

Elevated expression of exosomal microRNA-21 as a potential biomarker for the early diagnosis of pancreatic cancer using a tethered cationic lipoplex nanoparticle biochip

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Abstract. Pancreatic cancer (PC) has a poor prognosis due to the lack of effective molecular biomarkers for early diagnosis. Recent studies have investigated the use of exosomal microRNAs (exmiRs) as diagnostic biomarkers in cancer. The present study examined exmiR-21, exmiR-10b and exmiR-212-3p expression in patients with PC and healthy individuals. The expression levels of exmiR-21, exmiR-10b and exmiR-212-3p were examined in the peripheral blood plasma of 36 patients with PC and 65 healthy controls, using tethered cationic lipoplex nanoparticle biochip. The levels of exmiR-21 in the plasma of 34 mice were also evaluated. The expression levels of exmiR-21 and exmiR-10b were significantly greater in patients with PC compared with the control group. The receiver operating characteristic (ROC) analysis indicated that exmiR-21 had better diagnostic performance ($P=0.0003$; AUC, 0.7171) compared with the other two exmiRs. The diagnostic value of exmiR-21 improved when combined with exmiR-10b ($P<0.0001$; AUC, 0.791). Furthermore, exmiR-21 was capable of distinguishing patients with early-stage PC from controls and advanced-stage PC ($P<0.05$, early stage vs. healthy; $P<0.001$, early stage vs. advanced stage). The results of the present study revealed that the plasma levels of exmiR-21 and exmiR-10b were upregulated in patients with PC. The ROC analyses indicated that exmiR-21 had the best diagnostic performance among the three exmiRs. Furthermore, exmiR-21 was capable of discriminating patients with early-stage PC from healthy controls. These findings indicate the potential of determining the expression of exmiR-21 from serum using a tethered

cationic lipoplex nanoparticle biochip as a novel non-invasive strategy for the early diagnosis of PC.

Introduction

Pancreatic cancer (PC) is one of the most devastating types of cancer worldwide. As the third leading cause of cancer-associated mortality in the USA, PC was estimated to cause 44,330 deaths in this country in 2018 (1). The number of PC cases is projected to increase, and to become the second leading cause of cancer-associated mortality before 2030 (2).

Due to the anatomical location of PC, it is usually asymptomatic in the early stages of disease. However, without the use of invasive procedures, current screening methods are unable to detect early-stage PC (3), and these procedures are difficult to perform at the early stage of disease. Thus, the development of non-invasive detection methods, such as liquid biopsy, has been increasingly emphasized in recent years, and the identification of new biomarkers that can distinguish early stage PC is urgently needed (4).

MicroRNAs (miRs) play critical roles in a number of biological processes, such as cell proliferation, cell differentiation and apoptosis, through their regulation of gene expression by translational repression or mRNA degradation, resulting in gene silencing. Recent studies have indicated that miRs may also function as potential biomarkers for cancer detection (5-8). Exosomes are a type of extracellular vesicle, which are enclosed by a lipid bilayer that can contain various components, including proteins, lipids, DNA, mRNAs, miRs and non-coding RNAs (9). A number of studies have demonstrated that miRs contained within exosomes (exmiRs) play an important role in the development, metastasis and drug resistance of tumors (10-12). The lipid bilayer structure of exosomes prevents the enclosed miRs from degradation, allowing the detection of accurate numbers of stable exmiRs. Thus, isolating exosomes and quantifying the encapsulated exmiRs has become a promising approach for the noninvasive detection of cancer biomarkers (13).

miRs can be quantitatively measured in serum by reverse transcription-quantitative (RT-q)PCR. However, this approach is expensive and time-consuming (14). The tethered cationic lipoplex nanoparticle (TCLN) biochip is a novel technology

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for the direct analysis of exmiRs within a single exosome. Molecular beacons in liposome nanoparticles serve as molecular probes for target miRs. When serum is applied to the biochip, the molecular beacons recognize target miRs in exosomes, and the interaction is read by a total internal reflection fluorescence microscope (15). The copy numbers of the exmiRs can then be obtained by the fluorescence signals of the molecular beacons (MBs) (15). Wu *et al* (13) demonstrated that the levels of exmiRs detected by the TCLN biochip were consistent with the widely used reverse transcription-quantitative PCR (RT-qPCR) method. In addition, the TCLN biochip indicated a higher sensitivity compared with RT-qPCR (13).

The aim of the present study was to evaluate the expression levels of circulating exmiR-21, exmiR-10b and exmiR-212-3p between patients with PC and healthy individuals using a TCLN biochip and to determine whether these exmiRs can distinguish early-stage PC.

Materials and methods

Patients and clinical samples. A total of 36 patients with PC and 65 healthy individuals were enrolled for the present study between November 2017 and December 2018. The 36 patients with PC were divided into two groups, early- and advanced-stage; patients with stages I and II disease were categorized as the early stage group (n=19), whereas patients with stages III and IV disease were categorized in the advanced-stage group (n=17). All participants were from the Sir Run Run Shaw hospital of Zhejiang University (Zhejiang, China). The detailed characteristics of the patients are provided in Table I. All patients were newly diagnosed with PC, which was confirmed by two blinded pathologists from the Sir Run Run Shaw hospital, following surgery or ultrasound endoscopic guided fine needle aspiration biopsy. The blood samples of patients treated with chemotherapy, radiotherapy or immunotherapy were excluded. Tumors were staged according to the 8th American Joint Committee on Cancer (AJCC) TNM Staging of Pancreatic Cancer (16).

The blood samples included in the present study were collected in vacuum blood tubes with EDTA anticoagulant before surgery and pharmacotherapy, and handled within 1 h following collection. The blood samples were subjected to centrifugation at 2,795 x g for 10 min at 4°C. The plasma was then stored at -80°C. Written informed consent was obtained from the participants before sampling. The studies were conducted in accordance with the International Ethical Guidelines for Biomedical Research Involving Human Subjects (CIOMS), and the research protocols were approved by the Clinical Research Ethics Committee of Sir Run Run Shaw Hospital of Zhejiang University (Zhejiang, China)

Animal experiments. A total of 34 mice were obtained from the animal unit of Zhejiang University (Zhejiang, China). BALB/C nude mice (4-6 weeks old) were used in all experiments. All mice were female, and their weight ranged from 20-26 g. The housing temperature was from 22-24°C, and mice were given access to food and water. The air remained clean by using ventilation devices for ~8-15 times per hour, and the indoor lighting was controlled for 12 h of light and dark. All animal experiments were performed in the animal unit of

Zhejiang University according to procedures authorized and specifically approved by the institutional Ethical Committee. Subcutaneous xenograft mouse model were established in 12 mice by injecting 1x10⁶ PANC-1-Luciferase cells (Stem Cell Bank, Chinese Academy of Science) in the shoulders, via a left subcostal incision, made in mice that were anesthetized with 3% isoflurane. The other 22 normal mice were used as the normal control and were injected with the same volume of normal saline. Subsequently, the blood of the 34 mice was collected, when the subcutaneous tumors in the 12 xenograft mice grew to a longitudinal diameter of 1 cm. Blood was collected from the venous plexus of the eye socket, in mice anesthetized with 3% isoflurane, and the plasma was then separated by centrifugation at 2,795 x g for 10 min at 4°C. The plasma was then stored at -80°C. All mice were sacrificed by cervical dislocation under 3% isoflurane anesthesia, following the collection of blood. The plasma samples of these mice were used for further analysis.

Rationale for selecting exmiRs. As an oncomiR, a number of studies have demonstrated that miR-21 plays an important role in cell proliferation, migration, invasion and survival by regulating the cell cycle, apoptosis and invasion-related genes (17-21). Previous studies have proven that exosomal miR-21 is upregulated at the early stage of lung cancer, breast cancer, liver cancer and other malignant tumors (22-24).

Thus, exmiR-21 was considered as a potential biomarker for the diagnosis of cancer in the present study. However, to the best of our knowledge, few studies have focused on the association between exmiR-21 and PC, particularly at the early stage of PC. miR-10b was associated with the invasive and metastatic properties of various human cancer types. Kim *et al* (25) identified that miR-10b promotes breast cancer cell proliferation, migration and invasion via the inhibition of transcription factor TBX5 expression, leading to the repression of the tumor suppressor genes (dual specificity tyrosine phosphorylation regulated kinase 1A and PTEN). Previous studies regarded miR-10b as a potential diagnostic biomarker of breast cancer, bladder cancer and lung cancer (26). Ayaz *et al* (27) reported that miR-212-3p was specifically expressed in laryngeal squamous cell carcinoma. Furthermore, a study revealed that miR-212-3p was upregulated in PC tissue. However, to the best of our knowledge, few studies have focused on exmiRs in the aforementioned cancer types. Given that, exmiRs (particularly tumor-derived exmiRs) play an important role in the early diagnosis of several types of cancer in humans.

Preparing cationic lipoplex nanoparticles containing MBs. Firstly, miR-21-, miR-10b- and miR-212-3p-specific probes with fluorescent signal markers were designed. BHQ1 and 6-carboxy-fluorescein were assembled at the 3' and 5' ends of the probe, respectively. The center of the probe is a specially designed single-stranded base sequence for identifying the target RNA. The specific molecular probe is coated in the cationic liposome nanoparticles and linked to the glass substrate to construct a PC-associated miRNA liposome nanoparticle chip. The probe sequences were as follows: MiR-21-MB, 5'-6FAM-TCAACATCAGTCTGATAAGCT ATTATCAGACTGA-BHQ1-3'; miR-10b-MB, 5'-6FAM-A TACCACACAAATTCGGTTCTACAACCGAATTTG

Table I. Characteristics of patients with pancreatic cancer.

Characteristics	Cases, n (%)
Sex	
Female	15 (41.7)
Male	21 (58.3)
Age (years)	
<60	8 (22.2)
60-69	19 (52.8)
≥70	9 (25.0)
Cigarette smoking	
No	28 (77.8)
Yes	8 (22.2)
Alcohol drinking	
No	7 (19.4)
Yes	29 (80.6)
Diabetes mellitus	
No	33 (91.7)
Yes	3 (8.3)
ABO blood type	
A	13 (36.1)
B	8 (22.2)
O	12 (33.3)
AB	3 (8.3)
Stage	
I	6 (16.7)
II	13 (36.1)
III	5 (13.9)
IV	12 (33.3)
BMI	
<19.57	8 (22.2)
19.57-23.71	19 (52.8)
>23.71	9 (25.0)

Continuous variables are converted into categorical variables. All variables are displayed as numbers and percentages. BMI, body mass index.

TG-BHQ1-3'; miR-212-3p-MB, 5'-6FAM-CGGCCGTGAC TGGAGACTGTTAAGTCTCCAGTCA-BHQ1-3'.

Exosome characterization and quantification. A commercial exosome extraction kit (Total Exosome Isolation kit; cat. no. 4484450; Invitrogen; Thermo Fisher Scientific, Inc.) was used to extract mouse serum and human plasma exosomes to detect their particle size and potential. The dynamic light scattering (DLS) technology was used for detection, with the use of a 640 nm laser and 30 frames per sec, which is also known as Photon Correlation Spectroscopy (PCS) or quasi-elastic scattering, is a method for measuring the variation of light intensity fluctuations over time (28). The Zetasizer Nano ZS90 instrument (Malvern Instruments, Inc.) was used for particle size analysis. The DLS technology has the advantages of accuracy, speediness and good repeatability, and has

become a more conventional characterization method in nanotechnology (29). DLS is widely used, including for particle science testing in the life science field; it is mature, stable and credible. As a gold standard, it is recognized by the US Bureau of Standards and incorporated into the US measurement standard system. The calculation method of DLS should be able to reflect the particle size distribution within the system better.

Western blotting. Cells or exosomes were lysed with RIPA buffer (Beyotime Institute of Biotechnology), supplemented with protease and phosphatase inhibitor cocktail (Beyotime Institute of Biotechnology) on ice for 15 min. The lysates were centrifuged at 12,000 x g for 5 min to pellet cell debris at 4°C. The protein concentrations were measured by the BCA protein assay kit (Pierce; Thermo Fisher Scientific, Inc.), according to the manufacturer's instructions. Subsequently, the supernatant was added to SDS/PAGE sample loading buffer (Beyotime Institute of Biotechnology) and boiled for 10 min at 95°C. A total mass of 20 µg protein was run on SDS-PAGE (10%) and transferred to polyvinylidene fluoride membrane (Bio-Rad Laboratories, Inc.). The membranes were blocked for 60 min in 5% (w/v) non-fat dry milk or bovine serum albumin for 1 h at room temperature and incubated with primary antibodies overnight at 4°C. The next day, the membranes were washed thrice in tris-buffered saline containing 0.1% (v/v) Tween-20 (TBST), incubated with horseradish-peroxidase conjugated secondary antibodies (Thermo Fisher Scientific, Inc.) for 1 h at room temperature, and then washed thrice in TBST. Bands were visualized using ChemiDoc Touch Imaging System (Bio-Rad Laboratories, Inc.) and the band intensities were quantified using the Image Lab 5.2.1 software (Bio-Rad Laboratories, Inc.). The primary antibodies were: Anti-CD9 (1:2,000; cat. no. ab92726; Abcam), anti-CD63 (1:1,000; cat. no. ab134045; Abcam) and anti-heat shock protein 70 (HSP70; 1:1,000; cat. no. ab181606; Abcam). The secondary antibody was: Goat anti-rabbit IgG (H+L) secondary antibody, HRP (1:10,000; cat. no. 31460; Invitrogen; Thermo Fisher Scientific, Inc.)

Transmission electron microscopy (TEM). To evaluate the morphology of isolated exosomes, the exosomes were observed under a transmission electron microscope. Firstly, carbon-coated 400-mesh copper grids (Xinxing Bairui Co. Ltd.) were absorbed in a 20-µl aliquot of exosomes (~2 µg) for 2 min and allowed to dry at room temperature. Subsequently, the exosomes were fixed in 1% (v/v) glutaraldehyde (EM-grade) for 5 min at room temperature by placing drops carefully on the dried preparation. Next, the grids were washed twice with water for 5 min and then contrast-stained with 2% uranyl acetate for 10 min at room temperature. Images were obtained by transmission electron microscopy (Tecnaï T10; FEI).

Statistical analysis. Statistical analyses were performed and graphs were constructed using GraphPad Prism (version 7.0; GraphPad Software, Inc.) and SPSS (version 24.0; IBM Corp.) statistical software. The data are presented as the mean ± standard deviation. Significant differences between the mean values of two groups were determined using the Student's t-test. Significant differences between the mean values of three or four groups were determined using the one-way

ANOVA, followed by Student-Newman-Keuls for further multiple comparisons. Clinicopathological diagnoses were used as the gold standard to assess diagnostic accuracy, using the receiver operating characteristic (ROC) curves generated with the GraphPad Prism 7.0 software. All diagnostic metrics (sensitivity and specificity) were calculated using the standard formulae (30). $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Exosome characterization. DLS was used to detect the size of particles. Based on the DLS analysis, purified particles had a size of 53.5 nm in human plasma samples and 139.9 nm in mouse plasma samples, which was in accordance with the described view that exosomes are spherical particles with sizes ranging between 30 and 150 nm (31). The ζ potential values of plasma exosomes from both humans and mice were negative, -5.80 and -1.49 mV (Fig. 1A and D), respectively. CD9, CD63 and HSP70 are considered as exosomal-specific markers; the presence of these markers in the purified particles was assessed to characterize the purified particles as exosomes. Western blot analysis demonstrated that the purified particles expressed these markers, and thus could be considered as exosomes (Fig. 1B and E). Additionally, electron micrographs showed that the collected products had a distinctive cup shape (Fig. 1C and F). Overall, DLS, western blot analyses and electron micrographs confirmed that the particles tested were exosomes. Furthermore, as shown in Fig. 1G, the fluorescence intensity distributions indicated that more exosomes with higher exmiR-21, exmiR-10b and exmiR-212-3p expression were present in the samples of patients compared with those from healthy controls. Furthermore, murine PC exosomes revealed much higher fluorescence signals compared with plasma exosomes from healthy mouse (Fig. 1H).

Differential exmiR expression in patients with PC and in healthy controls. The expression levels of exmiR-21, exmiR-10b and exmiR-212-3p were examined in the plasma of 36 patients with PC and 65 healthy individuals. The results showed that the expression levels of exmiR-21 and exmiR-10b were significantly increased in the plasma of patients with PC compared with the healthy control group (Fig. 2A and B). However, no significant difference ($P = 0.0698$) was detected in the expression of exmiR-212-3p between the two groups (Fig. 2C).

ROC curve analysis. In order to evaluate the diagnostic performance of the three exmiRs, the ROC curve analysis was performed. The results revealed that exmiR-21 and exmiR-10b, but not exmiR-212-3p, could differentiate patients with PC from healthy individuals (Fig. 2D and E). The exmiR with the greatest area under the curve (AUC) was exmiR-21 ($P = 0.0003$; AUC, 0.7171); the AUC of exmiR-10b was 0.6543 ($P = 0.0105$). The ROC analysis indicated that the diagnostic accuracy of exmiR-21 was superior to the other miRs. ExmiR-21 and exmiR-10b were also combined in further ROC analysis, which indicated that exmiR-10b could improve the diagnostic performance of exmiR-21 ($P < 0.0001$; AUC, 0.791; Fig. 2).

ExmiR-21 was significantly upregulated in mice with PC and is a sensitive diagnostic marker. The expression of exmiR-21 was also examined using the TCLN biochip in the plasma of 12 mice with PC and 22 healthy mice. The level of exmiR-21 in the plasma of mice with PC was significantly higher ($P = 0.0001$) compared with the healthy controls (Fig. 3A). The AUC of exmiR-21 in mice was 0.8636 ($P = 0.0005$; Fig. 3B).

ExmiR-21 was significantly higher in patients with early-stage PC compared with healthy controls. The 36 patients with PC were stratified to examine the associations between the plasma levels of the three exmiRs and the PC stage, based on TNM staging. Detecting PC at an early stage (stages I or II) (32) may provide a window of opportunity when the disease can be adequately treated and stopped (33). The levels of exmiR-21 were significantly different in the early-stage group compared with the healthy control and patients with advanced-stage PC (Fig. 2F). In contrast, the expression levels of exmiR-10b and exmiR-212-3p showed no significant differences between the early-stage and the control groups (Fig. 2G and H).

Evaluating levels of the selected exmiRs with clinical parameters of patients with PC. Additionally, various clinical characteristics of the 36 patients with PC, such as age, sex, history of smoking or drinking, diabetes mellitus, blood type, cancer stage and body mass index were analyzed. However, the levels of the three exmiRs showed no significant differences according to these characteristics (Fig. 4).

Discussion

One of the most important causes of the high mortality of patients with PC is the lack of robust and sensitive biomarkers for early diagnosis. The majority of patients with PC are diagnosed at a stage too late for surgical treatment (32). Thus, the development of new diagnostic methods that can detect early-stage PC is necessary to improve the survival of patients with PC.

The present study demonstrated that exmiR-21 and exmiR-10b were expressed at significantly higher levels in patients with PC compared with the healthy controls. Among the three exmiRs, exmiR-21 demonstrated the optimal diagnostic performance and, when combined with exmiR-10b, the diagnostic performance of exmiR-21 increased. The plasma of 34 mice with PC were also examined, which demonstrated elevated levels of exmiR-21 expression compared with controls. Furthermore, the expression of the three exmiRs in patients with early- or late-stage PC was also investigated in the present study. The results indicated that exmiR-21 level may predict the stage of PC and suggested that elevated exmiR-21 expression may serve as a potential biomarker for the early diagnosis of PC.

Numerous studies have reported an association between the overexpression of miR-21 and miR-10b in the plasma of patients with PC (34,35). miR-21 and miR-10b have been associated with multiple cancer types, such as gastric and esophageal cancer (36), lung cancer (37) and breast cancer (38). However, few studies have examined exmiRs in plasma samples of patients with PC. Several studies have reported that exmiR-21 (4,39) and exmiR-10b (39,40) were significantly

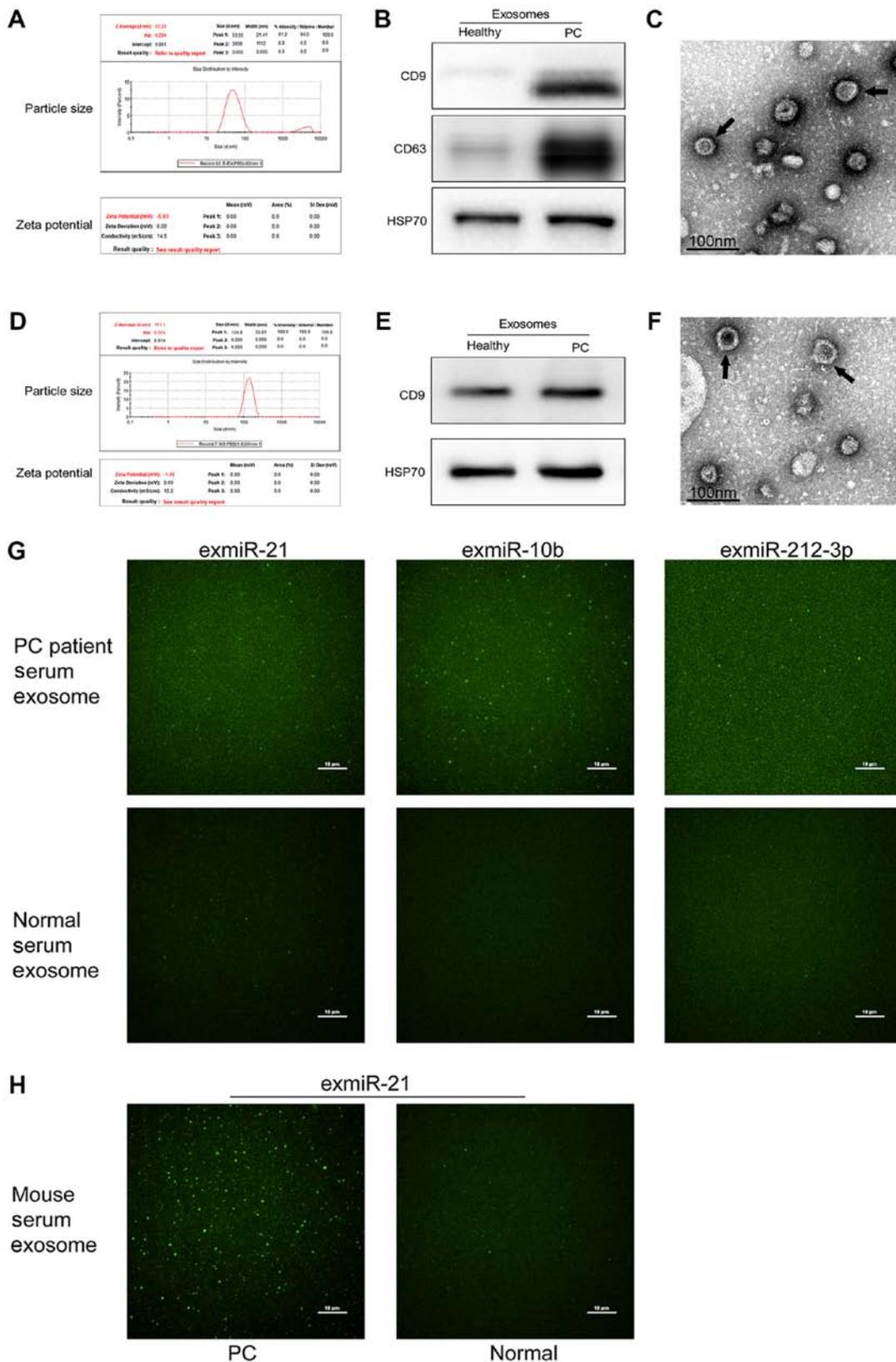


Figure 1. Characterization and quantification of exosome levels. (A) The particle size of exosomes in human plasma samples was 53.5 nm and the ζ potential was -5.8 mV. (B) Western blotting demonstrated the expression of specific exosome marker proteins (CD9, CD63 and HSP70) in exosomes derived from samples of patients with PC and healthy controls. (C) Transmission electron microscopy images of exosomes derived from humans. Black arrows point out the typical exosomes that have a cup shape and a double-layer membrane structure. (D) The particle size of exosomes in the plasma samples of mice was 139.9 nm and the ζ potential was -1.49 mV. (E) Western blotting demonstrated the expression of specific exosome marker proteins (CD9 and HSP70) in exosomes derived from plasma samples of mice with PC and healthy controls. (F) Transmission electron microscopy images of exosomes derived from mice. Black arrows point out the typical exosomes that have a cup shape and a double-layer membrane structure. (G) Total internal reflection fluorescence microscopy images of exmiR-21, exmiR-10b and exmiR-212-3p expression in human serum exosomes. (H) A set of TIRF microscopy images of exmiR-21 fluorescent signals from exosomes isolated from mice serum samples. PC, pancreatic cancer; HSP70, heat shock protein 70; exmiR, exosomal microRNA; TIRF, total internal reflection fluorescence.

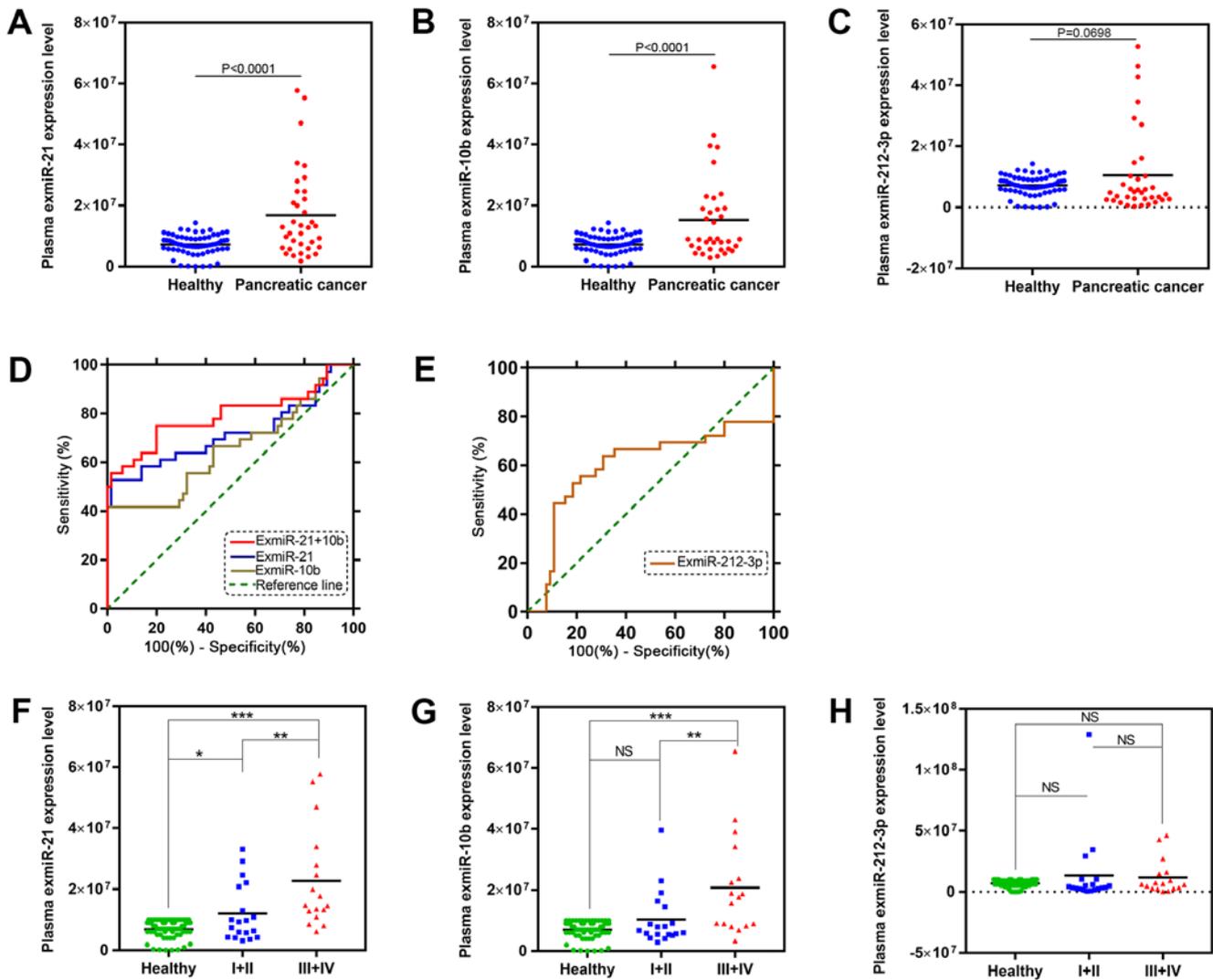


Figure 2. Comparing the expression pattern and diagnostic values of selected plasma exmiRs between patients with PC and control. The relative expression levels of (A) the exmiR-21, (B) exmiR-10b and (C) exmiR-212-3p. ROC analyses were performed among the candidate exmiRs: (D) Relative plasma exmiR-21, exmiR-10b and exmiR-21 combined with exmiR-10b levels in patients with PC (n=36) and healthy controls (n=65). (E) ROC curve of relative plasma exmiR-212-3p levels in patients with PC (n=36) and healthy controls (n=65). The patients with PC were further divided into two groups, early and advanced. Patients at stages I and II (according to the 8th American Joint Committee on Cancer) were divided into the early group, and the other patients at stages III and IV were in the advanced group. The relative expression level of (F) exmiR-21, (G) exmiR-10b and (H) exmiR-212-3p was compared between the different groups of individuals. *P<0.05. **P<0.001. ***P<0.0001. NS, no significant difference. PC, pancreatic cancer; exmiR; exosomal microRNA; ROC, receiver operating characteristic.

upregulated in patients with PC, and the results of the present study are consistent with these findings. The diagnostic performance of these exmiRs were also compared, which revealed that exmiR-21 showed the best diagnostic performance among these three exmiRs. Furthermore, the diagnostic performance of exmiR-21 was increased when combined with exmiR-10b. Wu *et al* (41) reported that miR-212-3p plays an important role in the development of gastric cancer. MiR-212-3p is one of the top differentially expressed miRs in PC (42), and a previous study confirmed that miR-212-3p is expressed at higher levels in PC tissues compared with controls (42). Furthermore, patients with PC and higher miR-212-3p expression had a decreased overall survival compared with those with low miR-212-3p expression. Therefore, miR-212-3p was selected as a candidate miR for analysis in the present study. However, no significant difference was observed in the expression of miR-212-3p between patients with PC and healthy controls,

and exmiR-212-3p exhibited the poorest diagnostic performance among the three exmiRs.

The results indicated that exmiR-21 could distinguish early-stage patients from both healthy individuals and patients with advanced-stage PC, indicating that exmiR-21 showed potential diagnostic value for early-stage PC.

Although the etiology of PC remains largely unclear, cigarette smoking, obesity, history of drinking alcohol and diabetes mellitus have been associated with an increased risk of PC (43). These risk factors were thus examined for any effect on the expression of the exmiRs in patients with PC in the present study. The expression levels of exmiRs were evaluated in subgroups, according to age, gender, body mass index, blood type (44), history of smoking, history of drinking and diabetes mellitus in the 36 patients with PC. No differences were observed in the levels of exmiRs in any of the subgroups. Further studies with a larger number of cases are

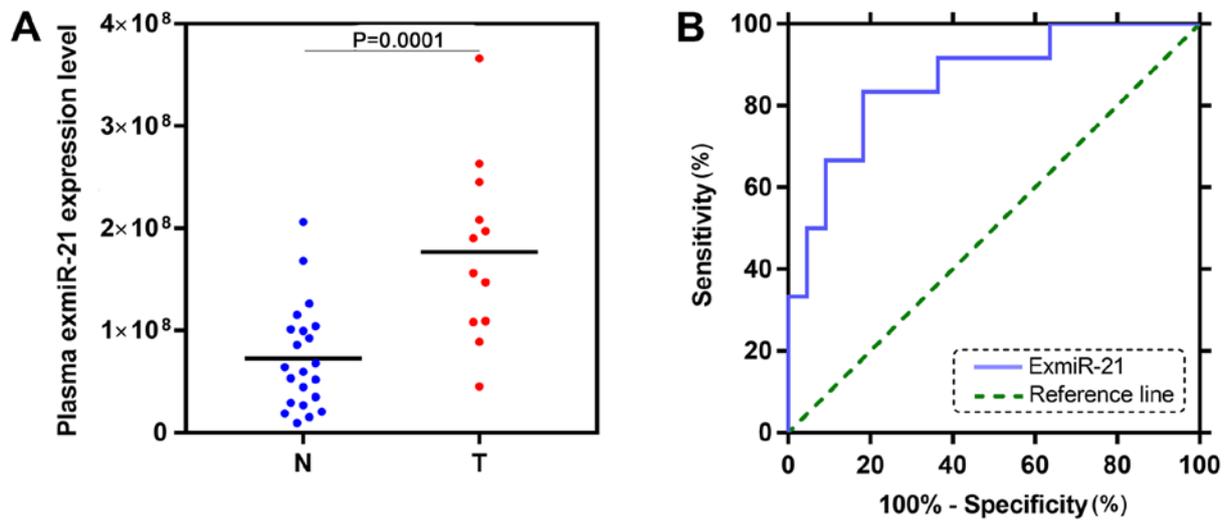


Figure 3. Expression level and diagnostic value of plasma exmiR-21 in mice. (A) Relative expression level of exmiR-21 in the plasma of mice with PC (n=12) and healthy mice (n=22). (B) ROC curve of relative plasma exmiR-21 in mice with PC and the healthy controls. P=0.0005; AUC, 0.8636. PC, pancreatic cancer; N, normal mice; T, mice with PC; exmiR, exosomal microRNA; ROC, receiver operating characteristics.

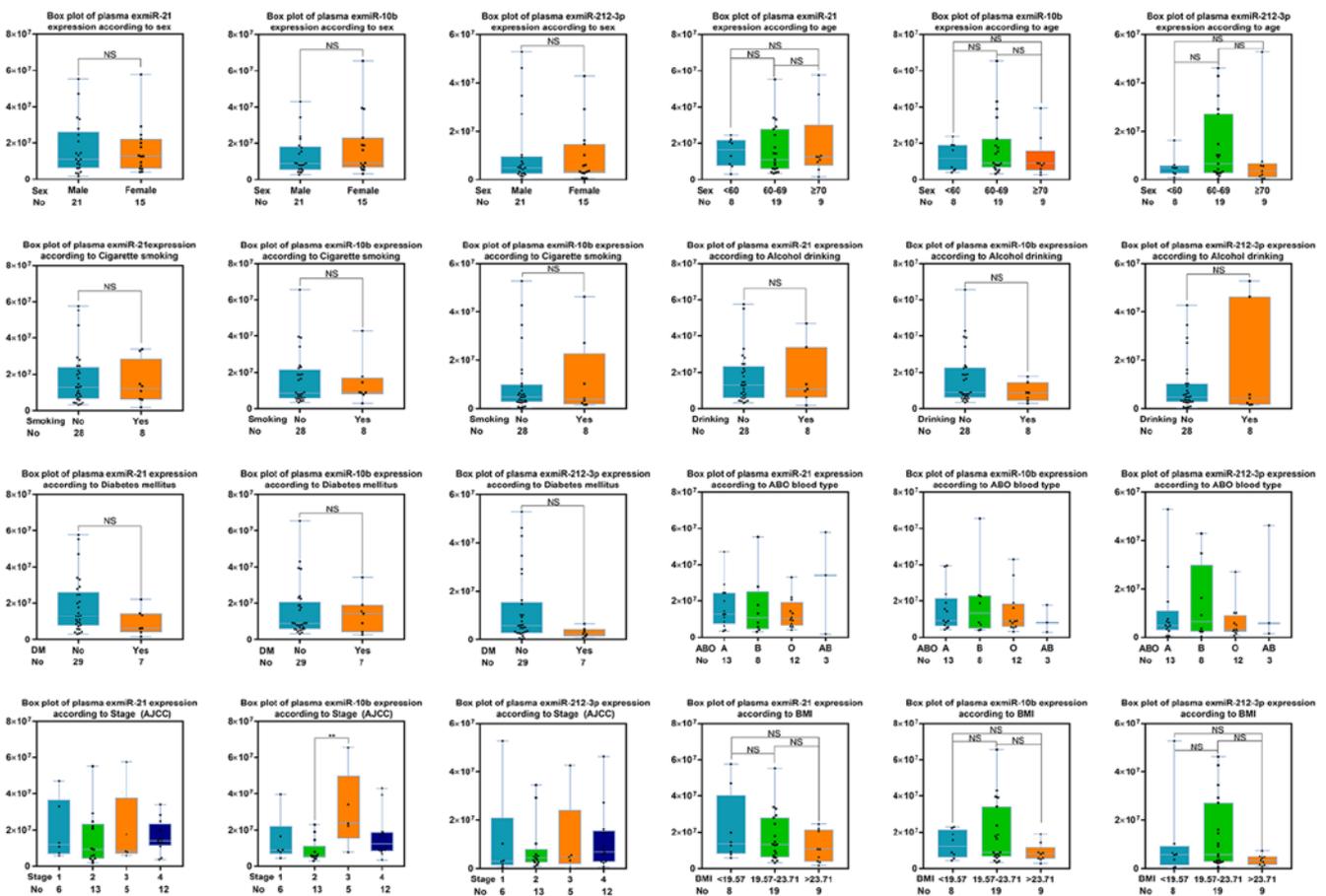


Figure 4. Association between the expression of exmiRs and clinicopathological parameters of patients with pancreatic cancer. **P<0.01. NS, no significant difference. ExmiR; exosomal microRNA; AJCC, American Joint Committee on Cancer; BMI, body mass index.

vital to evaluate the expression levels of these exmiRs based on clinical parameters.

To the best of our knowledge, the present study is the first to use the TCLN biochip to detect the expression levels of these candidate exmiRs in patients with PC, healthy indi-

viduals and mice, with the aim of investigating the diagnostic value of plasma exmiRs in PC. Compared with RTq-PCR, which involves cumbersome and time-consuming methods for exosome miR extraction and detection, the TCLN biochip is a novel and efficient method of detecting RNA targets for

the early diagnosis of cancer (6). Using the TCLN biochip, the levels of exmiRs can be easily determined in various cancer types. In the present study, the utility of using the TCLN biochip to evaluate exmiR-21 levels was demonstrated to distinguish patients with early-stage PC from healthy individuals. Although further large-scale studies are required to validate the present findings, these results suggest that this method will increase the number of patients with PC diagnosed at the early-stages, giving the opportunity to be treated with surgery and prevent disease progression.

The present study has several limitations. Due to the limited number of patients with PC in the study, a training and validation cohort could not be set. However, the plasma levels of exmiR-21 in 34 mice were validated. The present study also only compared the expression levels of the three exmiRs in patients with PC and healthy individuals; benign pancreatic disease could be a valuable inclusion (such as chronic pancreatitis, pancreatic neuroendocrine neoplasms or cystic tumors). Bloomston *et al* (45) suggested that miR-21 was uniquely upregulated in patients with PC compared with normal individuals and patients with chronic pancreatitis. Abue *et al* (46) reported that the miR-483-3p and miR-21 expression levels were significantly higher in PDAC than in normal individuals and patients with intraductal papillary mucinous neoplasm. Additionally, the present study only compared the preoperative patients with healthy individuals. Future studies should examine the levels of these exmiRs before and after treatment, such as surgery, radiotherapy, chemotherapy and molecular targeted therapy. As aforementioned, further studies with larger samples of both patients with PC and healthy individuals will be important to verify the reliability of the diagnostic performance of exmiR-21. Furthermore, other exmiRs that may have higher sensitivity and specificity in PC could also be considered. Therefore, RNA sequencing-based screening to identify PC-specific exmiRs will be conducted, in order to focus on exmiRs with the highest levels of upregulation, for further analysis.

The results of the present study demonstrated that plasma levels of exmiR-21 and exmiR-10b were increased in patients with PC compared with controls. ROC analyses demonstrated that exmiR-21 had the best diagnostic performance among the three exmiRs. ExmiR-21 was also found to discriminate patients with early-stage PC from both healthy controls and patients with advanced PC. These findings suggest that evaluating exmiR-21 using the TCLN biochip may be a useful non-invasive strategy for diagnosing early stage PC.

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Availability of data and materials

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

Authors' contributions

LC and SJ designed the study, performed the experiments, wrote the manuscript and prepared the figures. XP and GD were involved in the study design, and performed the experiments and data analysis. SZ collected the samples and characteristics of healthy individuals and patients. MW assisted with performing the experiment and verifying the results. All authors reviewed the manuscript, and approved the final version of the published manuscript.

Ethics approval and consent to participate

The research protocol was reviewed and approved by the Research Ethics Committee of Sir Run Run Shaw Hospital, School of Medicine, Zhejiang University (approval; no. 20170222-16). All participants or their guardians provided written informed consent for scientific research statement. All animal experiments were carried out in the animal unit of Zhejiang University according to procedures authorized and specifically approved by the institutional Ethical Committee.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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