Long non-coding RNA TMPO-AS1 serves as a tumor promoter in pancreatic carcinoma by regulating miR-383-5p/SOX11

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Abstract. The dysregulation of lncRNA TMPO antisense RNA 1 (TMPO-AS1) has been detected in various malignant tumors. However, the role of lncRNA TMPO-AS1 remains unclear in pancreatic carcinoma. The present study aimed to elucidate the functional mechanism of TMPO-AS1 in pancreatic carcinoma. In the present study, RT-qPCR, western blotting, MTT, Transwell, luciferase reporter and xenograft assays were used to investigate the role of lncRNA TMPO-AS1 in pancreatic carcinoma. Upregulation of lncRNA TMPO-AS1 was revealed in pancreatic carcinoma tissues and cells. Furthermore, knockdown of TMPO-AS1 restrained cell proliferation and motility in pancreatic carcinoma. In addition, microRNA (miR)-383-5p acted as a 'sponge' for lncRNA TMPO-AS1. The expression levels of lncRNA TMPO-AS1 and miR-383-5p were mutually inhibited in pancreatic carcinoma. Moreover, miR-383-5p was revealed to directly target SRY-related high-mobility group box 11 (SOX11). Notably, SOX11 could promote the occurrence of pancreatic carcinoma by interacting with the lncRNA TMPO-AS1/miR-383-5p axis. In conclusion, upregulation of IncRNA TMPO-AS1 promoted tumor growth, cell migration and invasion in pancreatic carcinoma by downregulating miR-383-5p and upregulating SOX11.

Introduction

Pancreatic carcinoma is a malignant tumor of the digestive tract, which is difficult to diagnose and treat. Approximately 90% of pancreatic cancers are ductal adenocarcinomas derived from ductal epithelium (1). In recent years, its morbidity and mortality have significantly increased. To make matters worse, the surgical mortality rate is high, and the cure rate is markedly low (2). At present, the cause of pancreatic cancer

is not clear. It has been reported that chronic pancreatitis and diabetes may be related to the occurrence of pancreatic cancer (3). In addition, pancreatic cancer is a highly malignant tumor with a markedly poor prognosis. The one-year survival rate for pancreatic cancer patients is 8%, and the five-year survival rate is 3%. The median survival time is only 2 to 3 months (4). Early diagnosis and early treatment is the key to improving the prognosis of pancreatic cancer patients. Therefore, it is urgent to explore new targets to improve the survival rate of pancreatic cancer patients.

Long noncoding RNAs (lncRNAs) have been reported to regulate the tumorigenesis of numerous human cancers, including pancreatic cancer. For example, lncRNA CASC2 suppressed the metastasis of pancreatic carcinoma cells by downregulating miR-21 (5). DUXAP8 is a pseudogene derived lncRNA that promotes the growth of pancreatic carcinoma cells through epigenetic silencing of CDKN1A and KLF2 (6). In the present study, the role of lncRNA TMPO antisense RNA 1 (TMPO-AS1) in pancreatic carcinoma was investigated, and previous studies have not been reported. The upregulation of lncRNA TMPO-AS1 has been detected in lung adenocarcinoma and prostate cancer (7,8). Functionally, IncRNA TMPO-AS1 was revealed to promote the progression of non-small cell lung cancer by regulating its natural antisense transcript TMPO (9). In addition, lncRNA TMPO-AS1 promoted the occurrence of osteosarcoma by regulating the miR-199a-5p/WNT7B axis (10). However, the regulatory mechanism of lncRNA TMPO-AS1 remains unknown in pancreatic cancer. Therefore, the present study aimed to clarify the function of lncRNA TMPO-AS1 in pancreatic cancer.

As is well known, competitive endogenous RNA (ceRNA) competes with miRNA target genes by binding microRNA (miRNA). lncRNAs can affect the expression of these target genes and ultimately regulate tumorigenesis by acting as ceRNAs (11). In the present study, it was predicted that miR-383-5p has a binding site with lncRNA TMPO-AS1. miR-383 has been demonstrated to suppress the development of pancreatic carcinoma by inhibiting the expression of GAB1 (12). In addition, SRY-related high-mobility group box 11 (SOX11) was predicted to be a potential target of miR-383-5p. The important role of SOX11 in human diseases has been reported. It has been revealed that SOX11 is associated with poor clinicopathological characteristics of human prostate cancer (13). miR-145 has been reported to suppress

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the growth of endometrial cancer by targeting SOX11 (14). However, the relationship between miR-383-5p and SOX11 in pancreatic cancer has not been investigated.

Therefore, the present study clarified the regulatory mechanism of the lncRNA TMPO-AS1/miR-383-5p/SOX11 axis. Concurrently, the functions of lncRNA TMPO-AS1, miR-383-5p and SOX11 were also investigated in pancreatic cancer. This research will help us better understand the functional mechanism of lncRNA TMPO-AS1 in pancreatic carcinoma.

Materials and methods

Clinical tissues. Thirty-eight patients with pancreatic carcinoma (33-81 years old; 24 males and 14 females) diagnosed at **Oingdao Hospital of Traditional Chinese Medicine (Oingdao** Hiser Hospital; Qingdao, China) were involved in the present study from January 2017 and November 2019. The inclusion criteria comprised of patients who were diagnosed with pancreatic carcinoma and had not received chemotherapy or other treatments. The exclusion criteria included patients with pancreatic carcinoma who had infections, immune system diseases, chronic diseases or other types of cancer concurrently, such as biliary tract or pulmonary infection, rheumatoid arthritis, uremia and leukemia. All participants only received surgery and provided written informed consents. The study was approved by the Institutional Ethics Committee of Qingdao Hospital of Traditional Chinese Medicine (Qingdao Hiser Hospital). The pancreatic carcinoma tissues and adjacent non-tumor tissues were frozen in liquid nitrogen and stored at -80°C for further experiments.

Cells culture and transfection. Pancreatic duct epithelial cell line HPDE6-C7 and pancreatic cancer cells SW1990, PANC-1 (ATCC) were grown in DMEM (Gibco; Thermo Fisher Scientific, Inc.) containing 10% fetal bovine serum (FBS; Invitrogen Thermo Fisher Scientific, Inc.). These cells were incubated at 37°C in an atmosphere with 5% CO₂.

The miR-383-5p mimic (5'-AGAUCAGAAGGUGAUUGU GGCU-3') and control miRNA mimic (mimic-NC, 5'-ACU ACUGAGUGACAGUAGA-3'), miR-383-5p inhibitor (5'-AGC CACAAUCACCUUCUGAUCU-3') and control miRNA inhibitor (inhibitor-NC, 5'-UUCUCCGAACGUGUCACGUTT -3') were purchased from Shanghai GenePharma Co., Ltd. The TMPO-AS1 or SOX11 siRNA (si-TMPO-AS1, 5'-AGGTAG AAACGCAGTTTAA-3'; si-SOX11, 5'-GCTAGGTTAGAA TACATTTAA-3') and non-silencing siRNA (si-NC, 5'-CAG UACUUUUGUGUAGUACAAA-3') oligonucleotide were also purchased from Shanghai GenePharma Co., Ltd. The cDNA encoding TMPO-AS1 was sub-cloned into the vector pcDNA3.1 (the pcDNA-TMPO-AS1 primers were used as follows: Forward, 5'-GGGGTACCGGGTTGGTGCGAG CTTCC-3' and reverse, 5'-CCGCTCGAGCACTGTTCAAAT TTAACA-3'; Invitrogen; Thermo Fisher Scientific, Inc.). The empty pcDNA3.1 vector (pcDNA-NC) was used as a control. Lipofectamine 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) was used for the transfection into SW1990 cells for 48 h (37°C) following to the manufacturer's protocol. Following transfection, the cells were cultured in a humidified incubator at 37°C and 5% CO₂ for 48 h before the initiation of further experiments. Reverse transcription-quantitative (RT-q)PCR. Total RNA was extracted from pancreatic cancer tissues and cells using TRIzol reagent (Invitrogen; Thermo Fisher Scientific, Inc.). The cDNA was synthesized using PrimeScript RT reagent kit according to the manufacturer's protocol (Takara Biotechnology Co., Ltd.; reverse transcription primer: 5'-GTC GTATCCAGTGCGTGTCGTGGAGTCGGCAAT TGCACT GGATAC GAC AGC CAC-3'; protocol: 30 min at 16°C, 30 min at 42°C, and 5 min at 85°C). RT-qPCR was performed using SYBR-Green or TaqMan kit (TaKaRa, Dalian, China) and primers. PCR was performed as follows: 25 cycles of 10 min at 98°C, 10 sec at 98°C, 10 sec at 55°C and 20 sec at 72°C, with a final extension at 72°C for 5 min. The relative expression of TMPO-AS1, miR-383-5p, SOX11 were quantified using the $2^{-\Delta\Delta cq}$ method (15) and they were normalized to GAPDH or U6. The primers used were: TMPO-AS1 forward, 5'-AACCCA GCCCACACACTAC-3' and reverse, 5'-GAATATGAGTGC CTGCAGAC-3'; miR-383-5p forward 5'-GGGAGATCAGAA GGTGATTGTGGCT-3' and reverse, 5'-CAGTGCGTGTCG TGGAGT-3'; U6 forward, 5'-GCTTCGGCAGCACATATA CTAAAAT-3' and reverse, 5'-CGCTTCACGAATTTGCGT GTCAT-3'; SOX11 forward, 5'-GGTGGATAAGGATTTGGA TTCG-3' and reverse, 5'-GCTCCGGCGTGCAGTAGT-3'; GAPDH forward, 5'-AGAAGGCTGGGGGCTCATTTG-3' and reverse, 5'-AGGGGCCATCCACAGTCTTC-3'.

Western blot analysis. Cells were lysed with RIPA cell lysis buffer (Pierce; Thermo Fisher Scientific, Inc.). In brief, the BCA Protein Assay Kit (Pierce; Thermo Fisher Scientific, Inc.) was used to quantify the protein concentration. Equal amounts of protein samples (30 μ g) were loaded onto 10% SDS-PAGE gel and then transferred onto PVDF membranes. After blocking with 5% skim milk at room temperature for 1 h, the membranes were incubated with SOX11 (rabbit polyclonal; dilution, 1:1,000; cat. no. ab170916) and GAPDH (rabbit monoclonal, dilution, 1:1,000; cat. no. ab181602) primary antibodies overnight at 4°C and then with the HRP-conjugated secondary antibody (mouse anti-rabbit, dilution, 1:2,000; cat. no. 58802; New England BioLabs, Inc.) for 2-3 h at room temperature. Finally, the images of the protein bands were captured by a Tanon detection system using ECL reagent (Thermo Fisher Scientific, Inc.).

MTT assay. Cell proliferation was measured by MTT assay. Briefly, the transfected SW1990 cells $(3x10^3 \text{ cells/well})$ were seeded into 96-well plates. These cells were incubated at 37°C in DMEM medium for 24, 48, 72 or 96 h, respectively. Then, the cells were incubated at 37°C for 4 h with fresh medium containing 10% MTT. Finally, DMEM culture medium was discarded and 200 μ l dimethyl sulfoxide was then added to each well to completely dissolve the crystals. The absorbance at 490 nm was detected by a spectrophotometer (Bio-Rad Laboratories, Inc.).

Transwell assay. Transwell chambers (8 μ m pore size; Corning, Inc.) were used to assess migration and invasion. SW1990 transfected cells (2x10⁴) were seeded in the upper chamber and DMEM medium with 10% fetal bovine serum was added in the lower chamber. Next, these cells were incubated at 37°C with 5% CO₂ for 18 h. Then the cells on the lower surface were fixed with 4% paraformaldehyde for 20 min at room



Figure 1. IncRNA TMPO-AS1 aggravates carcinogenic behaviors of pancreatic carcinoma. (A) IncRNA TMPO-AS1 expression in pancreatic carcinoma tissues and normal tissues (n=38). (B) IncRNA TMPO-AS1 expression in SW1990, PANC-1 and HPDE6-C7 cells (C) IncRNA TMPO-AS1 expression in SW1990 cells with its siRNA. (D-F) Cell proliferation, migration and invasion in SW1990 cells with TMPO-AS1 siRNA. *P<0.05, **P <0.01. TMPO-AS1, TMPO antisense RNA 1; si-, small interfering; NC, negative control.

temperature and stained with 0.1% crystal violet overnight at 4°C after the cells that did not migrate or invade were removed by a cotton swab. For the invasion assay, the Transwell chambers were precoated with Matrigel (BD Biosciences). Images were captured under an inverted microscope (magnification, x200; Olympus Corporation).

Bioinformatics analysis. To identify the putative miRNA target, we utilized online miRNA target analysis tools TargetScan (http://www.targetscan.org/) and starBase version 2.0 (http://starbase.sysu.edu.cn/n) to perform the prediction. The list of potential targets and binding sites were available by searching the database.

Dual luciferase reporter assay. SW1990 cells were seeded in 24-well plates and incubated for 24 h prior to transfection.

The 3'-UTR of wild-type or mutant TMPO-AS1 or SOX11 was inserted into the pmiR-GLO vector (Promega Corporation). The vector was then transfected into SW1990 cells with miR-383-5p mimics or miR-NC using Lipofectamine 2000 (Invitrogen; Thermo Fisher Scientific, Inc.). After 48 h, the luciferase activity was measured by the dual-luciferase reporter gene assay system (Promega Corporation). The firefly luciferase activities were normalized to *Renilla* luciferase activity.

Xenograft assays. All animal experiments were approved by the Ethics Committee of Animal Experiments of Qingdao Hospital of Traditional Chinese Medicine (Qingdao Hiser Hospital; Qingdao, China) (approval no. ANI2019-56; approval date: March 6, 2019). SW1990 cells were transfected with sh-TMPO-AS1 or empty vector (50 ng/well; Shanghai



Figure 2. IncRNA TMPO-AS1 and miR-383-5p expression levels are reciprocally inhibited in pancreatic carcinoma. (A) The binding site between lncRNA TMPO-AS1 and miR-383-5p. (B) Luciferase reporter assay. (C) TMPO-AS1 expression was detected in SW1990 cells with TMPO-AS1 siRNA or vector compared with si-NC or pcDNA-NC. (D) miR-383-5p expression was detected in SW1990 cells with TMPO-AS1 siRNA or vector compared with si-NC or pcDNA-NC. (E) miR-383-5p expression was detected in SW1990 cells with miR-383-5p mimics or inhibitor compared with mimic-NC or inhibitor-NC. (F) lncRNA TMPO-AS1 expression was assessed in SW1990 cells containing miR-383-5p mimics or inhibitor compared with mimic-NC or inhibitor-NC. (**P<0.01. TMPO-AS1, TMPO antisense RNA 1; miR-383-5p, microRNA-383-5p; wt, wild-type; mut, mutant; si-, small interfering; NC, negative control.

GenePharma Co., Ltd.) using Lipofectamine 2000 (Invitrogen; Thermo Fisher Scientific, Inc.). Next, transfected SW1990 cells (2x10⁶ cells in 100 μ l DMEM without FBS) were injected subcutaneously into the upper right flank of four female nude mice (4-week-old Balb/c/nu mice; 30 g; Changzhou Kaiwen Laboratory Animal Co., Ltd.). The mice were maintained under specific pathogen-free conditions and housed in ventilated cages with free access to food and water. Tumor growth was examined every 3 days, and the tumor volume (V) was determined by measuring the length (L) and width (W) of the tumor with a caliper and calculated using the formula $V=(L \times W^2) \times 0.5$. Subsequently, 18 days later, the tumors were removed and weighed. All animals were euthanized by intraperitoneal injection of 1% sodium pentobarbital (150 mg/kg) anesthetic. Animal death criteria included cardiac arrest, continued absence of spontaneous breathing for 2-3 min, extreme pupil dilation and no blink reflex in animals.

Statistical analysis. Data are expressed as the mean \pm standard deviation (SD) and analyzed by SPSS 19.0 (IBM Corp.) or Graphpad Prism 6 (Graphpad Software, Inc.). Paired Student's t-test was used for comparisons between tumor and adjacent

non-tumor samples of the same individuals, while unpaired Student's t-test was used for other comparisons between two groups. One-way ANOVA followed by Bonferroni's multiple comparisons post hoc test was used to calculate the difference between multiple groups. P<0.05 was considered to indicate a statistically significant difference.

Results

lncRNA TMPO-AS1 aggravates carcinogenic behaviors of pancreatic carcinoma. Compared with normal tissues, TMPO-AS1 expression was upregulated in pancreatic carcinoma tissues (Fig. 1A). Similarly, the expression of lncRNA TMPO-AS1 in SW1990 and PANC-1 pancreatic cancer cells was higher than that in HPDE6-C7 cells (Fig. 1B). Based on the aforementioned results, SW1990 cells were used to investigate the function of TMPO-AS1 in pancreatic carcinoma due to the significant difference of TMPO-AS1 expression. Next, TMPO-AS1 siRNA was transfected into SW1990 cells. The transfection efficiency was detected by RT-qPCR (Fig. 1C). An MTT assay revealed that downregulation of TMPO-AS1 inhibited cell proliferation in SW1990 cells (Fig. 1D). A Transwell



SW1990

Figure 3. miR-383-5p restrains pancreatic carcinoma cell proliferation and motility by interacting with lncRNA TMPO-AS1. (A) miR-383-5p expression in pancreatic carcinoma tissues and normal tissues (n=38). (B) miR-383-5p was negatively correlated with lncRNA TMPO-AS1 expression in pancreatic carcinoma tissues. (C) miR-383-5p expression in SW1990 cells with miR-383-5p mimics and TMPO-AS1 vector compared with mimic-NC or miR-383-5p mimics. (D-F) Cell proliferation, migration and invasion in SW1990 cells with miR-383-5p mimics and TMPO-AS1 vector compared with mimic-NC or miR-383-5p mimics. **P<0.01. TMPO-AS1, TMPO antisense RNA 1; miR-383-5p, microRNA-383-5p; NC, negative control.

assay revealed that knockdown of TMPO-AS1 restrained cell migration and invasion in SW1990 cells (Fig. 1E and F). The aforementioned results indicated that the upregulation of IncRNA TMPO-AS1 promoted cell proliferation and motility in pancreatic carcinoma.

IncRNA TMPO-AS1 and miR-383-5p expression levels are reciprocally inhibited in pancreatic carcinoma. The starBase version 2.0 (http://starbase.sysu.edu.cn/n) predicted that lncRNA TMPO-AS1 has a binding site with miR-383-5p (Fig. 2A). A dual luciferase reporter assay revealed that miR-383-5p mimics reduced the luciferase activity of wt-TMPO-AS1 and did not affect the luciferase activity of mut-TMPO-AS1 in SW1990 cells (Fig. 2B). This indicated that lncRNA TMPO-AS1 directly targeted miR-383-5p. Subsequently, the transfection efficiency of TMPO-AS1



Figure 4. miR-383-5p directly targets SOX11. (A) The binding sites between miR-383-5p and SOX11. (B) Luciferase reporter assay. (C) SOX11 expression was detected in SW1990 cells with miR-383-5p mimics or inhibitor compared with mimic-NC or inhibitor-NC. (D) SOX11 expression in pancreatic carcinoma tissues and normal tissues (n=38). (E) miR-383-5p was negatively correlated with SOX11 expression in pancreatic carcinoma tissues. (F) lncRNA TMPO-AS1 expression was positively correlated with SOX11 in pancreatic carcinoma tissues. **P<0.01. miR-383-5p, microRNA-383-5p; SOX11, SRY-related high-mobility group box 11; wt, wild-type; mut, mutant; NC, negative control.

vector and siRNA was detected in SW1990 cells by RT-qPCR (Fig. 2C). In addition, it was revealed that the expression of miR-383-5p in SW1990 cells was downregulated by the TMPO-AS1 vector and upregulated by TMPO-AS1 siRNA (Fig. 2D). Next, the transfection efficiency of miR-383-5p mimics or inhibitor was detected in SW1990 cells by RT-qPCR (Fig. 2E). Concurrently, miR-383-5p mimics in SW1990 cells also reduced TMPO-AS1 expression, while miR-383-5p inhibitor increased the expression of TMPO-AS1 in SW1990 cells (Fig. 2F). Collectively, the expression levels of lncRNA TMPO-AS1 and miR-383-5p exhibited mutual inhibition in pancreatic carcinoma.

miR-383-5p restrains pancreatic carcinoma cell proliferation and motility by interacting with lncRNA TMPO-AS1. In addition, downregulation of miR-383-5p was revealed in pancreatic carcinoma tissues compared with normal tissues (Fig. 3A). A negative correlation between miR-383-5p and TMPO-AS1 expression was revealed in pancreatic carcinoma tissues (Fig. 3B). Then, the miR-383-5p mimics and lncRNA TMPO-AS1 vector were transfected into SW1990 cells to explore the role of miR-383-5p and its interaction with lncRNA TMPO-AS1. Functionally, upregulation of TMPO-AS1 attenuated the inhibitory effect of miR-383-5p on cell proliferation in SW1990 cells (Fig. 3C). The results also revealed that the increased expression of miR-383-5p induced by its mimics was reduced by the lncRNA TMPO-AS1 vector (Fig. 3D). The TMPO-AS1 vector also reversed the inhibition of cell migration and invasion induced by miR-383-5p overexpression in SW1990 cells (Fig. 3E and F). These results indicated that overexpression of miR-383-5p inhibited cell proliferation and motility in pancreatic carcinoma. Moreover, the upregulation of TMPO-AS1 abolished the antitumor effects of miR-383-5p in pancreatic carcinoma.

miR-383-5p directly targets SOX11. Next, potential targets of miR-383-5p were predicted in the TargetScan database (http://www.targetscan.org). It was revealed that SOX11 has a binding site with miR-383-5p (Fig. 4A). It was also revealed that miR-383-5p mimics reduced the luciferase activity of wt-SOX11, indicating that SOX11 is a direct target of miR-383-5p (Fig. 4B). RT-qPCR and western blotting revealed



Figure 5. SOX11 is involved in the progression of pancreatic carcinoma. (A) SOX11 expression in SW1990 cells with SOX11 siRNA. (B-D) Cell proliferation, migration and invasion in SW1990 cells with SOX11 siRNA. **P<0.01. SOX11, SRY-related high-mobility group box 11; si-, small interfering; NC, negative control.

that miR-383-5p mimics inhibited SOX11 expression, while miR-383-5p inhibitors promoted SOX11 expression in SW1990 cells (Fig. 4C). In order to further confirm the relationship between SOX11 and miR-383-5p or TMPO-AS1, the expression level of SOX11 in pancreatic carcinoma tissues was first detected. Compared with normal tissues, SOX11 was revealed to be upregulated in pancreatic carcinoma tissues (Fig. 4D). Notably, miR-383-5p was revealed to negatively regulate SOX11 expression in pancreatic carcinoma tissues (Fig. 4E). Conversely, SOX11 expression was positively correlated with TMPO-AS1 expression in pancreatic carcinoma tissues (Fig. 4F). Therefore, SOX11 was revealed to be a direct target of miR-383-5p and TMPO-AS1 could positively regulate the expression of SOX11 in pancreatic carcinoma.

SOX11 is involved in the progression of pancreatic carcinoma. To explore the role of SOX11 in pancreatic carcinoma, SOX11 siRNA was transfected into SW1990 cells. RT-qPCR and western blotting revealed that SOX11 siRNA reduced its expression in SW1990 cells (Fig. 5A). In particular, it was determined that SOX11 downregulation inhibited cell proliferation (Fig. 5B). In addition, knockdown of SOX11 also suppressed cell migration and invasion in SW1990 cells (Fig. 5C and D). Collectively, SOX11 was revealed to play a carcinogenic role in pancreatic carcinoma by promoting cell proliferation, migration and invasion.

lncRNA TMPO-AS1 regulates pancreatic carcinoma tumor growth in vivo. In order to verify whether TMPO-AS1 expression regulates pancreatic carcinoma tumorigenesis *in vivo*, xenograft analysis was performed. Consistent with previous *in vitro* experiments, tumor growth in the pcDNA-TMPO-AS1 group was significantly slower than the pcDNA-NC group (Fig. 6A and B). In addition, it was also revealed that lncRNA TMPO-AS1 expression in the mice tissues of the pcDNA-TMPO-AS1 group was higher than that in the pcDNA-NC group (Fig. 6C). After nude mice were sacrificed and tumors were removed, and the tumor weight in the pcDNA-TMPO-AS1 group was revealed to be lower than that in the pcDNA-NC group (Fig. 6D). These



Figure 6. Effect of downregulated TMPO-AS1 on tumorigenesis in vivo. (A) Tumors from nude mice after injection of SW1990 cells with pcDNA-TMPO-AS1 or pcDNA-NC. (B) The tumor volume was measured every 3 days after injection. (C) The expression of TMPO-AS1 in mice tissues in the pcDNA-TMPO-AS1 group and pcDNA-NC group. (D) After 18 days of injection, tumors were removed from the nude mice and tumor weights were measured. **P<0.01. TMPO-AS1, TMPO antisense RNA 1; NC, negative control.

data demonstrated that lncRNA TMPO-AS1 downregulation significantly inhibited the growth of pancreatic carcinoma tumor xenograft.

Discussion

Recently, it has been discovered that an increasing number of lncRNAs play important roles in pancreatic carcinoma, including MACC1-AS1 and PVT1 (16,17). In particular, the upregulation of lncRNA PANDAR has been identified in pancreatic carcinoma and increased expression of lncRNA PANDAR promoted cell proliferation and inhibited cell apoptosis in pancreatic carcinoma (18). In the present study, lncRNA TMPO-AS1 was also upregulated in pancreatic cancer tissues and cells. The upregulation of TMPO-AS1 promoted cell viability, migration and invasion in pancreatic carcinoma. Notably, lncRNA TMPO-AS1 promoted the development of pancreatic carcinoma by regulating the miR-383-5p/SOX11 axis. These results indicated that lncRNA TMPO-AS1 acts as a tumor promoter in pancreatic carcinoma.

Consistent with our results, high expression of TMPO-AS1 was also detected in colorectal cancer and lung adenocarcinoma (19,20). In addition, TMPO-AS1 was revealed to promote the progression of cervical cancer by sponging miR-577 and upregulating RAB14 (21). In the present study, miR-383-5p was confirmed to act as a sponge for lncRNA TMPO-AS1. The overexpression of miR-383-5p inhibited the progression of pancreatic carcinoma. It has been reported that miR-383 suppressed the development of pancreatic carcinoma (12), similar to our results. In addition, it was revealed that lncRNA TMPO-AS1 functioned as an oncogene in pancreatic carcinoma by sponging miR-383-5p. Similarly,

LINC01128 accelerated the progression of cervical cancer by regulating the miR-383-5p/SFN axis (22). In the present study, lncRNA TMPO-AS1 was revealed to promote the progression of pancreatic carcinoma by mediating the miR-383-5p/SOX11 axis. Furthermore, SOX11 was confirmed to be a direct target of miR-383-5p.

It has been demonstrated that SOX11 is overexpressed in endometrial cancer and mantle cell lymphoma (23,24). In the present study, SOX11 was also upregulated in pancreatic carcinoma. In addition, knockdown of SOX11 inhibited cell proliferation, migration and invasion in pancreatic carcinoma. Similar to our study, it has been reported that the downregulation of SOX11 hindered the proliferation, migration, and invasion of thyroid tumor cells (25). It was revealed that miR-383-5p restrained the progression of pancreatic carcinoma by targeting SOX11. miR-145 also targeted SOX11 to suppress the growth of endometrial cancer (14). In addition, circular RNA CEP128 was proposed as a sponge of miR-145-5p, which promoted the progression of bladder cancer by regulating SOX11 (26). In the present research, IncRNA TMPO-AS1 was revealed to promote the development of pancreatic carcinoma by sponging miR-383-5p and upregulating SOX11. However, there is a limitation in the present study. Our results were only investigated in SW1990 cells, and the results should be verified in an additional cell line in the future.

In conclusion, the present study demonstrated that lncRNA TMPO-AS1 has carcinogenic effects in pancreatic carcinoma. In particular, upregulation of lncRNA TMPO-AS1 promoted cell proliferation, migration and invasion in pancreatic carcinoma by downregulating miR-383-5p and upregulating SOX11. This research may provide a new direction for the treatment of pancreatic carcinoma.

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Availability of data and materials

The datasets used during the present study are available from the corresponding author upon reasonable request.

Authors' contributions

FX and TZ designed the experiments. XS and SZ conducted the experiments. MN, and YC performed the animal experiments and prepared the figures. YW analyzed the data. FX and TZ wrote and edited the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The study was approved by the Institutional Ethics Committee of Qingdao Hospital of Traditional Chinese Medicine (Qingdao Hiser Hospital; Qingdao, China) (approval no. HUN2017-12; approval date: January 5, 2017). All participants provided written informed consents. All animal experiments were approved by the Ethics Committee of Animal Experiments of Qingdao Hospital of Traditional Chinese Medicine (Qingdao Hiser Hospital; Qingdao, China) (approval no. ANI2019-56; approval date: March 6, 2019).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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