinoma cell lines (Figure 3A). A greater amount of VEGF, which suppresses the differentiation and maturation of DCs, was secreted from Panc-1 and QGP-1 compared with KP-3L (P < 0.05). PGE₂ was secreted from KP-3L at a higher level than from the other cell lines. IL-21 mRNA, that likely suppressed the maturation of DCs and failed to induce antigen-specific T cell proliferation, was not expressed in any tumor cells by RT-PCR (Figure 3B). Moreover, to confirm the maturation of fused DCs, the expression of MHC class II and costimulatory molecules, such as CD80, CD86, and CD40, on DC/QGP-1 or DC/KP-3L was investigated by flow cytometry. The expression of these molecules was increased, consistent with the immunophenotype of mature DCs, and no suppression of the maturation of fused DCs was observed.

**Secretion of immunoregulatory cytokines from PBMCs cocultured with fused DCs**

ELISA was performed to assess interferon-γ, IL-10, TGF-β₁, and PGE₂ secretion in the supernatants of PBMCs 48 hours after the second stimulation by DCs fused with QGP-1, as a cell line that induced a low level of cytotoxicity, or DCs fused with KP-3L, as a cell line that induced a high level of cytotoxicity. As a control, cytokines in the supernatants of PBMCs cocultured with DCs alone were assessed. Interferon-γ secretion in PBMCs cocultured with DC/QGP-1 or DC/KP-3L was significantly higher than that in DCs alone (P < 0.05; Figure 4). There was no significant difference between DC/QGP-1 and DC/KP-3L. IL-10 secretion in PBMCs cocultured with DC/QGP-1 was significantly higher than that with DC/KP-3L and DCs alone (P < 0.05). The level of TGF-β₁ secretion was not significantly different among the three experimental groups. PGE₂ secretion in the PBMCs cocultured with DC/QGP-1 or DC/KP-3L was increased significantly, compared with that with DCs alone. There was no significant difference between DC/QGP-1 and DC/KP-3L. To investigate whether PBMCs cocultured with DC/QGP-1 secreted IL-10, cocultured PBMCs were immunofluorescently stained with anti-IL-10 antibody. The number of IL-10⁺ cells in PBMCs cocultured with DC/QGP-1 was significantly increased compared with that with DC/KP-3L and DCs alone (P < 0.05; Figure 5).

**Expansion of CD4⁺CD25⁺high Tregs**

The relative expansion of Tregs in PBMCs cocultured with DC/QGP-1 and DC/KP-3L for 14 days was compared.
with that in the PBMCs before coculture. CD4+CD25<sup>high</sup> Tregs in PBMCs cocultured with DC/QGP-1 were significantly increased compared with those in PBMCs cocultured with DC/KP-3L and DCs alone by flow cytometry (CD4+CD25<sup>high</sup>) (Figure 6) and immunocytochemistry (FOXP3) (Figure 7).

**Expression of MHC class I in pancreatic carcinoma cell lines**

The expression of MHC class I in carcinoma cell lines was evaluated by flow cytometry. Almost all tumor cells in the cell lines were positive for MHC class I antigen (Figure 8A) and there was no significant difference among cell lines. By contrast, from the perspective of mean fluorescent intensity, MHC class I antigen in KP-1NL and QGP-1 was significantly lower compared with that in KP-3L ($P < 0.05$; Figure 8B).

**Expression of Fas ligand in pancreatic carcinoma cells**

Fas ligand was not detectable in carcinoma cells by flow cytometry (data not shown).
The current study firstly investigated whether DCs fused with pancreatic carcinoma cells induced cytotoxicity against the tumor targets in four human pancreatic carcinoma cell lines, and examined the contribution of immunosuppressive factors against the cytotoxicity of PBMCs cocultured with fused DCs among cell lines.

This study demonstrated that DC/Panc-1, DC/KP-1NL, and DC/KP-3L significantly induced cytotoxicity against the tumor targets compared with DCs alone, excluding DC/QGP-1. When the cytotoxicity induced by fused DCs was compared among the pancreatic carcinoma cell lines, the level of cytotoxicity in QGP-1 was significantly lower compared with that in the other cell lines.

Recent studies have demonstrated that the mechanisms of tumor escape from immune recognition/destroy are likely to be multifactorial, such as the downregulation of MHC class I molecules, loss of tumor antigens, defective death receptor signaling, production of immunosuppressive cytokines, and the existence of suppressive cells. To reveal the factors that suppressed cytotoxicity of PBMCs cocultured with fused DCs, we examined these factors among pancreatic carcinoma cell lines and demonstrated that significantly higher levels of VEGF were detected in the supernatants of Panc-1 and QGP-1 compared with KP-3L. The previous study indicated that caudal related homeobox gene 2 (CDX2), an intestine-specific tumor suppressor gene, was expressed at a high level in QGP-1 cells, and cell proliferation and cyclin D1 mRNA

Discussion

The current study firstly investigated whether DCs fused with pancreatic carcinoma cells induced cytotoxicity against the tumor targets in four human pancreatic carcinoma cell lines, and examined the contribution of immunosuppressive factors against the cytotoxicity of PBMCs cocultured with fused DCs among cell lines.

Figure 3 (A) Immunoregulatory cytokine secretion from pancreatic carcinoma cell lines and dendritic cells (DCs). Forty-eight hours after suspending carcinoma cells and DCs at a density of 2 × 10^5/mL in the 6-well plate, the supernatants of the medium were harvested, and quantified for secreted interleukin-10 (IL-10), transforming growth factor (TGF)-β1, vascular endothelial growth factor (VEGF), and prostaglandin E (PGE) by enzyme linked immunosorbent assay (ELISA). Results are expressed as the mean ± standard deviation (SD) of four independent experiments, and compared by Student’s t-test. *P < 0.05. (B) IL-21 mRNA by reverse transcription-polymerase chain reaction (RT-PCR).

Notes: The expression level of IL-21 mRNA was assessed in four pancreatic carcinoma cell lines. The positive control for the expression of IL-21 mRNA was RNA from normal peripheral blood mononuclear cells (PBMCs) treated in anti-CD3-coated plates for 24 hours. The result is representative of three independent experiments.
Figure 4 Immunoregulatory cytokine secretion from peripheral blood mononuclear cells (PBMCs) cocultured with dendritic cells (DCs) fused with QGP-1 (DC/QGP-1), DCs fused with KP-3L (DC/KP-3L), or DCs alone.

Notes: Forty-eight hours after the second stimulation with fused DCs or DCs alone, the supernatants of PBMCs were harvested and quantified for secreted interferon (IFN)-γ, interleukin-10 (IL-10), transforming growth factor (TGF)-β1, and prostaglandin E2 (PGE2) by enzyme linked immunosorbent assay (ELISA). Results are expressed as the mean ± standard deviation (SD) of six independent experiments, and compared by Student’s t-test. *P < 0.05.

level were inhibited significantly after CDX2 transfection in pancreatic cancer cells, suggesting that CDX2 might play a role in inhibiting cell proliferation and repressing cyclin D1 transcriptional activity. However, in our study there were not any significant differences in induction of cytotoxicity among DCs alone, nonfused DCs, and fused DCs (DC/QGP-1) (Figure 2). Thus, the immunosuppressive potential of DC/QGP-1 may be due to only “active” in the context of professional antigen presentation from immune cells rather than an intrinsic property of QGP-1. VEGF is produced by many tumors and is important not only for tumor vascularization, but is also a key factor produced by solid tumors to inhibit immune recognition and to prevent DC maturation and maturation. Furthermore, to examine whether increased production of VEGF was associated with the suppression of cytotoxicity against pancreatic carcinoma cell lines, this study investigated the expression of MHC class II and costimulatory molecules in DC/QGP-1 or DC/KP-3L by flow cytometry. The expression of these molecules was similar to the expression of surface antigens in mature DCs. DC/QGP-1 and DC/KP-3L matured sufficiently to induce the immune response. Independently, expression of MHC class I antigens in KP-1NL and QGP-1 was significantly downregulated compared with KP-3L, when these results were expressed as fluorescent intensity. Downregulation of MHC class I was recognized in QGP-1 as well as KP-1NL, which induced a higher degree of cytotoxicity. MHC class I molecule downregulation occurs frequently in many cancers, and the altered phenotypes of MHC class I antigen expression can allow tumors to avoid recognition or survive attack by cytotoxic CD8+ T-cells. Kasuya et al have also demonstrated that the hypervascular pancreatic tumor, QGP-1 secreted a higher level of VEGF under a hypoxic environment than the hypovascular pancreatic ductal cell lines (BxPC-3 and AsPC-1).
These findings did not demonstrate that high secretion of VEGF and downregulation of MHC class I could be related to the immunosuppression of DC/QGP-1, despite a significant difference between QGP-1 and KP-3L. Furthermore, there was the elevated cytotoxicity of the nonfused DCs/tumor cell mixture (Figure 2). Nonfused DCs/tumor cell mixture may recognize a low level of soluble antigens secreted from tumor cells or surface antigens expressed weakly on the cell surface of tumor cells.

Recent studies have reported that CD4⁺CD25⁺Tregs are immunoregulatory and are important in immunological tolerance to self-antigens.¹⁹⁻²² The present study demonstrated that CD4⁺CD25⁺Tregs in PBMCs cocultured with DC/QGP-1 were significantly increased compared with those in PBMCs cocultured with DC/KP-3L or DCs alone by flow cytometry (CD4⁺CD25⁺) and immunocytochemistry (FOXP3). To investigate whether Tregs induced an immunosuppressive cytokine profile, the secretion of IL-10 and TGF-β1 was subsequently analyzed in the supernatants of PBMCs cocultured with fused DCs. There was no significant difference in the level of TGF-β1 secretion among the three experimental conditions. Although a recent study demonstrated the induction of Treg function in CD4⁺ T cells by PGE₂ secreted from carcinoma cells and the present study revealed that PGE₂ was secreted to a much higher level in the supernatants of KP-3L but not QGP-1 in the supernatants of PBMCs cocultured with DC/QGP-1 or DC/KP-3L compared with that in DCs alone, there was no significant difference between DC/QGP-1 and DC/KP-3L. Cytokines (IL-10 and TGF-β), immature myeloid DCs, CpG-ODN stimulated plasmacytoid DCs, and immunomodulatory drugs (vitamin D₃) can induce CD4⁺CD25⁺ T-cells to differentiate into CD4⁺CD25⁺ T cells.¹⁹⁻²² In this study, IL-10 secretion in PBMCs cocultured with DC/QGP-1 or DC/KP-3L was significantly higher than that with DC/KP-3L or DCs alone, suggesting that the source of increasing IL-10 may include DCs functionally altered by fusion mechanism and IL-10 secreting Tregs such as CD4⁺CD25⁺ T cells and CD4⁺ type 1 Tregs.⁴⁰ Subsequently, increasing IL-10 may be related to the expansion of Tregs and the induction of the immunosuppressive response. Leao et al.⁴⁴ demonstrated that effective depletion of Treg cells allows the recruitment of mesothelin-specific CD8⁺ T cells to the antitumor immune response against a mesothelin-expressing mouse pancreatic adenocarcinoma. Yamamoto et al.⁴⁴ also reported that Treg depletion combined with DC/tumor cell fusion hybrid vaccine enhanced the efficacy of immunotherapy in pancreatic cancer by activating CTLs. Similar effectiveness of Treg depletion in DC vaccination has been indicated in eradication of lymphoma cells in mice.⁴⁵

Although the overall 5-year survival for patients of pancreatic adenocarcinoma is only 1%–4%, that of malignant pancreatic endocrine carcinoma was 49%.⁴³ DC-based vaccination against pancreatic endocrine carcinoma with liver metastases revealed a decreased tumor marker level and a tumor regression of metastases.⁴⁴ However, information
about effects of DC-based immunotherapy against pancreatic endocrine carcinoma is quite limited. Thus, it is critical to compare the difference in induction of cytotoxicity between pancreatic ductal carcinoma and endocrine carcinoma, as shown in this study.

In conclusion, the present study firstly demonstrated that there were significant differences in the induction of CTL and CD4^+CD25^{high} Tregs, and the amount of IL-10 among DCs fused with different pancreatic carcinomas. These findings suggest that the reduced cytotoxicity of DC/QGP-1 may be related to the increased secretion of IL-10 and the extensive induction of CD4^+CD25^{high} Tregs.

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Disclosure
The authors report no conflicts of interest in this work.

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