Identifying key factors for the effectiveness of pancreatic cancer screening: A model-based analysis

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Abstract
Pancreatic cancer (PC) survival is poor, as detection usually occurs late, when treatment options are limited. Screening of high-risk individuals may enable early detection and a more favorable prognosis. Knowledge gaps prohibit establishing the effectiveness of screening. We developed a Microsimulation Screening Analysis model to analyze the impact of relevant uncertainties on the effect of PC screening in high-risk individuals. The model simulates two base cases: one in which lesions always progress to PC and one in which indolent and faster progressive lesions coexist. For each base case, the effect of annual and 5-yearly screening with endoscopic ultrasonography/magnetic resonance imaging was evaluated. The impact of variance in PC risk, screening test characteristics and surgery-related mortality was evaluated using sensitivity analyses. Screening resulted in a reduction of PC mortality by at least 16% in all simulated scenarios. This reduction depended strongly on the natural disease course (annual screening: −57% for “Progressive-only” vs −41% for “Indolent Included”). The number of screen and surveillance tests needed to prevent one cancer death was impacted most by PC risk. A 10% increase in test sensitivity reduced mortality by 1.9% at most. Test specificity is important for the number of surveillance tests. In conclusion, screening reduces PC mortality in all modeled scenarios. The natural disease course and PC risk strongly determines the effectiveness of screening. Test sensitivity seems of lesser influence than specificity. Future research should gain more insight in PC pathobiology to establish the true value of PC screening in high-risk individuals.

KEYWORDS
microsimulation model, pancreatic cancer, screening

Abbreviations: EUS, endoscopic ultrasonography; HGD, high-grade dysplasia; IGD, intermediate-grade dysplasia; LGD, low-grade dysplasia; LYG, life-years gained; MISCAN, Microsimulation Screening Analysis; MRI, magnetic resonance imaging; NNS, number needed to screen; NNSurv, number needed to surveil; NNT, number needed to treat; PanIN, pancreatic intraductal neoplasia; PC, pancreatic cancer.

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1 | INTRODUCTION

Pancreatic cancer (PC) has one of the poorest survival rates of all human cancers and is ranked among the top five of cancer-related deaths. Although the general incidence is low (lifetime risk of 1.3%), it is substantially increased in certain high-risk groups, with relative risks up to 132 in Peutz Jeghers Syndrome, for instance. Approximately 10% of PC cases are found in individuals with an inherited increased risk for this fatal disease. This high-risk population includes carriers of PC-prone gene mutations (eg, BRCA2, CDKN2A and LKB1) and first-degree relatives of familial PC patients. Familial PC patients have at least (a) one first-degree relative with PC, (b) one second-degree relative <50 years at time of PC diagnosis or (c) two relatives with PC without a known gene mutation.

Several studies have shown that screening individuals at high risk for PC can detect precursor lesions and early stage cancer. However, we lack solid evidence that the benefits of screening (reduced PC-related morbidity and mortality) outweigh its harms, such as patient burden, overdiagnosis and overtreatment. Determination of the effectiveness of screening requires large-scale and prolonged follow-up.

Computer simulation models have proven their value in estimating the long-term impact of screening using short-term indicators such as PC incidence and detection rates. The Microsimulation Screening Analysis (MISCAN) model, for example, has been used for the evaluation and optimization of screening programs for cancer of the cervix, breast, colon, prostate, lung and esophagus. This model has also been applied to test hypotheses on disease development and test characteristics.

In order to create an accurate simulation model, detailed information on the disease is necessary. Current knowledge gaps in PC screening include the natural disease course, test characteristics (sensitivity, specificity) for different disease stages, and the PC risk level. Decision-analytic modeling can be used to explore such uncertainties and quantify the impact of individual factors on the effect of screening. Identification of the factors with the strongest impact on screening may guide future research.

In our study, we aim to identify key parameters that define the effect of PC screening and to analyze their impact on the effect of PC screening in a high-risk population using the microsimulation model MISCAN. Consequently, we will identify areas to which further research should be directed.

2 | METHODS

2.1 | MISCAN-pancreas model

The MISCAN model generates a large simulated study population with individual life histories. Each person can develop a preinvasive lesion that can evolve into PC, from which that person may die at a certain point in time. This simulation results in an age- and time-specific output of disease incidence and mortality. The simulated population then undergoes PC screening, which will change life histories in case screening detects preinvasive lesions or early-stage cancer that can be surgically resected. These changes constitute the effects of PC screening and are represented by the number of prevented cancer cases and a potential shift in stage distribution.

The stochastic model underlying the simulation is composed by input parameters. These parameters relate to the demographic characteristics (eg, the life table), epidemiology and natural disease course (eg, duration of the different disease stages) and screening characteristics (eg, the sensitivity of the screening tests). Since evidence-based data are scarce in the setting of PC screening of high-risk individuals, many assumptions are based on consensus recommendations, as stated by the international Cancer of the Pancreas Screening-consortium.

2.2 | Base case analyses

2.2.1 | PC risk and natural disease course

We assume a lifetime risk of 7.5% for developing clinical PC, based on the current knowledge of PC risk in high-risk individuals. In the model, the disease is subdivided into preinvasive and invasive stages. Preinvasive lesions (ie, pancreatic intraductal neoplasia (PanIN) and intraductal papillary mucinous neoplasm) are subdivided into three consecutive stages, based on grade of dysplasia: low-, intermediate- and high-grade (respectively LGD, IGD and HGD). Invasive lesions are graded according to TNM stages: I, II and III/IV. Stages III and IV are simulated together because treatment and survival are comparable. While preinvasive lesions can only be detected by screening or surveillance, invasive stages can also be detected clinically (ie, because of symptoms).

The natural disease course of PC, in particular the duration of the different preinvasive stages, is unknown. Therefore, we model two...
base cases with contrasting disease pathways, to look at the effect of different progression patterns on screening outcomes. In the first base case, we simulate a path where all lesions progress from preclinical disease to clinical cancer, in an average of 14.3 years, based on the scarce evidence available.\textsuperscript{7} We assume durations are exponentially distributed (Weibull distribution, Shape 1) and that the duration of each disease stage is correlated with the next: in other words, when a lesion is fast growing in one stage, it does so in the next. Durations varied based on random selection in the distribution. There are no indolent lesions in this base case.

Since the average dwelling time of 14.3 years is debated, we model a second base case, in which both faster progressive and indolent (slow developing) lesions are simulated. In case of slow developing disease, lethal PC will never occur in the course of a lifetime. Preinvasive stages will at most progress to preclinical cancer Stage I. This is assumed to take over 30 years. In this base case, the preinvasive lesions that do progress to

| Value | Lifetime PC risk | Treatment mortality | Progressive-only pathway | Indolent Included pathway |
|-------|------------------|---------------------|--------------------------|--------------------------|
| Stage | Mean duration of progressive stages (years) | Stage Mean duration of indolent stages (years) | Stage Probability of being clinically diagnosed, before moving to the next stage\textsuperscript{12 b} |
| Preinvasive stage LGD | 3.33 | n.a. | 5.1% |
| Preinvasive stage IGD | 3.33 | n.a. | 13.1% |
| Preinvasive stage HGD | 3.33 | n.a. | 100% |
| Preclinical cancer Stage I | 2 | n.a. | 5-year relative survival\textsuperscript{12 b,c} |
| Preclinical cancer Stage II | 2 | n.a. | Clinical cancer Stage I |
| Preclinical cancer Stage III/IV | 1 | n.a. | Clinical cancer Stage II |
| Estimated mean total preclinical | 14.3 | 4.8 | Clinical cancer Stage III/IV |
| Screen test sensitivity\textsuperscript{d} | Preinvasive stage LGD | 60% | 90% |
| Preinvasive stage IGD | 60% | 93% |
| Preinvasive stage HGD | 75% | 99% |
| Preclinical cancer Stage I | 90% | Screen test specificity\textsuperscript{d} |
| Preclinical cancer Stage II | 93% | Any pancreatic lesion |
| Preclinical cancer Stage III/IV | 99% | 90% |

Notes: If someone has survived the first 5 years after diagnosis, we assumed lifelong PC survival. Detection (and associated management of preinvasive lesions) was assumed to lead to a 100% cure rate. For resection, we assumed a 3% mortality risk. Stage-specific survival of screen-detected invasive cancer was based on observed survival of clinically detected cancer in the Netherlands.

\textsuperscript{a}Optimized parameter.

\textsuperscript{b}Weighted averages are used to combine cancer Stage Ia and Ib into Stage I, and Ila and Ilb into Stage II.

\textsuperscript{c}A linear distribution of PC mortality was assumed over 5 years.

\textsuperscript{d}Combined MRI/EUS.
preclinical cancer, progress much faster, in an average time of 5 years, compared to the other base case. More detailed information on the assumed natural disease courses is provided in Table 1 and in the Appendix.

For both base cases, we use PC lifetime risk as a calibration target. We have created a high-risk population by multiplying the PC lifetime risk of the general population by 10. Figure 1 shows the prevalence of each disease stage by age, for both the “Progressive-only” and “Indolent Included” pathways.

### FIGURE 1
Estimated age-specific prevalence of different disease stages in high-risk individuals when modeling only progressive lesions (panel A), or modeling both indolent and faster progressive lesions (panel B), in the absence of screening.

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### 2.3 Assumptions for screening, surveillance and treatment

For both base cases, we simulated a hypothetical cohort of 1 000 000 high-risk individuals from birth until death with and without screening. When screening is implemented, these individuals are offered yearly or 5-yearly screening from the ages of 50 to 75. Individuals are assumed to adhere to all screening rounds. The screen test characteristics are equivalent to a combination of both an endoscopic ultrasonography (EUS) and a magnetic resonance imaging (MRI), as they provide the most accurate pancreatic imaging. The screen test is considered positive when a preinvasive or invasive lesion is detected, rightly or wrongly. Surveillance tests are defined as additional tests, performed every 6 months after a positive screen test until the lesion was resected (Figure 2). Test characteristics of the surveillance test are assumed identical to those of the screen test.

As explained in the abovementioned paragraph on natural disease course, disease development occurs in different stages. We assume test characteristics differ between these stages, based on preliminary results of a PC screening study cohort (Figure 2). We assume that 90% of normal pancreata (no lesions) are correctly identified on imaging and 10% will be diagnosed as false positive. These individuals will be referred to more intensive follow-up (surveillance test) or surgical resection. Figure 2 shows screen (and surveillance) test sensitivities for each disease stage. For individuals with a preinvasive lesion harboring LGD, IGD or HGD, tests are assumed to be positive in 60%, 60% and 75% of cases, respectively, and detection will result in surveillance or surgery. In case of preclinical cancer Stage I, II or III/IV, sensitivity is assumed to be 90%, 93% and 99%, respectively. Detection of preclinical cancer Stages I and II results in surgery. Individuals who are diagnosed with preclinical cancer Stage III/IV will receive palliative care.

Not all lesions detected by screening will develop into clinical cancer when left untreated. When detected, these clinically irrelevant findings may lead to overtreatment. In case of a positive screen test, individuals either undergo surgery (Figure 2, transitions 2 and 3) or surveillance. The proportion of patients that undergo resection is determined by the test sensitivity and the probability that a positive-tested patient undergoes resection, right or wrongfully so. This probability is based on different factors in the clinical process: risk of malignancy for the different precursor lesions, patient symptoms, changes over time, additional tests and multidisciplinary meetings (eg, test interpretation).

For example, when 90% of individuals with preclinical cancer Stage I will have a positive screening test and 90% of these patients will undergo resection, 81% of all individuals with preclinical cancer Stage I will undergo resection. Some individuals return straight back to regular screening, based on the interpretation of the positive test result. Detection and associated management of preinvasive lesions was assumed to lead to a 100% cure rate (ie, no cancer development). However, such patients remain at risk for development of new pancreatic lesions. For resection, we assume a baseline 3% mortality risk.

### 2.4 Outcome parameters and sensitivity analyses

For each of the six scenarios (two base cases with and without annual or 5-yearly screening), the model counts all relevant events, that is, the number of screening and surveillance tests, resections, cancer diagnoses and cancer deaths. From this, we can calculate cancer incidence and mortality rates, life-years gained (LYG) and interval-cancer rates (in the first 5 years and in the total period after screening). An interval cancer is defined as a clinically detected cancer after a negative screen test. For each scenario with screening, we calculate the number needed to screen (NNS), number needed to surveil (NNSurv) and number needed to treat to prevent one cancer death (NNT) as measures of effectiveness. We evaluated these measures of
effectiveness in terms of preventing cancer death instead of detecting one cancer because of the deadly nature of PC.

Besides assuming different disease pathways and screen intervals, we estimate the impact of changes in other variables as PC risk, test performance and treatment-related mortality on the effect of screening by sensitivity analyses (Figure 3 and Appendix Tables 2 and 3).

The lifetime risk for developing PC of 7.5% in the base cases is halved and doubled in the sensitivity analyses. Test specificity is decreased from 90% to 85% and increased to 95% and 100%. The assumed test sensitivity is varied by 5% (ie, 95% and 105% of its base case value) and by 10% (ie, 90% and 110% of base case value) for each disease stage, up to a maximum of 100%. Finally, we increase the treatment mortality risk from 3% to 5%.

3 | RESULTS

3.1 | Base case analyses; Progressive-only pathway

Without screening, 921 PC cases and 751 PC deaths per 10 000 individuals were found using the model including only progressive disease. The mortality rate was 108 deaths per 100 000 life years (LYs).

Annual screening led to a 49% decrease in number of cancer cases, from 921 to 466 cases, and prevented 432 cancer deaths per 10 000 persons simulated. This corresponds with a 57% reduction in mortality rate, from 108 to 46 deaths per 100 000 LYs (Table 2).

Five-yearly screening resulted in less resected lesions harboring HGD, as compared to annual screening (207 vs 233). The number of resected cancer Stages I, II and III were 256, 40 and 2 with 5-yearly screening and 252, 40 and 2 with annual screening. The NNT was 2.7 for yearly screening and 3.3 for 5-yearly screening.

3.2 | Base case analyses; Indolent Included pathway

Without screening, the model including both indolent and faster progressive disease, simulated 918 PC cases and 753 PC deaths per 10 000 individuals. The mortality rate was 108 deaths per 100.000 LYs.

Annual screening led to a 28% decrease in number of cancer cases, from 918 to 659 cases, and prevented 309 cancer deaths per

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FIGURE 2  Screening strategy evaluated in the MISCAN-pancreas model. Transition 1 ("continue screening") is the effect of a (false-) negative test result. Values in table are for the base case situation. *1- or 5-year interval. **In case of cancer Stage III/IV, patients only receive palliative care (no resection) [Color figure can be viewed at wileyonlinelibrary.com]
10 000 persons simulated. This corresponds with a 41% reduction in mortality rate, from 108 to 64 deaths per 100 000 LYs.

Five-yearly screening reduced mortality by 18%, compared to no screening. The NNS with yearly and 5-yearly screening were 600 and 326, respectively. Also, in this base case, the level of resected less invasive lesions (LGD and IGD) was higher when screening takes place annually, but to a lesser extent than in the Progressive-only model. The number of resected HGD lesions was also lower in case of 5-yearly screening compared to annual screening (204 vs 254). The number of resected cancer Stages I, II and III were 203, 74 and 4 with 5-yearly screening and 276, 66 and 5 with annual screening. The NNT was higher in case of 5-yearly screening, as compared to annual screening (6.0 vs 4.1).

### 3.3 Sensitivity analyses

The effect of different sensitivity analyses on NNS for both pathways in case of annual screening are visualized in Figure 3. An overview of all sensitivity analyses on different outcome measures is provided in Table 3 (annual screening) and in Appendix Tables 2 and 3 (5-yearly screening).

#### 3.3.1 PC risk

For Progressive-only disease and annual screening, doubling the PC base case risk lowered the NNS from 431 to 225. If the base case PC risk was decreased by half, the NNS becomes 844. Similar changes were seen in the Indolent Included pathway (Figure 3: Appendix Table 2 and 3).

#### 3.3.2 Screen test performance

In the Progressive-only pathway, lowering the test specificity from 90 to 85% led to an increase of the number of surveillance tests to prevent one cancer death (NNSurv), from 47 to 67, in case of annual
screening. Increased specificity up to 100% decreased the NNSurv from 47 to 5. In the Indolent Included pathway, lowering of the test specificity from 90 to 85% led to an increase in NNSurv from 65 to 93, in case of annual screening. With 5-yearly screening, the observed effect was smaller in both pathways.

For both pathways, an assumed 10% increase in test sensitivity for each disease stage lowered the PC incidence and mortality by <10%. The effect was smallest in case of annual screening.

### 3.3.3 Treatment mortality

Varying treatment mortality from 3% to 5% resulted in an increase in PC-related deaths of 4 to 8% for both disease pathways in both annual and 5-yearly screening. It also modestly increased the NNS; for annual screening in the Progressive-only pathway from 431 to 450, and in the Indolent Included pathway from 600 to 638.

### 4 DISCUSSION

Results from our model show that PC screening in high-risk individuals is promising and may be worthwhile. Based on the results of our exploratory study, we conclude that the characteristics of the natural disease course (ie, probability of progression and average duration of preclinical disease stages) largely determines the efficacy of PC screening. Another important parameter we identified is the level of risk for developing PC. This is highly relevant, given the various genetic backgrounds of high-risk individuals and the associated variable risk levels.
The sensitivity of the screening test has a much smaller effect on mortality reduction. The specificity of the test, however, is of particular importance for the number of surveillance tests (associated with burden) needed to prevent one PC death.

The influence of the test sensitivity on the NNS was negligible. This is partly caused by the fact that the probability that someone is referred for resection after a positive test was not varied (Appendix Table 1). Consequently, the change in resections (and LYG) was smaller than the change in screenpositives. In case of a screening interval of 5 years and a 10% lower sensitivity, 8% more screening tests were needed to prevent one PC death compared to the base case sensitivity (Progressive-only pathway). For the same comparison, the mortality rate was 4% lower. Furthermore, the low impact of the test sensitivity is influenced by the screening interval. Screening with a low test sensitivity can still be effective if the screen frequency is high. A screen test with a higher sensitivity could lead to a less intensive screening program with lower costs and a lower burden. Also, the specificity of PC screening tests could be much improved to decrease the number of false positive referrals and overtreatment. So, test development is still of much importance.

Microsimulation modeling is driven by available evidence and supplemented by assumptions. Ideally, these assumptions are based on high-quality evidence. In case of screening for PC in high-risk individuals, scientific data are still scarce. This paucity of high-quality evidence can partly be explained by the relatively short period of time in which the effects of PC screening have been studied (the first report dates from 1999). Also, as opposed to other types of cancer, in which histologic samples from suspicious lesions can be easily obtained, this is not the case for pancreatic lesions. EUS-guided tissue

| TABLE 3 | Results of the sensitivity analyses of the two base cases for annual screening |
|-----------------------------------------------|
| **Incidence (per 100 000 LYs)** | **Incidence reduction (%)** | **Mortality (per 100 000 LYs)** | **Mortality reduction (%)** | **NNS** | **NNT** | **NNSurv** |
| Population without screening | 132 | n.a. | 108 | n.a. | n.a. | n.a. |
| **Progressive-only pathway** | | | | | | |
| Base case | 67 | 49.2% | 46 | 57.4% | 431 | 2.9 | 47 |
| Risk | 34 | 50% | 23 | 58.2% | 844 | 2.8 | 87 |
| Doubled | 130 | 48.2% | 89 | 57.0% | 225 | 3.0 | 26 |
| Sensitivity | +5% | 66 | 50.0% | 45 | 58.3% | 427 | 2.8 | 46 |
| +10% | 64 | 51.5% | 44 | 59.3% | 422 | 2.8 | 46 |
| −5% | 68 | 48.5% | 46 | 57.4% | 437 | 2.9 | 47 |
| −10% | 69 | 47.7% | 47 | 56.5% | 443 | 2.9 | 48 |
| Specificity | 85% | 66 | 50% | 45 | 58.3% | 429 | 2.8 | 67 |
| 95% | 67 | 49.2% | 46 | 57.4% | 433 | 2.8 | 26 |
| 100% | 68 | 48.5% | 46 | 57.4% | 435 | 2.9 | 5 |
| Treatment mortality | 5% | 67 | 49.2% | 48 | 55.6% | 450 | 3.0 | 49 |
| **Indolent Included pathway** | | | | | | |
| Base case | 94 | 28.8% | 64 | 40.7% | 600 | 4.1 | 65 |
| Risk | 48 | 29.4% | 33 | 40.0% | 1178 | 4.1 | 122 |
| Doubled | 182 | 27.5% | 123 | 40.6% | 314 | 4.3 | 37 |
| Sensitivity | +5% | 93 | 29.5% | 63 | 41.7% | 587 | 4.1 | 64 |
| +10% | 92 | 30.3% | 62 | 42.6% | 575 | 4.1 | 62 |
| −5% | 96 | 27.3% | 65 | 39.8% | 615 | 4.2 | 66 |
| −10% | 97 | 26.5% | 66 | 38.9% | 630 | 4.2 | 68 |
| Specificity | 85% | 94 | 28.8% | 63 | 41.7% | 596 | 4.1 | 93 |
| 95% | 95 | 28.0% | 64 | 40.7% | 605 | 4.1 | 36 |
| 100% | 96 | 27.3% | 64 | 40.7% | 613 | 4.2 | 7 |
| Treatment mortality | 5% | 94 | 28.8% | 66 | 38.9% | 638 | 4.4 | 69 |

Abbreviations: LYs, life years; NNS, number needed to screen to prevent one cancer death; NNSurv, number needed to surveil; NNT, number needed to treat to prevent one cancer death.

aReduction of the PC incidence compared to no screening when PC risk is halved: 68, or doubled: 251.
bMortality reduction compared to no screening in case of halved PC risk: 55, or doubled PC risk: 207.
sampling, in particular of smaller lesions, is challenging and carries a risk of post-procedural pancreatitis.

The design of this model and its input parameters are based on assumptions and data that are consistent with current knowledge in this field. Importantly, when new data becomes available, it can be incorporated in our model to improve its predictive value. Another strength of our study is that we have used a formal microsimulation model (in which the assumptions are well defined, can be reproduced and of which outputs can be compared to observations), and performed comprehensive sensitivity analyses, through which we have tested the robustness of our results. The MISCAN microsimulation model was developed by the Department of Public Health of the Erasmus MC (Rotterdam, the Netherlands) and has been extensively validated for the evaluation of multiple cancer screening programs (cancer of the breast, cervix, colorectal, lung, prostate and oesophagus).\textsuperscript{2,7}

In this exploratory analysis, the only included negative aspect of treatment was a 3% to 5% surgical mortality risk. Therefore, a potential limitation is that we omitted other negative aspects, such as morbidity and loss in quality of life, as a result of screening, surveillance or resection. These aspects need consideration, since pancreatic surgery is associated with significant morbidity (40%-60%),\textsuperscript{10,11} such as delayed gastric emptying, wound infections and pancreatic fistulae, in addition to diabetes mellitus and/or exocrine insufficiency as late complications. Integrating these harms in the model will likely result in a less favorable effect. Furthermore, the assumed 100% attendance rate for PC screening might have led to an overestimation on the population effect. However, literature shows that high risk individuals are likely to participate in PC screening.\textsuperscript{14} Also, we did not consider systematic false-negative test results in our model. The sensitivity of a test is important, but repeated testing in a screening program generally enables missed lesions to be detected in a next screening-round. However, some lesions may be systematically missed due to their size or (pre-existing) parenchymal changes.

The NNS estimated for breast, cervical, colorectal and prostate cancer screening ranges from 1000 to 2000.\textsuperscript{15,16} Our estimation that approximately 500 persons need to be screened to prevent one PC death is significantly lower. Although many of the factors that influence the NNS in our model are based on assumptions, none of the sensitivity analyses resulted in a NNS of >1500. Moreover, the screening instruments (ie, EUS, MRI) used to detect (preinvasive) PC are more invasive than those used in currently implemented screening programs. Also, they are more expensive (eg, approximately €700 for EUS\textsuperscript{17} as compared to the €60 spend by other screening programs).\textsuperscript{5,7,18}

Few studies estimating the (cost-)effectiveness of PC screening have been published.\textsuperscript{17,19-22} Pandharipande et al\textsuperscript{21} concluded that screening with MRI of individuals with even modestly increased risk may incur life expectancy gains. Rulyak et al\textsuperscript{20} found that EUS screening in high-risk individuals increases patient life expectancy and is cost-effective compared to no screening. Both studies involved a one-time screening at the age of 50. Rubenstein et al\textsuperscript{17} concluded that for men with features of chronic pancreatitis, who are at high-risk for developing PC, the most effective management strategy is no intervention. This may be due to the assumed high morbidity rates and loss in quality of life after total pancreatectomy.

To conclude, we showed that the natural disease course of PC and its precursor lesions is one of the determining factors of the success of pancreatic screening in high-risk individuals. The risk for developing PC in the target population is another factor that plays a major role. Test sensitivity has a minor influence. Both base cases show that under plausible assumptions PC screening might be promising in a high-risk population. The current study underlines the importance of continued research pertaining the development of PC and differential risks within specific target populations. This is of interest not only to improve test strategies based on imaging, but also on biomarkers in serum and secretin stimulated pancreatic juice.

CONFLICT OF INTEREST
The authors disclosed no potential conflicts of interest.

DATA AVAILABILITY STATEMENT
All data on which the model inputs of this analysis are based are available upon reasonable request to the corresponding author: b. koopmann@erasusmc.nl.

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**SUPPORTING INFORMATION**

Additional supporting information may be found online in the Supporting Information section at the end of this article.

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