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DEPEND study protocol: Early detection of patients with pancreatic cancer: a pilot study to evaluate the utility of faecal elastase-1 and 13C-mixed triglyceride breath test as screening tools in high-risk individuals.

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DEPEND study protocol: Early detection of patients with pancreatic cancer: a pilot study to evaluate the utility of faecal elastase-1 and ¹³C-mixed triglyceride breath test as screening tools in high-risk individuals.

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Abstract

Introduction Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The incidence of PC is increasing, with little or no improvement in overall survival and the best chance for long-term survival in patients with PC relies on early detection and surgical resection. In this study, we propose the use of a diagnostic algorithm that combines tests of pancreatic exocrine function (Faecal elastase-1 test (FE-1) and the ^{13}C -mixed triglyceride (^{13}C -MTG) breath test) to identify patients with PC that urgently needs imaging studies.

Methods and analysis This prospective pilot (proof of concept) study will be carried out on 25 patients with resectable PC, 10 patients with chronic pancreatitis and 25 healthy volunteers. This study will construct a predictive algorithm for PC, utilising two tests of pancreatic exocrine function, FE-1 test and the ^{13}C -MTG breath test. Continuous flow isotope ratio mass spectrometry in the ^{13}C -MTG breath test will be used to analyse enriched $^{13}\text{CO}_2$ in exhaled breath samples. The additional predictive benefit of other potential biomarkers of PC will also be considered. Potential biomarkers of PC showing abilities to discriminate between patients with PC from healthy subjects or patients with chronic pancreatitis will be selected by metabolomic analysis.

Ethics and dissemination The study was approved by the North of Scotland Research and Ethics Committee on 1 October 2020 (reference: 20/NS/0105, favourable opinion granted). The results will be disseminated in presentations at academic national/international conferences and publication in peer-review journals.

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Strength and limitations of this study

- This pilot study is aiming to be the first study to construct a predictive algorithm for the early detection of pancreatic cancer (PC) based on tests of pancreatic exocrine function.
- PC patients with potentially curative pancreatic resection will be recruited in this study.
- We expect poor apparent ability for the algorithm to discriminate between PC and chronic pancreatitis patients due to limited sample size.
- Participants’ return of faecal specimens may be limited by embarrassment, hesitancy or inability of the study participants to provide a sample.

INTRODUCTION

Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The UK has one of the worst survival rates in Europe, with an average life expectancy at diagnosis of just 4–6 months and 20% survival at 1 year. In addition, only 3% of people survive for 5 years or longer and this figure has not improved four decades¹. Since complete surgical resection remains the only potentially curative option for PC, it is crucial to identify PC at an early stage, when it is still resectable. Unfortunately, only 10–20% of patients with PC are diagnosed at a stage when curative surgery remains an option².

The early symptoms of PC are usually vague and include weight loss, upper abdominal pain, backache, fatigue, diarrhoea and nausea. In 93% of patients, these symptoms may have been present for up to two years before diagnosis of PC³. Recent studies show that around 40% of patients diagnosed with PC require three or more visits to their GP before they are referred to a specialist^{4 5} thereby leading to a delay in the diagnosis. There is also an increased risk of developing PC in those with new-onset diabetes mellitus (NODM) i.e. less than three years since diagnosis⁶ and as many as 25% of PC cases are believed to present with NODM⁷.

Recent survey of GPs commissioned by Pancreatic Cancer Action⁸ suggested that the delay between not having access to a reliable diagnostic referral pathway and a confirmed diagnosis of PC is a major barrier to the early detection of PC. As the symptoms of PC are often non-specific and unlikely to prompt further clinical investigation, there is an urgent need to have a reliable and cheap screening test that can be used to identify high-risk individuals who require urgent imaging studies such as CT scan to confirm or exclude the

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presence of a pancreatic mass ⁹. Having such a test with reliable sensitivity and negative predictive value (NPV), would enable the GP to select patients for urgent referral to further investigations of PC. A high NPV value would enable the GP to reassure the patient that they did not need further imaging studies. Our innovative approach is to use tests of pancreatic exocrine function combined as a screening test in patients presenting with the vague symptoms that may be associated with PC.

Although pancreatic exocrine insufficiency (PEI) is a well-known complication of pancreatic cancer ^{10 11}, its presence is frequently overlooked in patients with advanced PC resulting in a decreased quality of life (QoL), malnutrition and morbidity ¹². Currently, the diagnostic tests of pancreatic exocrine function such as faecal elastase-1 (FE-1) or the ¹³C-mixed triglyceride (¹³C-MTG) breath test, may be used to detect PEI in patients after pancreatic surgery and chronic pancreatitis ¹³⁻¹⁶.

FE-1 test is a simple indirect and non-invasive method for assessing pancreatic secretion, which measures faecal concentrations of elastase-1, a proteolytic enzyme produced exclusively by pancreatic acinar cells, which binds to bile salts and passes through the gut with minimal degradation. FE-1 level correlate well with output of other pancreatic enzymes, such as amylase, lipase and trypsin ^{17 18} and have been shown to detect PEI in patients with advanced PC and following pancreatic surgery ¹⁴⁻¹⁶.

The ¹³C-MTG breath test assesses pancreatic lipase, it involves the ingestion of a ¹³C-labeled mixed triglyceride (1,3-distearoyl, 2- (carboxyl-¹³C)-octanoyl glycerol) mixed with a test meal. The ¹³C-MTG contains a ¹³C-labeled medium-chain fatty acid (¹³C-octanoate) at the Sn-

2 position, and two long-chain fatty acids (stearic acid) at the Sn-1 and Sn-3 positions of the glycerol backbone of the triacylglycerol. The ingested ^{13}C -MTG then undergoes the lipolysis of the two-long chain fatty acids from ^{13}C -MTG by pancreatic lipase to produce free fatty acids (stearic acid and ^{13}C -octanoate) and monoacylglycerol. After the intestinal absorption, the ^{13}C -free fatty acids are oxidised in the liver producing $^{13}\text{CO}_2$, which is then released on breath. Therefore, the $^{13}\text{CO}_2$ concentration in breath reflects pancreatic lipase activity. The ^{13}C -MTG breath test can be an alternative or complementary to the FE-1 test for PEI. The ^{13}C -MTG breath test has been validated in patients with chronic pancreatitis¹⁹ and patients with pancreatic cancer following surgical resection of the cancer^{20,21}, but not prior to surgery in patients with resectable PC.

In this study, we will validate FE-1 test and ^{13}C -MTG breath test in patients with resectable PC (at an early stage) and test their performance in discriminating between patients with PC and healthy subjects. These tests can provide a non-invasive risk stratification tool to identify patients at risk of having PC when they present with NODM, weight loss or vague upper GI symptoms. Furthermore, this will provide a rational approach to patients identified to have PEI to be selected to have pancreatic imaging such as computed tomography (CT) scan to confirm the presence of PC and having a pancreatic resection earlier (Figure 1). In addition, we will also be identifying other potential biomarkers of PC through unbiased multiomics analysis of blood samples (proteomics and metabolomics). The resulting data will be compared to the healthy control group and patients with chronic pancreatitis aiming to identify signals that are distinct to PC.

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METHODS AND ANALYSIS

Main centre

NIHR Southampton Biomedical Research Centre, Southampton Centre for Biomedical Research, Southampton General Hospital, Southampton, UK.

Dates of the study

From 16th March 2021 to 16th March 2022.

Design

This is a single cross-sectional pilot study of patients with PC, chronic pancreatitis and healthy controls, matched for age and sex. The primary objective is to determine if ¹³C-mixed triglyceride (¹³C-MTG) breath test and faecal elastase-1 (FE-1) concentrations combined can discriminate between the groups. Patients with PC will be recruited at the pre-operative stage within two weeks of initial diagnosis. The secondary objective of this study is the analysis of multiomic (proteomic and metabolomic) profiles and comparison between the groups.

Inclusion criteria

Male and female participants aged from 30-85 within the Wessex pancreatic cancer catchment area will be suitable for inclusion. In the cancer cohort, patients with a diagnosis of resectable pancreatic ductal adenocarcinoma (PDAC) (stage 1-2) will be included. Patients with chronic pancreatitis and healthy subjects will also be recruited as control groups and matched for age and sex to the cases of PDAC.

Exclusion criteria

Patients and controls with autoimmune or other chronic inflammatory conditions, or those who have previously had pancreatic disease or liver dysfunction will be excluded.

Intervention(s) or method

Each participant will be given a participant information sheet explaining the research and will sign an informed consent form. Participants are required to fast overnight for at least 12 hours prior to undergoing the ^{13}C -mixed triglyceride (^{13}C -MTG). At the Clinical Research Facility (CRF), full infection control precautions (related to COVID-19 pandemic) will be adhered to throughout the visit. Participants will have their temperature checked on arrival to ensure they are not pyrexial as part of screening for Covid-19, which will exclude them from the study. Anthropometric measurements (height, weight and waist circumference) of the participants will be recorded. Baseline blood samples will be taken prior to undergoing the ^{13}C -MTG breath test. The data collected will include the following: Full blood count (FBC), Urea and Electrolyte blood test (U&E), Liver function tests (AST, ALT, ALP, albumin and GGT) /clotting studies, glycated hemoglobin (HbA1c %), Lipid profile, Vitamins and micronutrients, Hyaluronic acid, Procollagen III amino terminal peptide (PIINP) and Tissue inhibitor of metalloproteinase-1 (TIMP-1). These tests will be analysed immediately after the blood has been drawn from the participants by the chemical pathology service laboratory at Southampton General Hospital. Further blood samples will be processed and stored for multiomics analysis. Participants will also be asked to provide spot stool samples for faecal elastase-1 (FE-1) concentrations by using a commercially available ELISA kit. Results of the FE-1 test will be expressed as $\mu\text{g/g}$ of stool.

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¹³C-MTG breath test protocol

Each participant will be asked to provide 2 baseline breath samples by blowing into 2 × 10ml exetainer tubes through a straw while wearing a facemask. After that, participants will be given an oral dose of 250mg of 2- [¹³C]-octanoyl-1,3-distearin (¹³C-MTG -Cambridge Isotope Laboratories, Andover, MA, USA) together with a solid test meal consisting of a crispbread or other gluten-free alternative, and 200ml of water. Post-prandial breath samples will be then collected every 30mins for 4 hours. No food or drinks except for water will be allowed during the duration of the breath test study. The collected breath samples will be analysed for the ratio of ¹³CO₂ to ¹²CO₂ using a Continuous Flow Isotope Ratio Mass Spectrometer (CF-IRMS SERCON Ltd, Crewe, UK) at the Southampton Centre for Biomedical Research. The increase in ¹³CO₂ content with regard to the baseline value of the initial breath sample is expressed as atom % excess and we will express the results of the ¹³C-MTG breath test as the cumulative percentage of ¹³C-label recovered on breath as ¹³CO₂ over 4 hours (cPDR over 4 hours).

Primary Outcome: Prediction of PC using a logistic model with measurement of FE-1 concentration expressed as µg/g of stool and ¹³C-MTG breath test expressed as cPDR over 4hrs as covariates.

Statistical analysis and data analysis plans

Sample size calculation

The sample size calculation for the proposed study is based on a previous study assessing pancreatic exocrine function using a ^{13}C -triglyceride breath test in healthy subjects and patients with a localised pancreatic mass ¹³. Using data from this study, we calculated that a sample size of 25 subjects in each group (patients with PC, patients with chronic pancreatitis and healthy control subjects) will give 100% power to detect a difference of 17.8% between patients with PC and healthy control group assuming a standard deviation of 10.5% in patients with PC at a significance level of 5% using a two-tailed test. The power calculation was carried out using IBM SPSS SamplePower version 3.

Statistical tests

Data will be presented as mean + SD, median + IQR when appropriate and 95% CI. Comparison of results of FE-1 and the ^{13}C -MTG breath test between each group will be evaluated by employing either the one way analysis of variance test (if the variables are normally distributed and parametric validity conditions are fulfilled) or the Mann Whitney U-test or Kruskal-Wallis test (if the variables are not normally distributed and non-parametric validity conditions are fulfilled). A multi-variable logistic regression analysis will be performed in order to determine the independent factors that are significantly associated with the presence of PDAC and logistic regression models will be constructed based on the identified independent factors. Furthermore, multi-variable regression analysis will be used to determine the optimal algorithm that combines the results of FE-1 test and ^{13}C -MTG breath test. We will also test whether potential biomarkers of PC identified by unbiased omics analysis improve the prediction of PDAC. A Receiver operating characteristic (ROC) curve will be constructed to evaluate the diagnostic accuracy (sensitivity, specificity,

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positive and negative predictive value, receiver operator curve analysis) of the algorithm for PDAC. Furthermore, the optimal cut off point for the algorithm to identify patients with PC will be estimated using the Youden index. A two-sided p value of <0.05 will be considered to indicate statistical significance. All analysis will be performed using SPSS (IBM, V.25.0) or Stata (Statacorp, v16.0).

Patient and public involvement statement

Patients, healthy subjects and the public were not involved in the study design, recruitment and conduct. We do not plan to inform the result to the study participants unless they apply for it.

ETHICS AND DISSEMINATION

Ethical approval has been granted by the North of Scotland Research and Ethics Committee, reference: 20/NS/0105, 1 October 2020. This manuscript reflects the latest protocol V.7 approved 29 October 2020.

The study will be conducted according to the principles of Good Clinical Practice, General data Protection Regulation and Data Protection Act 2018 for Health and care Research. The sponsor and study team will ensure approval of the study protocol, participant information sheets, consent forms, letters to general practitioners and supporting documents by the appropriate regulatory body and research and ethics committee prior to participant recruitment. Documents will be stored securely with restricted access for at least 10 years. Written, informed consent will be obtained from each patient and an identification number provided. Any published data will not contain personal identifiable data. The results of this study will be disseminated by presentation at academic national/international conferences and publication in peer-review journals.

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Authors' contributions: ZH and PA have designed the study concept and formulated the initial idea; ZH is the Chief investigator and responsible for protocol review. DM and PA wrote the initial protocol. CB, PA, SW and ZH were involved in statistical analysis plan. DM and ZH was responsible for organising the study operation requirements, identifying and discussion with eligible participants. DM is responsible for patient recruitment and consent. DM and PA prepared the test meal and obtained the samples from the participants, VG is sorting the standard operating procedures (SOPs) and logistics with biobanking and sample process. PA is responsible for the analysis of the breath samples and the interpretation of the results from the breath test. JW is managing the study team and logistics related to clinical trial involvement. PA, DM, CB, SW, VG, JW and ZH were involved in drafting of the protocol.

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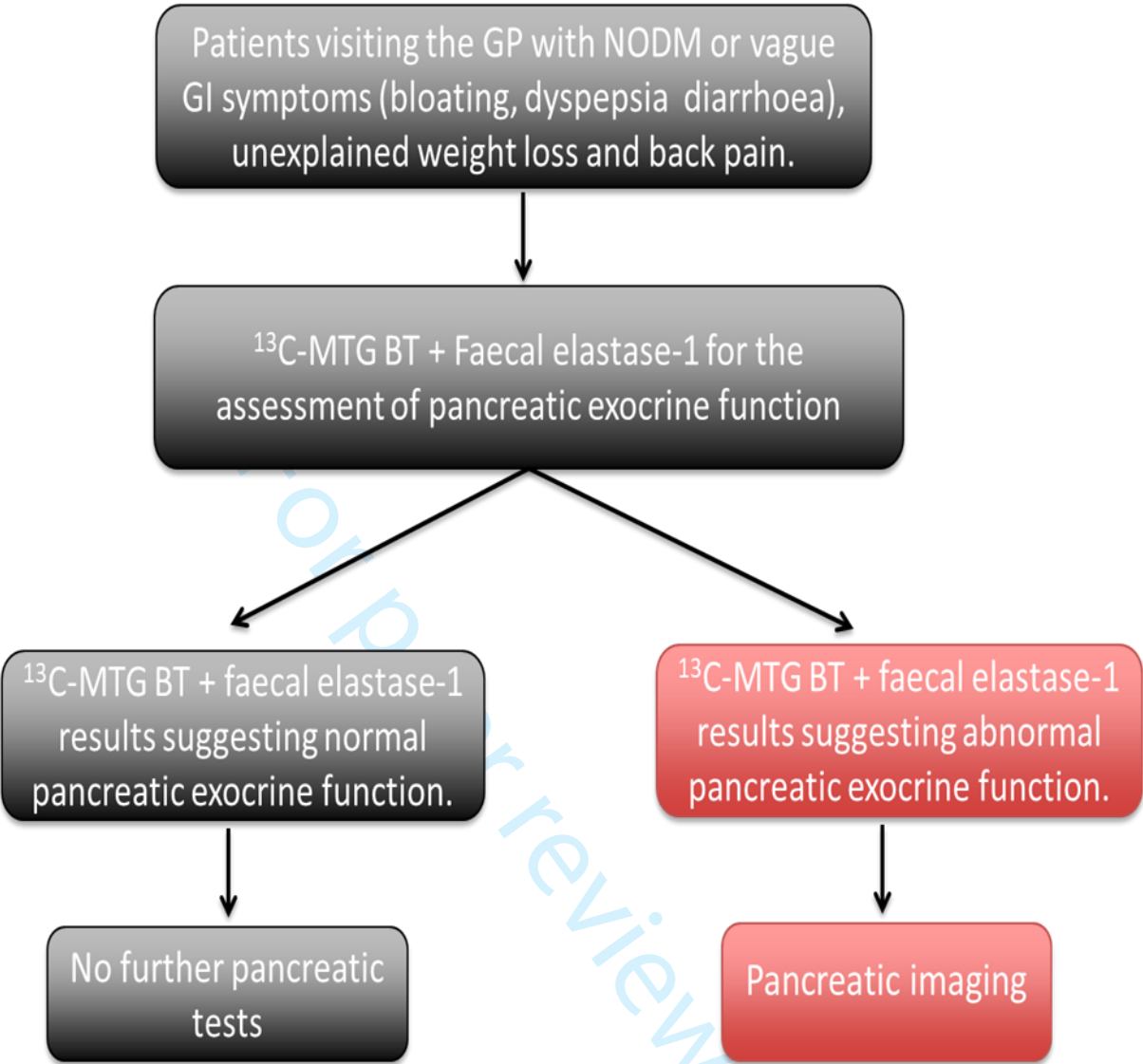
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Figure 1 Proposed diagnostic tool consisting of faecal elastase-1 test and the ^{13}C -mixed triglyceride breath test to screen for pancreatic exocrine dysfunction in patients 'at-risk' of pancreatic cancer.

For peer review only



Abbreviations: NODM- New Onset Diabetes Mellitus, ¹³C-MTG BT- ¹³C-mixed triglyceride breath test.

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DEPEND study protocol: Early detection of patients with pancreatic cancer: a pilot study to evaluate the utility of faecal elastase-1 and ¹³C-mixed triglyceride breath test as screening tools in high-risk individuals.

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Abstract

Introduction Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The incidence of PC is increasing, with little or no improvement in overall survival and the best chance for long-term survival in patients with PC relies on early detection and surgical resection. In this study, we propose the use of a diagnostic algorithm that combines tests of pancreatic exocrine function (Faecal elastase-1 test (FE-1) and the ^{13}C -mixed triglyceride (^{13}C -MTG) breath test) to identify patients with PC that urgently needs imaging studies.

Methods and analysis This prospective pilot (proof of concept) study will be carried out on 25 patients with resectable PC, 10 patients with chronic pancreatitis and 25 healthy volunteers. This study will construct a predictive algorithm for PC, utilising two tests of pancreatic exocrine function, FE-1 test and the ^{13}C -MTG breath test. Continuous flow isotope ratio mass spectrometry in the ^{13}C -MTG breath test will be used to analyse enriched $^{13}\text{CO}_2$ in exhaled breath samples. The additional predictive benefit of other potential biomarkers of PC will also be considered. Potential biomarkers of PC showing abilities to discriminate between patients with PC from healthy volunteers or patients with chronic pancreatitis will be selected by metabolomic analysis.

Ethics and dissemination The study was approved by the North of Scotland Research and Ethics Committee on 1 October 2020 (reference: 20/NS/0105, favourable opinion granted). The results will be disseminated in presentations at academic national/international conferences and publication in peer-review journals.

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Strength and limitations of this study

- This pilot study is aiming to be the first study to construct a predictive algorithm for the early detection of pancreatic cancer (PC) based on tests of pancreatic exocrine function.
- PC patients with potentially curative pancreatic resection will be recruited in this study.
- In this study, we would expect chronic pancreatitis to also have positive findings using the exocrine function tests. However, we are not proposing to use the faecal elastase-1 test and ¹³C-mixed triglyceride breath test to differentiate between chronic pancreatitis and pancreatic cancer at this stage.
- Participants’ return of faecal specimens may be limited by embarrassment, hesitancy or inability of the study participants to provide a sample.

59 INTRODUCTION

60 Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The UK
61 has one of the worst survival rates in Europe, with an average life expectancy at diagnosis of
62 just 4–6 months and 20% survival at 1 year. In addition, in the UK, only about 7% of people
63 survive for 5 years or longer¹. Since complete surgical resection remains the only potentially
64 curative option for PC, it is crucial to identify PC at an early stage, when it is still resectable.
65 Unfortunately, only 10–20% of patients with PC are diagnosed at a stage when curative
66 surgery remains an option².

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68 The early symptoms of PC are usually vague including weight loss, upper abdominal pain,
69 backache, fatigue, diarrhoea and nausea. In 93% of patients, these symptoms may have
70 been present for up to two years before diagnosis of PC³. Recent studies show that around
71 40% of patients diagnosed with PC require three or more visits to their GP before they are
72 referred to a specialist^{4,5} thereby leading to a delay in the diagnosis. There is also an
73 increased risk of developing PC in those with new-onset diabetes mellitus (NODM) i.e. less
74 than three years since diagnosis⁶ and as many as 25% of PC cases are believed to present
75 with NODM⁷.

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77 Recent survey of GPs commissioned by Pancreatic Cancer Action⁸ suggested that the delay
78 between not having access to a reliable diagnostic referral pathway and a confirmed
79 diagnosis of PC is a major barrier to the early detection of PC. As the symptoms of PC are
80 often non-specific and unlikely to prompt further clinical investigation, there is an urgent
81 need to have a reliable and cheap screening test that can be used to identify high-risk
82 individuals who require urgent imaging studies such as CT scan to confirm or exclude the

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3 83 presence of a pancreatic mass ⁹. Having such a test with reliable sensitivity and negative
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5 84 predictive value (NPV), would enable the GP to select patients for urgent referral to further
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8 85 investigations of PC. A high NPV value would enable the GP to reassure the patient that they
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10 86 did not need further imaging studies. Our innovative approach is to use tests of pancreatic
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13 87 exocrine function combined as a screening test in patients presenting with the vague
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15 88 symptoms that may be associated with PC.
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20 90 Although pancreatic exocrine insufficiency (PEI) is a well-known complication of pancreatic
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23 91 cancer ^{10 11}, its presence is frequently overlooked in patients with advanced PC resulting in a
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25 92 decreased quality of life (QoL), malnutrition and morbidity ¹². Currently, the diagnostic tests
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28 93 of pancreatic exocrine function such as faecal elastase-1 (FE-1) or the ¹³C-mixed triglyceride
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30 94 (¹³C-MTG) breath test, may be used to detect PEI in patients after pancreatic surgery and
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32 95 chronic pancreatitis ¹³⁻¹⁶.
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37 97 FE-1 test is a simple indirect and non-invasive method for assessing pancreatic secretion,
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40 98 which measures faecal concentrations of elastase-1, a proteolytic enzyme produced
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43 99 exclusively by pancreatic acinar cells, which binds to bile salts and passes through the gut
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45 100 with minimal degradation. FE-1 level correlate well with output of other pancreatic
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47 101 enzymes, such as amylase, lipase and trypsin ^{17 18} and have been shown to detect PEI in
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49 102 patients with advanced PC and following pancreatic surgery ¹⁴⁻¹⁶.
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54 104 The ¹³C-MTG breath test assesses pancreatic lipase, it involves the ingestion of a ¹³C-labeled
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57 105 mixed triglyceride (1,3-distearoyl, 2- (carboxyl-¹³C)-octanoyl glycerol) mixed with a test
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59 106 meal. The ¹³C-MTG contains a ¹³C-labeled medium-chain fatty acid (¹³C-octanoate) at the Sn-

2 position, and two long-chain fatty acids (stearic acid) at the Sn-1 and Sn-3 positions of the glycerol backbone of the triacylglycerol. The ingested ^{13}C -MTG then undergoes the lipolysis of the two-long chain fatty acids from ^{13}C -MTG by pancreatic lipase to produce free fatty acids (stearic acid and ^{13}C -octanoate) and monoacylglycerol. After the intestinal absorption, the ^{13}C -free fatty acids are oxidised in the liver producing $^{13}\text{CO}_2$, which is then released on breath. Therefore, the $^{13}\text{CO}_2$ concentration in breath reflects pancreatic lipase activity. The ^{13}C -MTG breath test can be an alternative or complementary to the FE-1 test for PEI. The ^{13}C -MTG breath test has been validated in patients with chronic pancreatitis¹⁹ and PC following surgical resection of the cancer^{20,21}, but not prior to surgery in patients with resectable PC.

Both FE-1 and the ^{13}C -MTG breath test are indirect tests of pancreatic exocrine function. To date, only one study has shown the performance of the ^{13}C -trioctanoin breath test with a sensitivity of 100% to detect the presence of PC in 14 patients¹³ while another study has shown that a FE-1 value of $\leq 20\mu\text{g/g}$ is an independent predictor of poor survival in patients with advanced PC²².

In this study, we will validate FE-1 test and ^{13}C -MTG breath test as potential diagnostic biomarkers of resectable PC and test their performance in discriminating between patients with PC and healthy subjects. We aim to assess the diagnostic performances of the exocrine function tests (as a combined diagnostic panel) in detecting PC. If so, then a logistic regression analysis will be performed that will integrate the results of FE-1 test and the ^{13}C -MTG breath test along with other potential biomarkers of PC as a diagnostic algorithm panel.

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131 These tests can provide a non-invasive risk stratification tool to identify patients at risk of
132 having PC when they present with NODM, weight loss or vague upper GI symptoms.
133 Furthermore, this will provide a rational approach to patients identified to have PEI to be
134 selected to have pancreatic imaging to confirm the presence of PC and having a pancreatic
135 resection earlier (Figure 1). In addition, we will be identifying other potential biomarkers of
136 PC through unbiased multi-omics analysis of blood samples (proteomics and metabolomics).

For peer review only

METHODS AND ANALYSIS

Main centre

NIHR Southampton Biomedical Research Centre, Southampton Centre for Biomedical Research, Southampton General Hospital, Southampton, UK.

Dates of the study

From 16th March 2021 to 16th March 2022.

Design

This is a single cross-sectional pilot study of patients with PC, chronic pancreatitis and healthy controls, matched for age and sex. The primary objective is to determine if ¹³C-mixed triglyceride (¹³C-MTG) breath test and faecal elastase-1 (FE-1) concentrations combined can discriminate between the groups. Patients with PC will be recruited at the pre-operative stage within two weeks of initial diagnosis. The secondary objective of this study is the analysis of multi-omic (proteomic and metabolomic) profiles and comparison between the groups.

Inclusion criteria

Male and female participants aged from 30-85 within the Wessex pancreatic cancer catchment area will be suitable for inclusion. In the cancer cohort, patients with a diagnosis of resectable pancreatic ductal adenocarcinoma (PDAC) (stage 1-2) will be included. Patients with chronic pancreatitis and healthy subjects will also be recruited as control groups and matched for age and sex to the cases of PDAC.

Exclusion criteria

Patients and controls with autoimmune or other chronic inflammatory conditions, or those who have previously had pancreatic disease or liver dysfunction will be excluded.

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3 163 **Intervention(s) or method**
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6 164 Each participant will be given a participant information sheet explaining the research and
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8 165 will sign an informed consent form. Participants are required to fast overnight for at least 12
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10 166 hours prior to undergoing the ¹³C-mixed triglyceride (¹³C-MTG). At the Clinical Research
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13 167 Facility (CRF), full infection control precautions (related to COVID-19 pandemic) will be
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15 168 adhered to throughout the visit. Participants will have their temperature checked on arrival
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18 169 to ensure they are not pyrexial as part of screening for Covid-19, which will exclude them
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20 170 from the study. Anthropometric measurements (height, weight and waist circumference) of
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23 171 the participants will be recorded. Baseline blood samples will be taken prior to undergoing
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25 172 the ¹³C-MTG breath test. The data collected will include the following: Full blood count
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28 173 (FBC), Urea and Electrolyte blood test (U&E), Liver function tests (AST, ALT, ALP, albumin
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30 174 and GGT) /clotting studies, glycated hemoglobin (HbA1c %), Lipid profile, Vitamins and
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33 175 micronutrients, Hyaluronic acid, Procollagen III amino terminal peptide (PIINP) and Tissue
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35 176 inhibitor of metalloproteinase-1 (TIMP-1). These tests will be analysed immediately after the
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38 177 blood has been drawn from the participants by the chemical pathology service laboratory at
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40 178 Southampton General Hospital. Further blood samples will be processed and stored for
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43 179 multi-omics analysis. Participants will also be asked to provide spot stool samples for faecal
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45 180 elastase-1 (FE-1) concentrations by using a commercially available ELISA kit. Results of the
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47 181 FE-1 test will be expressed as µg/g of stool.

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¹³C-MTG breath test protocol

Each participant will be asked to provide 2 baseline breath samples by blowing into 2 × 10ml exetainer tubes through a straw while wearing a facemask. After that, participants will be given an oral dose of 250mg of 2-¹³C-octanoyl-1,3-distearin (¹³C-MTG -Cambridge Isotope Laboratories, Andover, MA, USA) together with a solid test meal consisting of a crispbread or other gluten-free alternative, and 200ml of water. Post-prandial breath samples will be then collected every 30mins for 4 hours. No food or drinks except for water will be allowed during the duration of the breath test study. The collected breath samples will be analysed for the ratio of ¹³CO₂ to ¹²CO₂ using a Continuous Flow Isotope Ratio Mass Spectrometer (CF-IRMS SERCON Ltd, Crewe, UK) at the Southampton Centre for Biomedical Research. The increase in ¹³CO₂ content with regard to the baseline value of the initial breath sample is expressed as atom % excess and we will express the results of the ¹³C-MTG breath test as the cumulative percentage of ¹³C-label recovered on breath as ¹³CO₂ over 4 hours (cPDR over 4 hours).

Primary Outcome: Prediction of PC using a logistic model with measurement of FE-1 concentration expressed as µg/g of stool and ¹³C-MTG breath test expressed as cPDR over 4hrs as covariates.

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Statistical analysis and data analysis plans

Sample size calculation

The sample size calculation for the proposed study is based on a previous study assessing pancreatic exocrine function using a ¹³C-triglyceride breath test in healthy subjects and patients with a localised pancreatic mass¹³. In this study, they used the ¹³C-trioctanoin breath test to assess pancreatic exocrine function in 14 patients with a localised pancreatic mass and 5 healthy control subjects. The results of the study showed means ± standard deviation (SD) of the recovery of ¹³CO₂ over 3hrs after undergoing the ¹³C-trioctanoin breath test of 42% ± 3.4% for the healthy controls and 24.2% ± 10.5% for patients with a localised pancreatic mass with an effect size or mean difference between both groups of 17.8% and a within group SD of 9.1% (based on the SD estimates of 3.4% and 10.5% for each group respectively).

Using data from this study, we calculated that a sample size of 25 subjects in each group (patients with PC, patients with chronic pancreatitis and healthy control subjects) will give 100% power to detect a difference of 17.8% between patients with PC and healthy control group assuming a standard deviation of 10.5% in patients with PC at a significance level of 5% using a two-tailed test. The power calculation was carried out using IBM SPSS SamplePower version 3.

231 **Statistical tests**

232 Data will be presented as mean + SD, median + IQR when appropriate and 95% CI.

233 Comparison of results of FE-1 and the ¹³C-MTG breath test between each group will be

234 evaluated by employing either the one way analysis of variance test (if the variables are

235 normally distributed and parametric validity conditions are fulfilled) or the Mann Whitney

236 U-test or Kruskal-Wallis test (if the variables are not normally distributed and non-

237 parametric validity conditions are fulfilled). A multi-variable logistic regression analysis will

238 be performed in order to determine the independent factors that are significantly

239 associated with the presence of PDAC and logistic regression models will be constructed

240 based on the identified independent factors. Furthermore, multi-variable regression analysis

241 will be used to determine the optimal algorithm that combines the results of FE-1 test and

242 ¹³C-MTG breath test. We will also test whether potential biomarkers of PC identified by

243 unbiased omics analysis improve the prediction of PDAC. A Receiver operating characteristic

244 (ROC) curve will be constructed to evaluate the diagnostic accuracy (sensitivity, specificity,

245 positive and negative predictive value, receiver operator curve analysis) of the algorithm for

246 PDAC. Furthermore, the optimal cut off point for the algorithm to identify patients with PC

247 will be estimated using the Youden index. A two-sided p value of <0.05 will be considered to

248 indicate statistical significance. All analysis will be performed using SPSS (IBM, V.25.0) or

249 Stata (Statacorp, v16.0).

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251 **Patient and public involvement statement**

252 Patients, healthy subjects and the public were not involved in the study design, recruitment

253 and conduct. We do not plan to inform the result to the study participants unless they apply

254 for it.

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3 255 **ETHICS AND DISSEMINATION**
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5 256 Ethical approval has been granted by the North of Scotland Research and Ethics Committee,
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7 257 reference: 20/NS/0105, 1 October 2020. This manuscript reflects the latest protocol V.7
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15 260 The study will be conducted according to the principles of Good Clinical Practice, General
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17 261 data Protection Regulation and Data Protection Act 2018 for Health and care Research. The
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20 262 sponsor and study team will ensure approval of the study protocol, participant information
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22 263 sheets, consent forms, letters to general practitioners and supporting documents by the
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25 264 appropriate regulatory body and research and ethics committee prior to participant
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27 265 recruitment. Documents will be stored securely with restricted access for at least 10 years.
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30 266 Written, informed consent will be obtained from each patient and an identification number
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32 267 provided. Any published data will not contain personal identifiable data. The results of this
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35 268 study will be disseminated by presentation at academic national/international conferences
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37 269 and publication in peer-review journals.

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289 Southampton, UK.

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291 participated in this study.

292 **Authors' contributions:** ZH and PA have designed the study concept and formulated the
293 initial idea; ZH is the Chief investigator and responsible for protocol review. DM and PA
294 wrote the initial protocol. CB, PA, SW and ZH were involved in statistical analysis plan. DM
295 and ZH was responsible for organising the study operation requirements, identifying and
296 discussion with eligible participants. DM is responsible for patient recruitment and consent.
297 DM and PA prepared the test meal and obtained the samples from the participants, VG is
298 sorting the standard operating procedures (SOPs) and logistics with biobanking and sample
299 process. PA is responsible for the analysis of the breath samples and the interpretation of
300 the results from the breath test. JW is managing the study team and logistics related to
301 clinical trial involvement. PA, DM, CB, SW, VG, JW and ZH were involved in drafting of the
302 protocol.

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305 **Competing interest:** None.

306 **Patient consent for publication:** Not required.

307 **Provenance and peer review:** Not commissioned; externally peer reviewed.

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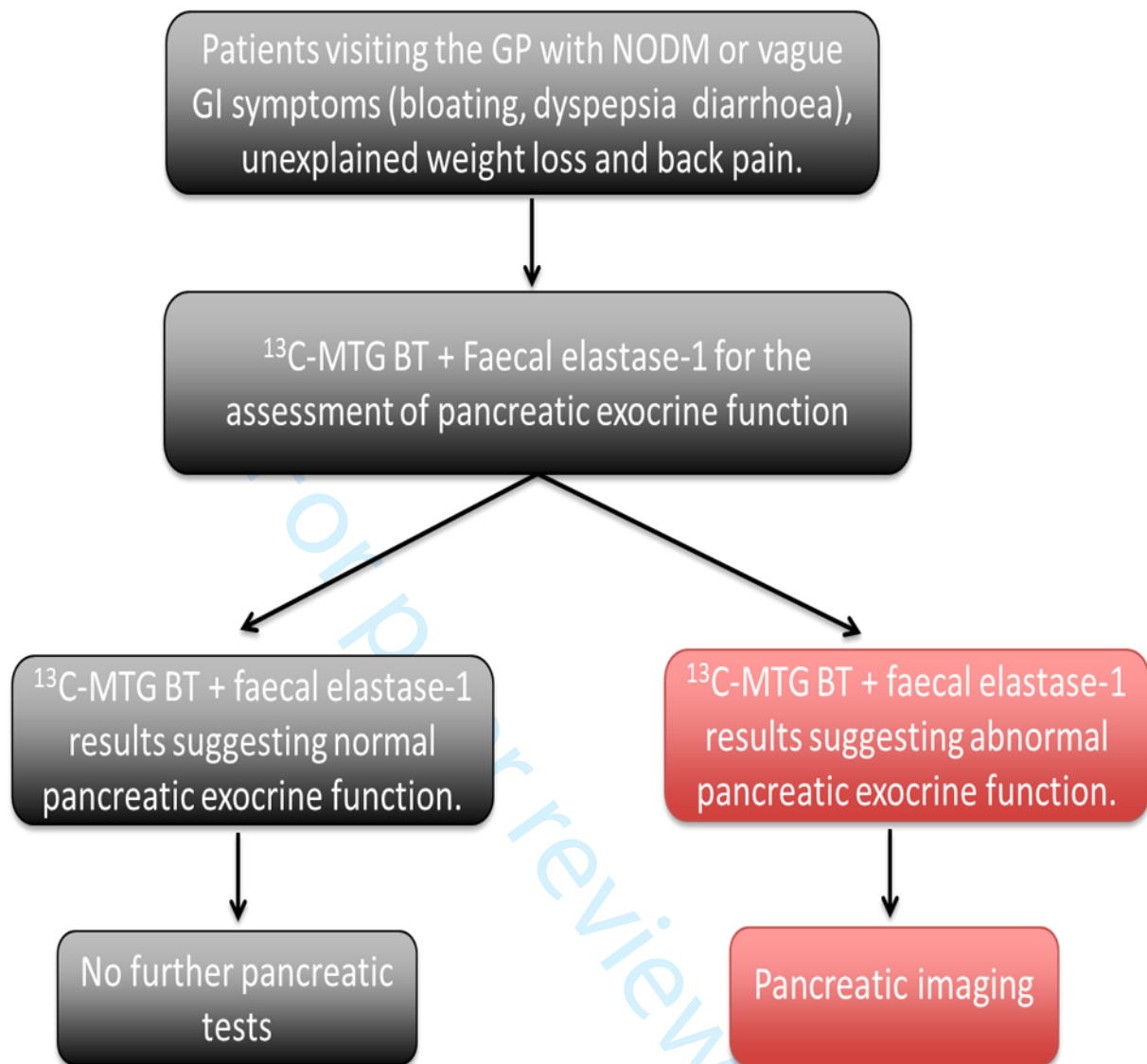
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Figure 1 Proposed diagnostic tool consisting of faecal elastase-1 test and the ¹³C-mixed triglyceride breath test to screen for pancreatic exocrine dysfunction in patients ‘at-risk’ of pancreatic cancer.

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Abbreviations: NODM- New Onset Diabetes Mellitus, ¹³C-MTG BT- ¹³C-mixed triglyceride breath test.

TRIPOD Checklist: Prediction Model Development

Section/Topic	Item	Checklist Item	Page
Title and abstract			
Title	1	Identify the study as developing and/or validating a multivariable prediction model, the target population, and the outcome to be predicted.	Page 1, lines 11-14
Abstract	2	Provide a summary of objectives, study design, setting, participants, sample size, predictors, outcome, statistical analysis, results, and conclusions.	Page 2, lines 22-43.
Introduction			
Background and objectives	3a	Explain the medical context (including whether diagnostic or prognostic) and rationale for developing or validating the multivariable prediction model, including references to existing models.	Pages 4-6, lines 57-130.
	3b	Specify the objectives, including whether the study describes the development or validation of the model or both.	Page 6, lines 124-130.
Methods			
Source of data	4a	Describe the study design or source of data (e.g., randomized trial, cohort, or registry data), separately for the development and validation data sets, if applicable.	Page 8, lines 145-152
	4b	Specify the key study dates, including start of accrual; end of accrual; and, if applicable, end of follow-up.	Page 8, lines 142-143.
Participants	5a	Specify key elements of the study setting (e.g., primary care, secondary care, general population) including number and location of centres.	Page 8, lines 139-141.
	5b	Describe eligibility criteria for participants.	Page 8, lines 153-161.
	5c	Give details of treatments received, if relevant.	n/a
Outcome	6a	Clearly define the outcome that is predicted by the prediction model, including how and when assessed.	Page 10, lines 203-205.
	6b	Report any actions to blind assessment of the outcome to be predicted.	n/a
Predictors	7a	Clearly define all predictors used in developing or validating the multivariable prediction model, including how and when they were measured.	n/a
	7b	Report any actions to blind assessment of predictors for the outcome and other predictors.	n/a
Sample size	8	Explain how the study size was arrived at.	Page 11, lines 215-222.
Missing data	9	Describe how missing data were handled (e.g., complete-case analysis, single imputation, multiple imputation) with details of any imputation method.	n/a
Statistical analysis methods	10a	Describe how predictors were handled in the analyses.	n/a
	10b	Specify type of model, all model-building procedures (including any predictor selection), and method for internal validation.	Page 12, lines 231-249.
	10d	Specify all measures used to assess model performance and, if relevant, to compare multiple models.	n/a
Risk groups	11	Provide details on how risk groups were created, if done.	n/a
Results			
Participants	13a	Describe the flow of participants through the study, including the number of participants with and without the outcome and, if applicable, a summary of the follow-up time. A diagram may be helpful.	n/a
	13b	Describe the characteristics of the participants (basic demographics, clinical features, available predictors), including the number of participants with missing data for predictors and outcome.	n/a
Model development	14a	Specify the number of participants and outcome events in each analysis.	n/a
	14b	If done, report the unadjusted association between each candidate predictor and outcome.	n/a
Model specification	15a	Present the full prediction model to allow predictions for individuals (i.e., all regression coefficients, and model intercept or baseline survival at a given time point).	n/a
	15b	Explain how to use the prediction model.	n/a
Model performance	16	Report performance measures (with CIs) for the prediction model.	n/a
Discussion			
Limitations	18	Discuss any limitations of the study (such as nonrepresentative sample, few events per predictor, missing data).	n/a

TRIPOD Checklist: Prediction Model Development

Interpretation	19b	Give an overall interpretation of the results, considering objectives, limitations, and results from similar studies, and other relevant evidence.	n/a
Implications	20	Discuss the potential clinical use of the model and implications for future research.	n/a
Other information			
Supplementary information	21	Provide information about the availability of supplementary resources, such as study protocol, Web calculator, and data sets.	n/a
Funding	22	Give the source of funding and the role of the funders for the present study.	Page 14, lines 303-304.

We recommend using the TRIPOD Checklist in conjunction with the TRIPOD Explanation and Elaboration document.

For peer review only

BMJ Open

DEPEND study protocol: Early detection of patients with pancreatic cancer: a pilot study to evaluate the utility of faecal elastase-1 and ¹³C-mixed triglyceride breath test as screening tools in high-risk individuals.

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DEPEND study protocol: Early detection of patients with pancreatic cancer: a pilot study to evaluate the utility of faecal elastase-1 and ¹³C-mixed triglyceride breath test as screening tools in high-risk individuals.

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Victoria M Goss ⁴, Jocelyn Walters ⁴, Zaed Z Hamady ^{1,2,3}.

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Abstract

Introduction Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The incidence of PC is increasing, with little or no improvement in overall survival and the best chance for long-term survival in patients with PC relies on early detection and surgical resection. In this study, we propose the use of a diagnostic algorithm that combines tests of pancreatic exocrine function (Faecal elastase-1 test (FE-1) and the ^{13}C -mixed triglyceride (^{13}C -MTG) breath test) to identify patients with PC that urgently needs imaging studies.

Methods and analysis This prospective pilot (proof of concept) study will be carried out on 25 patients with resectable PC, 10 patients with chronic pancreatitis and 25 healthy volunteers. This study will construct a predictive algorithm for PC, utilising two tests of pancreatic exocrine function, FE-1 test and the ^{13}C -MTG breath test. Continuous flow isotope ratio mass spectrometry in the ^{13}C -MTG breath test will be used to analyse enriched $^{13}\text{CO}_2$ in exhaled breath samples. The additional predictive benefit of other potential biomarkers of PC will also be considered. Potential biomarkers of PC showing abilities to discriminate between patients with PC from healthy subjects or patients with chronic pancreatitis will be selected by metabolomic analysis.

Ethics and dissemination The study was approved by the North of Scotland Research and Ethics Committee on 1 October 2020 (reference: 20/NS/0105, favourable opinion granted). The results will be disseminated in presentations at academic national/international conferences and publication in peer-review journals.

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Strength and limitations of this study

- We will use a non-invasive ¹³C-breath test to construct a predictive algorithm for detecting pancreatic cancer (PC).
- PC patients with potentially curative pancreatic resection will be recruited in this study.
- The predictive algorithm may have poor ability to discriminate between PC and chronic pancreatitis patients.
- Participants’ return of faecal specimens may be limited by embarrassment, hesitancy or inability of the study participants to provide a sample.

65 INTRODUCTION

66 Pancreatic cancer (PC) is the fifth leading cause of cancer related death in the UK. The UK
67 has one of the worst survival rates in Europe, with an average life expectancy at diagnosis of
68 just 4–6 months and 20% survival at 1 year. In addition, in the UK, only about 7% of people
69 survive for 5 years or longer¹. Since complete surgical resection remains the only potentially
70 curative option for PC, it is crucial to identify PC at an early stage, when it is still resectable.
71 Unfortunately, only 10–20% of patients with PC are diagnosed at a stage when curative
72 surgery remains an option².

73
74 The early symptoms of PC are usually vague and include weight loss, upper abdominal pain,
75 backache, fatigue, diarrhoea and nausea. In 93% of patients, these symptoms may have
76 been present for up to two years before diagnosis of PC³. Recent studies show that around
77 40% of patients diagnosed with PC require three or more visits to their GP before they are
78 referred to a specialist^{4 5} thereby leading to a delay in the diagnosis. There is also an
79 increased risk of developing PC in those with new-onset diabetes mellitus (NODM) i.e. less
80 than three years since diagnosis⁶ and as many as 25% of PC cases are believed to present
81 with NODM⁷.

82
83 Recent survey of GPs commissioned by Pancreatic Cancer Action⁸ suggested that the delay
84 between not having access to a reliable diagnostic referral pathway and a confirmed
85 diagnosis of PC is a major barrier to the early detection of PC. As the symptoms of PC are
86 often non-specific and unlikely to prompt further clinical investigation, there is an urgent
87 need to have a reliable and cheap screening test that can be used to identify high-risk
88 individuals who require urgent imaging studies such as CT scan to confirm or exclude the

presence of a pancreatic mass⁹. Having such a test with reliable sensitivity and negative predictive value (NPV), would enable the GP to select patients for urgent referral to further investigations of PC. A high NPV value would enable the GP to reassure the patient that they did not need further imaging studies. Our innovative approach is to use tests of pancreatic exocrine function combined as a screening test in patients presenting with the vague symptoms that may be associated with PC.

Although pancreatic exocrine insufficiency (PEI) is a well-known complication of pancreatic cancer^{10 11}, its presence is frequently overlooked in patients with advanced PC resulting in a decreased quality of life (QoL), malnutrition and morbidity¹². Currently, the diagnostic tests of pancreatic exocrine function such as faecal elastase-1 (FE-1) or the ¹³C-mixed triglyceride (¹³C-MTG) breath test, may be used to detect PEI in patients after pancreatic surgery and chronic pancreatitis¹³⁻¹⁶.

FE-1 test is a simple indirect and non-invasive method for assessing pancreatic secretion, which measures faecal concentrations of elastase-1, a proteolytic enzyme produced exclusively by pancreatic acinar cells, which binds to bile salts and passes through the gut with minimal degradation. FE-1 level correlate well with output of other pancreatic enzymes, such as amylase, lipase and trypsin^{17 18} and have been shown to detect PEI in patients with advanced PC and following pancreatic surgery¹⁴⁻¹⁶.

The ¹³C-MTG breath test assesses pancreatic lipase, it involves the ingestion of a ¹³C-labeled mixed triglyceride (1,3-distearoyl, 2- (carboxyl-¹³C)-octanoyl glycerol) mixed with a test meal. The ¹³C-MTG contains a ¹³C-labeled medium-chain fatty acid (¹³C-octanoate) at the Sn-

2 position, and two long-chain fatty acids (stearic acid) at the Sn-1 and Sn-3 positions of the glycerol backbone of the triacylglycerol. The ingested ^{13}C -MTG then undergoes the lipolysis of the two-long chain fatty acids from ^{13}C -MTG by pancreatic lipase to produce free fatty acids (stearic acid and ^{13}C -octanoate) and monoacylglycerol. After the intestinal absorption, the ^{13}C -free fatty acids are oxidised in the liver producing $^{13}\text{CO}_2$, which is then released on breath. Therefore, the $^{13}\text{CO}_2$ concentration in breath reflects pancreatic lipase activity. The ^{13}C -MTG breath test can be an alternative or complementary to the FE-1 test for PEI. The ^{13}C -MTG breath test has been validated in patients with chronic pancreatitis¹⁹ and patients with pancreatic cancer following surgical resection of the cancer^{20,21}, but not prior to surgery in patients with resectable PC.

Therefore, in order to improve the prognosis of patients with pancreatic cancer, it is essential to detect tumors at early stages, when they are more likely to be resectable. In this study, we will validate FE-1 test and ^{13}C -MTG breath test as potential early diagnostic biomarkers of resectable pancreatic cancer and test their performance in discriminating between patients with PC and healthy subjects. These tests can provide a non-invasive risk stratification tool to identify patients at risk of having PC when they present with NODM, weight loss or vague upper GI symptoms. Furthermore, this will provide a rational approach to patients identified to have PEI to be selected to have pancreatic imaging such as computed tomography (CT) scan to confirm the presence of PC and having a pancreatic resection earlier (Figure 1). In addition, we will also be identifying other potential biomarkers of PC through unbiased multiomics analysis of blood samples (proteomics and metabolomics). The resulting data will be compared to the healthy control group and patients with chronic pancreatitis aiming to identify signals that are distinct to PC.

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3 137 **METHODS AND ANALYSIS**
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5 138 **Main centre**
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7 139 NIHR Southampton Biomedical Research Centre, Southampton Centre for Biomedical
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9 140 Research, University Hospital Southampton NHS Foundation Trust, Southampton, UK.
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13 141 **Dates of the study**

14 142 From 16th March 2021 to 16th March 2022.
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18 144 **Design**

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20 145 This is a single cross-sectional pilot study of patients with PC, chronic pancreatitis and
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22 146 healthy controls, matched for age and sex. The primary objective is to determine if ¹³C-
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24 147 mixed triglyceride (¹³C-MTG) breath test and faecal elastase-1 (FE-1) concentrations
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26 148 combined can discriminate between the groups. Patients with PC will be recruited at the
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28 149 pre-operative stage within two weeks of initial diagnosis. The secondary objective of this
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30 150 study is the analysis of multiomic (proteomic and metabolomic) profiles and comparison
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32 151 between the groups.
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37 152 **Inclusion criteria**

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40 153 At the University Hospital Southampton NHS Foundation Trust, on average, 120 patients
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42 154 with PC are treated with surgery while 350 patients with PC are treated with chemotherapy
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44 155 every year. Male and female participants aged from 30-85 within the Wessex pancreatic
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46 156 cancer catchment area will be suitable for inclusion. In the cancer cohort, patients with a
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48 157 diagnosis of resectable pancreatic ductal adenocarcinoma (PDAC) (stage 1-2) will be
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50 158 included. Patients with chronic pancreatitis and healthy subjects will also be recruited as
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52 159 control groups and matched for age and sex to the cases of PDAC.
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3 162 **Exclusion criteria**
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6 163 Patients and controls with autoimmune or other chronic inflammatory conditions, or those
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8 164 who have previously had pancreatic disease or liver dysfunction will be excluded.
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Intervention(s) or method

Each participant will be given a participant information sheet explaining the research and will sign an informed consent form. Participants are required to fast overnight for at least 12 hours prior to undergoing the ¹³C-mixed triglyceride (¹³C-MTG). At the Clinical Research Facility (CRF), full infection control precautions (related to COVID-19 pandemic) will be adhered to throughout the visit. Participants will have their temperature checked on arrival to ensure they are not pyrexial as part of screening for Covid-19, which will exclude them from the study. Anthropometric measurements (height, weight and waist circumference) of the participants will be recorded. Baseline blood samples will be taken prior to undergoing the ¹³C-MTG breath test. The data collected will include the following: Full blood count (FBC), Urea and Electrolyte blood test (U&E), Liver function tests (AST, ALT, ALP, albumin and GGT) /clotting studies, glycated hemoglobin (HbA1c %), Lipid profile, Vitamins and micronutrients, Hyaluronic acid, Procollagen III amino terminal peptide (PIINP) and Tissue inhibitor of metalloproteinase-1 (TIMP-1). These tests will be analysed immediately after the blood has been drawn from the participants by the chemical pathology service laboratory at Southampton General Hospital. Further blood samples will be processed and stored for multiomics analysis. Participants will also be asked to provide spot stool samples for faecal elastase-1 (FE-1) concentrations by using a commercially available ELISA kit. Results of the FE-1 test will be expressed as µg/g of stool.

¹³C-MTG breath test protocol

Each participant will be asked to provide 2 baseline breath samples by blowing into 2 × 10ml exetainer tubes through a straw while wearing a facemask. After that, participants will be given an oral dose of 250mg of 2-¹³C-octanoyl-1,3-distearin (¹³C-MTG -Cambridge Isotope Laboratories, Andover, MA, USA) together with a solid test meal consisting of a crispbread or other gluten-free alternative, and 200ml of water. Post-prandial breath samples will be then collected every 30mins for 4 hours. No food or drinks except for water will be allowed during the duration of the breath test study. The collected breath samples will be analysed for the ratio of ¹³CO₂ to ¹²CO₂ using a Continuous Flow Isotope Ratio Mass Spectrometer (CF-IRMS SERCON Ltd, Crewe, UK) at the Southampton Centre for Biomedical Research. The increase in ¹³CO₂ content with regard to the baseline value of the initial breath sample is expressed as atom % excess and we will express the results of the ¹³C-MTG breath test as the cumulative percentage of ¹³C-label recovered on breath as ¹³CO₂ over 4 hours (cPDR over 4 hours).

Primary Outcome: Prediction of PC using a logistic model with measurement of FE-1 concentration expressed as µg/g of stool and ¹³C-MTG breath test expressed as cPDR over 4hrs as covariates.

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Statistical analysis and data analysis plans

Sample size calculation

The sample size calculation for the proposed study is based on a previous study assessing pancreatic exocrine function using a ¹³C-triglyceride breath test in healthy subjects and patients with a localised pancreatic mass ¹³. In this study, they used the ¹³C-trioctanoin breath test to assess pancreatic exocrine function in 14 patients with a localised pancreatic mass and 5 healthy control subjects. The results of the study showed means ± standard deviation (SD) of the recovery of ¹³CO₂ over 3hrs after undergoing the ¹³C-trioctanoin breath test of 42% ± 3.4% for the healthy controls and 24.2% ± 10.5% for patients with a localised pancreatic mass with an effect size or mean difference between both groups of 17.8% and a within group SD of 9.1% (based on the SD estimates of 3.4% and 10.5% for each group respectively).

Using data from this study, we calculated that a sample size of 25 subjects in each group (patients with PC, patients with chronic pancreatitis and healthy control subjects) will give 100% power to detect a difference of 17.8% between patients with PC and healthy control group assuming a standard deviation of 10.5% in patients with PC at a significance level of 5% using a two-tailed test. The power calculation was carried out using IBM SPSS SamplePower version 3.

234 Statistical tests

235 Data will be presented as mean + SD, median + IQR when appropriate and 95% CI.

236 Comparison of results of FE-1 and the ¹³C-MTG breath test between each group will be

237 evaluated by employing either the one way analysis of variance test (if the variables are

238 normally distributed and parametric validity conditions are fulfilled) or the Mann Whitney

239 U-test or Kruskal-Wallis test (if the variables are not normally distributed and non-

240 parametric validity conditions are fulfilled). A multi-variable logistic regression analysis will

241 be performed in order to determine the independent factors that are significantly

242 associated with the presence of PDAC and logistic regression models will be constructed

243 based on the identified independent factors. Furthermore, multi-variable regression analysis

244 will be used to determine the optimal algorithm that combines the results of FE-1 test and

245 ¹³C-MTG breath test. We will also test whether potential biomarkers of PC identified by

246 unbiased omics analysis improve the prediction of PDAC. A Receiver operating characteristic

247 (ROC) curve will be constructed to evaluate the diagnostic accuracy (sensitivity, specificity,

248 positive and negative predictive value, receiver operator curve analysis) of the algorithm for

249 PDAC. Furthermore, the optimal cut off point for the algorithm to identify patients with PC

250 will be estimated using the Youden index. A two-sided p value of <0.05 will be considered to

251 indicate statistical significance. All analysis will be performed using SPSS (IBM, V.25.0) or

252 Stata (Statacorp, v16.0).

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254 Patient and public involvement statement

255 Patients, healthy subjects and the public were not involved in the study design, recruitment

256 and conduct. We do not plan to inform the result to the study participants unless they apply

257 for it.

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3 258 **ETHICS AND DISSEMINATION**
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5 259 Ethical approval has been granted by the North of Scotland Research and Ethics Committee,
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8 260 reference: 20/NS/0105, 1 October 2020. This manuscript reflects the latest protocol V.7
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10 261 approved 29 October 2020.
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15 263 The study will be conducted according to the principles of Good Clinical Practice, General
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17 264 data Protection Regulation and Data Protection Act 2018 for Health and care Research. The
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20 265 sponsor and study team will ensure approval of the study protocol, participant information
21
22 266 sheets, consent forms, letters to general practitioners and supporting documents by the
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25 267 appropriate regulatory body and research and ethics committee prior to participant
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27 268 recruitment. Documents will be stored securely with restricted access for at least 10 years.
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30 269 Written, informed consent will be obtained from each patient and an identification number
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32 270 provided. Any published data will not contain personal identifiable data. The results of this
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35 271 study will be disseminated by presentation at academic national/international conferences
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37 272 and publication in peer-review journals.
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Authors' contributions: ZH and PA have designed the study concept and formulated the initial idea; ZH is the Chief investigator and responsible for protocol review. DM and PA wrote the initial protocol. CB, PA, SW and ZH were involved in statistical analysis plan. DM and ZH was responsible for organising the study operation requirements, identifying and discussion with eligible participants. DM is responsible for patient recruitment and consent. DM and PA prepared the test meal and obtained the samples from the participants, VG is sorting the standard operating procedures (SOPs) and logistics with biobanking and sample process. PA is responsible for the analysis of the breath samples and the interpretation of the results from the breath test. JW is managing the study team and logistics related to clinical trial involvement. PA, DM, CB, SW, VG, JW and ZH were involved in drafting of the protocol.

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Competing interest: None.

Patient consent for publication: Not required.

Provenance and peer review: Not commissioned; externally peer reviewed.

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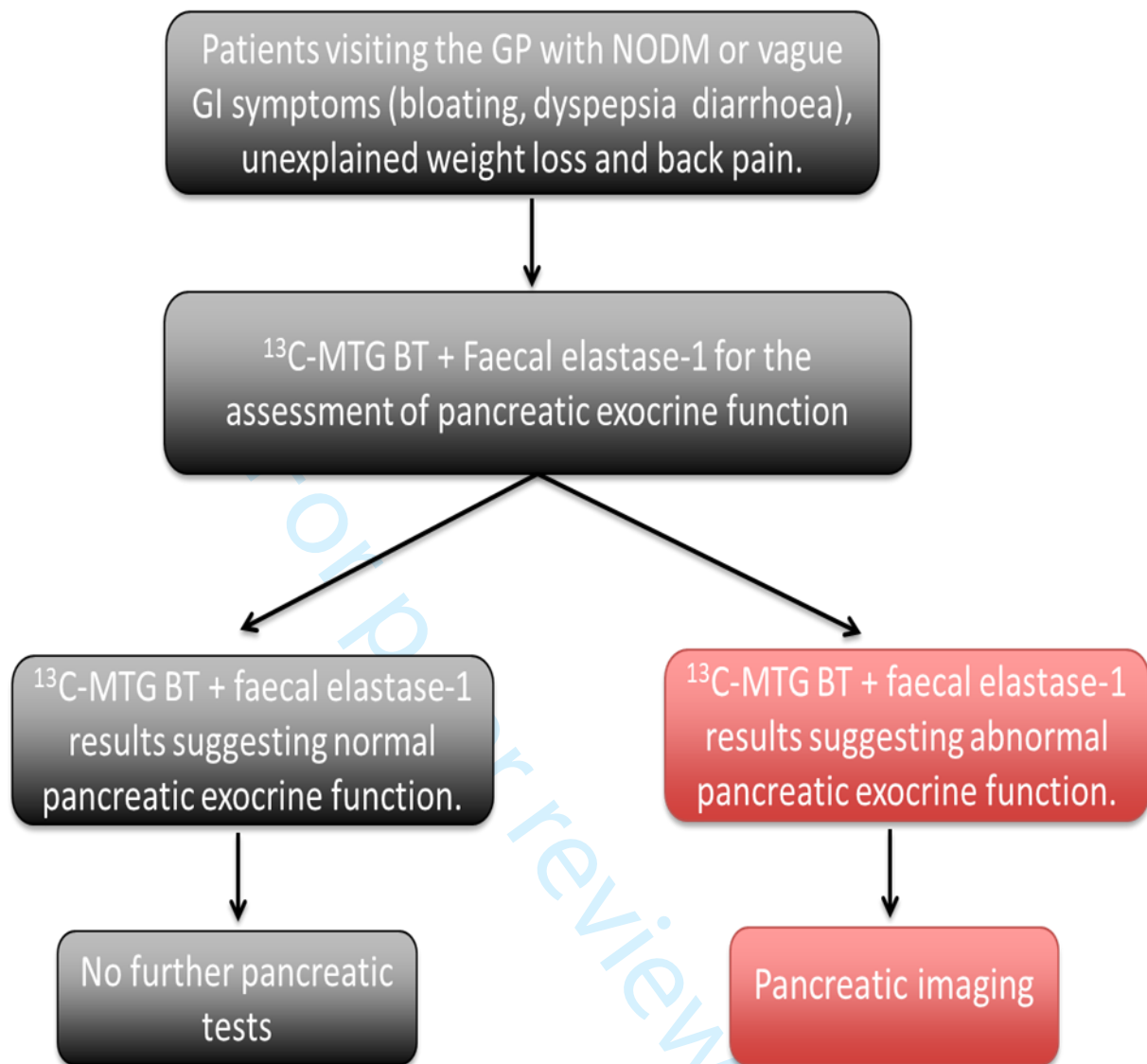
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Figure 1 Proposed diagnostic tool consisting of faecal elastase-1 test and the ¹³C-mixed triglyceride breath test to screen for pancreatic exocrine dysfunction in patients ‘at-risk’ of pancreatic cancer.

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Abbreviations: NODM- New Onset Diabetes Mellitus, ¹³C-MTG BT- ¹³C-mixed triglyceride breath test.

TRIPOD Checklist: Prediction Model Development

Section/Topic	Item	Checklist Item	Page
Title and abstract			
Title	1	Identify the study as developing and/or validating a multivariable prediction model, the target population, and the outcome to be predicted.	Page 1, lines 11-14
Abstract	2	Provide a summary of objectives, study design, setting, participants, sample size, predictors, outcome, statistical analysis, results, and conclusions.	Page 2, lines 22-43.
Introduction			
Background and objectives	3a	Explain the medical context (including whether diagnostic or prognostic) and rationale for developing or validating the multivariable prediction model, including references to existing models.	Pages 4-6, lines 57-130.
	3b	Specify the objectives, including whether the study describes the development or validation of the model or both.	Page 6, lines 124-130.
Methods			
Source of data	4a	Describe the study design or source of data (e.g., randomized trial, cohort, or registry data), separately for the development and validation data sets, if applicable.	Page 8, lines 145-152
	4b	Specify the key study dates, including start of accrual; end of accrual; and, if applicable, end of follow-up.	Page 8, lines 142-143.
Participants	5a	Specify key elements of the study setting (e.g., primary care, secondary care, general population) including number and location of centres.	Page 8, lines 139-141.
	5b	Describe eligibility criteria for participants.	Page 8, lines 153-161.
	5c	Give details of treatments received, if relevant.	n/a
Outcome	6a	Clearly define the outcome that is predicted by the prediction model, including how and when assessed.	Page 10, lines 203-205.
	6b	Report any actions to blind assessment of the outcome to be predicted.	n/a
Predictors	7a	Clearly define all predictors used in developing or validating the multivariable prediction model, including how and when they were measured.	n/a
	7b	Report any actions to blind assessment of predictors for the outcome and other predictors.	n/a
Sample size	8	Explain how the study size was arrived at.	Page 11, lines 215-222.
Missing data	9	Describe how missing data were handled (e.g., complete-case analysis, single imputation, multiple imputation) with details of any imputation method.	n/a
Statistical analysis methods	10a	Describe how predictors were handled in the analyses.	n/a
	10b	Specify type of model, all model-building procedures (including any predictor selection), and method for internal validation.	Page 12, lines 231-249.
	10d	Specify all measures used to assess model performance and, if relevant, to compare multiple models.	n/a
Risk groups	11	Provide details on how risk groups were created, if done.	n/a
Results			
Participants	13a	Describe the flow of participants through the study, including the number of participants with and without the outcome and, if applicable, a summary of the follow-up time. A diagram may be helpful.	n/a
	13b	Describe the characteristics of the participants (basic demographics, clinical features, available predictors), including the number of participants with missing data for predictors and outcome.	n/a
Model development	14a	Specify the number of participants and outcome events in each analysis.	n/a
	14b	If done, report the unadjusted association between each candidate predictor and outcome.	n/a
Model specification	15a	Present the full prediction model to allow predictions for individuals (i.e., all regression coefficients, and model intercept or baseline survival at a given time point).	n/a
	15b	Explain how to use the prediction model.	n/a
Model performance	16	Report performance measures (with CIs) for the prediction model.	n/a
Discussion			
Limitations	18	Discuss any limitations of the study (such as nonrepresentative sample, few events per predictor, missing data).	n/a

TRIPOD Checklist: Prediction Model Development

Interpretation	19b	Give an overall interpretation of the results, considering objectives, limitations, and results from similar studies, and other relevant evidence.	n/a
Implications	20	Discuss the potential clinical use of the model and implications for future research.	n/a
Other information			
Supplementary information	21	Provide information about the availability of supplementary resources, such as study protocol, Web calculator, and data sets.	n/a
Funding	22	Give the source of funding and the role of the funders for the present study.	Page 14, lines 303-304.

We recommend using the TRIPOD Checklist in conjunction with the TRIPOD Explanation and Elaboration document.

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