Review Article ZIP4: a promising early diagnostic and therapeutic targets for pancreatic cancer

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Abstract: Pancreatic cancer is an aggressive and metastatic tumor that lacks effective early detection and treatment methods. There is an urgent need to further understand its underlying molecular mechanisms and identify new biomarkers for early detection. Zinc, a critical trace element and catalytic cofactor, is tightly regulated within cells. ZIP4, a zinc transporter protein significantly overexpressed in human pancreatic cancer, appears to play a pivotal role in tumor development by modulating intracellular zinc concentration. This review highlights the role of ZIP4 in tumorigenesis, including its impact on pancreatic cancer growth, proliferation, migration, and drug resistance. ZIP4 exerts its effects by regulating zinc dependent transcriptional factors like CREB, STAT3, and ZEB1, resulting in upregulation of Cyclin D1, TP53INP1, ITGA3, CD44, ENT1 proteins, and miR-373. Moreover, ZIP4 mediates the miR373-PHLPP2-AKT signaling axis, which increases TGF-β expression. Coupled with CREB-activated macrophage catabolism-related genes SDC1 and DNM2, ZIP4 promotes cancer cachexia and supports amino acids to tumor cells under metabolic stress. Furthermore, ZIP4 facilitates bone resorption by osteoclasts via the RANKL-activated NF-KB pathway. A deeper understanding of these mechanisms may unveil potential targets for early diagnosis, prognosis assessment, and dietary recommendations for pancreatic cancer. These findings hold clinical significance not only for pancreatic cancer but also for other malignancies exhibiting heightened ZIP4 expression.

Keywords: Pancreatic cancer, ZIP4, Zinc, mechanisms, tumorigenesis

Introduction

Pancreatic cancer, notorious for its devastating impact, holds the highest mortality rate among cancers [1]. This grim prognosis is primarily due to its dense, proliferative stroma, a highly hypoxic and immunosuppressive tumor microenvironment, and the absence of effective screening tools and biomarkers. The maximum 5-year survival rate remains at a mere 10%, and by 2030, pancreatic cancer is projected to become the second leading cause of cancerrelated death in the United States [2]. Zinc, an essential trace element, serves as an intracellular secondary messenger that regulates multiple signaling cascades, significantly influencing protein structure, enzyme activity, and gene regulation. Alterations in zinc concentrations in both the serum and malignant tissues of cancer patients have been consistently reported, underscoring its crucial role in cancer biology [3]. The intracellular free zinc concentration must be tightly regulated, typically maintained at pico-to-low nanomolar levels. Zinc transporters are pivotal in preserving intracellular zinc homeostasis by facilitating the uptake of extracellular zinc and its release into the cytoplasm [4]. Emerging evidence emphasizes the crucial role of zinc homeostasis and transport in cancer development, particularly in breast and pancreatic cancers [5, 6]. Notably, ZIP4 stands out as the predominant upregulated zinc transporter protein and the most abundantly secreted exosomal protein in highly malignant pancreatic cancers [7, 8]. This review focuses on the tumor-promoting effects of ZIP4 on pancreatic cancer proliferation, metastasis, drug resistance, and cachexia. By unraveling the underlying pathological mechanisms, we aim to identify novel therapeutic targets and biomarkers for early detection, treatment, tumor staging, and prognosis assessment in pancreatic cancer.

The pathophysiological role of zinc

Zinc, an essential trace element and catalytic cofactor, plays crucial physiological and pathological roles in maintaining normal cell growth and proliferation through its interaction with various transcription factors and metalloenzymes containing zinc finger motifs [9, 10]. Zinc deficiency can induce apoptosis, hinder growth, delay bone development, lead to osteoporosis, impaired DNA synthesis, and immune system malfunction [11, 12]. Dysregulated zinc transporter proteins often disrupt the function of zinc finger transcription factors, including cyclic adenosine monophosphate response element binding protein (CREB), CREB binding protein, and metal-responsive transcription factor-1, resulting in abnormal cellular activity, disrupting the balance of normal growth, and promoting uncontrolled transcription and cell proliferation. Additionally, zinc accelerates the malignant development of tumors as it is a component of vital enzymes, such as carbonic anhydrase and matrix metalloproteinases, both of which are involved in hypoxia, cell proliferation, angiogenesis, and metastasis [13, 14]. However, excessive intracellular zinc is cytotoxic and can induce apoptosis, necessitating a homeostatic system to balance intracellular zinc uptake, storage, and efflux [15, 16]. Two solute linkage carrier gene families, SLC30 and SLC39, encode zinc transporters Zinc Transporter (ZnT) and Zrt- and Irt-like Protein (ZIP), respectively, which play antagonistic roles in regulating zinc availability [17]. ZnTs reduce intracellular zinc by facilitating its efflux from cells or sequestering it into intracellular vesicles, while ZIPs increase intracellular zinc by promoting extracellular uptake and vesicular zinc release into the cytoplasm [4].

Altered zinc transporter expression has been associated with various cancers. Studies have reported reduced zinc concentrations in the serum and malignant tissues of patients with prostate cancer or hepatocellular carcinoma [18, 19]. Breast cancer patients exhibit reduced serum zinc levels but increased intracellular zinc in malignant tissues due to low-level expression of ZnT1 [19-21]. Investigations also link breast cancer-associated proteins like ZIP6 to lymph node metastasis and estrogen-positive breast cancer [22], while ZIP10 has been implicated in breast cancer cell invasion [23].

Differences in serum and tumor tissue zinc concentrations across various cancers suggest tissue-specific zinc uptake mechanisms [4], collectively highlighting the crucial role of zinc transporter in cancer development. Among these transporters, ZIP4, encoded by the SLC39A4 gene, regulates intracellular zinc levels by absorbing dietary zinc and releasing it via vesicular compartments within intestinal epithelial cells [9]. The role of ZIP4 in tumors has gained attention (Table 1). Microarray analyses revealed overexpression of ZIP4 mRNA in human pancreatic cancer tissues. Li M et al. confirmed this through real-time Reverse Transcription Polymerase Chain Reaction (RT-PCR) and immunohistochemical (IHC) staining, showing ZIP4 mRNA levels were approximately 5.5-fold higher in pancreatic cancer cell lines and tissues compared to human pancreatic ductal epithelial cells and surrounding normal tissues. Notably, ZIP4 expression was also elevated in pancreatic cancer tissues compared to chronic pancreatitis tissues [5]. Furthermore, ZIP4 emerged as the most abundantly secreted exosomal protein in highly malignant pancreatic cancers [8]. These findings suggest ZIP4 is a potential contributor to pancreatic cancer pathogenesis and may serve as a valuable diagnostic marker. Overexpressed ZIP4 supplies additional zinc to fuel tumor cell growth and plays a multifaceted role in pancreatic cancer, including proliferation, invasion, metastasis, tumor malignancy grade, cancer cachexia, bone loss, and drug resistance.

ZIP4 promotes pancreatic cancer growth and proliferation and inhibits apoptosis

Zinc has been demonstrated to significantly impact cell proliferation by impacting DNA synthesis and cell cycle progression through deoxythymidine kinase and insulin-like growth factor-I [24]. In conditions of zinc deficiency, overexpressed ZIP4 exacerbates pancreatic cancer cell proliferation. Knockdown of ZIP4 significantly decreases tumor weight and volume in both subcutaneous and in situ models of the ASPC-1 pancreatic cancer cell line [25]. Zhang Y et al. using gene microarray, Reverse Transcription Quantitative Polymerase Chain Reaction, and western blot analysis, unraveled the signaling mechanism underlying ZIP4 mediated pancreatic cancer growth and proliferation [25]. They found that ZIP4 overexpres-

Cancer	Expression of ZIP4	Signal pathway	Outcome	Reference
Hepatocellular carcinoma		ZIP4 regulates the expression of MMP2 and MMP9; down-regulates the expression of pro-apoptotic genes such as cystatinase-3, cystatinase-9, Bax; promotes the expression of Bcl-2.	ZIP4 promotes migration, invasion and inhibits apoptosis in hepatocel- lular carcinoma; and may promote re-entry of hepatocellular carcinoma cells into the cell cycle after release from GO/G1 blockade.	$[76]$
Ocarian cancer		ZIP4 is involved in cellular activities related to HGSOC cancer stem cells through the ZIP4-HDAC4-VEGFA/ZIP4- NOTCH3-Jag-1 axis.	ZIP4 is functionally involved in CSC-associated cellular activities, includ- ing loss-of-nest resistance, colony formation, sphere formation, drug resistance, and LPA-induced EOC side populations in HGSOC cells.	[77, 78]
Glioma		ZIP4 levels are significantly associated with key genes for cell growth and angiogenesis in gliomas, such as VEGFA, MMP9, PDGFA, IGFBP2, IL-6 and IL-8.	High ZIP4 expression is significantly associated with higher glioma grade and shorter overall survival.	$[79]$
Gallbladder cancer		ZIP4 up-regulate CDK4 and c-MET.	ZIP4 promotes Gallbladder cancer cell proliferation and migration and inhibits apoptosis.	[80]
Nasopharyngeal carcinma		ZIP4 activates PI3K/Akt signaling pathway in Nasopha- ryngeal carcinma.	ZIP4 induces the EMT and promotes migration and invasion.	[81]
Non-small cell lung cancer		ZIP4 activates the Snail-N-cadherin signaling pathway and promoting the EMT process.	ZIP4 promoted cell migration, invasion, and metastasis, and its expression level was negatively correlated with overall survival and progression-free survival in NSCLC patients.	[82]
Prostate cancer		NA	ZIP4 has an inhibitory effect on the proliferation and invasion of prostate cancer.	[83]
Breast cancer		NA	ZIP4 promotes tumorigenicity of breast cancer stem cells.	[84]

Table 1. The role of ZIP4 in varies tumors

MMP, matrix metalloproteinase; Bcl-2, B-cell lymphoma-2; HGSOC, High-Grade Serous Ovarian Cancer; HDAC4, Histone Deacetylase 4; VEGFA, Vascular Endothelial Growth Factor A; NOTCH3, Notch homolog 3; Jag-1, Jagged-1; CSC, Cancer Stem Cells; LPA, Lysophosphatidic Acid; EOC, Epithelial Ovarian Cancer; IGFBP2, Insulin-Like Growth Factor Binding Protein 2; IL-6, Interleukin-6; IL-8, Interleukin-8; CDK4, Cyclin-Dependent Kinase 4; c-MET, cellular-mesenchymal to epithelial transition factor; PI3K, Phosphatidylinositol 3-Kinase; EMT, Epithelial-Mesenchymal Transition; NSCLC, Non-Small Cell Lung Cancer; NA, Not Applicable.

sion leads to elevated cyclin D1 levels, a critical regulator of the G1 to S-phase cell cycle transition, thereby promoting cell proliferation. ZIP4 overexpression also upregulates IL-6 expression via CREB activation, subsequently activating signal transducer and activator of transcription 3 (STAT3), leading to cyclin D1 elevating, and promoting pancreatic cancer cell proliferation and tumor progression. Additionally, ZIP4 overexpression significantly boosts microR-NA-373 (miR-373) expression (by 6- to over 200-fold), which suppresses tumor-suppressor molecules like CD44, large tumor suppressor homologue 2 (LATS2), and TP53-inducible nuclear protein 1 (TP53INP1), further driving cancer cell proliferation [26]. Silencing CD44, LATS2, and TP53INP1 resulted in severe ascites, increased tumor weight and metastasis to the peritoneum, liver, spleen, and colon. Most tumors showed poor differentiation with tumor areas exceeding 80% compared to controls. PH domain leucine-rich repeat protein phosphatase 2 (PHLPP2), a phosphatase involved in AKT signaling, is a target gene of miR-373 in pancreatic cancer. It inhibits CREB phosphorylation, establishing a CREB-miR-373-PHLPP2 feed-forward loop that activates the miR-373- PHLPP2-AKT-CyclinD1 signaling axis, driving pancreatic cancer growth [27]. Silencing miR-373 or blocking ZIP4 resulted in G0/G1 phase arrest, a decrease in the S-phase cell population, and significant reductions in tumor weight and size. Beyond promoting proliferation, ZIP4 also inhibits apoptosis. Under zinc-deficient conditions, ZIP4 knockdown increases susceptibility to apoptosis, as shown by flow cytometry analysis. ZIP4 overexpression prevents apoptosis by inhibiting the cleavage of caspase-7, caspase-9, and poly ADP-ribose polymerase, which are key components of apoptotic signaling. This resistance is partially mediated by the caspase-9 activator cytochrome C and the mitochondrial pathway pro-apoptotic protein Bax [28]. ZIP4 may also interfere with other apoptotic pathways, including the tumor necrosis factor (TNF) pathway via caspase-8 and the caspase-independent pathway managed by apoptosis-inducing factor. Impaired apoptosis contributes to both tumor progression and chemoresistance [29, 30]. Defective apoptosis can lead to tumor progression and chemoresistance [31]. Given the reduced zinc levels in the serum and tumor tissues of pancreatic cancer patients, overexpressed ZIP4 enables pancreatic cancer cells to extract zinc from limited sources, supporting excessive proliferation and inhibiting apoptosis [32]. By balancing proliferation and apoptosis, ZIP4 plays a pivotal role in pancreatic cancer growth, highlighting the importance of zinc transport proteins in cancer cells (Figure 1).

ZIP4 promotes pancreatic cancer cell invasion and migration

Distant metastasis within 24 months post-surgery occurs in approximately 60% of pancreatic cancer patients [33, 34], and is a major cause of death [35]. ZIP4 has been demonstrated as a crucial regulator of pancreatic cancer invasion and migration. Silencing ZIP4 in the ASPC-1 pancreatic cancer cell line notably reduced migration (~68%) and invasion (~81%), as confirmed by cell migration and invasion assays [25].

The degradation of the basement membrane is a key step in facilitating tumor cell migration and invasion, which begins with the breakdown of tight junctions between cells [36]. In normal epithelial cells, peripheral proteins like Zonula Occludens-1 (ZO-1) and transmembrane proteins like claudin-1 maintain cell-to-cell adhesion, preventing cancer cell migration [37]. Zinc has been shown to regulate the integrity of these tight junction proteins [38]. IHC staining analysis reveals that Zinc Finger E-Box Binding Homeobox 1 (ZEB1), a major transcription factor driving epithelial-mesenchymal transition (EMT), is overexpressed in pancreatic cancer tissues, while ZO-1 and claudin-1 levels are reduced compared to benign tissues [39]. Further studies indicate that overexpression of ZIP4 in pancreatic cancer cell lines upregulates the mesenchymal marker ZEB1 and reduces the epithelial markers ZO-1 and claudin-1. Through a ZEB1-dependent mechanism, ZIP4 amplifies downstream targets such as Focal Adhesion Kinase (an intracellular non-receptor tyrosine kinase associated with adhesion, migration and invasion) and Paxillin phosphorylate by downregulating claudin-1 and ZO-1, enhancing tumor invasion and metastasis independently of CREB [39].

EMT is a critical process for tumor metastasis, as it induces phenotypic changes that enable cancer cells to migrate, invade the basement membrane, and acquire stem cell-like charac-

Figure 1. The molecular signaling mechanisms and pathways of ZIP4 promoting pancreatic cancer cell proliferation, migration and drug resistance. (1) ZIP4 exerts a pro-muscle atrophic effect by phosphorylating STAT5 through the miR373-PHLPP2-AKT signaling axis to promote the expression of TGF-β. (2) ZIP4 increases the release of pancreatic cancer extracellular vesicles through the CREB-regulated RAB27B HSP70 and HSP90 stimulated the p38 MAPK signaling pathway and induced the expression of F-box protein and UBR2 in myotubes, leading to myofibrillar myosin heavy chain loss and myotube thinning. Meanwhile, ZIP4 activation of SDC1 and DNM2 via CREB and TRAIL secretion mediated through GSK3b could provide the required amino acid supply for tumor cell growth. (3) ZIP4 promotes the binding of RANKL to RANK on osteoclast membranes and facilitates bone resorption through activation of the NF-κB pathway, leading to decreased mineral density of bone tissue, increased bone crystallinity and bone strength loss.

teristics [40, 41]. ZEB1, apart from its role as a transcriptional repressor, can transition to a transcriptional activator by interacting with coactivators (Lef1, YAP1, P300, and Smad) regulating tumor progression via pathways like Hippo and Wnt [42-44]. The Hippo pathway plays an essential role in regulating cell shape, organ size, EMT, and tumorigenesis. YAP1, the major downstream effector of the Hippo pathway, has been shown to promote tumor growth in various cancers, including pancreatic, lung, and colon cancers. LATS2 can directly phosphorylate YAP1, inhibiting its activity through protein degradation. ZIP4 upregulates YAP1 expression by inhibiting LATS2 via miR-373, thereby activating downstream oncogenes [45- 47]. YAP1 also acts as a co-activator by forming a complex with ZEB1, binding to the ITGA3 promoter region to enhance EMT plasticity, cellular adhesion, sphere formation, and tumor sphere formation [48]. ZEB1's dual role - lower expression contributing to tumorigenesis and higher expression promoting metastasis - suggests its potential as a marker to distinguish between tumorigenesis and the EMT/metastasis stage [49]. Moreover, ZIP4 overexpression significantly elevates the Vascular Endothelial Growth Factor (VEGF), Neuropilin-1 (NRP-1, a receptor or co-receptor for specific isoforms of VEGF), matrix metalloproteinase (MMP)-9 and MMP-2 expression, which are critical molecules in angiogenesis, invasion, and metastasis within the ZIP4-initiated signaling cascade [50]. The specific regulatory mechanisms for NRP-1, VEGF, MMP-9, and MMP-2 in response to ZIP4 necessitate further investigation to comprehensively understand their role in ZIP4-mediated pancreatic cancer cell invasion and migration.

ZIP4 and drug resistance

Breakthroughs in pancreatic cancer treatment remain elusive, with gemcitabine-based monotherapy or combinations serving as the standard for advanced cases. Despite extensive exploration, no combination therapy has significantly outperformed gemcitabine alone [51]. Disease progression often leads to gemcitabine resistance, culminating in cancer recurrence [52]. Liu M et al. demonstrated that ZIP4 expression influences both the growth and chemoresistance of pancreatic cancer using a three-dimensional tumor culture model based on spheroids. When ZIP4 was overexpressed, MIA-PaCa-2 cell line enhanced resistance to cisplatin, 5-Fluorouracil (5-FU), and gemcitabine. suggesting ZIP4 as a potential predictor of chemoresistance in pancreatic cancer patients [51]. Their subsequent investigation delineated the regulatory cascade through which ZIP4 modulates drug resistance: ZIP4 phosphorylates STAT3, which subsequently upregulates ZEB1. ZEB1 increases the expression of integrin subunits ITGA1 and ITGB3, leading to the formation of integrin α3β1. This integrin complex downregulates equilibrium transporter protein 1 (ENT1) via c-Jun N-terminal Kinase mitogen-activated protein kinase (JNK MAPK) activation [51], reducing cellular uptake of gemcitabine and contributing to drug resistance [53]. Elevated ENT1 expression in pancreatic cancer typically indicates responsiveness to gemcitabine therapy, but ENT1 downregulation or deletion leads to gemcitabine resistance [54, 55]. Additionally, ZEB1 inhibits miR-203, further contributing to drug resistance in pancreatic cancer [56].

Beyond integrin α3β1, integrin α2β1 enhances pancreatic cancer cell resistance to 5-FU through B-cell lymphoma 2 (Bcl-2) upregulation. ITGB1 promotes gemcitabine tolerance by activating Cdc42, while collagen/ITGA1 signaling plays a crucial role in gemcitabine tolerance [57, 58] (Figure 2). These findings reveal unconventional pathways regulating metastasis and chemoresistance in pancreatic cancer and

offer potential targets for future therapeutic strategies.

ZIP4 and cancer cachexia

Cachexia, characterized by progressive muscle wasting, fat loss, reduced appetite, and weight loss [59-61], affects over 80% of pancreatic cancer patients and is resistant to refractory to conventional nutritional support [62, 63]. This significant increase in muscle protein catabolism not only compromises patients' quality of life but also impairs their tolerance and responsiveness to chemotherapy, thus contributing to tumor progression [64, 65]. ZIP4, along with its upstream and downstream regulatory molecules, plays a crucial role in pancreatic cancer malignancy. Studies have shown that ZIP4 enhances muscle proteolysis, evidenced by significantly reduced tibialis anterior muscle mass and cross-sectional area in animal models with high ZIP4 expression compared to ZIP4 knockdown groups [27, 66]. ZIP's pro-muscle atrophy effect is mediated through the phosphorylation of STAT5 via the miR373-PHLPP2-AKT signaling axis, which promotes the expression of the soluble malignant plasma factor Transforming Growth Factor-β (TGF-β) [27]. CircANAPC7, a ZIP4-induced miR373 sponge, not only suppresses ZIP4-mediated cell proliferation through the miR373-PHLPP2-AKT-CyclinD1 signaling axis but also effectively ameliorates muscle atrophy in pancreatic cancer by inhibiting the miR373-PHLPP2-AKT-STAT5-TGF-β signaling axis. Mice with high circANAPC7 expression exhibited increased levels of myofibrillar protein and myosin heavy chain, alongside reduced expression of muscle atrophy markers such as Ubiquitin Protein Ligase E3 Component N-Recognin 2 (UBR2) protein, Muscle Atrophy F-box protein-1, and Muscle RING-finger protein-1. These mice also demonstrated improved grip strength, increased muscle weight, and enhanced cross-sectional area of muscle fibers [27]. This was supported by Kaplan-Meier survival curves, wherein circANAPC7 significantly increased median survival in pancreatic cancer xenograft mice compared to controls, underscoring the role of circANAPPC7 and ZIP4 in pancreatic cancer growth and malignancy.

Additionally, Yang J et al. demonstrated that ZIP4 enhances the release of heat shock proteins (HSP) HSP70 and HSP90 from pancreatic

ing pathways through zinc finger transcription factors CREB, STAT3, and ZEB1 leading to cyclinD1 up-regulate and through miR-373 negatively regulating tumor suppressor molecules TP53INP1, LATS2 and CD44 to promote pancreatic cancer proliferation and migration. And ZIP4 promotes the activation of MAP kinase JNK by integrin α3β1 through ZEB1 down-regulation of ENT1 leading to gemcitabine resistance.

cancer extracellular vesicles (EVs) through CREB-regulated RAB27B, a GTPase necessary for controlling EV release [66]. These EV-associated proteins act as danger-associated molecular patterns, activating toll-like receptor 4 and stimulating the p38 MAPK pathway as well as the expression of F-box and UBR2 protein in myotubes, leading to thinning of myotubes and loss of myofibrillar myosin heavy chain [66]. Moreover, acetyl coenzyme A synthetase short-chain family member 2 (ACSS2), an enzyme involved in lipid synthesis, is overexpressed in pancreatic intraepithelial neoplasia lesions and pancreatic cancer tissues. ACSS2 promotes ZIP4 transcription by modifying the Ets Variant 4 promoter with H3K27ac histones. ZIP4, in turn, induces macropinocytosis through CREB-activated macropinocytosis catabolismassociated genes like Dynamin-2 (DNM2) and Syndecan-1 (SDC1), facilitating amino acids supply for tumor cell growth under metabolic stress states [67]. Notably, the downregulation

of ZIP4 had a less pronounced effect on macropinocytosis compared to ACSS2 downregulation, suggesting that additional downstream signals of ACSS2, independent of ZIP4, may play a role in this process. ZIP4 also contributes to pancreatic cancer cachexia by secreting tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) 10 through the glycogen synthase kinase-b (GSK3b) pathway, leading to muscular atrophy [67].

In addition to its role in cachexia, ZIP4 has been implicated in bone loss associated with malignancy. ZIP4 promotes bone resorption by binding to Receptor Activator of Nuclear Factor Kappa-Β Ligand (RANKL), a member of the TNF superfamily, which interacts with RANK on osteoclast membranes. This activates osteoclasts, and initiates the NF-κB pathway, resulting in reducing bone mineral density, increasing bone crystallinity, and compromising bone strength in situ xenograft mouse models [68].

Zinc and ZIP4 as therapeutic targets

Studies have shown that serum zinc levels are significantly reduced in pancreatic cancer patients, and zinc content in pancreatic tumor tissue is lower compared to adjacent normal tissue [32]. Furthermore, monitoring changes in trace element concentrations during pancreatic cancer treatment revealed that patients with decreasing zinc levels had higher mortality rates than those whose levels remained stable or increased [69]. Patients with larger reductions in zinc levels during treatment exhibited higher mortality rates compared to those with smaller reductions, and individuals whose zinc status shifted from normal to deficient had an increased risk of death [69]. The decline in zinc levels has been associated with cancer progression and poor outcomes, aligning with the overexpression of ZIP4 in human pancreatic cancer.

Numerous studies have highlighted the inhibitory impact of ZIP4 silencing on pancreatic cancer growth in subcutaneous and in situ xenograft models in nude mice [25, 39, 51]. ZIP4 silencing significantly reduced primary tumor weights, jaundice, peritoneal dissemination, hepatic and lung metastases, colon/intestinal obstruction, ascites, and weight loss, while also improving survival in *in situ* xenograft nude mice [25]. Histological analyses also indicated a reduction in tumor grade following ZIP4 silencing, potentially increasing tumor responsiveness to subsequent treatments. Consequently, combining ZIP4 short hairpin RNA with chemotherapy or radiotherapy may enhance the effectiveness of tailored therapies. Additionally, identifying ZIP4 expression profiles before and after pancreatic cancer treatment could guide the selection of initial and sequential ZIP4 therapies [25]. Tan X et al. were the first to demonstrat the utility of exosomal ZIP4 as a novel biomarker for pancreatic cancer diagnosis, leveraging clinical blood samples from healthy individuals and patients with malignant pancreatic cancer, benign pancreatic disease, or biliary tract disease. ZIP4 effectively distinguished malignant pancreatic cancer patients from both healthy controls (AUC= 0.8931) and patients with benign pancreatic diseases (AUC=0.89) [8], highlighting ZIP4's potential as a valuable diagnostic marker for pancreatic cancer.

Conclusion and perspectives

Pancreatic cancer is diagnosed late in over 80% of cases, hindering timely intervention due to the absence of distinct symptoms, rendering surgery unfeasible in most instances [70]. Approximately 15% of newly diagnosed cases present with non-metastatic and resectable tumors, with recurrence occurring within a year for most patients [71, 72]. This highlights the urgent need for novel biomarkers that facilitate early diagnosis and treatment of pancreatic cancer. This review outlines various mechanisms wherein ZIP4 contributes to pancreatic cancer development and proposes avenues for further exploration. (1) We have summarized how ZIP4 promotes pancreatic cancer growth, proliferation, migration, chemoresistance, and cachexia. Some of its tumor-promoting effects are attributed to the disruption of tight junctions, promotion of EMT, and resistance to gemcitabine chemotherapy. However, studies have shown that ZIP4 also regulates the expression of MMP and increases resistance to other chemotherapeutic agents, such as 5-FU and cisplatin, although the exact mechanisms remain unclear. Therefore, further investigation is needed, such as how ZIP4 regulates MMP to promote tumor metastasis and the ZIP4's synergistic or inhibitory effects in combination with gemcitabine-based therapies; (2) ZIP4's biolog-

ical effects are mediated through major transcriptional activators including CREB, STAT3, and ZEB1, making them potential therapeutic targets. However, effective methods for precisely targeting ZIP4 or its downstream transcriptional activators and delivering such therapies need further development; (3) Given that zinc levels in ZIP4-overexpressing tumor cells remain lower than those in surrounding normal pancreatic tissue, we hypothesize that this may be due to excessive zinc consumption required for tumor survival and growth. However, the exact reasons for this phenomenon remain to be explored. It's generally believed that zinc can exert anti-tumor activity by improving immune function, reducing oxidative stress, and promoting DNA damage repair [73, 74]. Thus, exploring strategies to harness high levels of zinc in pancreatic cancer to induce anti-tumor activity might be more beneficial than directly silencing ZIP4; (4) Given ZIP4's role in zinc uptake, heavy meat diets, particularly red and double-cooked meat, cause excessive zinc absorption in humans, which a recognized risk factor for pancreatic cancer development [75]. This suggests that dietary recommendations could be valuable for pancreatic cancer patients. Carefully designed, large-scale cohort studies are necessary to evaluate zinc status and its implications for early diagnosis and prognosis in patients with pancreatic cancer. In conclusion, Zinc and ZIP4 may serve as promising diagnostic, therapeutic, and prognostic biomarkers for pancreatic cancer. Targeting ZIP4 with innovative treatments may offer promising treatment strategies for not only pancreatic cancer but also other malignancies displaying elevated ZIP4 expression.

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Disclosure of conflict of interest

None.

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