Hypoxia-mediated immune evasion of pancreatic carcinoma cells

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Abstract. Hypoxia is one of the characteristics of human and animal tumors. To investigate the association between hypoxia and the immune evasion of cancer cells, the present study examined paraffin sections of pancreatic tissues from patients with pancreatic carcinoma, chronic pancreatitis and normal pancreatic tissue and established a series of PANC-1 cell lines, which were cultured under various hypoxic and normoxic conditions. The results demonstrated that the expression of hypoxia-inducible 1α (HIF- 1α) in pancreatic carcinoma was significantly higher compared with that in the chronic pancreatitis and normal pancreatic tissues, which revealed that a hypoxic microenvironment existed in pancreatic carcinoma. HIF-1a was inversely correlated with major histocompatibility complex class I chain-related (MIC) genes, which indicated that hypoxia was involved in tumor immune evasion. The cell experiments demonstrated that the mechanism involved shedding of the MIC from the membrane of the pancreatic carcinoma cells, which then formed soluble (s)MIC. The sMIC genes downregulated natural killer (NK) group 2, member D and the cytotoxic activity of NK cells. Depending on its activity, the nitric oxide-cyclic guanosine monophosphate-protein kinase G signaling pathway can either increase or inhibit immune evasion of pancreatic cancer cells.

Introduction

Through mutation, tumor cells may express novel antigens, which are recognized by the immune system. The innate and acquired immune systems are subsequently activated, resulting in tumor cell death. However, there are exceptions, in which tumor cells modify the surface antigens and alter the microenvironment to evade the immune response (1).

There is substantial evidence that hypoxia is present in malignant solid tumors and is associated with increased tumor growth, invasion, metastatic potential and resistance to therapy (2). Fink *et al* (3) indicated that liver cancer cells exposed to hypoxia were more tolerant to natural killer (NK) cells compared with those cultivated under normoxic conditions.

Hypoxia-inducible factor-1 (HIF-1) is a sequence-specific DNA-binding transcription factor, which is regulated by hypoxia and transcriptionally activates the expression of several genes (4). HIF-1 consists of HIF-1 α and HIF-1 β (5). HIF-1 β is stable intracellularly, while HIF-1 α is associated with cellular O₂ (6). The major histocompatibility complex class I chain-related (MIC) gene has seven family members; however, only the MICA and MICB genes encode proteins (7). The MICA/B proteins are located in the cell membrane and act as ligands of the NK group 2, member D (NKG2D). These proteins are rarely expressed by normal cells; however, they are expressed in a variety of malignant diseases (8-10). NKG2D is well-established as a receptor expressed on NK cells. Following ligand binding, it transfers the signal downstream to activate the cytotoxic activity of NK and T-cells (11-13).

Hypoxia in pancreatic cancer widely exists; therefore, a series of experiments was designed in order to determine whether pancreatic cancer cells resisted the cytotoxic effect of NK cells and whether the resistance was associated with hypoxia. The present study also aimed to elucidate whether the resistance observed was associated with the expression of NKG2D on NK cells and the molecule MIC on the pancreatic cancer cell membrane.

Materials and methods

Materials. Mouse anti-human MICA/B (1:150) and NKG2D (1:100) monoclonal antibodies were obtained from Santa Cruz Biotechnology, Inc, (Dallas, TX, USA). Rabbit anti-human HIF-1 α polyclonal antibody was obtained from Boshide Bio, Inc. (Wuhan, China). Glyceryl trinitrate (GTN) was obtained from Sabex (Boucherville, QC, Canada), KT5823 and NG-mono-methyl-L-arginine (L-NMMA) were obtained from Biyuntian (Shanghai, China) and MTT and 8-bromoguanosine cyclic monophosphate (8-Br-cGMP) were obtained from Sigma-Aldrich (St. Louis, MO, USA).

Patients and samples. Pathological specimens were obtained from 42 patients undergoing surgical resection for pancreatic carcinoma and nine patients undergoing surgical resection for chronic pancreatitis at Xiangya Hospital (Changsha, China) between April 2010 and April 2012. In addition, eight normal pancreatic tissue samples were obtained from the Xiangya

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Sample			HIF-1a				MIC A/B			
	n	-	+	++	+++	P-value	0	1	2	P-value
Pancreatic carcinoma	42	10	2	12	18		4	13	25	
Chronic pancreatitis	9	7	2	0	0	< 0.001	7	2	0	<0.001
Normal pancreas	8	0	0	0	0		7	1	0	

Table I. Expression of HIF-1 α and MICA/B in pancreatic carcinoma, chronic pancreatitis and normal pancreatic tissues.

Transplant Center (Changsha, China) or from autopsy. The general sample information for the 59 total samples, graded as previously described (14,15), is shown in Table I. The study was approved by the ethics committee of Xiangya Hospital, Central South University, Changsha, China.

Cells. The PANC-1 cell line, purchased from the Resource Center of Shanghai Institutes of Biological Sciences (Shanghai, China), was maintained in a monolayer culture in Dulbecco's modified Eagle's medium supplemented with 20% fetal bovine serum (Gibco-BRL, Invitrogen Life Technologies, Carlsbad, CA, USA). NK cells were purchased from the Resource Center of Shanghai Institutes of Biological Sciences (Shanghai, China). NK cells were cultured in RPMI-1640 containing 10% fetal bovine serum, penicillin (100 U/ml) and streptomycin (100 μ g/ml). Interleukin (IL)-2 (1000 U/ml) was also added to the medium. Following 72 h of incubation the NK cells were able to be used as effector cells.

Culture conditions. In order to establish hypoxic conditions, the cells were placed in airtight chambers, which were flushed with a gaseous mixture of 4.5% CO₂, 95% N₂ and 0.5% O₂ with optional addition of GTN (10 nM), 8-Br-cGMP (10 nM) or KT5823 (10 μ M). Another group of cells was incubated with 20% O₂ and optional addition of L-NMMA (5 μ g/ml).

Immunohistochemical analysis. The tumor samples were fixed with 10% paraformaldehyde (Blue Star, Shanghai, China) and embedded in paraffin (Behai Chemical Industry Company, Changsha, China). Sections (4 μ m) were then cut from the samples using a Leica RM2135 microtome (Leica Biosystems, Bensheim, Germany) and were adhered to microscope slides. The sections were dewaxed and washed three times with phosphate-buffered saline (PBS; Sigma-Aldrich). For non-specific inhibition, each section was incubated in 10% normal goat serum (Boster, Wuhan, China) for 30 min at 37°C and then with primary antibodies, rabbit anti-human polyclonal HIF-1 α and the mouse anti-human monoclonal MIC A/B, overnight at 4°C. The sections were washed three times with PBS and were incubated with secondary antibody linked with biotin and then with horseradish peroxidase (HRP)-marked anti-biotin (Boster) for 30 min at 37°C. Subsequently, the sections were incubated in freshly prepared diaminobenzidine (Boster) and subsequently counterstained with hematoxylin (Boster). The sections were then observed under an optical microscope (CX41; Olympus Corp., Tokyo, Japan).

Reverse transcription quantitative polymerase chain reaction (RT-qPCR) analysis. The total RNA was extracted using TRIzol reagent. RT was performed using 1 μ g total RNA and oligo(dT) primers (Sangon Biotech Co., Ltd, Shanghai, China) which were as follows: homo-MICA forward, 5'-AGGTACATCTGGATGGTCAG-3' and reverse, 5'-TTGTCTTCATGGATCTCACA-3' with an amplified fragment of 232 bp; homo-MICB forward, 5'-CTTCGTTACAACCT CATGGT-3' and reverse, 5'-ATATGAGTCAG GGTCCTCCT-3' with an amplified fragment of 227 bp and homo-GAPDH forward, 5'-ACCACAGTCCATGCCATCAC-3' and reverse, 5'-TCCACCACCCT GTTGCTGTA-3' with an amplified fragment of 450 bp. The PCR cycling conditions were as follows: 95°C for 5 min, followed by 40 cycles of 94°C for 30 sec, annealing at 54°C for 30 sec and 72°C for 30 sec, with a final extension at 72°C for 5 min. Conditions were then maintained at 4°C.

ELISA analysis. A total of two non-overlapping epitope antibodies were used for ELISA analysis. The polystyrene board plates (Boster) were coated with the mouse anti-human monoclonal MIC A/B immunoglobulin G (IgG) overnight at 4°C, then inhibited with 5% fetal bovine serum for 2 h at 37°C and washed with PBS. Subsequently, the cell culture medium samples were added and incubated for 1 h at 37°C. The plates were then washed with PBS and incubated with goat anti-mouse IgG antibody linked with HRP for 1 h at 37°C. TMB chromogenic agent and TMB terminated liquid (Boster) were added and, following reaction, the absorbance was measured at 450 nm using a DU640 ultraviolet spectrophotometer (Beckman Coulter, Miami, FL, USA).

Flow cytometric analysis. The cultured cells were collected and resuspended in PBS at a density of 1x10⁶/ml. Subsequently, antibody linked with fluorochrome (mouse anti-human monoclonal MIC A/B IgG) was added and incubated for 30 min at 4°C in the absence of visible light. Following washing three times with PBS, the cells were detected by fluorescence-activated cell sorting (FACS) using a FACSCalibur instrument (BD Biosciences, Franklin Lakes, NJ, USA).

MTT analysis. The cells were cultured in 96-well plates. MTT (20 μ l of 5 mg/ml) was added to each well and the cells were incubated for 4 h at 37°C. Following incubation, 100 μ l

	HIF-1α				MICA/B			
Cases (n)	Positives (n)	Positive rate (%)	χ^2	P-value	Low	High	χ^2	P-value
18	14	77.8	0.025	>0.05	7	11	0.010	>0.05
24	18	75.0	0.025	20.05	10	14	0.019	20.05
32	24	75.0	0.010	> 0.05	13	19	0.111	> 0.05
10	8	80.0	0.010	>0.03	4	6		>0.03
28	16	57.1	4 0 2 5	-0.05	6	22	10.280	<0.05
14	13	92.9	4.023	<0.05	11	3	10.309	<0.05
12	9	75.0			2	10		
12	8	66.7	1.116	>0.05	3	9	9.141	< 0.05
18	15	88.3			12	6		
32	25	78.1	0.010	>0.05	12	20	0 111	> 0.05
10	7	70.0	0.010		5	5	0.111	>0.05
23	21	91.3	4 602	<0.05	9	14	0.014	>0.05
19	11	57.9	4.093	<0.05	8	11	0.014	×0.05
	Cases (n) 18 24 32 10 28 14 12 12 18 32 10 23 19	Cases (n)Positives (n)181424183224108281614131291281815322510723211911	HIF-1 α Cases (n)Positives (n)Positive rate (%)181477.8241875.0322475.010880.0281657.1141392.912975.0181588.3322578.110770.0232191.3191157.9	HIF-1 α Cases (n)Positives (n)Positive rate (%) χ^2 181477.80.025241875.00.010322475.00.01010880.00.010281657.14.025141392.91.1161588.366.71.116181588.30.010232578.10.010232191.34.693191157.94.693	HIF-1 α Cases (n)Positives (n)Positive rate (%) χ^2 P-value181477.80.025>0.05241875.00.010>0.05322475.00.010>0.0510880.00.010>0.05281657.14.025<0.05	HIF-1 α HIF-1 α LowCases (n)Positives (n)Positive rate (%) χ^2 P-valueLow181477.80.025>0.05710241875.00.010>0.05134322475.00.010>0.051310880.00.010>0.056141392.94.025<0.05	HIF-1 α MICACases (n)Positives (n)Positive rate (%) χ^2 P-valueLowHigh181477.80.025>0.05711241875.00.010>0.05131910880.00.010>0.05146281657.14.025<0.05	$\begin{array}{c c c c c c c c c c c c c c c c c c c $

Table II. Expression	on levels of HIF-	-1α and MICA/B	according to	clinicopa	thological p	parameters.
			0		0	

HIF-1a, hypoxia-inducible factor 1a; MICA/B, major histocompatibility complex class I chain-related A/B; TNM, tumor-node-metastasis.

Table III. Correlations between HIF-1 α and MICA/B.

HIF-1α	Low (0/1)	High (2)	r	P-value
Low (-/+)	0	12	0.500	.0.001
High (++/+++)	17	13	-0.522	<0.001
Total	17	25		

HIF-1 α , hypoxia-inducible factor 1 α ; MICA/B, major histocompatibility complex class I chain-related A/B.

MTT solution was added and the cells were agitated on an orbital shaker for 1 min. The absorbance was then read at 570 nm using an Elx-800 microplate reader (Winooski, VT, USA).

Statistical analysis. All statistical analyses were performed using the SPSS 13.0 statistical software package (SPSS, Inc., Chicago, IL, USA). Differences between groups were compared using single factor analysis of variance and Student's t-test. P<0.05 was considered to indicate a statistically significant difference between values. Table IV. Expression of sMICA/B under different culturing conditions.

Conditions/ supplements	Absorbance	sMICA (pg/ml)	sMICB (pg/ml)
20% O ₂	0.598	127.14	3.664
	0.776	169.92	5.219
	0.654	140.60	3.495
20% O ₂ + L-NMMA	1.113	250.91	9.274
	1.429	326.86	7.077
	1.127	254.28	6.165
$0.5\% \mathrm{O_2}$	1.556	357.38	5.590
	1.388	317.00	7.517
	1.360	310.27	8.125
0.5% O ₂ + GTN	0.868	192.03	6.030
	0.796	174.73	7.348
	0.881	195.16	7.719

Cells were cultured in Dulbecco's modified Eagle's medium containing 20% fetal bovine serum. HIF-1 α , hypoxia-inducible factor 1 α ; sMICA/B, soluble major histocompatibility complex class I chain-related A/B; GTN, glyceryl trinitrate; L-NMMA, NG-mono-methyl-L-arginine.





Figure 1. Expression levels of HIF-1 α and MIC in pancreatic carcinoma, chronic pancreatitis and normal pancreatic tissues. HIF-1 α in (A and B) pancreatic carcinoma tissue, (C and D) chronic pancreatitis tissue, (E and F) chronic pancreatitis tissue, and (G and H) normal pancreas tissue. (I and J) MIC in pancreatic carcinoma tissue. Magnification: A, C E G and H, x100; B, D, F, H and J, x400. Dark granules indicate positive staining. HIF-1 α , hypoxia-inducible factor 1 α ; MIC, major histocompatibility complex class I chain-related.

Figure 2. Protein expression of MIC in PANC-A cells under different culture conditions. (A) Cells cultured in 0.5% O₂ (grey) and 20% O₂. (B) Cells cultured in 0.5% O₂ (grey) and 0.5% O₂ + glyceryl trinitrate. (C) Cells cultured in 0.5% O₂ (grey) and 0.5% O₂ + 8-bromoguanosine cyclic monophosphate. (D) Cells cultured in 20% O₂ and 20% O₂ + NG-mono-methyl-L-arginine, a nitric oxide synthase inhibitor (grey). (E) Cells cultured in 20% O₂ and 20% O₂ + KT5823, a protein kinase G inhibitor (grey). HIF-1 α , hypoxia-inducible factor 1 α ; mMIC, membrane-bound major histocompatibility complex class I chain-related.

Results

Expression levels of HIF-1 α and MICA/B in pancreatic carcinoma, chronic pancreatitis and normal pancreatic tissues. As shown in Table I, protein expression of HIF-1 α was observed in 32 out of 42 pancreatic carcinoma samples and 2 out of 9 chronic pancreatitis samples; however, HIF-1 α protein was not detected in the normal pancreatic samples. In addition, protein expression of MICA/B was observed in 38 out of 42 pancreatic

carcinoma samples and 2 out of 9 chronic pancreatitis samples, but was detected in only one normal pancreatic tissue sample. The expression levels of HIF-1 α and MICA/B were either positively or negatively correlated to tumor-node-metastasis (TNM) staging; however, no correlation with the pathological type was observed (P>0.05; Table II). According to Spearman's rank correlation coefficient, the two proteins were negatively correlated (P<0.001; Table III). Immunohistochemistry also

Pre-incubation	Cytotoxic activity of NK cells (%) (NK:PANC-1 cell ratio)						
conditions	6.25:1	12.5:1	25:1	50:1			
20% O ₂	32.4±5.1	35.7±4.5	40.7±5.9	43.9±5.5			
$20\%~O_2 + L\text{-NMMA}$	19.1±3.4	22.1±4.2	25.4±4.4	33.3±5.1			
0.5% O ₂	15.8±3.3	17.7±2.9	25.2±4.0	29.1±4.2			
$0.5\% O_2 + GTN$	16.6±2.8	21.8±3.1	29.5±4.1	38.2±5.7			

Table V. Cytotoxic activity of NK cells on PANC-1 cells.

Values are presented as the mean ± standard deviation.NK, natural killer; GTN, glyceryl trinitrate; L-NMMA, NG-mono-methyl-L-arginine.



Figure 3. Expression of sMICA/B in medium under different culture conditions. (A) sMICA and (B) sMICB under different culture conditions. sMIC; soluble major histocompatibility complex class I chain-related; GTN, glyceryl trinitrate; L-NMMA, NG-mono-methyl-L-arginine.

revealed that, in the pancreatic carcinoma cells, the expression of MIC on the membrane was reduced; however, it was detected in the interstitial tissue (Fig. 1).

Effects of hypoxia on membrane-bound (m) and soluble (s) MICA/B. As demonstrated previously, $HIF-1\alpha$, regulated by hypoxia, was increased in pancreatic carcinoma tissues, while mMIC was decreased with accumulation in interstitial



Figure 4. Expression of NK group 2, member D in NK cells co-incubated with PANC-1 cells at a ratio of 1:1 for 24 h. (A) PANC-1 cells pre-incubated in 0.5% O_2 and 20% O_2 (grey). (B) PANC-1 cells pre-incubated under 0.5% O_2 and 0.5% O_2 + glyceryl trinitrate (grey). (C) PANC-1 cells pre-incubated under 20% O_2 (grey) and 20% O_2 + NG-mono-methyl-L-arginine; NK, natural killer.



Figure 5. Cytotoxic activity of NK cells. (A) PANC-1 cells pre-cultured in 0.5% O_2 and 0.5% O_2 + GTN. (B) PANC-1 cells pre-cultured in 20% O_2 and 20% O_2 + L-NMMA. *P<0.05 vs. the other group at the same proportion. NK, natural killer; GTN, glyceryl trinitrate; L-NMMA, NG-mono-methyl-L-arginine.

tissue (16). To assess whether hypoxia reduced mMIC and increased sMIC, the PANC-1 pancreatic carcinoma cell line was cultured in 0.5% O₂. The results revealed that the protein expression of mMIC decreased compared with that in cells cultured in 20% O_2 (Fig. 2). However, no clear differences were detected in the mRNA expression levels of mMICA/B between the two groups of cells. Furthermore, the addition of a single dose of either GTN (10 nM) or 8-Br-cGMP (10 nM) to the cells incubated for 24 h in 0.5% O2 was sufficient to prevent the reduction in mMIC. Inhibiting nitric oxide (NO) synthesis with the NO synthase inhibitor L-NMMA (5 μ g/ml) or suppressing protein kinase G (PKG) activity with the PKG inhibitor KT5823 (10 μ M) reduced the expression of mMIC in the cells cultured in 20% O₂ (Fig. 2). By contrast, sMICA increased ~2-fold (P<0.05) in 0.5% O2, which was prevented by co-incubation with GTN (10 nM; Fig. 3). Co-incubation with L-NMMA (5 μ g/ml) also increased the levels of sMICA (Table IV, Fig. 3). No significant differences were observed in sMICB (Table IV; Fig. 3).

Effect of hypoxia on the protein expression of NKG2D. To assess the effect of hypoxia on the protein expression of NKG2D, PANC-1 cells which were pre-incubated in 0.5% O_2 and 20% O_2 with or without GTN and L-NMMA were collected and co-incubated with NK cells. As Fig. 4 shows, compared with the cells incubated in 20% O_2 , those cultured in 0.5% O_2 had lower protein expression levels of NKG2D (40.5 vs. 70.3%). The addition of GTN promoted the expression of NKG2D to 62.5%, while L-NMMA downregulated the expression to 48.4% (Fig. 4).

Effect of hypoxia on the cytotoxic activity of NK cells. To determine the effect of hypoxia on the cytotoxic activity of NK cells, the PANC-1 cells were pre-cultured under different conditions and then co-incubated with NK cells in varying proportions (1:6.25, 1:12.5, 1:25 and 1:50) for 24 h. An MTT assay revealed that cells cultured in 0.5% O_2 were more tolerant to the NK cells compared with those incubated in 20% O_2 . GTN increased the cytotoxic activity and L-NMMA decreased the cytotoxic activity of the NK cells (Table V; Fig. 5).

Discussion

The present study demonstrated an increase in the levels of sMIC, resulting in a decrease in the protein expression of NKG2D, which was key in regulating the hypoxia-mediated immune evasion of human pancreatic carcinoma cells. Furthermore, the results of the present study clearly suggested that NO signaling was essential in this process.

Pathological slides revealed that HIF-1 α was highly expressed in pancreatic carcinoma tissues, which indicated that hypoxia was present in the pancreatic carcinoma. In addition, MIC decreased compared with the chronic pancreatitis and normal pancreatic tissues, suggesting that hypoxia may downregulate MIC.

To mimic the hypoxic microenvironment experienced by cancer cells, the present study cultured the PANC-1 cell line in 0.5% O_2 , with control cells cultured in 20% O_2 . The results demonstrated that the expression of HIF-1 α was high and that of MIC was low (data not shown). However, RT-qPCR revealed no clear differences in the mRNA expression of MICA/B between cells cultured in 0.5% O₂ and 20% O₂, raising the question of whether this was due to downregulation of MIC translation or detachment of MIC to the extracellular space. ELISA demonstrated that the level sMIC, particularly sMICA, increased in the cell culture medium exposed to 0.5% O₂, which had a negative effect on the expression of NKG2D and the cytotoxic activity of NK cells. The results indicated that hypoxia did not inhibit the gene expression of MIC, but induced the detachment of MIC from the cell membrane and downregulated the expression of NKG2D and the cytotoxicity of NK cells. This resulted in the reduced immunogenicity of cancer cells and subsequent evasion of the immune system. Matrix metalloproteinases (MMP) may be important molecules in this regulation, as it has been reported that MMP-2 and -9 are often upregulated in tumors (17). Salih et al (18) investigated the addition of MMP inhibitor to the medium, which revealed marked increases in mMIC. Another study confirmed that the hypoxia-induced detachment of mMIC is regulated by MMPs (19). However, the reason for downregulation in the expression of NKG2D and NK cytotoxicy remain to be elucidated. One theory is that the binding of sMIC and NKG2D causes the internalization of NKG2D. Doubrovina et al (20) demonstrated that following binding to soluble MIC, NKG2D is decreasingly located at the membrane and is increasingly present in the cytoplasm. Groh et al (21) also observed that sMIC induced the internalization and degradation of NKG2D on T cells.

Studies have also demonstrated that hypoxia markedly inhibits the cellular NO-cGMP-PKG pathway (22-25). Inhibiting the pathway with drugs has a mimetic effect of hypoxia (26-29). In the present study, GTN and 8-Be-cGMP were used as NO-cGMP-PKG pathway stimulators, and L-NMMA and KT5823 were used as inhibitors. Treatment with GTN and 8-Be-cGMP reduced sMICA and promoted the expression of NKG2D and NK cytotoxicity, whereas treatment with L-NMMA and KT5823 had an opposite effect. It is understood that the NO-cGMP-PKG pathway begins with NO synthesis. As a soluble gas molecule, NO passes through the cell membrane and activates soluble guanylyl cyclase (sGC), which catalyzes the conversion of GTP to cGMP. cGMP then acts as a secondary messenger, activating PKG and amplifying NO signals to downstream effectors (26). According to the results of the present study, activation of the NO-cGMP-PKG pathway weakened the hypoxia-mediated immune evasion of pancreatic carcinoma cells.

All the experiments in the present study were performed *in vitro* and, as there is greater interest in the immune response *in vivo*, subsequent investigation aims to establish a nude mouse model to study the effect of NO-mimetic drugs on tumorigenicity.

In conclusion, the results of the present study indicated that the NO-cGMP-PKG signaling pathway regulated the immune escape observed in pancreatic cancer. Activation of this pathway can reverse the immune escape, whereas its inhibition promoted the immune escape. The hypoxic environment, which the pancreatic cancer cells existed in, was closely associated with this signaling pathway. Further research on NO-cGMP-PKG signal pathway and hypoxia environment will provide novel directions for the development of immunotherapies for pancreatic cancer.

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