Urinary volatile organic compound analysis for the diagnosis of cancer: a systematic literature review and quality assessment

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Search Strategy

Database: Embase Classic+Embase <1947 to 2019 December 13> Search Strategy: Volatile Organic Compounds/ or volatile organic compound*.mp. (20445) biomarkers.mp. or BIOMARKERS/ (385783) 3 Metabolomics/ or metabo*omics.mp. (35735) Magnetic Resonance Spectroscopy/ or magnetic resonance spectrometry.mp. (100046) 5 mass spectrometry.mp. or Mass Spectrometry/ (463935) metabolic profiling.mp. (4883) 7 metabotyping.mp. (71) urine.mp. or URINE/ (564820) 1 or 2 (405092) 3 or 4 or 5 or 6 or 7 (569457) 8 and 9 and 10 (5080) 12 prostate.mp. or PROSTATE/ (347922) prosta*.mp. (583298) gastric.mp. or GASTRIC/ (369992) gastro*.mp. (894775) esophageal.mp. or ESOPHAGEAL/ (149069) 17 esophago*.mp. (53403) colorectal.mp. or COLORECTAL/ (261334) rectal*.mp. (153592) bowel.mp. or BOWER/ (249315) 21 intestinal.mp. or INTESTINAL/ (386822) intestin*.mp. (799633) pancreas.mp. or PANCREAS/ (394885) pancrea*.mp. (520745) carcinoma.mp. or CARCINOMA/ (1218118) ca*.mp. (24378921) 27 12 or 13 (583298) 14 or 15 (1143350) 16 or 17 (180363) 18 or 19 or 20 or 21 or 22 (1231013) 23 or 24 (520745) 31 25 or 26 (24378921) 32 27 or 28 or 29 or 30 or 31 (3084057) 32 and 33 (2400354) 35 11 and 34 (689)

limit 35 to english language (676)

37 limit 36 to human (571)

Database: Ovid MEDLINE(R) ALL <1946 to December 13, 2019> Search Strategy:

- 1 Volatile Organic Compounds/ or volatile organic compound*.mp. (14685)
- 2 biomarkers.mp. or BIOMARKERS/ (491461)
- 3 Metabolomics/ or metabo*omics.mp. (25507)
- 4 Magnetic Resonance Spectroscopy/ or magnetic resonance spectrometry.mp. (147500)
- 5 mass spectrometry.mp. or Mass Spectrometry/ (309505)
- 6 metabolic profiling.mp. (3623)
- 7 metabotyping.mp. (55)
- 8 urine.mp. or URINE/ (357244)
- 9 1 or 2 (505235)
- 10 3 or 4 or 5 or 6 or 7 (454801)
- 11 8 and 9 and 10 (4492)
- 12 prostate.mp. or PROSTATE/ (189208)
- 13 prosta*.mp. (364594)
- 14 gastric.mp. or GASTRIC/ (287312)
- 15 gastro*.mp. (522330)
- 16 esophageal.mp. or ESOPHAGEAL/ (137920)
- 17 esophago*.mp. (37648)
- 18 colorectal.mp. or COLORECTAL/ (154456)
- 19 rectal*.mp. (114296)
- 20 bowel.mp. or BOWER/ (151056)
- 21 intestinal.mp. or INTESTINAL/ (403809)
- 22 intestin*.mp. (499655)
- 23 pancreas.mp. or PANCREAS/ (137239)
- 24 pancrea*.mp. (325641)
- 25 carcinoma.mp. or CARCINOMA/ (805969)
- 26 ca*.mp. (18137476)
- 27 12 or 13 (364594)
- 28 14 or 15 (714689)
- 29 16 or 17 (150433)
- 30 18 or 19 or 20 or 21 or 22 (799204)
- 31 23 or 24 (325641)
- 32 25 or 26 (18137476)
- 33 27 or 28 or 29 or 30 or 31 (2026101)
- 34 32 and 33 (1468790)
- 35 11 and 34 (449)
- 36 limit 35 to english language (435)
- 37 limit 36 to human (315)

PRISMA Checklist

Section/topic	#	Checklist item	Reported on page #
TITLE			
Title	1	Identify the report as a systematic review, meta-analysis, or both.	1
ABSTRACT			
Structured summary	2	Provide a structured summary including, as applicable: background; objectives; data sources; study eligibility criteria, participants, and interventions; study appraisal and synthesis methods; results; limitations; conclusions and implications of key findings; systematic review registration number.	1; no registration number
INTRODUCTION			
Rationale	3	Describe the rationale for the review in the context of what is already known.	1-2
Objectives	4	Provide an explicit statement of questions being addressed with reference to participants, interventions, comparisons, outcomes, and study design (PICOS).	2
METHODS	-		
Protocol and registration	5	Indicate if a review protocol exists, if and where it can be accessed (e.g., Web address), and, if available, provide registration information including registration number.	13
Eligibility criteria	6	Specify study characteristics (e.g., PICOS, length of follow-up) and report characteristics (e.g., years considered, language, publication status) used as criteria for eligibility, giving rationale.	13-14
Information sources	7	Describe all information sources (e.g., databases with dates of coverage, contact with study authors to identify additional studies) in the search and date last searched.	13
Search	8	Present full electronic search strategy for at least one database, including any limits used, such that it could be repeated.	13-14, Supplementary
Study selection	9	State the process for selecting studies (i.e., screening, eligibility, included in systematic review, and, if applicable, included in the meta-analysis).	13-14
Data collection process	10	Describe method of data extraction from reports (e.g., piloted forms, independently, in duplicate) and any processes for obtaining and confirming data from investigators.	13-14

11	List and define all variables for which data were sought (e.g., PICOS, funding sources) and any assumptions and simplifications made.	14
12	Describe methods used for assessing risk of bias of individual studies (including specification of whether this was done at the study or outcome level), and how this information is to be used in any data synthesis.	13-14
13	State the principal summary measures (e.g., risk ratio, difference in means).	14
14	Describe the methods of handling data and combining results of studies, if done, including measures of consistency (e.g., I2) for each meta-analysis.	N/A
15	Specify any assessment of risk of bias that may affect the cumulative evidence (e.g., publication bias, selective reporting within studies).	13-14
16	Describe methods of additional analyses (e.g., sensitivity or subgroup analyses, meta-regression), if done, indicating which were pre-specified.	N/A
17	Give numbers of studies screened, assessed for eligibility, and included in the review, with reasons for exclusions at each stage, ideally with a flow diagram.	2, Figure 1
18	For each study, present characteristics for which data were extracted (e.g., study size, PICOS, follow-up period) and provide the citations.	2, Table 1
19	Present data on risk of bias of each study and, if available, any outcome level assessment (see item 12).	6, Table 2, Table S1
20	For all outcomes considered (benefits or harms), present, for each study: (a) simple summary data for each intervention group (b) effect estimates and confidence intervals, ideally with a forest plot.	8-10, Figure 2, 3 and 4, Table 3, Table S2
21	Present results of each meta-analysis done, including confidence intervals and measures of consistency.	N/A
22	Present results of any assessment of risk of bias across studies (see Item 15).	6, Table 2, Table S1
23	Give results of additional analyses, if done (e.g., sensitivity or subgroup analyses, meta-regression [see Item 16]).	N/A
24	Summarize the main findings including the strength of evidence for each main outcome; consider their relevance to key groups (e.g., healthcare providers, users, and policy makers).	11-13, Table 4
25	Discuss limitations at study and outcome level (e.g., risk of bias), and at review-level (e.g., incomplete retrieval of identified research, reporting bias).	13
	12 13 14 15 16 17 18 19 20 21 22 23	12 Describe methods used for assessing risk of bias of individual studies (including specification of whether this was done at the study or outcome level), and how this information is to be used in any data synthesis. 13 State the principal summary measures (e.g., risk ratio, difference in means). 14 Describe the methods of handling data and combining results of studies, if done, including measures of consistency (e.g., I2) for each meta-analysis. 15 Specify any assessment of risk of bias that may affect the cumulative evidence (e.g., publication bias, selective reporting within studies). 16 Describe methods of additional analyses (e.g., sensitivity or subgroup analyses, meta-regression), if done, indicating which were pre-specified. 17 Give numbers of studies screened, assessed for eligibility, and included in the review, with reasons for exclusions at each stage, ideally with a flow diagram. 18 For each study, present characteristics for which data were extracted (e.g., study size, PICOS, follow-up period) and provide the citations. 19 Present data on risk of bias of each study and, if available, any outcome level assessment (see item 12). 20 For all outcomes considered (benefits or harms), present, for each study: (a) simple summary data for each intervention group (b) effect estimates and confidence intervals, ideally with a forest plot. 21 Present results of each meta-analysis done, including confidence intervals and measures of consistency. 22 Present results of any assessment of risk of bias across studies (see Item 15). 23 Give results of additional analyses, if done (e.g., sensitivity or subgroup analyses, meta-regression [see Item 16]). 24 Summarize the main findings including the strength of evidence for each main outcome; consider their relevance to key groups (e.g., healthcare providers, users, and policy makers).

Conclusions	Provide a general interpretation of the results in the context of other evidence, and implications for future research.								
FUNDING									
Funding	27	Describe sources of funding for the systematic review and other support (e.g., supply of data); role of funders for the systematic review.	15						

Supplementary Materials

1/(Total number of VOCs identified in each study)

Total number of VOCs identified in each cancer type

x

Total number of studies that identified this VOC

Figure S1. Equation for weighted means of each identified volatile organic compound (VOC).

 Table S1. Quality assessment of metabolic metadata based on CAWG-MSI guidelines.

	Author	
	Biological replicates	Sample preparation
	Biofluid harvesting	
	Storage condition	
	Extraction method	
	Extract concentration, dilution, resolubilisation process	
	Extract enrichment	
	Extract cleanup	
	Extract storage and/or relocation	
	Instrument description	Experimental analysis
	Auto-injector	
	Separation column	
	Technique specific sample preparation	
	Separation parameters	
	Instrument description	Instrumental
	Sample introduction & delivery	performance
	Ionisation source	
	Mass analyser description and acquisition mode	
	Data acquisition parameters	
	m/z calibration standard	Method validation
	Accuracy of m/z calibration	
	Mass resolution	
	Ion source optimisation parameters	
	Chromatographic resolution	
	QC samples	
	Internal Standards used	
	Accuracy of Internal standard	
	Precision of internal standard	
	Retention time markers	
	Accuracy and precision for replicated analyses	
	Accuracy and precision for validation samples	
	Cycles per column/injector/septum/blank	
Description, quantifier of IS metabolite	Relative quantification	Metabolite identification
Method for assessing instrument		
Replicate analyses		
Calibration curves for each metabolite	Absolute quantification	
Range of standards used		
Quantification of method accuracy		
Quantification of method precision		
ПОО		
ПОО		
Recovery/ stability		
Data file format/ conversion methods	Post-acquisition data pre-processing	Data pre-processing
Data-preprocessing methods		
Level		Metabolite identification

Lima A.R. 2019	Υ	Y	Υ	Υ	N	Υ	N	Υ	Y	Υ	Y	Υ	Y	Υ	Y	Υ	Υ	Υ	N	N A	N	Υ	N	Υ	Υ	N	N	Z	Ν	Υ	N	Υ	Y	N								N	Y	2
Struck-Lewicka W. 2014	Υ	Y	Y	Y	Y	N	N	N	Y	N	Y	Y	Y	Y	N	Y	Υ	Υ	N	N A	N	Y	N	Y	Y	N	N	Υ	N	N	N	Y	Υ	N								N	Υ	2
Gao Q. 2019	Υ	Y	Y	Υ	N	Y	N	Υ	Y	N	Y	Y	Y	Y	N	Υ	Υ	Υ	N	N A	N	N	N	N	Y	N	N	N	N	Υ	N	Y	N	N								N	Y	2
Jimenez-Pacheco A. 2018	Υ	Y	Υ	Υ	N	Y	N	Υ	Υ	Υ	Y	N	Y	Υ	N	Υ	Υ	Υ	N	N A	N	N	N	N	Ν	N A	N A	N	N	N	N	N	N	N								N	N	2
Khalid T. 2015	Υ	Y	Υ	Υ	N	Υ	N	Υ	Υ	N	Υ	N	Y	Υ	N	Υ	Υ	Υ	N	N A	N	N	N	N	N	N A	N A	N	N	N	N	N	Υ	Υ								N	N	2
Spanel P. 1999	Υ	Y	Υ	N	N	N	N	N	Υ	N	N	N	N	Υ	Y	Υ	Υ	Υ	N	N A	N	N	N	N	N	N A	N A	N	N	N	N				N	N	N	N	N	N	N	N	N	1
Chen Y. 2016	Υ	Y	Υ	Υ	Υ	N	N	N	Υ	N	Υ	Υ	Y	Υ	N	Υ	Υ	Υ	N	N A	N	N	N	N	Υ	N	N	N	N	Υ	N				Υ	Υ	Υ	Υ	Υ	Υ	Υ	N	N	1
Navaneethan U. 2015	Υ	Y	Υ	Υ	N	N	N	N	N	N	N	N	N	Υ	Y	Υ	Υ	Υ	N	N A	N	N	N	N	N	N A	N A	N	Υ	N	N				N	N	N	N	N	N	Υ	N	Υ	2
Panebianco C 2017	Υ	Y	Υ	Υ	N	Υ	N	N	Υ	Υ	Υ	N	Y	Υ	Υ	N	N	N	N	N A	N	N	N	N	N	N A	N A	N	N	N	N	N	Υ	N								N	Y	2
Arasaradnam RP 2014	Υ	Y	Υ	Υ	N	N	N	N	Υ	Υ	Υ	N	Y	Υ	N	Υ	Υ	Υ	N	N A	N	Υ	N	N	N	N A	N A	N	N	N	N	N	Υ	N								N	Y	N A
Huang J. 2013	Υ	Y	Υ	N	N	N	N	N	N	N	N	N	N	Υ	Y	Υ	Υ	Υ	N	N A	N	N	N	N	N	N A	N A	N	N	N	N				N	N	N	Υ	N	N	N	N	N	2
Rozhentsov A.A. 2014	Y	N	Y	Υ	N	Υ	N	Y	N	N	N	N	Y	N	N	N	N	N	N	N A	N	N	N	N	N	N A	N A	N	N	N	N				N	N	N	N	N	N	N	N	N	N A
Silva C.L. 2011	Υ	Y	Y	Υ	N	Υ	N	Υ	Y	N	Y	N	Y	Y	Y	Y	Υ	Υ	N	N A	N	Υ	N	N	N	N A	N A	N	N	N	N	N	N	N								N	N	2

Y: yes, N: no, NA: not applicable. QC: quality control; IS: internal standard; LLOQ: lower limits of quantification; LLOD: lower limits of detection

Table S2. List of biomarkers noted to be significantly increased and/or decreased in cancers and associated metabolic pathway.

Compound name	Chemical Classes	Metabolic pathway	Prostate Cancer	Gastrointestinal Cancer
Ethanol	Alcohol	Glycolysis / Gluconeogenesis		
Acetaldehyde	Aldehyde	Pyruvate metabolism; Glycolysis / Gluconeogenesis; Glycerophospholipid metabolism		
Phenylacetic acid	Aromatic compound	Phenylalanine metabolism		↑
Butyric acid	Fatty acid	Butanoate metabolism	\	
Methylglyoxal	Ketoaldehyde	Glycine, serine and threonine metabolism; Pyruvate metabolism	↓	
Acetic acid		Glyoxylate and dicarboxylate metabolism; Pyruvate metabolism; Glycolysis / Gluconeogenesis	\	↑
Hydrogen sulfide	Organic acid	Sulfur metabolism		↑
Propionic acid		Propanoate metabolism	\downarrow	