



Primary and Secondary Prevention of Pancreatic Cancer

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Abstract

Purpose of Review Pancreatic cancer remains one of the most lethal of malignancies with 5-year survival of only 8%. A number of reasons account for the high fatality rate including few known modifiable risk factors, no effective screening tools, and lack of early diagnostic symptoms. Therefore, in this review, we aim to summarize existing evidence from major studies concerning (1) risk factors for risk assessment and risk stratification, and (2) screening modalities and early detection markers to better understand the ways to prevent pancreatic cancer or identify it at earlier stages. Improvements in primary and secondary prevention of pancreatic cancer are critical to reduce the morbidity and mortality of this deadly disease.

Recent Findings We searched the published literature and identified studies of pancreatic cancer risk published prior to September 30, 2018, with an emphasis on manuscripts publicized during the last 5–10 years. Known and suspected risk factors include familial and genetic risk, smoking, obesity, alcohol, poor diet including sugary sweetened beverages, diabetes, and periodontal disease. Recent advances have identified potential early detection markers (e.g., ctDNA, circulating cancer cells, metabolites, and miRNA).

Summary Currently, pancreatic cancer has few known and suspected risk factors, and risk assessment tools have limited utility given their modest discriminatory power. Although emerging evidence suggests blood-based biomarkers may be useful as early detection markers, findings need to be confirmed in prospective studies. Due to the rarity of disease, future studies should consider a two-tiered approach in which risk assessment is used to identify high-risk individuals for screening, and then effective imaging and biomarkers in pathways known to affect pancreatic cancer risk are employed; these combination approaches may reduce false positives and mortality compared with just risk assessment or screening alone.

Keywords Pancreatic cancer · Risk factors · Early detection

Introduction

Pancreatic cancer remains one of the most lethal of malignancies. An estimated 232,306 cases of pancreatic cancer occur globally each year, and 227,023 die from the disease [1]. Since 2004, incidence rates have increased by 1.5% per year in the

USA [2]. This is extremely alarming given that half of the individuals diagnosed with pancreatic cancer die within 6 months and that the 5-year survival for pancreatic cancer is 8% [3]. While the 5-year survival rate improves to 37.4% in patients presenting with stage 1 or localized disease, only 10% of patients are identified at this early stage [4]. The majority of patients (53%) are diagnosed with distant, metastatic cancer, and have a 5-year survival of 2.9% [5]. A number of reasons exist for the late diagnosis and high fatality rate, including few known modifiable risk factors, no effective screening tools, and lack of early diagnostic symptoms unique to pancreas cancer. Thus, approaches to prevent disease or identify it at earlier stages (e.g., stage 1a) are critical to reduce the morbidity and mortality of this deadly disease. Therefore, in this review, we aim to review the recent literature with regard to the (1) identification for risk factors for risk assessment and risk stratification (Table 1 and Fig. 1) and (2) identification of screening modalities and early detection markers.

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Table 1 A summary of epidemiologic studies examining the association between risk factors and pancreatic cancer risk

Risk factor	Study, country, year	Study population	Study design	Main findings
Familial and genetic risk	Molina-Montes E et al., Europe, 2018	1431 cases and 1090 controls	Case-control	Pancreatic cancer is associated with 2 first degree relatives of pancreatic cancer (OR = 3.88; 95% CI 2.97–9.72) Pancreatic cancer is associated with family history of cancer (OR = 1.30; 95% CI 1.13–1.54) Pancreatic cancer is associated with family history of diabetes (OR = 1.24; 95% CI: 1.01–1.52)
	Ghadirian P et al., Canada, 2002	174 cases and 136 controls	Case-control	Pancreatic cancer is associated with family history of pancreatic cancer (relative risk = 5.0; $p = 0.01$)
	Fehringer G et al., Canada, 2014	690 cases and 736 control	Case-control	Pancreatic cancer is associated with family history of pancreatic cancer (OR = 1.68; 95% CI 1.03–2.72) Pancreatic cancer is associated with one sibling history of pancreatic cancer (OR = 5.22; 95% CI 2.03–13.41)
	Permuth-Wey J et al., 2009	2617 cases and 6284 controls in 7 case-control study 3951 cases and 1,104,508 participants in 2 cohort study	Meta-analysis	Pancreatic cancer is associated with family history of pancreatic cancer (summary RR = 1.80; 95% CI 1.48–2.12)
Smoking	Klein AP et al., Europe, 2018	11,537 cases and 17,107 controls from Pancreatic Cancer Cohort Consortium (PanScan) and the Pancreatic Cancer Case-Control Consortium (PanC4)	Genome-wide meta-analysis	New genome-wide significant loci: rs13303010 at 1p36.33 (NOC2L, OR = 1.26; 95% CI 1.19–1.35, $P = 8.36 \times 10^{-14}$) rs2941471 at 8q21.11 (HNF4G, OR = 0.89; 95% CI 0.85–0.93, $P = 6.60 \times 10^{-10}$) rs4795218 at 17q12 (HNF1B, OR = 0.88; 95% CI 0.84–0.92, $P = 1.32 \times 10^{-8}$)
	Zheng Z. et al., China, 2016	323 case-control pairs	Case-control	Pancreatic cancer is associated with smoking (OR = 1.78, 95% CI 1.02–3.10) but frequency and quantities were not statistically significant
	Wang Y. et al., China, 2014	307 cases and 1228 controls	Case-control	Current smokers vs. never smokers (OR = 1.71; 95% CI 1.25–2.35) Pancreatic cancer is associated with smoking intensity (≥ 20 cigarettes/day) (OR = 2.18; 95% CI 1.66–3.69)
	Bosetti C. et al., 2012	6507 cases and 12,890 controls from 12 case-control studies within the International Pancreatic Cancer Case-Control Consortium (PanC4)	Pooled Analysis	Current smokers vs. never smokers (OR = 2.2; 95% CI 1.7–2.8) Pancreatic cancer is associated with smoking intensity (≥ 35 cigarettes/day) (OR = 3.4; P for trend < 0.0001)
Obesity	Zou L. et al., 2014	9044 cases and 32,039 controls from 30 retrospective studies 8962 cases and 2,075,947 participants in 12 prospective studies	Meta-analysis	Compared to non-smokers: 10 cigarettes/day (OR = 1.5; 95% CI 1.4–1.6) 20 cigarettes/day (OR = 1.9; 95% CI 1.8–2.0) 30 cigarettes/day (OR = 2.0; 95% CI 1.9–2.1) 40 cigarettes/day (OR = 2.1; 95% CI 1.9–2.3)
	Jiao L. et al., 2010	2454 cases and 943,759 participants from 7 cohort studies	Pooled analysis	Overweight vs. normal weight (RR = 1.13; 95% CI 1.03–1.23) Obese vs. normal weight (RR = 1.19 95% CI: 1.05–1.35)
	Arislan AA et al., 2010	2170 cases and 2209 controls from PanScan consortium	Pooled analysis	BMI: highest vs lowest BMI quartile (OR = 1.33; 95% CI 1.12–1.58) (OR = 1.33; 95% CI 1.04–1.69) among men (OR = 1.34; 95% CI 1.05–1.70) among women Waist-to-hip ratio (WHR): highest vs. lowest WHR quartile among women (OR = 1.87; 95% CI 1.31–2.69); but not significant among men
	Genkinger JM et al., 2011	2135 cases and 846,340 participants from 14 cohort studies	Pooled analysis	BMI: Early adulthood BMI ≥ 25 kg/m ² vs. BMI 21–22.9 kg/m ² (RR = 1.30, 95% CI 1.09–1.56) WHR: highest vs lowest WHR quartile (RR = 1.35; 95% CI 1.03–1.78)
Alcohol	Genkinger JM et al., 2015	1947 cases and 647,478 participants from 20 prospective in the National Cancer Institute BMI and Mortality Cohort Consortium	Pooled analysis	WHR (HR = 1.09; 95% CI 1.02–1.17 per 0.1 increase, pancreatic cancer mortality). Waist circumference (HR = 1.07; 95% CI 1.00–1.14 per 10 cm increase, pancreatic cancer mortality) BMI in early adulthood (HR = 1.18; 95% CI 1.11–1.25 per 5 kg/m ² increase, pancreatic cancer mortality)
	Genkinger JM et al., 2009	2187 cases and 862,664 participants from 14 cohort studies	Pooled analysis	Pancreatic cancer is associated with alcohol (RR = 1.22; 95% CI 1.03–1.45, > or = 30 vs. 0 g/day)

Table 1 (continued)

Risk factor	Study, country, year	Study population	Study design	Main findings
Diet	Lucenteforte et al., 2012	5585 cases and 11,827 controls from PanC4	Pooled analysis	No association between light-to-moderate alcohol consumption (≤ 4 drinks per day) and pancreatic cancer risk. Pancreatic cancer is associated with higher consumption level (drinking ≥ 9 drinks per day) (OR = 1.6; 95% CI 1.2–2.2)
	Michaud et al., 2010	1530 cases and 1530 controls from PanScan	Pooled analysis	No association between total alcohol intake (≥ 60 g/day vs. 0–5 g/day) and pancreatic cancer risk (OR = 1.38; 95% CI 0.86–2.23)
	Inoue-Choi M. et al., USA, 2011	256 cases and 34,642 participants from Iowa Women's Health Study	Cohort	No association between dietary intake of any nutrients or food groups and pancreatic cancer
	Nöthlings U. et al., 2008	610 cases and 183,513 participants from Multiethnic Cohort (MEC) study 517 cases and 424,978 participants from EPIC study	Cohort	No significant association between food group or food item pattern and pancreatic cancer risk in the overall cohort
Physical activity	Bosetti et al., 2013	326 cases and 652 controls	Case-control	Pancreatic cancer is associated with “animal products” pattern (OR = 2.03; 95% CI 1.29–3.19) and “starch-rich” pattern (OR = 1.69; 95% CI: 1.02–2.79). Pancreatic cancer is inversely associated with “vitamins and fiber” pattern (OR = 0.55; 95% CI 0.35–0.86). No association between “unsaturated fats” pattern (OR = 1.13; 95% CI 0.71–1.78) and pancreatic cancer
	Chan et al., 2013	532 cases and 1701 controls	Case-control	Pancreatic cancer is inversely associated with the prudent dietary pattern (greater intake of vegetables, fruit, fish, poultry, whole grains, and low-fat dairy) (OR = 0.51; 95% CI 0.31–0.84, p-trend = 0.001 among men); (OR = 0.51; 95% CI: 0.29–0.90, p-trend = 0.04 among women). Pancreatic cancer is associated with the western dietary pattern (higher intake of red and processed meats, potato chips, sugary beverages, sweets, high-fat dairy, eggs, and refined grains) (OR = 2.4; 95% CI 1.3–4.2, p trend = 0.008 among men) but was not associated with risk among women.
	Michaud et al., 2005	366 cases and 124,672 participants from 2 cohorts	Pooled analysis	No associations between the prudent pattern (RR = 1.32; 95% CI 0.66–2.63) or the western pattern (RR = 0.91; 95% CI 0.57–1.47) and pancreatic cancer.
	Behrens et al., 2015	10,501 cases	Meta-analysis	High vs. low levels of physical activity (RR = 0.93; 95% CI: 0.88–0.98) among 22 cohort studies and (RR = 0.78; 95% CI 0.66–0.94) among 8 case-control studies.
Physical activity	Brenner et al., 2014	826 cases and 930 controls	Case-control	Pancreatic cancer is associated with regular leisure-time physical activity (OR = 0.65; 95% CI: 0.52–0.87)
	Calton et al., 2008	70 cases and 33,530 participants	Cohort	The RRs for increasing quartiles of total physical activity were 1.0, 0.80, 0.66, 0.52 (95%CI 0.26–1.05, p trend = 0.05)
	Hanley et al., 2001	312 cases and 2919 controls	Case-control	Pancreatic cancer is inversely associated with composite moderate and strenuous physical activity among men (OR = 0.53; 95% CI 0.31–0.90). No association between physical activity and pancreatic cancer among women, though a tendency for reduced risk with increasing levels of strenuous activity was suggested (p for trend = 0.06)
	Berrington de Gonzalez et al., 2006	324 cases and 438,405 participants	Cohort	No significant decreased risk with total physical activity (RR = 0.82; 95% CI 0.50–1.35)
Physical activity	Nothlings et al., 2007	472 cases and 167,430 participants	Cohort	No association between physical activity and pancreatic cancer
	Patel et al., 2005	242 cases and 145,627 participants	Cohort	No association between physical activity and pancreatic cancer

Table 1 (continued)

Risk factor	Study, country, year	Study population	Study design	Main findings
Reproductive factors	Stolzenberg-Solomon et al., 2008	654 cases and 495,035 participants	Cohort	No association between physical activity and pancreatic cancer (RR = 1.20; 95% CI 0.63–2.27)
	Andersson et al., 2018	110 cases and 17,035 participants	Cohort	No association between physical activity and pancreatic cancer
	Prizment et al., 2007	228 cases and 37,459 participants	Cohort	No association between age at menopause or first childbirth, parity and breastfeeding history and pancreatic cancer
	Skinner et al., 2003	243 cases and 115,474 participants	Cohort	Pancreatic cancer is inversely associated with age at menopause (HR = 0.61, 95% CI 0.40–0.94 for menopause at 45–49 years; HR = 0.75, 95% CI 0.51–1.09 for 50–54 years; and HR = 0.35, 95% CI 0.18–0.68 for menopause at 55 years or more; <i>P</i> trend = 0.005) No associations between age at first birth, number of births, age at menarche, or use of hormones and pancreatic cancer
	Navarro Silvera et al., 2016	187 cases and 89,835 participants	Cohort	Pancreatic cancer is inversely associated with ≥ 5 births vs. nulliparous (RR = 0.58, 95% CI 0.34–0.98)
Medical history	Luo et al., 2016	2535 cases from 5 case-control studies and 6 cohort studies	Meta-analysis	Pancreatic cancer is associated with postmenopausal (OR = 2.44, 95% CI 1.45–4.09) No associations between age at first live birth, parity, age at menarche, use of oral contraceptive, and use of hormone replacement therapy (HRT) and pancreatic cancer. However, among parous women, risk increased with increasing parity.
	Stevens et al., 2009	1182 cases and 995,192 participants	Cohort	Pancreatic cancer is associated with age at first birth (RR = 1.21, 95% CI 1.01–1.45, highest versus lowest categories)
	Lujan-Barroso et al., 2016	2838 cases and 4748 controls from Panc4	Pooled analysis	No significant association between age at menarche, number of children, age at first birth, breast-feeding, type of menopause, age at menopause, or time since menopause and pancreatic cancer
	Kirkegard et al., 2018	41,669 acute pancreatitis and 208,340 comparison. 937 pancreatic cases through follow up period	Cohort	Pancreatic cancer is inversely associated with hysterectomy (OR = 0.78, 95% CI 0.67–0.91)
	Raimondi et al., 2010	12 case-control studies and 10 cohort studies	Meta-analysis	Pancreatic cancer is associated with acute pancreatitis (HR = 2.02, 95% CI 1.57–2.61)
	Pang et al., 2017	595 cases and 512,000 participants for cohort study	Cohort and meta-analysis	Pancreatic cancer is associated with all types of pancreatitis (RR = 5.1, 95% CI: 3.5–7.3 for unspecified pancreatitis; RR = 13.3, 95% CI: 6.1–28.9 for chronic pancreatitis; RR = 69.0, 95% CI 56.4–84.4 for hereditary pancreatitis.
	Bosetti et al., 2014	8305 cases and 13,987 controls	Pooled analysis	Pancreatic cancer is associated with diabetes (HR = 1.87, 95% CI 1.48–2.37 for cohort study; HR = 1.52, 95% CI 1.43–1.63 for meta-analysis)
	Haugvik et al., 2015	827 cases and 2407 controls	Meta-analysis	Pancreatic cancer is associated with diabetes 20 or more years after diabetes diagnosis (OR = 1.30, 95% CI 1.03–1.63)
	Walker et al., 2015	536 cases and 869 controls	Case-control	Pancreatic cancer is associated with diabetes (OR = 2.74, 95% CI 1.63–4.62)
	Lu et al., 2015	529 cases and 5000 controls	Nested case-control	No association between metformin and pancreatic cancer
Wang et al., 2014	10 cohort studies and 3 case-control studies	Meta-analysis	No association between metformin and pancreatic cancer	
Hamada et al., 2018	583 cases and 113,059 participants	Cohort	Pancreatic cancer is inversely associated with metformin use (RR = 0.63, 95% CI 0.46–0.86)	

Table 1 (continued)

Risk factor	Study, country, year	Study population	Study design	Main findings
	Chiu et al., 2011	190 cases and 760 control	Case-control	No association between statin use and pancreatic cancer (HR = 0.98, 95% CI 0.82–1.16)
	Walker et al., 2015	536 cases and 869 controls	Case-control	No association between statin use and pancreatic cancer (OR = 0.87, 95% CI 0.56–1.36) Pancreatic cancer is inversely associated with statin use among men (OR = 0.50, 95% CI 0.32–0.79); no association between statin use and pancreatic cancer among women (OR = 0.86, 95% CI 0.52–1.43)
	Cui et al., 2012	7807 cases and 1,692,863 participants from 16 studies	Meta-analysis	No significant association between statin use and pancreatic cancer (RR = 0.89, 95% CI 0.74–1.07)
	Archibugi et al., 2018	11,975 cases and 3,433,175 controls from 27 studies	Meta-analysis	Pancreatic cancer is inversely associated with statin use (OR = 0.70, 95% CI 0.60–0.82)
	Ben et al., 2012	943 cases and 1128 controls	Case-control	Pancreatic cancer is associated with hepatitis B (OR = 1.60, 95% CI 1.15–2.24)
	Xu et al., 2013	3758 cases and 744,120 controls	Meta-analysis	Pancreatic cancer is associated with hepatitis B (OR = 1.20, 95% CI 1.01–1.39), Pancreatic cancer is associated with hepatitis C (OR = 1.26, 95% CI 1.03–1.50)
	Krull et al., 2016	116 cases and 20,360 participants	Cohort	No significant association between hepatitis B (OR = 1.22, 95% CI: 0.81–1.84) or hepatitis C (OR = 0.69, 95% CI: 0.28–1.69) and pancreatic cancer
	Risch et al., 2014	761 case and 794 controls	Nested case-control	Pancreatic cancer is inversely associated with CagA seropositive individuals (OR = 0.68, 95% CI 0.54–0.84)
	Huang et al., 2017	448 cases and 448 controls	Nested case-control	No association between <i>H. pylori</i> seropositivity (OR = 0.96, 95% CI: 0.70–1.31) or CagA seropositivity (OR = 1.07, 95% CI: 0.77–1.48)
	Liu et al., 2017	65,155 participants	Meta-analysis	No association between <i>H. pylori</i> infection (OR = 1.09, 95% CI 0.81–1.47) or CagA seropositivity (OR = 1.18, 95% CI 0.80–1.72) and pancreatic cancer
	Schulte et al., 2015	580 cases and 626 controls	Nested case-control	No association between <i>H. pylori</i> seropositivity (OR 1.00, 95% CI: 0.74–1.35) or CagA seropositivity (OR 0.74, 95% CI 0.48–1.15) and pancreatic cancer
	Fan et al., 2018	361 cases and 371 controls	Nested case-control	Pancreatic cancer is associated with <i>Porphyromonas gingivalis</i> (OR = 1.60, 95% CI 1.15–2.22), <i>Aggregatibacter actinomycetemcomitans</i> (OR = 2.20, 95% CI 1.16–4.18) and pancreatic cancer
	Huang et al., 2016	126 cases and 19,924 participants	Cohort	Pancreatic cancer is associated with unacceptable dental plaque (HR = 2.1, 95% CI: 1.0–4.7)
	Michaud et al., 2013	405 cases and 416 controls	Nested case-control	Pancreatic cancer is associated with Porphyromonas gingivalis ATTC 53978 (OR = 2.14, 95% CI 1.05–4.36; > 200 ng/ml vs ≤ 200 ng/ml)
	Michaud et al., 2007	216 cases and 51,529 participants	Cohort	Pancreatic cancer is associated with history of periodontal disease (RR = 1.64, 95% CI 1.19–2.26)

Risk Factors for Pancreatic Cancer

Familial and Genetic Risk

Genetic variation, both familial and sporadic, plays an important role in pancreatic cancer. Family history of any cancer has been associated with a 15–30% higher pancreatic cancer risk [6, 7]; risk is stronger when considering only a family history of pancreatic cancer, with risk ratios of 1.68 for any relative, 3.88 for at least two first-degree relatives, and up to fivefold for when an individual has an affected sibling [6, 8]. A meta-analysis consisting of seven case-control studies and two cohort studies reported an association between having a family history of pancreatic cancer and pancreatic cancer risk (summary RR = 1.80, 95% CI 1.48–2.12). Increased risks were observed when considering the number of first degree relatives affected (summary RR = 4.6, 95% CI 0.5, 16.4; summary RR = 6.4, 95% CI 1.8–16.4; summary RR = 32.0, 95% CI 10.2–74.7, for one relative, two relatives, or three relatives affected, respectively) [9].

The associations between family history and pancreatic cancer risk suggest that there are genes of varying penetrance that influence the pancreatic carcinogenesis. Several genes have been implicated in pancreatic cancer risk (e.g., BRCA1, BRCA2, PALB2, ATM, CDKN2A, APC, MLH1, MSH2, MSH6, PMS2, PRSS1, and STK11), as well as the ABO genotype [10•]. A recent meta-analysis conducted with the largest pancreatic cancer GWAS that included up to 11,537 cases and 17,107 controls observed several new genome-wide significant loci. Specifically, SNPs located on the *NOC2L* gene were statistically significantly associated with pancreatic cancer risk (OR = 1.26, 95% CI 1.19–1.35, $P = 8.36 \times 10^{-14}$) [11•]. Additionally, genetic syndromes have been shown to be associated with a 4–40% increased pancreatic cancer risk such as familial atypical multiple mole melanoma, Peutz-Jeghers syndrome, hereditary pancreatitis, hereditary nonpolyposis colon cancer, and multiple endocrine neoplasia type 1 syndrome [12]. However, family history and/or genetic predisposition is reported to account for only 5–10% of all pancreatic cancer cases in the US; estimates have remained stable over the last few decades [10•, 13, 14].

Lifestyle Factors

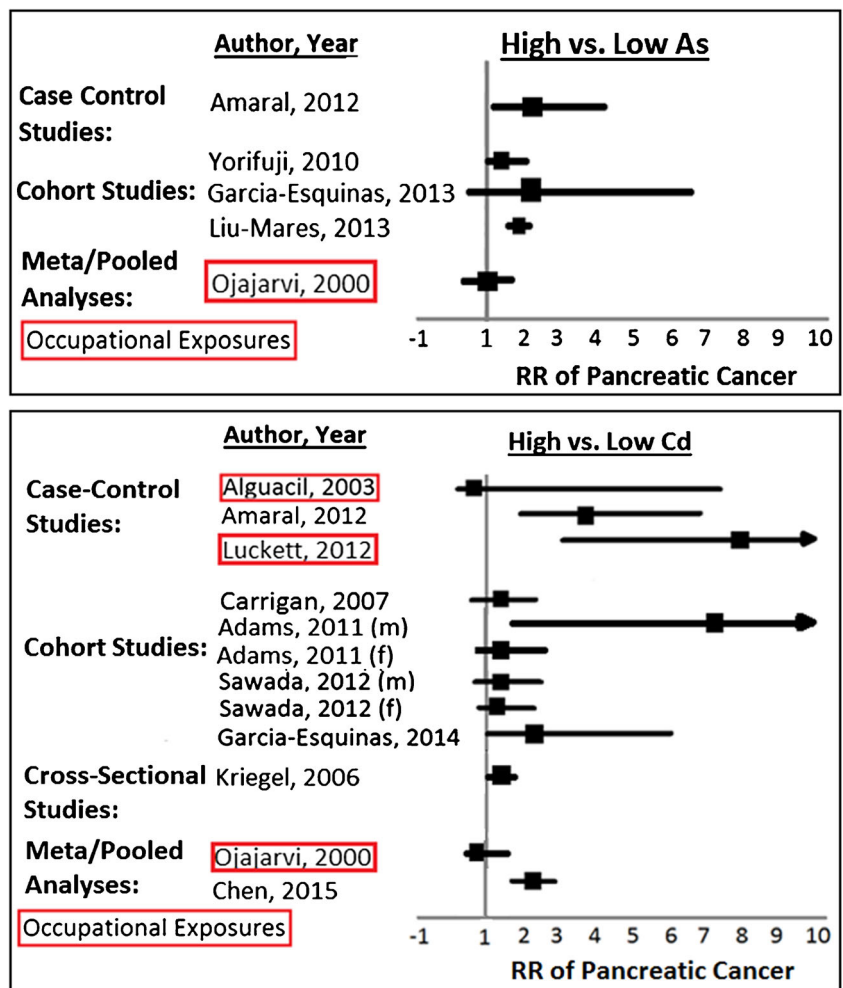
Smoking Tobacco use is one of the strongest and most consistent lifestyle risk factors for pancreatic cancer with an estimated population attributable fraction of 11–32% [15•]. Nicotine derivatives in cigarette smoke can promote carcinogenesis of the pancreas by inducing cellular damage, formation of DNA adducts, or interfering with physiological pathways [16]. Two recent case-control studies [17, 18], a pooled analysis of 12 case-control studies from Panc4 consortium [19], as well as a large meta-analysis of 30 case-control and 12 cohort studies

[20], reported a twofold higher pancreatic cancer risk for current compared with never smokers, and the risk increased up to threefold with ≥ 35 cigarettes per day.

Overall and Central Obesity The World Cancer Research Fund (WCRF) and American Institute for Cancer Research (AICR) estimate that healthy weight is the second most important factor, besides not smoking, for cancer prevention; excess weight is estimated to be attributable for up to 15% of all pancreatic cancer cases [21]. Excess body fat has been implicated in pancreatic cancer risk due to modification of insulin, hormonal, and inflammation pathways [22–25]. Prior research has shown that overall obesity, as measured by BMI, and central obesity, as measured by waist circumference or waist-to-hip ratio, are positively associated with pancreatic cancer incidence [26, 27, 28•]. In four pooled analyses [26, 27, 28•, 29], a significant 8–14% higher pancreatic cancer risk and mortality were observed for a 5 kg/m² increase in BMI at baseline (usually measured in mid to late adulthood) and a 55% higher risk when examining > 35 compared with 18.5–24.9 kg/m². A slightly stronger 18–20% higher pancreatic cancer risk was observed for a 5 kg/m² increment in BMI at younger ages (usually retrospectively assessed for ages 18–21) [28•, 29]. In only one [28•] of three pooled analyses [26, 28•, 29] was waist circumference positively associated with pancreatic cancer; however, all three pooled analyses reported statistically significant positive associations for waist-to-hip ratio [26, 28•, 29]. In 2018, the expert panel of the WCRF/AICR report stated that there was convincing evidence that greater body fatness is associated with higher pancreatic cancer risk [30•].

Alcohol Heavy alcohol drinking has been hypothesized to be associated with higher pancreatic cancer risk. Alcohol may promote carcinogenesis through several mechanisms, such as the creation of acetaldehyde, an alcohol metabolism byproduct; upregulation of immunosuppressive and inflammatory pathways; activation of phase I cytochrome P450 biotransformation enzymes; and folate depletion that can interfere with DNA processes [31–34]. Most case-control studies have observed no association with alcohol intake [35–54], while a number of case-control studies found positive [55–60] and inverse [61–63] associations. Additionally, inconsistent associations have been reported with pancreatic cancer risk from 12 prospective studies [64–70, 71•, 72•, 73•, 74, 75]. In a pooled analysis of 14 prospective cohort studies, a 22% (95% CI 3–45%) higher pancreatic cancer risk was observed for alcohol intake ≥ 30 g(g) (equivalent to \geq two drinks per day) compared with 0 g/day [71•]; whereas no statistically significant association was noted in PanScan or PanC4 case-control consortia for that contrast [72•, 73•]. Yet, a 60% increased risk was noted for those consuming more than

Fig. 1 Risk estimates for studies examining arsenic and cadmium exposure with pancreatic cancer risk. The meta-analyses included are Ojajarvi, 2000 (92 occupational studies); Chen, 2015: (five published studies and one de novo study). The black squares and horizontal lines correspond to the study-specific relative risks and 95% confidence intervals. The area of the black squares is proportional to the inverse of the sum of the between-studies variance and the study-specific variance. The red boxes represent studies in which the exposure was occupational exposure to arsenic or cadmium



nine drinks compared with <one drink/day in PanC4 [72•]. Based on this evidence, the WCRF/AICR expert panel concluded that the current data on alcohol intake were too inconsistent to reach a judgment [30•].

Diet Dietary factors have long been hypothesized to be associated with pancreatic cancer risk due to the large geographic variation in incidence rates worldwide [76]. To date, most research, including consortia studies, has focused on individual foods and nutrients including fruits and vegetables [77], dairy products [78], sugar-sweetened beverages [79], fats, meats, and protein (all individual studies cited in WCRF/AICR 2018 report) [80]. In 2018, an expert panel for the WCRF/AICR stated the evidence was suggestive for a positive association between consumption of red meat, processed meat, foods containing saturated fatty acids, and foods containing fructose and pancreatic cancer risk [30•]. However, as pancreatic cancer is believed to be a disease of multifactorial origins, it is also critical to understand risk in the context of multiple, simultaneous dietary, and lifestyle factors. Few studies

have examined associations between dietary and lifestyle patterns and indices, which capture multiple exposures simultaneously, and pancreatic cancer risk. In studies examining data-driven dietary patterns (e.g., prudent pattern) identified within their own cohorts; results have been inconsistent [48, 81–85], whereas indices that are a priori (e.g., Alternative Healthy Eating Index) have been associated with lower pancreatic cancer risk for those who adhered to a healthier dietary and lifestyle [86–89]. Yet, case-control studies pose the potential for information and selection biases, or and cohort studies had limited case numbers and statistical power [81, 82, 85, 86].

Physical Activity Physical activity, through reducing insulin resistance, adiposity, DNA damage, and inflammation, has been posited to lower pancreatic cancer risk [90]. Prior case-control [91, 92] and a cohort study [93], as well as two large meta-analyses [90, 94], have suggested that physical activity is inversely associated with pancreatic cancer risk with risk estimates ranging from 7 to 35% lower risk when comparing high to low levels of physical activity, while several

prospective cohort studies have found nonsignificant or no association [95–98]. Some studies have suggested that risk may be limited to leisure-time physical activity only [91, 94]. The WCRF in 2018 stated that the evidence of a protective effect of physical activity on pancreatic cancer risk is too limited and inconsistent to draw a conclusion [30].

Reproductive Factors

Due to suggested hormonal effects on pancreatic carcinogenesis in rat and human tissue models [99–102], reproductive and hormonal factors have been hypothesized to play a role in pancreatic cancer risk. Results have been inconsistent when examining age at menarche, oral contraceptive use, parity, age at first birth, menopausal status, and hormone replacement therapy (HRT) use as possible risk factors of pancreatic cancer. Although individual studies have reported null [103, 104], inverse [105], and positive associations with increasing parity [106], a meta-analysis of six cohorts and five case-control studies reported a 21% higher pancreatic cancer risk for highest vs. lowest categories of age at first birth [107]. Overall, there was no association with hormone use [104–106] and age at menopause [103, 108]. However, one cohort study observed a protective effect of older age of menopause on pancreatic cancer risk (HR = 0.35 (95% CI 0.18–0.68) comparing menopause at ≥ 55 to < 45 years) [104], and for women who have had a hysterectomy (OR = 0.78; 95% CI 0.67–0.91) in the Panc4 case-control consortium [109]. Overall, there is conflicting evidence of reproductive factors and pancreatic cancer risk in women that warrants further research.

Medical History

Pancreatitis Pancreatitis, the acute or chronic inflammation of the pancreas [110], has been implicated in pancreatic cancer incidence. Specifically, a twofold higher risk has been observed in patients diagnosed with acute pancreatitis [110, 111], while a meta-analysis of six cohort studies and one case-control reported a 13-fold higher pancreatic cancer risk in those diagnosed with chronic pancreatitis [112].

Diabetes and Metformin Type 2 diabetes mellitus, a disease that can occur when the body develops insulin resistance or not enough is produced, has been associated with an excess pancreatic cancer risk; however, it has also been posited that diabetes may be a consequence or early manifestation of pancreatic cancer. Two meta-analyses and one pooled analysis observed a 50–90% higher pancreatic cancer risk in patients who have a history of diabetes [113–115]; risk was similar even for individuals who had diabetes for up to 20 years prior to diagnosis [114]. Medications, like metformin, used to treat type II diabetes, has been hypothesized to have been

protective due to its hypoglycemic and hypoinsulinemic effects [116], but results have been heterogeneous. A case-control and nested case-control study have observed a nonsignificant positive association between metformin use and pancreatic cancer risk [116, 117] while a meta-analysis of 10 cohort studies and three case-control studies observed an inverse association for metformin use in individuals with type 2 diabetes (summary RR = 0.63, 95% CI 0.46, 0.86) [118].

Statins Statins are traditionally used for the treatment and prevention of cardiovascular disease and have been studied in relation to cancer due to their possible anti-neoplastic properties [119]. One case-control and one cohort studies observed no association with statin use and pancreatic cancer risk [119, 120], while a 34% (95% CI 8–53%) lower risk was observed in a case-control study; the statistically significant inverse association was limited to men (OR_{men} = 0.50; 95% CI 0.32–0.79; OR_{women} = 0.86; 95% CI 0.52–1.43) [121]. Two meta-analyses observed a suggestive (summary RR = 0.89, 95% CI 0.74, 1.07) [122] and statistically significant protective effect of statin use on pancreatic cancer risk (pooled OR = 0.70; 95% CI 0.60–0.82) [123].

Hepatitis B, Hepatitis C, and *Helicobacter pylori* (*H. pylori*) Infection Hepatitis B, hepatitis C, and *H. pylori* infections have been investigated due to their ability to be detected and replicated within the pancreas [124, 125], their association with pancreatitis [126, 127], and their ability to enhance inflammatory responses which may promote pancreatic carcinogenesis [128], respectively. A meta-analysis of five case-control and three cohort studies [129] observed a 20–60% higher pancreatic cancer risk with a previous hepatitis B infection, while a large Japanese cohort observed a null association [130]. For hepatitis C, a 26% higher risk was observed in the meta-analysis [129], while subsequent to the meta-analysis, a suggestive inverse association (OR = 0.69, 95% CI 0.28–1.69) was observed in the large Japanese cohort [130]. For *H. pylori*, a nested case-control study [128] and two meta-analyses [131, 132] observed no statistically significant association with *H. pylori* infection. In contrast, a case-control study examining CagA genes, found in some *H. pylori* strains, found a lower pancreatic cancer risk in CagA seropositive individuals (OR = 0.68; 95% CI 0.54–0.84) and a nonsignificant increased risk for CagA negative, *H. pylori*-positive individuals when compared with those who were seronegative for both *H. pylori* and CagA [133].

Periodontal Disease The oral microbiome has recently been hypothesized to be involved in immune response and carcinogen metabolism [134]. Poor oral health has been associated with up to a twofold higher pancreatic cancer risk in multiple prospective studies [134, 135, 136, 137]. *Porphyromonas gingivalis*, has specifically been implicated with reported

significant odds ratios of 1.60 (presence vs. absence) [134] and 2.14 (presence vs. absence of antibodies) [136•].

Environmental (Other “Environmental” Exposures Are Discussed under the Lifestyle Section) and Occupational Exposures

Metals and Metalloids Although the International Agency for Research on Cancer (IARC) concluded there are sufficient evidence to classify inorganic arsenic (As) and cadmium (Cd) as class I human carcinogens [138, 139•], these statements refer to other cancers. Few studies have examined these metals and metalloids with pancreatic cancer risk; the associations have been inconsistent with studies reporting null [140–144] and positive associations [145–154]. Furthermore, a recent study by Antwi et al. reported significant associations between exposure to asbestos, benzene, and chlorinated hydrocarbons and an increased pancreatic cancer risk with ORs ranging from 1.21 to 1.70 [155]. Given that most studies were small, had limited power, were retrospective or restricted to populations exposed to high occupational levels, future high-quality prospective studies with direct measurements of metal exposure are needed.

Summary of Risk Factors

As many of the suspected risk factors for pancreatic cancer may be modifiable, primary prevention by reducing harmful exposures and increasing preventative exposures over time may help to reduce incidence and mortality rates of this highly fatal cancer. Throughout the last 30 years, smoking rates have decreased in the USA [156] and worldwide [157], while obesity and diabetes rates have increased globally [158–160]. Changes in these key risk factors for pancreatic cancer, accounting for latency, may have a strong impact on future incidence and mortality of pancreatic cancer.

Screening

Primary Prevention of Pancreatic Cancer: Existing Pancreatic Cancer Risk Models Have Modest Discrimination

Given the high fatality rate and no current effective chemopreventive agents or screening tools for pancreatic cancer, prevention through identification of novel risk factors and behavioral modification of these factors offers the most promising approach to reducing incidence and mortality. As described above, these established or suspected risk factors [79, 161–164], the majority of which confirm risks < 1.5–2-fold [163], are insufficient, even jointly, for early detection or risk stratification. Currently, a few validated risk assessment

models integrating established risk factors were developed for primary prevention [165•, 166•, 167, 168]. The PancPro model includes the number of family members affected, their relationship and age at diagnosis; the AUC was 0.61 (95% CI 0.51 to 0.71) for any family history, which increased to 0.75 (95% CI 0.68 to 0.81) when the relationship and age at onset were included [166•]. The Klein model includes smoking, diabetes, alcohol use, family history of pancreatic cancer, body mass index, ABO genotype, and three risk alleles; the AUC range from 0.57 for genetic factors, 0.58 for non-genetic factors to 0.61 for genetic and non-genetic factors [165•]. A third model included five SNPs, smoking, and family history of cancer (AUC = 0.63, 95% CI 0.60–0.66) [167]. The last model included age, height, BMI, fasting glucose, urine glucose, smoking, and age at smoking initiation, and drinking habits showed similar c-statistics for men and women, 0.81 (95% CI:0.80–0.83) and 0.80 (95% CI:0.79–0.82), respectively [168]. The use of current risk assessment models has limited utility in the general population due to the low incidence of the disease [165•] and the modest discriminatory power for all models evaluated [169].

No Effective Screening Modality for Pancreatic Cancer Exists

Currently, the US Preventive Services Task Force does not recommend screening for pancreatic cancer in asymptomatic individuals [170]. Further, due to the low prevalence of the disease, no effective screening tool, and lack of effective treatments after diagnosis, population-level screening has the potential to cause significant harm that may outweigh the benefits [170]. However, screening recommendations for higher risk individuals have been proposed. The International Consortium for Pancreatic Cancer Screening (CAPs) recommends that individuals who are first-degree relatives (FDRs) of patients with pancreatic cancer from a familial pancreatic cancer kindred with at least two affected FDRs or patients with Peutz-Jeghers syndrome and p16, BRCA2, and hereditary non-polyposis colorectal cancer (HNPCC) mutation carriers with ≥ 1 affected FDR should undergo initial screening using endoscopic ultrasonography (EUS) and/or magnetic resonance cholangiopancreatography (MRI) [171•]. No consensus was reached for the age to initiate screening or stop surveillance, the optimal screening modalities, and intervals for follow-up imaging, and which screening abnormalities were of sufficient concern for surgery to be recommended. In contrast to CAPs, The American College of Gastroenterology (ACG) [172•] recommended endoscopic ultrasound (EUS) and/or magnetic resonance imaging (MRI) of the pancreas annually starting at age 50 years or 10 years younger than the earliest age of pancreatic cancer in the family. Further, they conditionally recommended, that patients with Peutz-Jeghers syndrome should start surveillance at age 35 years [172•].

Given the rareness of the disease, the lack of consensus regarding screening recommendations, a two-tiered approach in which risk assessment models are employed to identify high-risk individuals for screening using additional biomarkers in pathways known to affect pancreatic cancer risk may reduce false positives and mortality compared with just risk assessment or screening alone. Recent modeling has supported this approach for other diseases [173–175]; thus research into less invasive biomarkers may provide an opportunity to improve primary and secondary prevention approaches.

Early Detection Using Blood Markers

As most people are diagnosed at the late stage, surgical resection is only possible for approximately 15–20% of patients [176]. A major focus of pancreatic cancer research is to develop effective early detection methods through biomarkers (which includes genetic markers discussed in the familial and genetic risk section) with sufficient sensitivity and specificity to accurately detect asymptomatic pancreatic adenocarcinoma at the early stage when treatment might be more effective and thereby increase the 5-year survival. Several novel candidate biomarkers have been proposed for earlier diagnosis, though none have been adopted into routine clinical use. Prior studies, conducted for other cancers, suggest that the inclusion of biomarker and genetic data may improve the performance of existing risk models; the inclusion of biomarkers into existing risk models depends on easily being able to obtain these measures. As such, tissue does not lend itself to a screening or risk assessment as tissue sampling of the pancreas is not trivial. A promising alternative is measurement in blood, a less invasive and more easily collected biospecimen. Below, we summarize the latest research on blood biomarkers for early detection.

CA19-9

To date, CA19-9, a type of carbohydrate secreted by exocrine epithelial cells and, more specifically, an isolated form of Lewis antigen, is currently the best serological pancreatic cancer biomarker that is approved by the FDA for pancreatic cancer management (e.g., prognostic marker). Yet it lacks the sensitivity and specificity to be utilized as a screening tool. Prior retrospective, cross-sectional or nested case-control studies have suggested that CA19-9 has a sensitivity and specificity of 68–74% when examining pancreatic cancer cases with healthy or non-cancer controls [177, 178]. Further complicating the use of CA19-9 as a screening tool, CA19-9 may also be elevated in non-malignant conditions, such as pancreatitis and biliary obstruction or other malignancies (e.g., colorectal cancer), and it can only be expressed in individuals with Lewis a+/b- or Lewis a+/b+ genotypes (5–10% of population are Lewis a-/b- genotype and cannot express CA19-9) [179].

Therefore, many efforts have been taken to improve the performance of the CA19-9 test. Like most complex diseases, the etiology of pancreatic cancers involves a number and combination of risk factors. Thus, a panel of multiple biomarkers may be necessary for use as a screening tool [180–183, 184•]; the combined effect of a panel may increase sensitivity and reduce false positives. To this end, one large prospective study, the Prostate, Lung, Colorectal, and Ovarian Cancer Screening Trial (PLCO), examined 67 biomarkers including CA 19-9, carcinoembryonic antigen (CEA), neuron-specific enolase, beta human chorionic gonadotropin, carcinoembryonic antigen-related cell adhesion molecule 1, and prolactin (which are significantly altered in sera) in combination. CA19-9 plus CEA had the highest diagnostic power of 0.66 in all possible two biomarker panels; no biomarkers were identified that performed significantly better than CA19-9 alone (AUC = 0.66) [185]. Given the low diagnostic power, CA19-9 alone or in combination is not effective as a screening tool.

Proteins and Proteomics

Aberrant glycosylation of glycoproteins has been correlated to several diseases including cancer; a number of studies have examined select proteins or glycoproteins with pancreatic cancer risk. In one cross-sectional study, the combination of α -1-antichymotrypsin (AACT), thrombospondin-1 (THBS-1), and haptoglobin (HPT) (AUC = 0.95, AUC = 0.85) outperformed CA 19-9 (AUC = 0.89) in distinguishing 37 pancreatic cancer cases from 30 healthy control and 112 non-cancer controls, respectively [186]. Other studies have observed strong AUCs for MUC5AC, a member of the mucin family, a heterogeneous group of 21 abundant, high molecular weight O-glycoproteins that can be either secreted or membrane bound; the AUC for the combination of MUC5AC with CA19-9 to differentiate pancreatic cancer cases from benign and chronic pancreatitis controls was statistically significantly greater (0.91, 0.86–0.95) compared with the AUC for the CA19-9 model alone (0.61, CI 0.86–0.95). Inclusion of MUC5AC with CA19-9 improved its specificity (from 43 to 83%) and sensitivity (from 79 to 83%) for differentiating pancreatic cancer cases from controls (e.g., healthy, benign gastrointestinal conditions, chronic pancreatitis) [187]. Whereas, a large prospective study, United Kingdom Collaborative Trial of Ovarian Cancer Screening (UKCTOCS, which profiled 225 serum proteins, found the combination of THBS-1 and CA19-9 achieved a significantly higher AUC of 0.85 ($P < 0.01$) than both markers alone [188]. In addition, plasma thrombospondin-2 (THBS-2) concentrations discriminated among all stages of pancreatic adenocarcinoma with a receiver operating characteristic (ROC) c-statistic of 0.76–0.88 [189]; the c-statistic improved to 0.96–0.97 with CA19-9. Further, the sensitivity was 87% and specificity was 98% for the combination of THBS-2 and CA19-9. A recent study identified CA19-9 and melanoma

inhibitory activity (MIA) or CA19-9 and macrophage inhibitory cytokine-1 (MIC-1) as best biomarkers to separate early stage pancreatic adenocarcinoma cases from chronic pancreatitis (AUC CA19-9 + MIA = 0.86 vs. AUC CA19-9 = 0.81) or IPMN (AUC CA19-9 + MIC-1 = 0.81 vs. AUC CA19-9 = 0.75) in 188 pancreatic adenocarcinoma cases and 220 non-cancer controls [190]. Future research should confirm these findings in larger prospective studies.

ctDNA

Nucleic acids are released due to apoptosis and necrosis of cells and circulate in the peripheral blood [191]. Circulating nucleic acids (DNA, mRNA, and microRNA (miRNA)) have been positively associated with tumor burden and malignant progression [191]. Thus, many attempts have been made to exploit ctDNA as a cancer biomarker for many tumor types including the pancreas. The major ctDNA biomarker of interest for pancreatic cancer is mutated KRAS, given it is the earliest genetic alteration, an important component in the pathogenesis of pancreatic cancer [192], and is mutated in > 90% of pancreas cancer patients [193]. Circulating mutated KRAS DNA was identified in 48% of individuals with localized pancreatic cancer and in 85% of patients with advanced disease in a cross-sectional study that included 155 pancreas cancer patients [194]. When KRAS mutations in ctDNA were combined with four protein markers (CA19-9, CEA, HGF, OPN), the sensitivity increased from 30% for detectable KRAS mutation alone to 64% with 99.5% specificity [195]. In a subsequent cross-sectional study, they tested CancerSEEK, a combined assay for genetic alterations and a panel of eight protein biomarkers, which detects 95% of pancreas cancer at 99% specificity [196]. However, it should be noted that these studies were cross-sectional and not prospective, and mutations in KRAS are not specific to pancreatic cancer but arise in many other cancers.

Circulating Cancer Cells

The presence of circulating tumor cells (CTCs) that have disseminated into peripheral blood is the first step during the formation of metastasis [197]. Therefore, the detection of pancreatic tumor cells in the peripheral circulation may be a useful tool for screening. The greatest challenge in the detection of CTCs is their rarity in the blood (~ 1 CTC per billion blood cells). In a cross-sectional study ($N = 25$ cases, 15 benign controls) [198], the positive expression rates of C-MET, h-TERT, CK20, and CEA in the pancreatic cancer group were 80% (20/25), 100% (25/25), 84% (21/25), and 80% (20/25), respectively, while in the benign disease control group the rates were 0% (0/15), 0% (0/15), 6.77% (1/15), and 0% (0/15), respectively. Several other studies also reported the presence of CTCs in peripheral blood from pancreatic cancer patients, but using

different platforms make it challenging to reach a consensus for clinical application [199, 200]. Although promising, given the small clinical sample and cross-sectional nature, future prospective research is warranted.

Circulating Exosomal DNA

Exosomes are 40–150 nm extracellular vesicles that contain DNA, RNA, and proteins [201]. All living cells, including cancer cells, generate exosomes, and cancer cells generate higher levels of exosomes than normal cells [202]. Exosomes arise from viable cancer cells and may reflect different biology than circulating cell-free DNA (cfDNA) shed from dying tissues [203]. Emerging research has focused on exosomes and their molecular contents as a potential cancer biomarker [204]. Allenson et al. [203] compared exosome-derived DNA to cfDNA to validate KRAS detection rates in liquid biopsies of patients with pancreatic adenocarcinoma using a discovery cohort of 88 pancreatic adenocarcinoma patients, 54 age-matched healthy controls, and a validation cohort of 39 cancer patients and 82 healthy controls. KRAS mutations in exoDNA, were identified in 7.4%, 66.7%, 80%, and 85% of age-matched controls, localized, locally advanced, and metastatic pancreatic adenocarcinoma patients, respectively. Comparatively, mutant KRAS cfDNA was detected in 14.8%, 45.5%, 30.8%, and 57.9% of these individuals. Similarly, KRAS mutations (39.6%) was observed in 48 pancreatic adenocarcinoma patients but only 2.6% in healthy subject [205]. Other studies have also observed higher glypican-1-positive (GPC1) exosome levels [206] in patients with pancreatic cancer than in controls; GPC1+ crExos (from pancreatic adenocarcinoma, chronic pancreatitis patients and healthy individuals) revealed a near perfect classifier with an AUC of 1.0 (95% CI 0.956–1.0). However, the sample size was small, and all studies were cross-sectional.

Antibody Arrays

Given antibodies are generated against certain tumor-associated antigens (e.g., mesothelin, TNP1) [207], antibody arrays may be useful as potential cancer biomarkers [208]. A three-protein (ERBB2, TNC, and ESR1) panel of plasma biomarkers was identified from 130 test set [209] and demonstrated an AUC of 0.68 and 0.86 when using prediagnostic and diagnostic specimens of pancreatic adenocarcinoma, respectively. When CA 19-9 was added to the panel, the AUC increased to 0.71 and 0.97 for prediagnostic and diagnostic specimens, respectively, suggesting the possibility for use as a diagnostic biomarker panel. Additional studies [210] suggest that IGFBP2 and IGFBP3 are statistically more effective (AUC = 0.94) than CA19-9 alone (AUC = 0.89) at discriminating pancreas cancer patients ($n = 101$) at an early stage from healthy controls ($n = 38$). Gerdtsson et al. [211] evaluated an

antibody array on human recombinant antibody targeting cytokines and estimated AUC values in the test sets ranged from 0.77 to 0.87 to distinguish pancreatic adenocarcinoma vs. individuals not known to have pancreatic cancer as controls.

Metabolites

Interested in whether altered metabolism may indicate sub-clinical pancreatic cancer, Mayers et al. [212•] collected pre-diagnostic plasma from pancreatic cancer cases ($N=453$) and matched controls ($N=900$) in a pooled analysis of individual-level data from four prospective cohort studies (median time between blood collected and diagnosis was 8.7 years). They discovered three metabolites (out of 133 studied), the branched chain amino acids (BCAAs) isoleucine, leucine, and valine were significantly associated with a future diagnosis of pancreatic adenocarcinoma. The result also confirmed that plasma BCAAs were elevated in mice with early-stage pancreatic cancers driven by mutant *Kras* expression. Although not the focus of this review, select urinary metabolites have also been identified as potential early detection markers, including Acetone, O-Acetylcarnitine, Dimethylamine, and Choline. [213]

miRNA

MicroRNAs (miRNAs) are small RNAs (22–25 nt) that negatively regulate gene expression by binding to complementary mRNA resulting in gene silencing, translational repression, or target degradation. The deregulation of some miRNAs has been identified as a mechanism responsible for cell transformation including pancreatic cancer development [214]. Previous studies have reported miR-21, miR-375, miR-196, miR-210, and miR-200 as potential miRNA candidates [214–217]. One small cross-sectional study ($n=48$ cases) conducted profiling of 45 miRNAs and suggested that MicroRNA-375 improves diagnosis of pancreatic adenocarcinoma in this study (70% accuracy) but did not outperform CA19-9 [218]. Lai et al. [219] found that exosomal glypican-1 (GPC1) is not diagnostic for pancreatic adenocarcinoma whereas the AUC for exosomal miR-10b, miR-21, miR-30c, miR-181a, and miR-let7a had 100% sensitivity and specificity with respect to their accuracy in distinguishing pancreatic adenocarcinoma from normal controls; only miR-106b and miR-483 failed to have an excellent AUC. However, their sample size is small, and the findings should be prospectively confirmed.

Conclusions

Given the late stage of diagnosis and the lack of effective treatment of pancreatic cancer, primary (reduction in exposure

to risk factors) and secondary prevention efforts (effective screening modalities) are the best approaches to reduce the morbidity and mortality from this disease. Currently, pancreatic cancer has few known and suspected risk factors, and risk assessment tools have limited utility given their modest discriminatory power range of 0.57–0.81 [220]. Although emerging evidence suggests blood-based biomarkers may be useful as early detection markers, findings need to be confirmed in prospective studies. Due to the rarity of disease, future studies should consider a two-tiered approach in which risk assessment is used to identify high-risk individuals for screening, and then effective imaging and biomarkers in pathways known to affect pancreatic cancer risk are employed; these combination approaches may reduce false positives and mortality compared with just risk assessment or screening alone.

Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

Human and Animal Rights and Informed Consent This article does not contain any studies with human or animal subjects performed by any of the authors.

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