Supplementary

Table S1 Comparison of PC derived EVs vs. other cancer biomarkers

| Potential biomarker | Sample type | Diagnostic value in PC patients | Advantages | Disadvantages | References |
|------------------------------------|-------------------------------|--|--|--|------------|
| Carbohydrate antigen 19-9 (CA19-9) | Serum | Sensitivity: 79-81%; Specificity: 82-90% | Relatively easy collection; reliable marker for treatment response and monitoring | Poor screening marker; elevated expression in benign jaundice, pancreatitis, ovarian cancer or other malignancies | (1) |
| Circulating tumour cells (CTCs) | Serum/plasma | Sensitivity: 75%; Specificity: 96.4%; AUC: 0.867; 95% CI: 0.798-0.935 | Correlated with poor prognosis | Low concentration in serum/plasma; Lack of evidence in large scale clinical setting; variable in isolation techniques | (2-4) |
| Cell free DNA (cfDNA) | Plasma | Combination of 5mC and 5hmC prediction model Sensitivity: 93.8%; Specificity: 95.5%; AUC: 0.99 | | Utility is limited to identifying existing mutation in clinical setting; Lacks evidence in large scale clinical setting | (5) |
| Extracellular vesicles (EVs) | Plasma/serum/pancreatic juice | GPC1+ study: Sensitivity: 95-100%; Specificity: 95-100% | Correlated to early detection, prognostic marker and potential tumour staging marker | Variability in isolation techniques; Lacks evidence in high quality isolation in clinical setting | (6-17) |

AUC, area under curve.

Table S2 Potential EV biomarkers in pancreatic cancer

| Type of EV cargo | EV content | Experimental Approach | Sample Type | PC patient sample size | Sensitivity and specificity | Relevance to PC | Reference |
|----------------------------|---|---|---|--|---|--|------------|
| Micro RNA (miRNA) | miR-1246, miR-3976, miR-4644, miR-4306 | RT-PCR, qRT-PCR | Serum samples and PC cell lines | miR-1246, miR-4644 and miR-4306: Patients: 12. miR-3976: Patients: 131 | miR-1246: Sensitivity: 66.7%, Specificity:100%, AUC: 0.814. miR-4644: Sensitivity: 75%, Specificity: 76.9%, AUC: 0.76 | Elevated expression | (13,15,18) |
| | miR-18a | qRT-PCR | Patient plasma samples | Patients: 36 | Not available | Elevated expression | (19,20) |
| | miR-17-5p | qRT-PCR | Patient serum samples | Patients: 22 | Sensitivity: 72.7%, Specificity: 92.6% | Correlated to advanced stage of PC | (21) |
| | miR-122-5p | qRT-PCR | Patient plasma samples | Patients: 216 | AUC: 0.81 | Diagnostic marker | (7) |
| | miR-let7a | LC-MS/MS, qRT-PCR | Patient plasma samples | Patients: 29 | Sensitivity: 100%, Specificity | Lower expression linked to PC progression | (22,23) |
| | miR-191, miR-451a and miR-21 | qRT-PCR | Patient plasma samples | Patients: 32 | miR-191: Sensitivity: 71.9%, Specificity: 84.2%. miR-451a: Sensitivity: 65.6%, Specificity: 85.7%. miR-21: Sensitivity: 80.7%, Specificity: 81.0% | Elevated expression | (24) |
| | miR-21 | Western Blotting, TCLN biochip | Patient plasma and mouse serum sample | es Patients: 36 | Sensitivity: 95.5%, Specificity: 81.5% | Elevated expression | (25) |
| | miR-451a | qRT-PCR | Patient serum samples | Patients: 6 | Sensitivity: 69.2%, Specificity: 70.8% | Elevated expression | (26) |
| | miR-196b/LCN2/TIMP1 | RT-PCR | Patient serum and duodenal juice | Patients: 50 | Sensitivity: 80%, Specificity: 80%, AUC: 0.93 | Elevated expression | (27) |
| | miR-214 | qRT-PCR | Patient plasma samples | Patients: 20 | Not available | Lower expression linked to better survival rate. Diagnostic marker | (28) |
| | Ratio of miR-3940-5p/miR-8069 | 3D digital PCR | Patient urine samples | Patients: 43 | Sensitivity: 58.1%, Specificity: 89.2% | Diagnostic marker | (29) |
| | miR-192-5p, miR-19a-3p, and miR-19b-3p | qRT-PCR | Patient serum samples | Patients: 159 | Not available | Elevated expression; Diagnostic and prognostic value | (30) |
| | miR-10b, miR-21, miR-30c, miR- 181a | LSPR-based assay | Patient plasma samples | Patients: 29 | Sensitivity: 100%, Specificity: 100% | Diagnostic marker | (22) |
| Circular RNA (circ RNA) | has_circ_0000896 and has_ circ_0000128 | qRT-PCR | Patient plasma and cell culture samples | Patients: 8 | Not available | Elevated expression | (31) |
| | circRNA-PDE8A | RNA binding protein immunoprecipitation assay, Biotinylated RNA pulldown assays | Patient plasma samples | Patients: 93 | Not available | Elevated expression; correlated PC progression | (32) |
| | circ-IARS | qRT-PCR | Patient plasma, tissue and cell culture samples | Patients: 92 | Not available | Elevated expression | (33) |
| | circRNA-0000069 | Flow-cytometry, Western blotting, RT- qPCR | Patient tissue and cell culture samples | Patients: 179 | Not available | Elevated expression | (34) |

Table S2 (continued)

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| Type of EV cargo | EV content | Experimental Approach | Sample Type | PC patient sample size | Sensitivity and specificity | Relevance to PC | Reference |
|------------------|---|--|---|------------------------|--------------------------------------|---|-----------|
| mRNA | CK18, CD63 | Next Generation sequencing and qRT-PC | RPatient plasma samples | Patients: 89 | AUC: 0.93 | Detected in PDAC patients | (35) |
| | FGA, KRT19, HIST1H2BK, ITIH2, MARCH2, CLDN1, MAL2 and TIMP1 | RNA sequencing analysis | Patient plasma samples | Patients: 284 | AUC: 0.949 | Detected in PDAC patients | (36) |
| Proteins | Glypican-1 | UPLC-MS, Western Blot Analysis, qRT- PCR | Patient serum and tissue samples, Anima studies, cell lines | I Patients: 190 | Sensitivity: 100%, Specificity: 100% | Diagnostic/Screening tool | (6) |
| | MIF | Proteomics, RNA sequencing, tissue processing, immunofluorescence, SDS- PAGE, Western Blot, flow cytometry | Human peripheral blood samples, animal studies and Cell lines | Patients: 18 | Not available | Initiates formation of pre- metastatic niche in the liver | (37) |
| DNA | NOTCH1, BRCA2 | Next generation sequencing | Patient pleural fluid, blood, plasma | Patient: 3 | Not available | Detected in patient exoDNA samples | (38) |
| | KRAS ^{G12D} and TP53 ^{R273H} | ddPCR | Patient serum samples | Patients: 171 | Not available | Elevated expression | (39) |
| | KRAS | ddPCR | Patient plasma samples | Patients: 194 | Not available | Detected in PC patients | (40) |
| | KRAS | ddPCR | Patient plasma samples | Patients: 88 | Not available | Elevated expression | (41) |

GPC1+, glypican-1; RT-PCR, reverse transcriptase polymerase chain reaction; qRT-PCR, quantitative real time polymerase chain reaction; LC-MS/MS, Liquid chromatography-mass spectrometry; TCLN biochip, tethered cationic lipoplex nanoparticle biochip; LSPR-based assay, localized surface plasmon resonance based assay; UPLC-MS, ultra performance liquid chromatography mass spectrometry; SDS-PAGE, sodium dodecyl-sulfate polyacrylamide gel electrophoresis; ddPCR, droplet digital polymerase chain reaction; MIF, migration inhibitory factor; AUC, area under curve.

Table S3 Preclinical and clinical trials of PC derived EVs

| Disease Model | Trial | Pharmacological drug | Administration | References |
|---------------|--------------------|-----------------------------|---------------------------|------------|
| Mouse | Preclinical | siPAK4 | Intratumoral injection | (42) |
| Mouse | Preclinical | Gemcitabine | Intravenous injection | (43) |
| Mouse | Preclinical | siKRASG12D | Intraperitoneal injection | (44) |
| Mouse | Preclinical | siKRASG12D | Intraperitoneal injection | (45) |
| Mouse | Preclinical | siKRASG12D and miRNA-145-5p | Intratumoral injection | (46) |
| Human | Clinical (Phase 1) | siRNA KrasG12D | Intravenous injection | (47) |

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