Novel crosstalk between KLF4 and ZEB1 regulates gemcitabine resistance in pancreatic ductal adenocarcinoma

ZHIYI WANG1, YUAN CHEN2, YANLIANG LIN3, XINXING WANG1, XIANPING CUI1, ZHENHAI ZHANG1, GUOZHE XIAN1 and CHENGKUN QIN1

Departments of 1Hepatobiliary Surgery, 2Paediatrics and 3Center Laboratory, Shandong Provincial Hospital Affiliated to Shandong University, Jinan, Shandong 250021, P.R. China

Received May 26, 2017; Accepted August 8, 2017

DOI: 10.3892/ijo.2017.4099

Abstract. Pancreatic ductal adenocarcinoma (PDAC) is one of the most lethal malignancies with broad resistance to chemotherapeutic drugs. Krüppel-like factor 4 (KLF4) is a candidate tumor suppressor in PDAC. However, the precise role of KLF4 in gemcitabine resistance of PDAC remains largely unclear. In this study, we demonstrated that gemcitabine inhibited KLF4 expression. Moreover, gemcitabine also reduced the levels of miR-200b and miR-183, but promoted ZEB1 expression in PDAC cells. KLF4 knockdown blocked the expression of miR-200b and miR-183, and inversely, KLF4 overexpression promoted the expression of miR-200b and miR-183, suggesting that KLF4 positively regulated the expression of miR-200b and miR-183. Moreover, KLF4 knockdown enhanced ZEB1 expression and gemcitabine resistance while KLF4 overexpression induced the opposite effect. ChIP assays verified that KLF4 positively regulated the expression of miR-200b and miR-183 by directly binding to their promoters. Then, miR-200b and miR-183 directly inhibited ZEB1 expression by targeting its 3'UTR region. ZEB1 knockdown attenuated gemcitabine resistance in PDAC cells. KLF4 overexpression promoted gemcitabine sensitivity of PDAC in vivo by negatively regulating ZEB1 expression. Our results revealed that novel crosstalk between KLF4 and ZEB1 regulated gemcitabine resistance in PDAC.

Introduction

Pancreatic ductal adenocarcinoma (PDAC) is a highly lethal malignancy with an estimated 5-year survival rate <5%, and by 2020, it is estimated to be the second leading cause of cancer-related mortality (1). Due to the lack of early detection methods and effective preventive strategies, PDAC is usually diagnosed at an advanced stage when it is deemed unresectable and exhibits poor response to chemotherapeutic drugs, including gemcitabine and 5-fluorouracil (2). Thus, it is essential to understand the underlying mechanisms for the chemoresistance of PDAC.

Krüppel-like factor 4 (KLF4) is a transcription factor that serves as an important regulator in cell differentiation, cell cycle, cell survival and apoptosis (3). KLF4 exerts seemingly paradoxical functions in different cell types. KLF4 is downregulated in bladder and gastric cancer cells, and enforced expression of KLF4 promotes apoptosis in these cells (4). In PDAC, KLF4 also functions as a tumor-suppressor (5). Conversely, KLF4 exhibits oncogenic functions in oral, skin squamous carcinoma cells and breast ductal carcinoma (6). To date, little is known about the precise role of KLF4 in gemcitabine resistance of PDAC.

The epithelial-mesenchymal transition (EMT)-activator ZEB1 is a transcriptional repressor of epithelial genes, such as E-cadherin (7). ZEB1 has been widely implicated in the invasiveness and metastasis of solid tumors, including lung adenocarcinoma, colorectal and breast cancers (8-10). Ectopic expression of ZEB1 also promotes drug-resistant phenotypes (7). However, the role of ZEB1 in gemcitabine resistance of PDAC remains unexplored.

In this study, we showed that gemcitabine treatment inhibited KLF4 expression and promoted ZEB1 expression. Further investigation demonstrated that KLF4 suppressed ZEB1 expression and gemcitabine resistance by directly regulating the expression of miR-200b and miR-183. In addition, ZEB1 knockdown attenuated gemcitabine resistance in PDAC cells.

Materials and methods

Antibodies and reagents. The following monoclonal antibodies (mAbs) and polyclonal antibodies (pAbs) were used: anti-KLF4 (pAb, ab106629), and anti-ZEB1 (pAb, ab155249) were purchased from Abcam (Cambridge, MA, USA); anti-GAPDH (pAb) was purchased from Anbo Biotechnology Co. (San Francisco, CA, USA); Cell counting kit-8 was purchased from Dojindo Laboratories (Tokyo, Japan). Gemcitabine was purchased from Eli Lilly (Indianapolis, IN, USA). ZEB1 siRNA, miR-200b
and miR-183 were synthesized by Biosune Biotechnology (Shanghai, China).

**Cell culture.** The human PDAC cell lines BxPC-3, Panc-1 and MIApaca-2 were purchased from the American Type Culture Collection (Manassas, VA, USA). Cells were cultured in RPMI-1640 medium (Hyclone, USA) supplemented with 10% fetal bovine serum (Hyclone), 100 U/ml penicillin, 100 U/ml streptomycin and 0.03% L-glutamine at 37°C in 5% CO2.

**Cell viability assay.** PDAC cells were seeded at a density of 1.0x10^4 cells/well in 96-well microplates with 100 µl RPMI-1640 medium. After grown to ~80% confluence, the cells were incubated with the indicated concentration of gemcitabine. After treatment for 72 h, the cells were incubated with 10% WST-8 dye for 2 h at 37°C. The absorbance was detected at 450 nm using a SpectraMax M2. IC50 values were determined using GraphPad Prism 6.0.

**Flow cytometric analysis.** PDAC cells were seeded at a density of 2x10^5 cells/well in 6-well microplates. After treatment, flow cytometric analysis was performed to determine cell apoptosis. In brief, after treatment with gemcitabine, the cells were digested using trypsin lacking EDTA and phenol red, and were incubated in binding buffer containing Annexin V-FITC (2.5 mg/ml) and propidium iodide (5 mg/ml) for 10 min in the dark at room temperature. Subsequently, the labeled cells were detected by a flow cytometer (Beckman Coulter, Chicago, IL, USA).

**Luciferase reporter assay.** Panc-1 and MIApaca-2 cells were co-transfected with increasing concentrations of pcDNA3.1 carrying KLF4 and 0.2 µg of luciferase reporter plasmid containing the miR-200b or miR-183 promoter using Lipofectamine 3000 (Roche); the pRL-SV40 plasmid (Promega) was transfected as a normalization control. After 24-h incubation, cell lysates were collected to determine luciferase activity using the Dual-Luciferase assay (Promega) according to the manufacturer's instructions. The primers were used as follows: 5′-ATATGCGGCCGCGGTGTGGGTAAGGTGCT-3′ and 5′-GCTTGGTACCAGGCCACCTTCTCTCTTCTTCTTCTC-3′ for KLF4; 5′-AGAAAGGGTGGGAAGGAGGACA-3′ and 5′-GGACTCTGCTGGAGAAGCTCAGTA-3′ for miR-200b; 5′-GAGACTCCGCTGGAGAAGCTCAGTA-3′ and 5′-CGGGAAGGTGCTTCTCTGCT-3′ for miR-183. The sequence of ZEB1-3′UTR containing the miR-200b or miR-183 seed regions were amplified using specific primers (forward, 5′-CGAGCTCAGTGCCATTTCTCAGTATT-3′ and reverse, 5′-CCGCTCGAGAAGTTCTTTTCTGAGTGGTGGG-3′). The product was digested using the restriction enzymes BamHI and AgeI, and was then linked to p-GC365-EGFP plasmid. The plasmids were transformed into E. coli DH5α. Positive clones were harvested and identified by sequencing. The recombinant plasmids were transfected to 293T cells by Lipofectamine 3000 (Invitrogen). The infection efficiency was determined by measuring fluorescence intensity under microscopy (Olympus 3.3RTV, Japan). After 48 h of transfection, the supernatant was collected by centrifugation at 4,000 g for 10 min at 4°C, followed by ultrafiltration centrifugation at 4000 g for 12 min. The concentrated solution was stored at -80°C for future experiments.

**Gene transfection.** BxPC-3, Panc-1 and MIApaca-2 cells were seeded at a density of 2x10^5 cells/well in 6-well microplates. After growing overnight, the cells were transfected with lentivirus carrying shRNAs against KLF4 or KLF4 cDNA (GeneChem, Shanghai, China). For ZEB1 knockdown, PDAC cells were transfected with siRNA against ZEB1 (GGTAGAGGTATATGTAATA) using Lipofectamine 3000 (Invitrogen). Western blot analysis. Western blot analysis was performed as previously described (11). Equal amounts of protein were loaded to sodium dodecyl sulfate polyacrylamide gels. The protein was subsequently transferred onto PVDF membranes. After blocked with 5% BSA, membranes were incubated with the indicated primary antibodies overnight at 4°C, followed by treatment with horseradish peroxidase-conjugated secondary antibodies for 1 h at room temperature. The protein signals were detected using the ECL kit and quantified using an Alpha Imager 2200 (Alpha Innotech Corp., USA).

**Northern blot analysis.** The total RNA was extracted using TRIzol reagent (Invitrogen). RNA (20 µg) was subjected to 15% urea-polyacrylamide gels, and then transferred to nylon membrane. The expression of miR-200b, miR-183 and U6 were detected using biotin-labeled oligonucleotide probes (5′-TCATCATTACCGGCGATATTA-3′, miR-200b; 5′-CGAGCTCAGTGCCATTTCTCAGTATT-3′, miR-183 and 5′-ATATGGAACGCTTCAGAATT-3′, U6). The signals of blots were measured using the Fujifilm LAS-4000 imaging system.

**Chromatin immunoprecipitation assay (ChIP).** ChIP was performed using a ChIP assay kit (Millipore) according to the manufacturer's instructions. In brief, Panc-1 cells were crosslinked with fresh 1% formaldehyde, followed by incubation in SDS lysis buffer containing 1% protease inhibitors.
The cell lysates were sonicated to shear crosslinked DNA to 200-1,000 bp in length. Protein/DNA complexes were immunoprecipitated with 3 µg anti-ZEB1 antibody or control IgG. The complexes were eluted from the antibodies, and were dissociated with 5 M NaCl. ChIP samples were analyzed with real-time quantitative PCR using primers specific for the promoters of miR-200b and miR-183. The primers are listed in Tables I and II.

**Quantitative RT-PCR.** Total RNA was extracted using TRizol reagent (Invitrogen). The expression of miR-200b and miR-183 were analyzed using TaqMan miRNA assays (Applied Biosystems, CA, USA) according to the manufacturer's instructions. U6 snRNA was used as an internal control.

**Immunohistochemistry.** Immunohistochemistry was performed as previously described (12). Briefly, tumor sections were fixed in formalin, embedded with paraffin, deparaffinized in xylene, and hydrated with ethanol. After antigen retrieval using microwave, the tissue sections were treated with 3% hydrogen peroxide and blocked with 10% goat serum. The sections were then stained with anti-KLF4 and anti-ZEB1 antibodies, followed by treatment with a biotinylated secondary antibody. The images were captured using a fluorescence microscope (Leica DM IRE2).

**Mouse xenograft models.** The orthotopic PDAC nude mouse model was established as previously described by us (12), with a minor modification. Briefly, 4- to 6-week-old BALB/c nude mice of both sexes were obtained from Weitonglihua Animal Center (Beijing, China) and maintained under specific pathogen-free conditions in the animal facility. All procedures were approved by the Institutional Laboratory Animal Care and Use Committee at Shandong Provincial Hospital. Panc-1 cells (1x10^7) in 100 µl PBS/Matrigel (1:1, v/v, BD Biosciences, USA) transplanted with control lentivirus or lentivirus carrying KLF4 cDNA were injected into both flanks of the mice. One week after injection, gemcitabine (80 mg/kg; q3 days) was administered via intraperitoneal injection for six weeks. The long (L) and short axes (S) of the tumors were measured weekly using Vernier calipers, and the tumor volumes were calculated as follows: V = (L x S^2) x π/6. The mice were sacrificed 24 h after the last injection, and the tumors were harvested for subsequent analysis.

**Statistical analysis.** Statistical analysis was carried out using the SPSS 18.0 software package. All data are presented as the mean ± standard deviation from at least three independent experiments. Student’s t-test was used for comparisons between two groups. One-way ANOVA was used for comparisons among multiple groups. p<0.05 was considered statistically significant.

**Results**

**Gemcitabine inhibits the expression of KLF4, miR-200b and miR-183 and promotes ZEB1 expression.** First, we measured the expression of KLF4 and ZEB1 in BxPC-3, Panc-1 and MIApaca-2 cells. Fig. 1A showed a higher level of KLF4 in BxPC-3 cells than those in Panc-1 and MIApaca-2 cells while lower levels of ZEB1 in BxPC-3 cells than those in Panc-1 and MIApaca-2 cells. Further investigation found...
that gemcitabine treatment reduced the expression of KLF4, miR-200b and miR-183 in a dose-dependent manner, whereas led to a dose-dependent increase in ZEB1 expression (Fig. 1D and E).

KLF4 attenuates ZEB1 expression and gemcitabine resistance by upregulation of miR-200b and miR-183. Next, we examined the correlation between KLF4 and ZEB1. KLF4 knockdown significantly promoted ZEB1 expression while KLF4 overexpression inhibited ZEB1 expression (Fig. 2A). KLF4 depletion inhibited gemcitabine-induced activation of caspase 3 while KLF4 overexpression promoted the activation of caspase 3 even in the absence of gemcitabine (Fig. 2B), suggesting that KLF4 downregulation conferred gemcitabine resistance of PDAC cells. KLF4 knockdown restrained the expression of miR-200b and miR-183 (Fig. 2C). In contrast, KLF4 overexpression elevated the levels of miR-200b and miR-183 (Fig. 2C). In KLF4-depleted cells, forced expression of miR-200b and miR-183 repressed the ZEB1 expression again (Fig. 2D). In KLF4-overexpressing cells, forced expression of anti-miR-200b and anti-miR-183 restored the ZEB1 expression (Fig. 2E). Restoration of miR-200b and miR-183 activated caspase 3 in KLF4-depleted cells whereas anti-miR-200b and anti-miR-183 significantly reduced caspase 3 activation in KLF4-overexpressing cells (Fig. 2F). Overexpression of miR-200b and miR-183 in KLF4-depleted cells promoted cell apoptosis in the presence of gemcitabine, but inhibition of miR-200b and miR-183 in KLF4-overexpressing cells blocked cell apoptosis induced by gemcitabine (Fig. 2G). These results suggested that KLF4 knockdown contributed to ZEB1 expression and gemcitabine resistance by suppressing miR-200b and miR-183.
KLF4 positively regulates the expression of miR-200b and miR-183 in PDAC cells. Since KLF4 is a transcriptional factor that regulates the expression of multiple genes, we hypothesized that KLF4 directly regulated the expression of miR-200b and miR-183. The putative binding sites of KLF4 located on the promoters of miR-200b and miR-183 were identified. Figure 2 demonstrates the effects of KLF4 on ZEB1 expression and gemcitabine resistance. (A) BxPC-3, Panc-1, and MIApaca-2 cells were transfected with lentivirus carrying shRNAs against KLF4 or KLF4 cDNA. The expression of KLF4 and ZEB1 were analyzed by western blot analysis. *p<0.01 vs. control; **p<0.001 vs. control. (B) The KLF4-depleted or KLF4-overexpressing cells were treated with 10 µM gemcitabine. The activation of caspase 3 was determined by western blot analysis. (C) BxPC-3, Panc-1, and MIApaca-2 cells were transfected with lentivirus shRNAs against KLF4 or KLF4 cDNA. The expression of miR-200b and miR-183 were analyzed by RT-PCR and northern blot analysis. (D) The KLF4-depleted cells were transfected with miR-200b and miR-183. The expression of ZEB1 was analyzed by western blot analysis. (E) The KLF4-overexpressing cells were transfected with anti-miR-200b and anti-miR-183. The expression of ZEB1 was analyzed by western blot analysis. (F) The KLF4-depleted cells were transfected with miR-200b and miR-183, followed by treatment with 10 µM gemcitabine. The KLF4-overexpressing cells were transfected with anti-miR-200b and anti-miR-183, followed by treatment with 10 µM gemcitabine. The activation of caspase 3 was determined by western blot analysis. Cell viabilities were determined by CCK8 assay (G). *p<0.01.
shown in Fig. 3A and D. Chromatin immunoprecipitation (ChIP) assays revealed that KLF4 could interact with site 4 and 5 in the miR-200b promoter and the site 4 in the miR-183 promoter (Fig. 3B and E). We subsequently constructed the pGL3-200-luc and pGL3-183-luc plasmids by inserting these sequences into the pGL3 luciferase reporter. Panc-1 and MIApaca-2 cells were co-transfected with increasing concentrations of pcDNA3.1 plasmid carrying KLF4 and 0.2 µg pGL3-200b-luc using Lipofectamine 3000. After 48-h transfection, luciferase activity was measured. ‘p<0.01 vs. control group; ‘p<0.05 vs. the group transfected with 0.1 µg pcDNA3.1 carrying KLF4. (D) The putative binding sites of KLF4 in miR-183 promoter are shown. (E) ChIP assays identified the binding sites of KLF4 in miR-183 promoter. (F) Panc-1 and MIApaca-2 cells were co-transfected with increasing concentrations of pcDNA3.1 plasmid carrying KLF4 and 0.2 µg pGL3-200b-luc using Lipofectamine 3000. After 48-h transfection, luciferase activity was measured. ‘p<0.05 vs. control group; ‘p<0.05 vs. the group transfected with 0.1 µg pcDNA3.1 carrying KLF4.

miR-200b and miR-183 directly targets ZEB1 in PDAC cells. We further explored whether miR-200b and miR-183 directly targeted the 3’UTR region of ZEB1. Targetscan (http://www.targetscan.org/) was used to predict the recognition sites of miR-200b and miR-183. As shown in Fig. 4A, four putative sites for miR-200b and two putative sites for miR-183 were found. The sequences containing these sites were cloned into the luciferase reporter plasmid to obtain the ZEB1-3’UTR-WT luc plasmid (wild-type), and the sequence containing the mutant sites was cloned to construct ZEB1-3’UTR-MT luc plasmid (mutant type). Subsequently, the Panc-1 cells were co-transfected with ZEB1-3’UTR-WT or ZEB1-3’UTR-MT and miR-200b or miR-183. Both miR-200b and miR-183 significantly inhibited luciferase activities in the cells carrying ZEB1-3’UTR-WT compared to those carrying negative control (NC) and ZEB1-3’UTR-MT (Fig. 4B). Furthermore, both miR-200b and miR-183 notably reduced mRNA and protein levels of ZEB1 in BxPC-3, Panc-1 and MIApaca-2 cells (Fig. 4C and D). Conversely, both anti-miR-200b and anti-miR-183 elevated mRNA and protein levels of ZEB1 (Fig. 4C and D). These results indicated that miR-200b and miR-183 directly targeted ZEB1.

ZEB1 knockdown contributes to gemcitabine sensitivity of PDAC cells. As shown in Fig. 5A, gemcitabine treatment promoted ZEB1 expression, which was consistent with the results in Fig. 1D. ZEB1 knockdown mildly activated caspase 3. A combination of ZEB1 knockdown and gemcitabine significantly increased cleaved caspase 3, suggesting that ZEB1 knockdown facilitated gemcitabine sensitivity of PDAC cells. Fig. 5B illustrated that gemcitabine treatment induced cell apoptosis in ZEB1-depleted cells. These data indicated that ZEB1 knockdown contributed to gemcitabine sensitivity of PDAC cells.

Negative association between KLF4 and ZEB1 regulates gemcitabine resistance of PDAC in vivo. We established xenograft tumors of PDAC to determine the effect of KLF4 overexpression on the ZEB1 expression and gemcitabine resistance of PDAC in vivo. The results showed that KLF4
overexpression reduced ZEB1 expression, consistent with the data in vitro (Fig. 6A-C). Linear regression analysis showed that KLF4 expression was negatively correlated to ZEB1 expression (Fig. 6D). Compared to the control group, KLF4 overexpression did not significantly affect tumor growth, whereas it enhanced gemcitabine sensitivity (Fig. 6E and F). These results suggested that KLF4 overexpression attenuated gemcitabine resistance of PDAC in vivo by negatively regulating ZEB1 expression.

Discussion

Chemotherapy resistance has been a major challenge in improving the overall survival of patients with PDAC. Gemcitabine is a standard clinical chemotherapeutic drug for advanced PDAC, but results in a progression-free survival interval ranging from 0.9 to 4.2 months only (12). Thus, understanding of the mechanism by which chemotherapy resistance occurs contributes to the development of novel therapeutic strategies for overcoming advanced PDAC. KLF4 has been identified as a tumor suppressor in diverse types of cancer (13-15), including PDAC (5). In the pancreas, KLF4 is predominantly expressed in PDAC and regulates the expression of cytokeratin-19, a specific marker for PDAC (16). In this study, we observed a higher level of KLF4 expression in BxPC-3 compared to those in Panc-1 and MIApaca-2. BxPC-3 cells were more sensitive to gemcitabine than Panc-1 and MIApaca-2. Fig. 1D illustrated that gemcitabine treatment

Figure 4. miR-200b and miR-183 directly targets ZEB1 in PDAC cells. (A) Targeting sequences of miR-200b and miR-183 in the 3’UTR region of ZEB1. (B) Panc-1 cells were co-transfected with miR-200b or miR-183 and ZEB1-3’UTR-WT or ZEB1-3’UTR-MT. The relative luciferase activities were determined using the Dual-Luciferase assay. *p<0.01 vs. NC group. (C) BxPC-3, Panc-1 and MIApaca-2 cells were transfected with miR-200b, miR-183, anti-miR-200b or anti-miR-183. The mRNA expression of ZEB1 was determined by RT-PCR. *p<0.01 vs. NC group. The protein expression of ZEB1 was determined by western blot analysis (D). *p<0.01 vs. NC group.
Figure 5. ZEB1 knockdown attenuates gemcitabine resistance in PDAC cells. BxPC-3, Panc-1 and MIApaca-2 cells were transfected with siRNA against ZEB1 for 48 h, followed by treatment with 10 µM gemcitabine. (A) The expression of ZEB1 and cleaved caspase 3 were analyzed by western blot analysis. (B) Cell viabilities were determined by CCK8 assay. *p<0.05; **p<0.01.

Figure 6. KLF4 reduces gemcitabine resistance of PDAC in vivo by negatively regulating ZEB1 expression. (A) Representative IHC staining for KLF4 and ZEB1 in tumor tissues from control mice or KLF4 expressing mice. (B) Western blot analysis of KLF4 and ZEB1 in tumor tissues from control mice or KLF4 expressing mice. (C) The relative expression of KLF4 and ZEB1 in tumor tissues from control mice or KLF4 expressing mice. (D) Linear regression analysis of KLF4 and ZEB1 expression was performed using GraphPad Prism 6.0 software. (E) Xenograft tumors from control mice and KLF4 expressing mice are shown. (F) Growth curves of tumors from control mice or KLF4 expressing mice after gemcitabine treatment. *p<0.05; **p<0.01; ***p<0.001.
miR-200 family members cooperate to affect stemness properties in PDAC cells by suppressing Bmi1 (19). miR-183 and miR-200b subsequently inhibited the Wnt/β-catenin signaling pathway (20,21). In this study, overexpression of miR-183 and miR-200b reduced gemcitabine resistance of PDAC cells while KLF4 overexpression reduced gemcitabine resistance. These results confirmed that the low level of KLF4 promoted gemcitabine resistance of PDAC cells. How does KLF4 regulate gemcitabine resistance of PDAC cells? Our results showed that gemcitabine treatment simultaneously triggered the downregulation of KLF4, miR-200b and miR-183 in PDAC cells. KLF4 silencing inhibited the expression of miR-200b and miR-183. In contrast, KLF4 overexpression promoted the expression of miR-200b and miR-183. ChIP assays revealed that KLF4 positively regulated the expression of miR-200b and miR-183 by directly binding to their promoters. These results suggested that lower expression of KLF4 conferred PDAC cells with greater gemcitabine resistance by upregulation of miR-200b and miR-183. Consistently, the expression of miR-200b has been found to be significantly downregulated in gemcitabine-resistant cells (18). The expression of miR-183 is related to expression of miR-200 family members (19). miR-183 and miR-200 family members cooperate to affect stemness properties in PDAC cells by suppressing Bmi1 (19), function as EMT inhibitors and accelerate epithelial differentiation by targeting the Wnt/β-catenin signaling pathway (20,21). In this study, overexpression of miR-183 and miR-200b reduced the expression of ZEB1 at both mRNA and protein levels in PDAC cells, even in KLF4-depleted cells that ZEB1 was strongly increased. Conversely, anti-miR-200b and anti-miR-183 increased the mRNA and protein levels of ZEB1, even in KLF4-overexpressing cells in which ZEB1 was significantly inhibited. Combined with the data from luciferase reporter assays, we confirmed that miR-200b and miR-183 targeted ZEB1 in PDAC cells. Congruously, ZEB1 has been found to be direct target of miR-200b in hepatocellular carcinoma, osteosarcoma and non-small cell lung cancer (22-24). Although it has been confirmed that ZEB1 represses the transcription of miR-183 in lung cancer (25,26), it is unclear whether miR-183 directly targets ZEB1. Our results indicated that KLF4 knockdown enhanced gemcitabine resistance by suppressing the miR-200b/miR-183/ZEB1 signaling pathway. ZEB1 is an EMT-activator that is correlated to tumor metastasis, drug resistance and poor prognosis in different tumor types (27). A series of studies have suggested that EMT promotes early-stage dissemination, invasion and metastasis of PDAC (28,29). A recent study demonstrates that EMT is dispensable for metastasis but increases chemoresistance in pancreatic cancer (30). Our results also demonstrated that ZEB1 knockdown promoted gemcitabine sensitivity of PDAC cells.

In conclusion, we demonstrated that gemcitabine inhibited the expression of KLF4, blocking KLF4-mediated downstream signals. Further investigation confirmed that KLF4 induced the expression of miR-200b and miR-183 by directly binding to the promoters of miR-200b and miR-183. Elevated miR-200b and miR-183 subsequently inhibited ZEB1 expression by directly targeting the 3'UTR of ZEB1. Inhibition of ZEB1 attenuated gemcitabine resistance. In addition, KLF4 overexpression enhanced gemcitabine sensitivity of PDAC in vivo by inhibiting ZEB1 expression. Taken together, our results suggested that novel crosstalk between KLF4 and ZEB1 regulates gemcitabine resistance in PDAC cells (Fig. 7).

Acknowledgements

This study was supported by the National Natural Science Foundation of China (grant nos. 81572272 and 81201778), Shandong Provincial Natural Science Foundation (grant no. ZR2013HQ026) and the Science and Technology Development Plan Project of Shandong Province (grant nos. 2013G0021810 and 2016 GSF201127).

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