<table>
<thead>
<tr>
<th>Title</th>
<th>Pages</th>
<th>Authors</th>
</tr>
</thead>
<tbody>
<tr>
<td>Retrospective Study Analysis of clinicopathological features and</td>
<td>445</td>
<td>Chen YR, Xu ZX, Jiang LX, Dong ZW, Yu PF, Zhang Z, Gu GL</td>
</tr>
<tr>
<td>prognostic factors of breast cancer brain metastasis</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Clinical study of standard residual liver volume and transient</td>
<td>459</td>
<td>Yue ZQ, Zhang P, Yan S, Ju LL, Wang HX, Yuan LX, Chen L, Wu JZ, Cao YL</td>
</tr>
<tr>
<td>elastography in predicting poor prognosis of patients after</td>
<td></td>
<td></td>
</tr>
<tr>
<td>hemihepatectomy</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Observational Study System  describing  surgical field extension</td>
<td>471</td>
<td>Sakamoto A, Noguchi T, Matsuda S</td>
</tr>
<tr>
<td>associated with flap reconstruction after resection of a</td>
<td></td>
<td></td>
</tr>
<tr>
<td>superficial malignant soft tissue tumor</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Basic Study Computational exploration of the significance of</td>
<td>479</td>
<td>Wang SL, Zhuo GZ, Wang LP, Jiang XH, Liu GH, Pan YB, Li YR</td>
</tr>
<tr>
<td>COPS6 in cancer: Functional and clinical relevance across tumor</td>
<td></td>
<td></td>
</tr>
<tr>
<td>types</td>
<td></td>
<td></td>
</tr>
<tr>
<td>META-ANALYSIS Circulating tumor cells as potential prognostic</td>
<td>504</td>
<td>Zhang ZH, Bao YW, Zhao YJ, Wang JQ, Guo JT, Sun SY</td>
</tr>
<tr>
<td>biomarkers for early-stage pancreatic cancer: A systematic review</td>
<td></td>
<td></td>
</tr>
<tr>
<td>and meta-analysis</td>
<td></td>
<td></td>
</tr>
<tr>
<td>SCIENTOMETRICS Bibliometric analysis of the global research status</td>
<td>518</td>
<td>Zhang YZ, Li MZ, Wang GX, Wang DW</td>
</tr>
<tr>
<td>and trends of mechanotransduction in cancer</td>
<td></td>
<td></td>
</tr>
<tr>
<td>CASE REPORT Autoimmune diabetes from pembrolizumab: A case report</td>
<td>535</td>
<td>Bhandari H, Khalid F, Bodla ZH, Muhammad T, Du D, Meghal T</td>
</tr>
<tr>
<td>and review of literature</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Calcitriol induced hypercalcemia - a rare phenomenon in lung cancer</td>
<td>544</td>
<td>Prakash A, Khalid F, Alahwan A, Bader H, Du D, Meghal T</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
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https://www.wjgnet.com/bpg/gerinfo/208

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https://www.wjgnet.com/bpg/GerInfo/239

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https://www.wjgnet.com/bpg/GerInfo/242

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Circulating tumor cells as potential prognostic biomarkers for early-stage pancreatic cancer: A systematic review and meta-analysis

Zi-Han Zhang, Yi-Wen Bao, Ya-Jun Zhao, Jian-Quan Wang, Jin-Tao Guo, Si-Yu Sun

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Abstract

BACKGROUND
Pancreatic cancer is difficult to be diagnosed early clinically, while often leads to poor prognosis. If optimal personalized treatment plan can be provided to pancreatic cancer patient at an earlier stage, this can greatly improve overall survival (OS). Circulating tumor cells (CTCs) are a collective term for various types of tumor cells present in the peripheral blood (PB), which are formed by detachment during the development of solid tumor lesions. Most CTCs undergo apoptosis or are phagocytosed after entering the PB, whereas a few can escape and anchor at distal sites to develop metastasis, increasing the risk of death for patients with malignant tumors.

AIM
To investigate the significance of CTCs in predicting the prognosis of early pancreatic cancer patients.

METHODS
The PubMed, EMBASE, Web of Science, Cochrane Library, China National Knowledge Infrastructure, China Biology Medicine, and ChinaInfo databases were searched for articles published through December 2022. Studies were considered qualified if they included patients with early pancreatic cancer, analyzed the prognostic value of CTCs, and were full papers reported in English or Chinese. Researches were selected and assessed using the Preferred Reporting Items for Systematic Reviews and Meta-Analyses protocol and the Newcastle-Ottawa Scale criteria. We used a funnel plot to assess publication bias.

RESULTS
From 1595 publications, we identified eight eligible studies that collectively enrolled 355 patients with pancreatic cancer. Among these original studies, two were carried out in China; three in the United States; and one each in Italy, Spain, and Norway. All eight studies analyzed the relevance between CTCs and the
prognosis of patients with early-stage pancreatic cancer after surgery. A meta-analysis showed that the patients
that were positive pre-treatment or post-treatment for CTCs were associated with decreased OS [hazard ratio (HR)
= 1.93, 95% confidence interval (CI): 1.197-3.126, \( P = 0.007 \)] and decreased relapse-free/disease-free/progression-
free survival [HR = 1.27, 95% CI: 1.137-1.419, \( P < 0.001 \)] in early-stage pancreatic cancer. Additionally, the results
suggest no statistically noticeable publication bias for overall, disease-free, progression-free, and recurrence-free
survival.

CONCLUSION
This pooled meta-analysis shows that CTCs, as biomarkers, can afford reliable prognostic information for patients
with early-stage pancreatic cancer and help develop individualized treatment plans.

Key Words: Pancreatic cancer; Surgery; Prognosis; Systematic review; Meta-analysis; Biomarkers

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Core Tip: There is no consensus regarding the prognostic value of circulating tumor cells (CTCs) in early-stage pancreatic
cancer after surgery. This is the first systematic review and meta-analysis to investigate the potential of CTCs in predicting
survival time in early pancreatic cancer. We pooled the analyses of the relationship between CTCs and overall/disease-
free/progression-free/relapse-free survival in related studies. Patients testing positive for CTCs pre- or intra-surgery may
have worse prognoses, requiring more intense chemotherapy and closer follow-up.

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INTRODUCTION
Pancreatic cancer is a fatal disease with poor prognosis. In 2018, pancreatic cancer was the seventh leading cause of
cancer-related mortality worldwide[1]. The number of annual diagnoses of and deaths related to pancreatic cancer in
China have exceeded those in the United States[2]. The incidence of pancreatic cancer continues to increase at a rate of
0.5%-1.0% each year, and it is projected to be the second deadliest cancer by 2030 in Western countries[3]. Despite
continuous advances in chemotheraphy, radiation, and surgical techniques; the prognosis of patients with pancreatic
cancer remains significantly poor. Approximately half of the patients experience reoccurrence within the first year,
mainly due to metastatic disease occurrence after surgical resection. Consequently, one of the most important challenges
is to identify a factor that can assess the survival outcome of patients with early pancreatic cancer before surgery, optimize treatment, and assist in the development of monitoring strategies.

Currently, technological innovations in imaging and endoscopy are being used to improve the diagnostic accuracy of
pancreatic cancer. Computed tomography (CT) and magnetic resonance imaging (MRI) remain first-line diagnostic
modalities for clinical suspicion. Endoscopic ultrasound (EUS) and EUS-guided fine-needle aspiration (EUS-FNA), also
play important roles in its diagnosis. However, imaging features of early pancreatic cancer are subtle, and the general
consensus is that it is difficult to detect early lesions[4]. Liquid biopsy is a new technology that detects biomarkers, such
as carcinoembryonic antigen, carbohydrate antigen 19-9 (CA19-9), or CA125, from nonsolid biological tissues, such as
blood[5], and has received increased attention because of its convenience and noninvasiveness[6]. However, these
biomarkers all lack sufficient sensitivity and specificity for diagnostic purposes.

With the development of medical testing technologies, circulating tumor cells (CTCs) have emerged as a popular
research topic over the last few decades. CTCs are a small number of tumor cells that detach from primary tumors,
circulate through the bloodstream, and are the main source of its dissemination and metastasis[7]. Unlike traditional
histopathological examinations, which are usually complex and risky for difficult-to-biopsy tumors such as pancreatic
cancer, CTCs are noninvasive. CTCs can provide real-time and comprehensive information about the tumor because they
are enriched in the bloodstream and originate from different regions of the original tumor or metastasis. Moreover, CTCs
can present large-scale health information, such as the expression of genes and proteins and alteration of cellular contents
and cell membranes, which are essential for improving diagnostic accuracy and developing individualized treatments[8].
Recently, an increasing number of studies have revealed that CTCs show promise for prognostic evaluation in several
tumors, including lung[9], renal[10], breast[11], gastric[12], and colorectal cancers[13].

Nonetheless, the prognostic effect of CTCs in early pancreatic cancer remains ambiguous, mainly because of the
increasing number of CTC isolation methods and different study designs. Therefore, this study aimed to perform a
structural meta-analysis of currently available evidence on the prognostic value of CTCs in early-stage pancreatic cancer.
MATERIALS AND METHODS

Search strategy
Following the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines, we conducted a comprehensive literature search of all studies published in database repositories before December 2022 that were related to the use of CTCs for the diagnosis of pancreatic cancer. The search was performed only in English and Chinese databases including PubMed, EMBASE, Web of Science, Cochrane Library, China National Knowledge Infrastructure (CNKI), China Biology Medicine (CBM), and Chinalinfo. Only studies published in Chinese and English were included in the analysis. Search terms included these keywords, MeSH (medical subject headings) terms, and their entry terms: “Pancreatic Neoplasms” (MeSH), “Neoplasm, Pancreatic,” “Pancreatic Neoplasms,” “Pancreas Neoplasms,” “Neoplasm, Pancreas,” “Neoplasms, Pancreas,” “Pancreas Neoplasms,” “Neoplasms, Pancreatic,” “Cancer of Pancreas,” “Pancreas Cancers,” “Cancer, Pancreas,” “Cancers, Pancreas,” “Pancreatic Cancer,” “Cancer, Pancreatic,” “Cancers, Pancreatic,” “Pancreatic Cancer,” “Cancer of the Pancreas,” “Neoplastic Cells, Circulating” (MeSH), “Neoplasms Circulating Cells,” “Circulating Neoplastic Cells,” “Cell, Circulating Neoplastic,” “Cells, Circulating Neoplastic,” “Circulating Neoplastic Cell,” “Neoplastic Cell, Circulating,” “Circulating Tumor Cells,” “Cell, Circulating Tumor,” “Cells, Circulating Tumor,” “Circulating Tumor Cell,” “Tumor Cell, Circulating,” “Cells, Neoplasms Circulating,” “Cell, Neoplasm Circulating,” “Neoplasms Circulating Cell,” “Circulating Cells, Neoplasm,” “Tumor Cells, Embolic,” “Cell, Embolic Tumor,” “Cells, Embolic Tumor,” “Embolic Tumor Cell,” “Tumor Cell, Embolic,” “Embolic Tumor Cells,” “Embolism, Tumor,” “Embolisms, Tumor,” “Tumor Embolism,” “Tumor Embolisms,” “Prognostic,” “Prognosis,” “Prognostic”. These terms were supplemented by the logical operators “and” and “or.” To expand the literature, we reviewed and evaluated the references of the included studies.

Selection criteria
Two independent investigators reviewed the article titles and abstracts according to inclusion and exclusion criteria to exclude irrelevant studies. Subsequently, the full texts of the included studies were analyzed to determine their suitability for meta-analysis. In case of any contradictions, a third reviewer was consulted for adjudication. The inclusion and exclusion criteria were formulated based on the PRISMA of Diagnostic Test Accuracy Studies guidelines. Inclusion criteria: (1) Studies included participants of any age with histologically or cytologically confirmed early pancreatic cancer (early pancreatic cancer was defined as pancreatic adenocarcinoma with a maximum tumor diameter of 4 cm, regional lymph node metastasis of no more than three nodes, and no distant metastasis); (2) studies that investigated pre- or intra-operative CTCs as a prognostic biomarker in blood for early pancreatic cancer patients’ survival results after surgery; (3) sufficient published data available for calculating hazard ratio (HR) and 95% confidence interval (CI) of overall survival (OS) and progression-free survival (PFS); (4) studies that were published in English or Chinese; and (5) studies that were reported as full paper publications. Exclusion criteria: (1) Patients were diagnosed with pancreatic cancer at advanced stages or studies that didn’t analyze the results of early pancreatic cancer patients independently; (2) studies that did not provide adequate data on the prognostic performance of CTCs for early pancreatic cancer patients; (3) patients didn’t receive surgical therapy; and (4) articles were published in languages other than English and Chinese. Overall, five non-human studies, case reports, comments, meta-analyses, reviews, and published clinical and treatment guidelines were excluded.

Data extraction
To decrease systematic errors during data collection, two reviewers independently selected the studies. Any conflicting cases were carefully reviewed, and a third reviewer was consulted to reach a consensus. Data extraction from the eligible papers included the following items: (1) General information about the article: first author, publication date, and country; (2) study information: number of patients, evidence of confirmed pancreatic cancer, CTCs separation solution, CTCs determination criteria, and follow-up time; and (3) data for the meta-analysis: HR with 95% CI for OS, PFS or disease-free survival (DFS), recurrence-free survival (RFS). For articles without HR and 95% CI, we used Engauge Digitizer 11.3 to calculate them based on the survival rate extracted from Kaplan-Meier curves.

Risk of bias
The quality of the included studies was assessed based on the Newcastle-Ottawa Scale (NOS) criteria for non-randomized studies, which included three key domains covering “selection,” “comparability,” and “outcome.” A star rating system was used to semi-quantitatively evaluate study quality, and those who met the standards for each item were awarded one or two stars. Scores range from zero to nine. A score equal to or greater than seven indicates high quality. This tool objectively evaluates the risk of bias and assesses concerns regarding its applicability. The quality assessment was performed by two independent reviewers. Any disagreements were discussed until an agreement was reached. Publication bias was investigated using a funnel plot, with P < 0.05 indicating a significant publication bias.

Statistical analysis
Review Manager 5.3 software (The Nordic Cochrane Center, The Cochrane Collaboration, Copenhagen, Denmark) was used to assess the pooled HR effect size. Heterogeneity between studies was assessed using Cochrane’s Q test and I² statistics. According to Higgins and Thompson, I² > 50% or P < 0.1 was viewed as consistent significant heterogeneity. A fixed-effect model was used when minor heterogeneity was observed. Otherwise, a random-effects model was used to calculate the pooled HR. Subgroup analyses based on ethnicity, separation solution, treatment, and follow-up time were performed to explore potential sources of heterogeneity. Finally, we conducted a sensitivity analysis to evaluate the effect
of a single study on overall outcomes.

RESULTS

Characteristics and quality of the included studies
A total of 1963 articles were collected, including 91 from CNKI, 273 from ChianInfo, 90 from CBM, 258 from PubMed, 55 from EMBASE, 31 from Cochrane, and 1165 from Web of Science. After deleting 368 duplicate references, we screened the titles and abstracts of 1595 residual articles. In total, 1551 articles were excluded (520 meta-analyses or reviews; 32 non-human studies; 15 articles published in languages other than English and Chinese; 161 meeting abstracts, case reports, guidelines, or letters; and 823 irrelevant studies). According to the inclusion and exclusion criteria, we read the full text of the 44 remaining studies, of which 36 articles were ultimately excluded (5 irrelevant studies and 31 articles with data deficiency). Consequently, eight studies involving 355 patients with early-stage pancreatic cancer were included in our meta-analysis.[14-21]. A flow diagram of the literature search and filtering process is shown in Figure 1.

The characteristics of these studies are summarized in Table 1. The overall research quality was rated as moderate by the NOS, with an average score of 6.75 (Table 2). Among these original studies, two were conducted in China; three in the United States; and one each in Italy, Spain, and Norway. The publication years were 2014 (n = 1), 2018 (n = 1), 2021 (n = 5), and 2022 (n = 1). All eight studies analyzed the correlation between CTCs and prognosis in patients with early-stage pancreatic cancer after surgery. While four studies included early pancreatic cancer patients only,[14,19-21], the remaining four studies contained early and advanced pancreatic cancer patients with early pancreatic cancer patients accounting for at least 30% of the total patients[15-18]. In total, five studies detected CTCs in the peripheral blood (PB), two studies detected CTCs in the PB and portal venous blood (PVB), and one study detected CTCs in the central venous catheter and portal blood. Blood samples were collected solely before surgery in four studies[16,17,21], before and after surgery in three studies[14,15,18], and during surgery in only one study[19]. Only pre- and intra-operative CTC data were included in this meta-analysis. Although eight studies applied different CTC enrichment and separation methods, including commercially available CTC detection kits,[14,17], dielectrophoresis-field flow fractionation (DEP-FFF)[15], and CTC isolation systems[18-21], most methods essentially rely on the density characteristics of CTCs as well as the epithelial and mesenchymal markers expressed by CTCs. Most studies applied DFS, OS, PFS, and RFS as survival outcomes, while only Hugenschmidt et al.[20] adopted cancer-specific survival as an outcome indicator. Seven of the eight studies reported the adjusted HR and 95%CI for the association between DFS/PFS/RFS and positive CTCs. For the one that did not, we calculated these values according to the Kaplan-Meier curves provided in the articles. In addition, only three of the eight included studies reported the adjusted HR and 95%CI for the association between OS and positive CTCs while another two studies provided Kaplan-Meier curves for OS.

The relationship between CTCs and prognosis of early-stage pancreatic cancer patients after surgery
We conducted a meta-analysis to assess the association between CTCs detected in the blood samples of patients with early-stage pancreatic cancer and their prognosis after surgery. The degree of heterogeneity among the five studies that provided the adjusted HR and 95%CI for OS was low (F = 41%, P = 0.15); therefore, we chose a fixed-effect model for analysis. The pooled analyses of the five studies, containing 169 patients, showed that positive detection of pre-treatment or post-treatment CTCs was associated with decreased OS (HR = 1.93, 95%CI: 1.20-3.13) that was statistically significant (Z = 2.69, P = 0.007) (Figure 2A). There was a moderate degree of heterogeneity among the seven studies available for the adjusted HR and 95%CI of DFS/PFS/RFS (F = 56%, P = 0.01); therefore, we chose a random-effects model for analysis. The results showed that positive pre-treatment or post-treatment CTCs were associated with decreased DFS/PFS/RFS (HR = 1.97, 95%CI: 1.20-3.25) that were statistically significant (Z = 2.67, P = 0.008) (Figure 2B).

Subgroup analysis
Because of the high heterogeneity in the DFS/PFS/RFS results, meta-regression was conducted to explore the sources of heterogeneity. No apparent deviation was found in the ethnicity, treatment, and follow-up time subgroups. Later analysis demonstrated that CTC-positive patients were associated with decreased DFS/PFS/RFS in subgroups that detected CTCs by the CellSearch system (HR = 2.62, 95%CI: 1.65-4.16, Z = 4.09, P < 0.001), and the heterogeneity between subgroups was low (P = 0.018) (Figure 3).

Sensitivity analysis
To explore the potential sources of this difference, we conducted sensitivity analysis by sequentially excluding each study. When the study by Xing et al.[14] was excluded, the heterogeneity of the remaining studies was significantly reduced (OS: F = 0%, P = 0.44; DFS/PFS/RFS: F = 0%, P = 0.76), and the pooled results for both OS and DFS/PFS/RFS were increased (OS: HR = 3.06, 95%CI: 1.59-5.86, Z = 3.36, P < 0.001; DFS/PFS/RFS: HR = 2.60, 95%CI: 1.78-3.812, Z = 4.001, P < 0.001) (Figure 4). Nevertheless, the direction of the pooled results for the OS and DFS/PFS/RFS subgroups were not affected, indicating a negative association between CTC positivity and lower OS or DFS/PFS/RFS.

Publication bias
We used a funnel plot to assess publication bias in this meta-analysis. The results showed no publication bias for OS (P = 0.653) and DFS/PFS/RFS (P = 0.117) (Figure 5).
Table 1 Characteristics of the included studies

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<tr>
<th>Ref.</th>
<th>Year</th>
<th>Country</th>
<th>Sample</th>
<th>Separation Solution</th>
<th>Markers and expression level on PC CTC</th>
<th>Cases</th>
<th>No-positive</th>
<th>Treatment</th>
<th>Outcome</th>
<th>Follow-up time</th>
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<tbody>
<tr>
<td>Xing et al[14]</td>
<td>2021</td>
<td>China</td>
<td>Peripheral blood</td>
<td>SE-iFISH</td>
<td>CD44+ CTC: DAPI +/CD45-/CD31 +/CD44 +/Vimentin (+ or -) with aneuploid CEP8</td>
<td>73</td>
<td>Not reported</td>
<td>Surgery</td>
<td>OS/DFS</td>
<td>Median 10.8 mo (1.2-31.8 mo)</td>
</tr>
<tr>
<td>Semaan et al[15]</td>
<td>2021</td>
<td>United States</td>
<td>Peripheral blood</td>
<td>DEP-FFF</td>
<td>pEMT-CTC: CD45+, EpCAM + and/or Pan-CK +, Vimentin +, DAPI +</td>
<td>31</td>
<td>28</td>
<td>Surgery/neoadjuvant treatment</td>
<td>OS/PPS</td>
<td>Median 15.4 mo (0.43-1.9 mo)</td>
</tr>
<tr>
<td>Court et al[16]</td>
<td>2018</td>
<td>United States</td>
<td>Peripheral blood</td>
<td>NanoVelcro chip</td>
<td>DAPI +/CD45 -/CK +; CD45 positivity greater than 2 × background</td>
<td>40</td>
<td>27</td>
<td>Surgery/chemotherapy</td>
<td>OS/RFS</td>
<td>≥ 24 mo</td>
</tr>
<tr>
<td>White et al[18]</td>
<td>2021</td>
<td>United States</td>
<td>Peripheral blood and portal venous blood</td>
<td>CellSearch</td>
<td>DAPI +/CK +/CD45 -</td>
<td>33</td>
<td>21</td>
<td>Surgery/neoadjuvant treatment</td>
<td>OS/RFS</td>
<td>Median 14.1 mo (0.86-1.97 mo)</td>
</tr>
<tr>
<td>Bissolati et al[19]</td>
<td>2014</td>
<td>Italy</td>
<td>Peripheral blood and portal venous blood</td>
<td>CellSearch</td>
<td>DAPI +/CK +/CD45 -</td>
<td>20</td>
<td>9</td>
<td>Surgery/chemotherapy</td>
<td>OS/PPS</td>
<td>Median 39.2 mo (36-45 mo)</td>
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</tbody>
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DEP-FFF: Dielectrophoresis-field flow fractionation; ICC: Immunocytochemistry; SE-iFISH: Subtraction enrichment and immunostaining-fluorescence in situ hybridization; PC: Pancreatic cancer; CTCs: Circulating tumor cells; LT-PCR: Ligand-targeted polymerase chain reaction; CK: Cytokeratin; DAPI: 4',6-diamidino-2-phenylindole; EpCAM: Epithelial cell adhesion molecule; CSS: Cancer specific survival; DFS: Disease-free survival; OS: Overall survival; PFS: Progression-free survival; pEMT: Partial epithelial-mesenchymal transition.

The relationship between CA19-9 and the prognosis of early-stage pancreatic cancer patients after surgery

In order to compare the potential of CTC and CA19-9 in predicting patient prognosis, we extracted the adjusted HR and 95%CI of DFS related to CA19-9 in the included articles. Only four studies had the required data[14,16-18]. The degree of heterogeneity of the adjusted HR and 95%CI for DFS was high ($I^2=73\%$, $P=0.01$), so we used a random-effects model for the analysis. The pooled analyses of the four studies, containing 171 patients, showed that the CA19-9 level in the PVB of early-stage pancreatic cancer patients was not an independent predictor of a shorter time to recurrence ($HR=1$, 95%CI: 1.00-1.00; $Z=0.36$, $P=0.72$) (Figure 6). Considering the impact of sampling time on the results, we excluded the study of White et al[18], which collected venous blood after resection. Though the heterogeneity of the remaining studies was significantly reduced ($I^2=48\%$, $P=0.15$), the combined HR was still 1 (95%CI: 1.00-1.00; $Z=2.20$, $P=0.03$) (Figure 7).
### Table 2 Quality assessment of the included studies according to the Newcastle-Ottawa Scale criteria for non-randomized studies

<table>
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<td>Selection of non-exposed</td>
<td>Ascertainment of exposure</td>
<td>Outcome not present at start</td>
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<tr>
<td>Xing et al[14]</td>
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<td>Semaan et al[15]</td>
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<td>Padillo-Ruiz et al[21]</td>
<td>-</td>
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<td>Court et al[16]</td>
<td>-</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>Cheng et al[17]</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>White et al[18]</td>
<td>-</td>
<td>1</td>
<td>1</td>
<td>1</td>
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<tr>
<td>Bissolati et al[19]</td>
<td>1</td>
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</tr>
<tr>
<td>Hugenschmidt et al[20]</td>
<td>-</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
</tbody>
</table>

1Meets the criteria of the scale. A maximum of one ‘star’ for each item within the ‘Selection’ and ‘Outcome’ categories; maximum of two ‘stars’ for ‘Comparability’. -Does not meet the criteria of the scale.

### DISCUSSION

In this meta-analysis, we found that CTC detected in patients with early pancreatic cancer is negatively correlated with the prognosis time after surgery. After data collection and filtering, eight studies from 2014 to 2022, including 355 patients with early-stage pancreatic cancer, were included in our analysis. Through data sorting and meta-analysis, we demonstrated that testing positive pre- or intra-treatment for CTCs was associated with decreased OS and RFS/DFS/PFS in early-stage pancreatic cancer. Nevertheless, our meta-analysis showed a high degree of heterogeneity in DFS/PFS/RFS. To determine the potential sources of the heterogeneity, we conducted a subgroup analysis. The pooled results were not affected by ethnicity, treatment, or follow-up time, whereas subgroup analysis by CTCs separation solution decreased the heterogeneity between the groups, indicating that this might have caused the heterogeneity. Studies that enumerated CTCs by the CellSearch system demonstrated a more obvious correlation between CTC-positive patients and decreased DFS/PFS/RFS. However, except for this system, all other separation solutions contained only one study; therefore, we were unable to obtain pooled results. Sensitivity analysis was conducted by using the “leave-one-out method.” Notably, when a study by Xing et al[14] was removed, the heterogeneity of the remaining studies was significantly reduced, and the pooled results for both OS and DFS/PFS/RFS were relatively elevated. We infer that the differences present in this study may be derived from the separation solutions employed as well as the stemness markers used to identify the stem cell-like phenotype of CTCs. Importantly, the results demonstrating the survival-jeopardizing effects of CTCs were maintained. We suggest that patients testing positive for CTCs pre- or intra-treatment may have a worse prognosis and
require more intense chemotherapy and closer follow-up. Furthermore, we collected data regarding CA19-9 in the included studies and four studies containing 171 patients, that met our requirements. Our meta-analysis showed that the preoperative CA19-9 level in the PVB of early-stage pancreatic cancer patients was not an independent predictor for shorter time of recurrence after resection.

The prevalence of pancreatic cancer has increased dramatically globally, and is expected to become a leading cause of cancer-related mortality[22]. Only 10%-15% of patients have localized pancreatic cancer suitable for surgery[3], which is the only potentially curative therapy known to date. Even for patients with a localized disease, a high proportion experience postoperative recurrence within 5 years, caused by local tumor recurrence or distant metastases. The 1- and 5-year survival rates are 63% and 17%, respectively[23]. Therefore, identifying high-risk populations for screening and prevention, early diagnosis, and establishing personalized treatment plans are currently the primary challenges[24].

Imaging, including EUS, CT and MRI, which can provide a convenient and noninvasive diagnosis, remains the first-line diagnostic modality for pancreatic cancer and is used to evaluate therapeutic efficacy in many organs[25-28]. However, cross-sectional imaging is limited in the visualization of small and metastatic tumors, which can frequently result in underestimation of the pancreatic cancer stage[29]. EUS-FNA or EUS-fine needle biopsy (EUS-FNB) can localize pancreatic lesions measuring < 3 cm, providing a minimally invasive tissue biopsy[30]. EUS-FNB is increasingly becoming a practical tool for diagnosing malignancy in various pancreatic solid lesions[31,32]. EUS-FNA combined with needle-based confocal laser endomicroscopy (nCLE) can achieve real-time imaging for in vivo tissue analysis[33]. However, these examinations requires anesthesia and may be accompanied by complications such as acute pancreatitis, tumor dissemination, and postoperative hemorrhage. Besides, nCLE is limited by the duration of the surgical time and the operability of the 19G FNA needle[33]. More recently, liquid biopsy has received a great deal of attention for its ability to assess a comprehensive cancer profile in a noninvasive and real-time manner[26]. Serum CA19-9 level is the only diagnostic biomarker approved by the United States Food and Drug Administration for pancreatic cancer. Moreover, CA19-9 can be an independent predictor of prognosis for pancreatic cancer patients[34]. An elevated CA19-9 level after

Figure 1 Flow diagram for searching and filtering eligible studies included in the meta-analysis.
A

<table>
<thead>
<tr>
<th>Study or subgroup</th>
<th>log[hazard ratio]</th>
<th>SE</th>
<th>Weight</th>
<th>IV, fixed, 95%CI</th>
<th>Hazard ratio IV, fixed, 95%CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Alexander Serman2021</td>
<td>2.14593128</td>
<td>1.18388675</td>
<td>4.3%</td>
<td>0.65 [0.84, 0.70]</td>
<td>-</td>
</tr>
<tr>
<td>Cheng Xing2021</td>
<td>0.11511281</td>
<td>0.36263032</td>
<td>45.6%</td>
<td>1.12 [0.55, 2.28]</td>
<td>-</td>
</tr>
<tr>
<td>Javier Padillo-Ruiz2021</td>
<td>1.18264702</td>
<td>0.38930008</td>
<td>39.6%</td>
<td>3.26 [1.52, 7.00]</td>
<td>-</td>
</tr>
<tr>
<td>Massimiliano Bissolati2014</td>
<td>1.31640923</td>
<td>1.11628455</td>
<td>4.8%</td>
<td>3.73 [0.42, 33.26]</td>
<td>-</td>
</tr>
<tr>
<td>Michael O. White2021</td>
<td>-0.28768207</td>
<td>1.02623894</td>
<td>5.7%</td>
<td>0.75 [0.10, 5.63]</td>
<td>-</td>
</tr>
<tr>
<td>Total (95% CI)</td>
<td>-</td>
<td>-</td>
<td>100.0%</td>
<td>1.93 [1.20, 3.13]</td>
<td>-</td>
</tr>
<tr>
<td>Heterogeneity: Chisq = 6.83, df = 4 (P = 0.15), I^2 = 41%</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Test for overall effect: Z = 2.69 (P = 0.007)</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
</tbody>
</table>

B

<table>
<thead>
<tr>
<th>Study or subgroup</th>
<th>log[hazard ratio]</th>
<th>SE</th>
<th>Weight</th>
<th>IV, random, 95%CI</th>
<th>Hazard ratio IV, random, 95%CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Michael O. White2021</td>
<td>0.90825958</td>
<td>0.5919223</td>
<td>11.4%</td>
<td>2.48 [0.78, 7.91]</td>
<td>-</td>
</tr>
<tr>
<td>Massimiliano Bissolati2014</td>
<td>-0.52733734</td>
<td>1.36131066</td>
<td>3.1%</td>
<td>0.59 [0.04, 8.50]</td>
<td>-</td>
</tr>
<tr>
<td>Javier Padillo-Ruiz2021</td>
<td>0.52442222</td>
<td>0.73826281</td>
<td>8.5%</td>
<td>1.28 [0.30, 5.49]</td>
<td>-</td>
</tr>
<tr>
<td>Harald Hugenschmidt2021</td>
<td>1.02961942</td>
<td>0.26155128</td>
<td>22.7%</td>
<td>2.80 [1.68, 4.68]</td>
<td>-</td>
</tr>
<tr>
<td>Hao Cheng2022</td>
<td>1.31774782</td>
<td>0.55327808</td>
<td>12.4%</td>
<td>3.74 [1.26, 11.05]</td>
<td>-</td>
</tr>
<tr>
<td>Colin M. Court2018</td>
<td>0.9439059</td>
<td>0.54525224</td>
<td>12.6%</td>
<td>2.57 [0.88, 7.48]</td>
<td>-</td>
</tr>
<tr>
<td>Cheng Xing2021</td>
<td>0.18913794</td>
<td>0.05820024</td>
<td>29.3%</td>
<td>1.21 [1.08, 1.35]</td>
<td>-</td>
</tr>
<tr>
<td>Total (95% CI)</td>
<td>-</td>
<td>-</td>
<td>100.0%</td>
<td>1.97 [1.26, 3.25]</td>
<td>-</td>
</tr>
<tr>
<td>Heterogeneity: Tau^2 = 0.22, Chisq = 16.92, df = 6 (P = 0.010), I^2 = 65%</td>
<td>-</td>
<td>-</td>
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<td>-</td>
<td>-</td>
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<tr>
<td>Test for overall effect: Z = 2.67 (P = 0.008)</td>
<td>-</td>
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</table>

Figure 2 Forest plots of the five studies show a shorter overall survival and disease-free survival/progression-free survival/recurrence-free survival in early pancreatic patients with positive pre-treatment or post-treatment circulating tumor cells detection. A: Overall survival; B: Disease-free survival/progression-free survival/recurrence-free survival. CI: Confidence interval.

Figure 3 Forest plot of the three studies that used the CellSearch system shows a decreased disease-free survival/progression-free survival/recurrence-free survival in early pancreatic patients that were positive pre-treatment or post-treatment for circulating tumor cells. CI: Confidence interval.

resection or during chemotherapy predicts a high probability of tumor recurrence or progression[35]. However, the prognostic ability of preoperative CA19-9 is still disputed, since it is mostly detected in advanced stages, it is neither sensitive nor specific enough to identify early-stage patients or for the differential diagnosis of patients at different stages; additionally, it is also positive in many other benign and malignant pancreatic diseases such as pancreatitis, cholestasis, and gastric cancer[7]. In this meta-analysis, the pooled results of the four studies, that we examined, demonstrated that the prognostic effect of CA19-9 in peripheral venous blood on early postoperative recurrence of early-stage pancreatic cancer is not as obvious as that of CTCs.

In the past decade, an increasing number of studies have examined the prognostic value of CTCs in cancers of various organs, as the formation of tumor metastases relies heavily on the survival of CTCs and their ability to mediate angiogenesis in target organs[36]. CTCs are tumor cells shed from both primary and secondary foci and are found circulating in the bloodstream; therefore, they can provide valuable information about primary tumors and secondary deposits. In addition, isolation and in vivo cultures of animal xenografts provide deeper information on individual tumor characteristics[37]. Prospective observational studies have also revealed that CTC numbers rarely drop to zero even after complete resection in both chemo-naïve and post-neoadjuvant patients and can be observed longitudinally before disease recurrence[38]. However, the deficiency of relevant studies and different research designs make the clinical significance of CTCs in early pancreatic cancer prognosis a controversial topic. A thorough analysis of the prognostic performance of CTCs is critical for monitoring and developing treatment strategies for patients with pancreatic cancer.

CTCs in the bloodstream are difficult to capture and identify because their concentration in the circulatory system is extremely low (1-10 cells/10 mL) in most cases[39]. Furthermore, CTCs are scattered among an enormous number of erythrocytes and leukocytes, posing tremendous challenges for their complete collection and accurate detection. To
Figure 4 Sensitivity analysis of the influence of each study on the pooled results for the overall survival and disease-free survival/progression-free survival/recurrence-free survival subgroups. A: Overall survival; B: Disease-free survival/progression-free survival/recurrence-free survival.
Figure 5 Funnel plots were generated for the included studies to determine whether or not publication bias was found for both the overall survival and disease-free survival/progression-free survival/recurrence-free survival subgroups. A: Overall survival; B: Disease-free survival/progression-free survival/recurrence-free survival.

Figure 6 Forest plots of the four studies show that the carbohydrate antigen 19-9 level in the venous blood of early-stage pancreatic cancer patients was not an independent predictor of a shorter time to recurrence. CI: Confidence interval.

Figure 7 Forest plots of the three studies, that detected the carbohydrate antigen 19-9 level in the venous blood before surgery, show that it was not an independent predictor of a shorter time to recurrence for early-stage pancreatic cancer patients. CI: Confidence interval.
metastasis and chemoresistance [45]. In our included studies, Court et al [16] used the microfluidic NanoVelcro CTC chip to evaluate the presence and number of CTCs. This platform greatly improves CTC capture and identification by utilizing anti-EpCAM-coated 3D-nanosubstrates in conjunction with microfluidic chaotic mixers and by adding tumor identification markers, which enhance the synergistic effects of cell–substrate contact frequency as well as its affinity [46]. This platform also allows seamless integration with laser capture microdissection for single CTC isolation [47]; moreover, the separated cells can be subjected to downstream molecular analyses [48]. However, an important limitation of this assay is that it is similar to the CellSearch system. In addition, using membrane filtration may reduce specificity, as some CTCs are found to be equal or smaller in size than nucleated blood cells [49]. The combination of specific mesenchymal markers may be a future direction for these assays. Considering their failure to detect EpCAM- or CK-negative CTCs, Lin et al [50] developed an integrated tumor cell surface molecule-independent SE-iFISH platform in 2015. Using fluorescence in situ hybridization with a specific chromosome centromere probe, this system can detect aneuploidy in the PB, which is a common manifestation of chromosome instability and malignant solid tumors. Moreover, Semaan et al [15] used an antigen-independent approach called DEP-FFF, which utilizes the physical properties of cells and allows the isolation of phenotypically distinct CTCs. In doing so, they obtained not only epithelial and mesenchymal CTCs, but also intermediate-state CTCs, which may show greater invasiveness and therapeutic resistance. Another valid method to detect the molecular characterization of CTCs is PCR. This is most likely due to the detection of multiple tumor markers, which could downstage the effect of the heterogeneity that exists in CTCs. Among the included studies, Cheng et al [17] used immunomagnetic depletion and ligand-targeted polymerase chain reaction to detect the expression rate of folate receptor + CTCs in patients with different stages of pancreatic cancer. Traditional genetic studies of CTCs are limited by the low specificity of the enrichment methods, which contributes to the presence of nuclear blood cells, necrotic cells, tumor-derived exosomes, and cellular fragments. Therefore, the obtained nucleic acids may not accurately reflect the hereditary properties of CTCs [42]. In recent years, single-cell separation and whole-genome amplification of CTCs have been developed to overcome this challenge. However, studies on pancreatic cancer are still lacking.

Although the pooled results of our meta-analysis indicated that CTCs were associated with shorter OS and DFS/PFS/RFS in patients with early pancreatic cancer after surgery, three of the eight included studies showed no significant correlation between the existence of CTCs and both OS and DFS/PFS/RFS. Colin et al [45] found that CTCs were independent predictors of RFS following surgery and could correctly identify patients with occult metastatic disease preoperatively. However, when the analysis was limited to the early-stage subset, CTC count was no longer associated with shorter RFS. One possibility is that they used only 4 mL of PB, which may be too small to detect CTCs in patients with early-stage pancreatic cancer. For example, White et al [18], the CTC number in the PB did not contribute to predicting the OS or RFS of patients after resection, and the CTC number in the PVB was not associated with RFS. However, their results showed complete collinearity between the number of CTCs detected in the PVB and the OS. They hypothesized that this was caused by pancreatic venous drainage, as well as the capture and dilution effects of the liver for CTCs in the portal circulation. Similarly, the research of Bissolati et al [19] showed no significant correlation between the number of CTCs and survival time. However, they found that patients with a positive intraoperative detection of CTCs in the PVB had a higher liver metastasis rate, indicating that CTCs are of great significance for guiding adjuvant chemotherapy and postoperative follow-up monitoring. Overall, these studies revealed that CTCs in the blood of patients with pancreatic cancer are difficult to detect, particularly in early-stage patients.

In contrast, despite the different separation technologies and biomarkers used, the prognostic value of CTCs as a survival indicator for patients with early-stage pancreatic cancer was demonstrated in the remaining five studies. Moreover, they indicated that CTCs could provide tumor characteristics. Xing et al [14] used CD44+ as a marker to detect CTCs, which represent a more stem-like phenotype and can cause tumor metastasis, promoting tumor growth, angiogenesis, and drug resistance. Semaan et al [15] detected the longitudinal characterization of CTC subtypes and described EMT-CTCs as a more aggressive phenotype that is more prone to therapeutic resistance. In addition, the preoperative CTCs level in the PVB of early-stage pancreatic cancer patients may be a better prognostic marker then the preoperative CA19-9 level. This may broaden the selectivity of biomarkers for pancreatic cancer.

Our study has several limitations. First, the small number of included studies as well as the numerous types of separation strategies and research designs limited the power of our analysis. Second, several studies did not provide HR and 95% CI directly; therefore, we estimated them based on Kaplan-Meier curves, which may cause deviations. Finally, four of the eight studies recruited patients with pancreatic cancer at all stages, and we only extracted data about patients with early-stage pancreatic cancer; hence, the information was not detailed.

**CONCLUSION**

This meta-analysis reveals the potential of CTCs as a prognostic biomarker for early-stage pancreatic cancer. Although we rigorously gathered and analyzed the data, the essential limitations of the included studies caused a high degree of heterogeneity and hindered deeper exploration, which reduced the confidence level of our study. To overcome these difficulties, large-scale multicenter cohort studies are urgently needed to explore the full potential of CTCs.
ARTICLE HIGHLIGHTS

Research background
Pancreatic cancer is a terribly invasive and poorly prognosis disease with a five-year survival less than 10%. Recently, more and more studies demonstrated that circulating tumor cells (CTCs) can be a significant prognostic marker of pancreatic cancer.

Research motivation
In this study, we conducted a meta-analysis to analyze the prognostic role of CTCs in patients with pancreatic cancer and investigated whether CTCs can provide prognostic information and assist develop personalized treatment plans.

Research objectives
Our research aims at exploring the predictive effect of CTCs on survival indicators of pancreatic cancer patients in different studies.

Research methods
A standardized literature search of databases was conducted for articles about CTCs published through December 2022. After screening based on inclusion and exclusion criteria, data relevant to prognosis were extracted for analysis. We used a fixed- or random-effect model to calculate the pooled hazard ratio (HR) and 95% confidence interval (CI) of overall survival (OS) and progression-free survival (PFS) according to the degree of heterogeneity.

Research results
Eight eligible studies with a total number of 355 patients with early-stage pancreatic cancer were included. This meta-analysis showed that positive pre-treatment or post-treatment CTCs was associated with shorter OS (HR = 1.93, 95%CI: 1.197-3.126, \( P = 0.007 \)) and decreased relapse-free/disease-free/PFS (HR = 1.27, 95%CI: 1.137-1.419, \( P < 0.001 \)) in patients with early-stage pancreatic cancer. While the CA19-9 level in the portal venous blood of early-stage pancreatic cancer patients showed no significant correlation with postoperative recurrence time of patients (HR = 1, 95%CI: 1.00-1.00, \( P = 0.03 \)).

Research conclusions
Our meta-analysis indicates that CTCs are closely related to the prognosis of early pancreatic cancer patients and can serve as a guiding indicator for developing patient important treatment plans.

Research perspectives
Researchers should extend follow-up time to observe the relationship between CTC and OS. Besides, large-scale multicenter cohort studies are urgently needed to explore the full potential of CTCs.

ACKNOWLEDGEMENTS
We thank Professor Si-Yu Sun, for his support and guidance, as well as all other doctors who participated in this study.

FOOTNOTES
Author contributions: Zhang ZH contributed to the data acquisition and analysis, and article drafting; Bao YW, Zhao YJ, and Wang JQ and Sun SY oversaw acquiring and analyzing data; Guo JT was the study supervisor and was responsible for the revision of the manuscript; all authors read and gave their final approval for this version of the manuscript to be submitted.

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PRISMA 2009 Checklist statement: The authors have read the PRISMA 2009 Checklist, and the manuscript was prepared and revised according to the PRISMA 2009 Checklist.

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Country/Territory of origin: China
REFERENCES


