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Observational Study

Identification of exhaled breath volatile organic compounds to distinguish between

pancreatic adenocarcinoma, pancreatic cystic neoplasm, and patients without

pancreatic lesions

Tiankanon K et al. PDAC exhaled breath volatile organic compound

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Abstract

BACKGROUND

Volatile organic compounds (VOC) are a promising potential biomarker that may be

able to identify the presence of cancers.

AIM

To identify exhaled breath VOCs that distinguish pancreatic ductal adenocarcinoma

(PDAC) from intraductal papillary mucinous neoplasm (IPMN) and healthy volunteers.

METHODS

We collected exhaled breath from histologically proven PDAC patients, radiological

diagnosis IPMN, and healthy volunteers using the ReCIVA® device during 10/2021-

11/2022. The VOCs were identified by thermal desorption-gas chromatography/field-

asymmetric ion mobility spectrometry and compared between groups.

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RESULTS

156 participants (44% male, mean age 62.6 ± 10.6) were enrolled (54 PDAC, 42 IPMN, and 60 controls). Among the 9 VOCs identified, two VOCs that showed the difference between groups were dimethyl sulfide [0.73 vs 0.74 vs 0.94 arbitrary units (AU), respectively; P = 0.008], and acetone dimers (3.95 vs 4.49 vs 5.19 AU, respectively; P < 0.001). After adjusting for the imbalance parameters, PDAC showed higher dimethyl sulfide levels than the control and IPMN groups, with adjusted odds ratio (aOR) of 6.98 (95%CI: 1.15-42.17) and 4.56 (1.03-20.20), respectively (P < 0.05 both). Acetone dimer levels were also higher in PDAC compared to controls and IPMN (aOR: 5.12 (1.80-14.57) and aOR: 3.35 (1.47-7.63), respectively (P < 0.05 both). Acetone dimer, but not dimethyl sulfide, performed better than CA19-9 in PDAC diagnosis (AUROC 0.910 vs 0.796). The AUROC of acetone dimer increased to 0.936 when combined with CA19-9, which was better than CA19-9 alone (P < 0.05).

CONCLUSION

The VOCs that potentially distinguish PDAC from IPMN and healthy participants were dimethyl sulfide and acetone dimer. Additional prospective studies are required to validate these findings.

Key Words: Volatile organic compound; Pancreas; Adenocarcinoma; Pancreatic Intraductal Neoplasms; Breathing

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Core Tip: Participants with pancreatic ductal adenocarcinoma (PDAC) exhibit distinct exhaled breath volatile organic compounds (VOC) from those with intraductal papillary

mucinous neoplasm (IPMN) or the general population. Dimethyl sulfide and acetone dimer were the VOCs shown to potentially identify PDAC from IPMN and healthy subjects.

INTRODUCTION

Pancreatic ductal adenocarcinoma (PDAC), or pancreatic cancer, is the most common pancreatic malignant lesion, accounting for the seventh highest total of cancer-related deaths worldwide^[1]. The incidence of PDAC is on the rise and is considered as one of the most deadly cancers^[1]. Due to the non-specific symptoms of PDAC, most patients are diagnosed at an advanced stage with a 5-year survival rate of less than 8%^[1-4]. Current diagnostic techniques rely on imaging, endoscopic ultrasonography, and biopsy to determine tumor histology. Unfortunately, the performance of these techniques is suboptimal for detecting tumors smaller than 2 cm in diameter. Additionally, biomarkers commonly used, particularly CA19-9, have shown poor performance^[5,6].

In addition to pancreatic cancer, another pancreatic lesion that is being detected more frequently due to advances in diagnostic imaging technology is intraductal papillary neoplasm (IPMN)^[7]. The reported prevalence of incidental pancreatic cysts, including IPMN, identified on multi-detector computed tomography (CT) and magnetic resonance imaging (MRI), is as high as 2.6% and 13.5%, respectively^[8,9]. These lesions carry the risk of malignant transformation to pancreatic cancer, which occurs 76% of the time in main duct type IPMN (MD-IPMN), 16% in branch duct type IPMN (BD-IPMN), and 46% in mixed type IPMN^[10]. Similar to PDAC, no biomarkers are currently available to predict the malignant transformation of IPMN.

Volatile organic compounds (VOC) are organic compounds generated by the metabolism of cells and released into the blood and other body fluids, with high vapor pressure under room temperature conditions^[11]. VOCs can be found in exhaled breath, urine, bile, feces, or saliva, and represent biochemical reactions caused by biological activities, such as apoptosis, oxidative stress, or inflammation^[11]. According to

accumulating evidence, the analysis of VOC profiles is a promising novel cancer biomarker to diagnose many types of cancer, such as lung, breast, gastrointestinal, and urological cancers^[12]. One of the simplest biological specimens to access is breath samples. Samples can be collected in large volume, completely noninvasively, and with high patient acceptability. Previous studies have shown satisfactory results in the ability of exhaled breath to distinguish VOC profiles between cancer and noncancer participants^[13-15]. We hypothesized that biomarkers for PDAC exist in exhaled breath and that these VOC profiles in PDAC patients differ from those with IPMN and those without pancreatic lesions. We also believe that VOC profiles may have the potential as an early detection biomarker and screening tool for patients at risk of this malignancy.

Although a few studies have suggested that VOCs could be employed as new biomarkers, the VOCs reported in each investigation varied. This was owing to the fact that the type and quantity of VOCs discovered depended on the methods employed^[16]. Because of their high resolution and sensitivity, gas chromatography-mass spectrometry (GC-MS) and selective ion flow tube mass spectrometry (SIFT-MS) were the most often used techniques^[12]. However, these techniques are expensive, time-consuming, and require expertise to operate, making them unsuitable for point-of-care screening programs^[17,18]. Field asymmetric ion mobility spectrometry (FAIMS), which has equivalent sensitivity to GC-MS and SIFT-MS but is less expensive and less technical, has recently been recommended as being more suitable for clinical study^[18,19]. To date, no study has been conducted to differentiate VOCs for PDAC using FAIMS. Our objective in this study was to identify VOCs of exhaled breath that could be used to distinguish PDAC from IPMN and controls without pancreatic lesions using the FAIMS technique, and to investigate the diagnostic performance of VOCs in comparison to CA19-9.

MATERIALS AND METHODS

We conducted a single-center, cross-sectional study between October 2021 and January 2023 at the Center of Excellence for Innovation and Endoscopy in Gastrointestinal

Oncology, Division of Gastroenterology, Department of Medicine, Chulalongkorn University. The study protocol was reviewed and approved by the Institutional Research Committee, Faculty of Medicine, Chulalongkorn University (IRB No.0482/65) and registered in the Thai Clinical Trials Registry (TCTR20211109002).

Participant population

Participants with cytological or histological confirmation of PDAC and radiological confirmation of IPMN were enrolled. The stages of PDAC were classified as early, locally advanced, and advanced stages according to the American Joint Committee on Cancer/Union for International Cancer Control TNM classification of malignant tumors[20]. IPMN patients were classified as MD-IPMN, BD-IPMN, and mixed-type IPMN according to the Sendai and Fukuoka consensus guidelines^[7]. Any PDAC or IPMN participant who had undergone previous treatment, including surgical intervention, pancreatic radiation, systemic chemotherapy, and endoscopic local ablations, were excluded from the study to avoid possible effects of the treatment. Healthy volunteers without pancreatic lesions confirmed by abdominal imaging, including transabdominal ultrasound, endoscopic ultrasonography, abdominal CT, or MRI, with a minimum age of 18 years were also recruited as the control group. Participants with a history of other malignancies, breathing through tracheostomy tubes, pregnant at the time of recruitment, concurrent active infection, history of infection within 3 wk prior to breath sample collection, diagnosed with cirrhosis, endstage renal disease, or participants with poorly controlled diabetes, defined as HbA1C of \geq 8%, were also excluded from the study.

Data collection

All eligible participants were provided with details of the study. Verbal and written informed consent were obtained from every participant before enrollment. Baseline characteristics including age, gender, smoking status, alcohol consumption, comorbidities, such as chronic viral hepatitis B and C, nonalcoholic fatty liver disease,

ischemic heart disease, diabetes, current medication, and previous history of endoscopic treatment were obtained from participants and electronic medical records. Laboratory data associated with pancreatic diseases including liver function test, amylase, lipase, and the tumor biomarker CA19-9 were also obtained prior to breath sampling. Information on tumor location, size, staging, and metastasis sites was gathered from electronic medical records.

Breath samples collection

All participants had been advised to fast, stop smoking, exercise, and withhold current medication for 6 h prior to breath sample collection to minimize confounders in exhaled breaths. Participants were instructed to breathe normally through a disposable face mask that covered both the nose and mouth for 2-3 min. The face mask was connected to the ReCIVATM breath sample system (Owlstone Medical, Cambridge, United Kingdom), which was coupled with software to monitor real-time progression of the breath sample collection (Figure 1A). During this process, a pure oxygen gas supply was provided through the face masks to avoid contamination of confounding surrounding gas into the system. Approximately 100 mL of exhaled air was collected and distributed equally into 4 thermal desorption (TD) tube (UnityTM-XR, Markes International Ltd, Llantrisant, United Kingdom) prior to VOC extraction process.

VOC extraction and measurement

The VOC extraction protocol was carried out using a previously validated method published in the previous study^[21]. Extraction of VOCs from TD tubes was performed using a thermal desorption unit (UnityTM-XR, Markes International Ltd, Llantrisant, United Kingdom). The collected TD tubes were heated from 10 °C to 250 °C in split flow mode to release the VOCs collected inside. The VOC profiling was then analyzed by gas chromatography (GC) using Thermo Scientific TRACE1310 GC system (Waltham, MA, United States), which is equipped with an HP-PLOT U gas chromatography column of 30 m length × 0.32 mm of inner diameter × 10 μm of film thickness (Agilent

Technologies, Santa Clara, CA, United States). The GC column was initially programmed to heat at 40 °C for 2 min and ramp to 130 °C at the rate of 10 °C/min for 10 min. Helium was used as the carrier gas at a flow rate of 1 mL/min during the VOC separation. The extracted VOCs then flowed through a transfer line at 130 °C to final extraction using a FAIMS system (Owlstone Medical Lonestar VOC Analyzer FAIMS system, Cambridge, United Kingdom) (Figure 1B). After final extraction, VOCs extracted from each participant's group were shown in the form of chromatograms (Figure 1C).

VOC identification

The identification was achieved by calibrating the retention time of standard solutions with the chromatogram and comparing them with the in-house VOC data library^[21]. The in-house VOC data library was created with XGBoost to identify important features of VOC profiles, and the algorithm also chose the optimal combination of VOCs to yield optimal statistical parameters. The XGBoost algorithm is one of the AI algorithms established in the previous VOC study^[21]. The XGBoost enabled the minimization of false predictions by continuously learning from the model that was previously constructed as a series and thus maximizing the prediction model's efficiency. System calibration was performed daily to ensure the accuracy of all instruments. The difference in types of VOCs extracted and relative amounts of the VOCs, expressed as arbitrary unit (AU), were analyzed and compared between groups.

Statistical analysis

The sample size was calculated to achieve 80% power with a two-sided type I error of 0.05 based on data from the previous VOC study to distinguish between PDAC and the control population^[15]. With a 1:1:1 ratio among PDAC, IPMN, and control groups, a minimum sample size of 41 in each group was estimated.

The descriptive statistics of the categorical variables were expressed as frequency (percentage) and the descriptive statistics of the continuous variables were expressed as

mean \pm SD or median and range, as appropriate. Pearson's Chi-square test was used to compare categorical variables. The Mann-Whitney U test or ANOVA was used to compare differences between continuous variables. Multivariate logistic regression analysis was performed to assess the association between the level of VOCs and the participant's diagnosis. A receiver-operating characteristic (ROC) curve was constructed to identify the optimal cutoff point for VOCs to distinguish PDAC from other diagnoses. Statistical significance was established at P < 0.05. All statistical tests were performed with SPSS version 22.0.0 (SPSS Inc, Chicago, IL, United States).

RESULTS

Baseline characteristics

A total of 156 participants (44% male) with a mean age of 62.6 ± 10.6 years were enrolled. Among these, 54 participants were PDAC patients, 42 were IPMN patients, and 60 were selected for the control group. PDAC patients consisted of 3 (5%) early stage, 27 (50%) locally advanced stage, and 24 (44%) advanced stage. The advanced-stage PDAC metastatic sites included 17 (32%) liver, 12 (22.6%) lymph nodes, 8 (15%) peritoneum, and 4 (7%) lungs. Regarding the participants in IPMN group, 5 (11.9%) participants were classified as MD-IPMN, 35 (83.3%) as BD-IPMN, and 6 (14.2%) were malignant IPMN.

Baseline characteristics in terms of age, sex, smoking and alcohol consumption, and participant co-morbidities were not different among the three groups. Patients with PDAC, however, had statistically higher levels of bilirubin, aspartate aminotransferase (AST), alanine aminotransferase (ALT), and alkaline phosphatase (ALP), and international normalized ratio (INR). A lower level of albumin compared to IPMN and control groups was also reported (Table 1).

Comparisons of VOCs profiles

A total of 9 VOCs from exhaled breath were identified (Table 2). Two VOCs, dimethyl sulfide and acetone dimer, demonstrated a statistically significant difference between

the PDAC, IPMN, and control groups (0.94 vs 0.74 vs 0.73 AU; P = 0.008 for dimethyl sulfide; 5.19 vs 4.49 vs 3.97 AU; P < 0.001 for acetone dimer, respectively).

After adjusting for imbalanced factors between groups, including levels of bilirubin, AST, ALT, ALP, INR, and albumin levels using multivariate logistic regression, the PDAC group showed a significantly higher level of dimethyl sulfide compared to IPMN with an adjusted odd ratio (aOR) of 4.56 (95%CI: 1.03-20.20, P = 0.046), and the control group with aOR of 6.98 (95%CI: 1.15-42.17, P = 0.034), respectively (Figure 2).

The acetone dimer also showed a significantly higher level in the PDAC group compared to the IPMN group (aOR 3.35; 95%CI 1.47-7.63, P = 0.015), and the control group (aOR 5.12; 95%CI 1.80-14.57, P = 0.002) (Figure 3). However, when comparing the levels of dimethyl sulfide and acetone dimer of the IPMN group with those of controls, no statistically different VOC levels were demonstrated.

In subgroup analyses, the levels of dimethyl sulfide were higher in the metastatic PDAC than in the localized PDAC group (0.96 vs 0.92 AU; P = 0.016) as shown in Table 3. However, the acetone dimer level does not differ between PDAC stages.

Performance of dimethyl sulfide and acetone dimer as a biomarker

The mean value of the PDAC biomarker, CA19-9, was significantly different between PDAC, IPMN and healthy participants (mean \pm SD of 307 \pm 467, 43 \pm 89, and 11 \pm 5.4, respectively; P < 0.001). When looking at the ROC curves, CA19-9 with a cutoff value of 28.75 was able to distinguish PDAC patients from non-PDAC participants with an AUROC of 0.796.

Dimethyl sulfide using a cutoff value of 0.78 AU distinguished PDAC from non-PDAC participants with an AUROC of 0.671 and 68.5% sensitivity, 59.8% specificity, 47.4% positive predictive value (PPV), 78.2% negative predictive value (NPV) and 62.8% accuracy. These results did not show any statistical difference compared to the performance of CA19-9 (P = 0.115).

Acetone dimer with a cutoff value of 4.97 AU, outperformed CA19-9 with an AUROC of 0.910 and 87.0% sensitivity, 92.2% specificity, 85.5% PPV, 93.1% NPV, and 90.4%

accuracy. When combined CA19-9 with acetone dimer level, at a cutoff point of 0.29 AU, the AUROC increased to 0.936 which was significantly better than the performance of CA19-9 alone (P = 0.020) and reported a sensitivity, specificity, PPV, NPV, and accuracy of 96.7%, 86.0%, 82.9%, 97.4%, and 90.4%, respectively. (Table 4 and Figure 4).

DISCUSSION

Our study reported that both exhaled breath VOCs dimethyl sulfide and acetone dimer showed statistically higher concentrations in the PDAC group compared to the non-PDAC and control groups. The significant discriminatory performance of VOCs, particularly acetone dimer, supported previous evidence that exhaled VOCs have the potential to be new biomarkers for PDAC detection^[13-15].

Our findings of higher acetone dimer levels in PDAC patients were consistent with the findings from previous research[14]. Acetone dimer levels in human secretions have been reported to be higher in a variety of digestive cancers, including colorectal cancer^[22,23], cancer^[22], $(HCC)^{[21,24]}$, gastric hepatocellular carcinoma cholangiocarcinoma^[25]. The dysregulation of ketone metabolism caused by cancer cells is thought to be one possible explanation for the increase in acetone levels. Acetone is a ketone compound derived from the spontaneous degradation of acetoacetate, a ketone body generated from acetyl-CoA, mainly obtained from the beta-oxidation of longchain fatty acids [22,26]. Because of their ability to divide uninhibitedly, cancer cells cause an abnormal increase in glucose metabolism and deplete the availability of glucose for the surrounding cells, shifting the energy source toward the lipid metabolism pathway, resulting in increased ketone body production^[27]. These extra ketone bodies can be metabolized by the mitochondria of both cancer and surrounding cells via the tricarboxylic acid cycle and provide energy to the cell in addition to the normal glucose pathways^[22].

Despite the findings that acetone dimer levels were higher in PDAC groups, acetone monomer levels failed to show significant differences. Acetone is typically reported without labeling the ionized form (whether it is acetone monomer or acetone dimer)

when using GC-MS or SIFT-MS methods^[14,28]. However, by utilizing FAIMS, we were able to discriminate between the two ionized forms since the FAIMS analyzer contains a beta radiation source (Ni-63) that emits electrons that ionize the air and moisture that flows through the analyzer. The process generates positive and negative hydronium ions, which react with molecules with a strong proton affinity, such as acetone, to produce distinct ion peaks of an acetone monomer or dimer^[29]. The amount of monomer or dimer produced varies by factors such as temperature and, more importantly, substance concentration. A previous study revealed that proton-bound dimers occur when the gas concentration is sufficiently high^[30]. As a result, the acetone dimer may have a stronger association with the amount of acetone generated than the monomer.

Another source of acetone is the alcohol dehydrogenase enzyme, which uses an oxidation-reduction reaction to convert isopropyl alcohol to acetone^[31]. Not surprisingly, previous research showed that acetone is directly related to isopropyl alcohol, and these two VOCs usually co-exist^[32,33]. While acetone levels were higher in PDAC, our study failed to show statistical differences in isopropyl alcohol levels between PDAC and non-PDAC participants. This suggests that, in contrast to other cancers, the alcohol dehydrogenase enzyme may not be the primary energy pathway in IPMN and PDAC.

Our findings also revealed that PDAC patients produced more dimethyl sulfide compared to non-PDAC patients and controls, which is one of the volatile sulfur compounds produced by the metabolism of sulfur-containing amino acids like methionine and cysteine^[34]. Dimethyl sulfide is normally detected in very low concentrations in normal population serum. Higher concentrations were found in patients with pulmonary hypertension, cirrhosis, and hepatocellular carcinoma^[24]. The proposed mechanism of dimethyl sulfide production is the methylation of methanethiol by thiol S-methyltransferase, which occurs endogenously or is synthesized by the microbiota in the intestinal tract (e.g., Streptococcus, Fusobacterium, Salmonella, Enterobacter, and Helicobacter)^[35]. A previous research reported significantly lower levels

of dimethyl sulfide in cholangiocarcinoma exhaled breath, which could reflect a reduced formation of the antioxidant glutathione linked to cancer pathways^[25]. In another study, dimethyl sulfide was found to be one of the discriminatory VOCs in HCC when compared to cirrhosis^[24].

To the best of our knowledge, no previous study has demonstrated statistical differences in dimethyl sulfide levels between PDAC and non-PDAC populations. In this study, we discovered a correlation between dimethyl sulfide levels and PDAC metastasis status. This finding suggested that dimethyl sulfide could be used as a biomarker for PDAC metastasis. However, more data on symptom-free survival and response to treatment after follow-up is required to confirm these findings. Based on the findings of the HCC study, one of our hypotheses is that the higher level of dimethyl sulfide in the metastasis group is due to liver lesions, as liver metastasis was the most common type of metastasis. Another possibility is that cancer cells' protein methylation has changed, or that the gut microbiota has changed as a result of duodenal invasion. While hypotheses can be offered, the reasons why PDAC patients demonstrated a higher level of dimethyl sulfide remain unanswered and necessitates further investigation.

Acetone dimer, at a cut-off value of 4.97, is the only VOC that demonstrated a better PDAC discriminating performance when compared to CA19-9, a well-known tumor marker for PDAC. Acetone dimer with AUROC of 0.910 in our study was consistent with previous reports of exhaled breath acetone shown to differentiate PDAC from controls with AUROCs of 0.757-0.901^[14,36]. We also showed that acetone dimer, when combined with CA19-9 at a cutoff value of 0.29, can improve the AUROC to 0.936. This intriguing finding highlighted not only the possibility of using acetone dimer as a sole biomarker, but also the importance of combining acetone dimer with the previous CA19-9 to improve diagnostic accuracy. External validation of our findings, however, is required before applying them to real-world clinical practice.

The strength of this study compared to previous research is the method used for alveolar air collection and the use of FAIMS in combination with the XGBoost algorithm VOC library for VOC profile analysis. Previous VOC studies employed a variety of methods for collecting breath samples, such as aluminum bags or the Bio-VOC sample device^[14,15], which potentially introduced ambient air into the samples collection more than alveolar air. We selected the ReCIVATM breath sample system with software to monitor the real-time progression of the breath sample collection with a pure oxygen gas supply to avoid contamination with confounding gas. We also collected the gas in thermal desorption tubes, which are more stable than aluminum bags.

A second difference compared to previous studies was the use of FAIMS instead of previously used complex analytic techniques for VOC extraction such as GC-MS, ion-molecule reaction mass spectrometry, or electrical nose^[13-15]. FAIMS, can distinguish VOCs by using specific mobilities in the electrical field of the substances^[37]. The advantages of this method include better VOC discrimination performance at low concentrations, which we believe is more appropriate in exhale breath samples, and a shorter runtime (approximately 10 min) compared to GC-MS (up to 45 min)^[19,37]. As a result, it is regarded as a promising method in real clinical practice. FAIMS does have limitations, specifically the difficultly in reproducing between different instruments, resulting in minor variations in waveform between units^[37]. To ensure the accuracy of the VOCs extracted, we created our own VOC library by standardizing gas processing and comparing VOC chromatograms on the same machine. We also used continuous pure oxygen gas flow during breath sample collection to prevent environmental VOC contamination.

Aside from the VOC collection and analysis method, another study strength was the homogeneity of non-PDAC participants. Most of the previous data used a combination of patients with benign pancreatic diseases including chronic pancreatitis, pancreatic cysts, pancreatic pseudocysts, IPMN, and healthy volunteers as the controls^[14,36,38]. The heterogeneity in benign pancreatic pathophysiology within the control groups may have affected the ability to detect differences in VOCs between groups in these studies. To prevent these confounding factors in our study, we inclusively enrolled only IPMN

participants who had potential malignant transformation to serve as a second comparison group and enrolled standard healthy volunteers for the control group.

Our study did have several limitations. First, our recruited PDAC population was primarily comprised of patients with locally advanced and advanced stage PDAC. This resulted from the limitation of current PDAC diagnosis, since most PDAC patients are diagnosed in the advanced stage. The VOCs collected in our study may not be representative of VOCs detected in the early-stages of PDAC. The study could not show if early stage PDAC VOC levels were significantly different than at later stages of the disease and might not be effective as a screening method. A larger longitudinal study with more early-stage PDAC patients is needed. Secondly, because of the small number of malignant IPMN in our study, a conclusion that the VOCs can distinguish malignant from benign IPMN is problematic. Further research with larger numbers of malignant IPMN is still needed. Third, we cannot conclude which PDAC was developed from IPMN in our study. However, no PDAC patients had any cystic component or other pancreatic cystic lesion that represented an area of current or previous IPMN. Fourth, this is a "proof-of-concept" study. We intended to identify PDAC VOCs from potentially malignant pancreatic lesions like IPMN. Other benign pancreatic tumors with no malignant potential such as serous cystic neoplasm, as well as other periampullary cancers such as distal cholangiocarcinoma, ampullary cancer, or duodenal cancer, were not included in the comparison group. The specific VOCs that distinguish PDAC from these populations require further investigation.

Multiple exogenous confounding factors such as environmental factors, occupational exposure, and underlying diseases could have possibly altered the VOCs profile in real-world practice. In addition, the VOC analysis in this study was derived from a single cohort, and the model performance was evaluated in a cohort of participants from the same center. The generalizability of our study results should be interpreted with caution. Lastly, the external validation of these findings using an independent cohort is required before applying this approach to clinical practice.

CONCLUSION

The use of alveolar VOC composition offers potential as a biomarker to help distinguish PDAC from IPMN and normal pancreas. The two VOCs that showed promising results were dimethyl sulfide and acetone dimer. The combination of acetone dimer with CA19-9 showed the highest AUROC to distinguish PDAC from the non-PDAC population. More prospective diagnostic research is required to validate our findings.

ARTICLE HIGHLIGHTS

Research background

In various gastrointestinal malignancies, including pancreatic adenocarcinoma, volatile organic compounds (VOC) have demonstrated a good result in distinguishing patients with cancer and those without cancer.

Research motivation

No previous research has been conducted to capture exhaled breath VOCs using thermal desorption-gas chromatography/field-asymmetric ion mobility spectrometry (TD-GC/FAIMS) methods in order to distinguish pancreatic ductal adenocarcinoma (PDAC). The control population in the previous study was heterogeneous and reduced the validity of the results.

Research objectives

To identify exhale breath VOCs that can distinguish PDAC patients from those with Intraductal papillary mucinous neoplasm (IPMN) and those with no pancreatic lesions using TD-GC/FAIMS.

Research methods

Exhaled breath collection using the ReCIVA® device and VOCs identification using TD-GC/FAIMS method.

Research results

Dimethyl sulfide and acetone dimer were higher in PDAC patients when compared to IPMN and healthy participants. Dimethyl sulfide levels have been linked to PDAC metastasis status. Combining acetone dimer with the previous biomarker, CA19-9, has been shown to improve PDAC diagnostic accuracy.

Research conclusions

Dimethyl sulfide and acetone dimer were the two VOCs that can potentially distinguish PDAC from IPMN and healthy participants.

Research perspectives

Further validation with a larger cohort using a longitudinal approach is needed to confirm this study findings.

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