

Delivering quality data in biomarker discovery

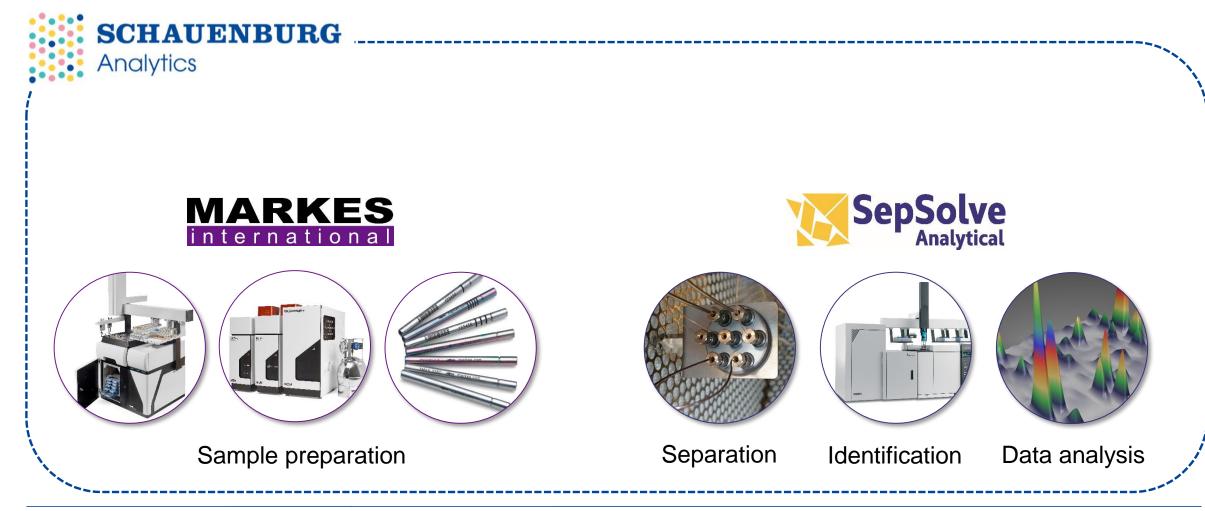
Laura McGregor Product Marketing Manager





Who are SepSolve Analytical?

Experts in analytical chemistry







Biomarker analysis & the challenges involved

Breath analysis workflow using thermal desorption (TD)-GC-MS

Typical errors and ways to overcome them

Beyond breath – investigating other sources of biomarkers

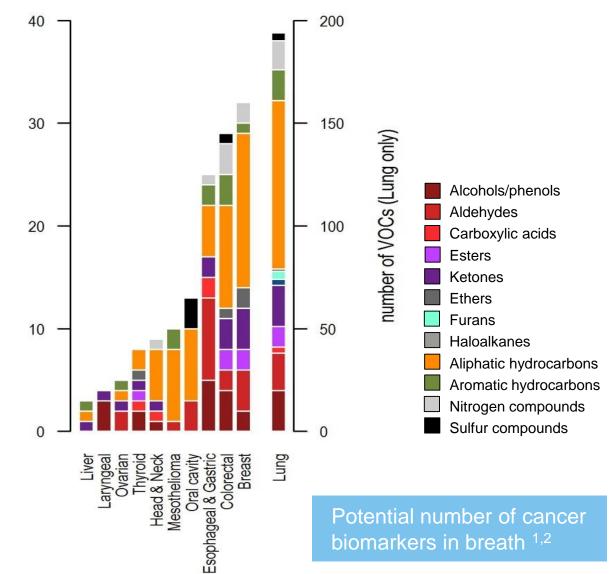


The diversity of the breath volatilome

- >800 VOCs have previously been reported from breath^{1, 2}
 - Many different types of compounds
 - Need to quantify at trace levels (sub-ppb)

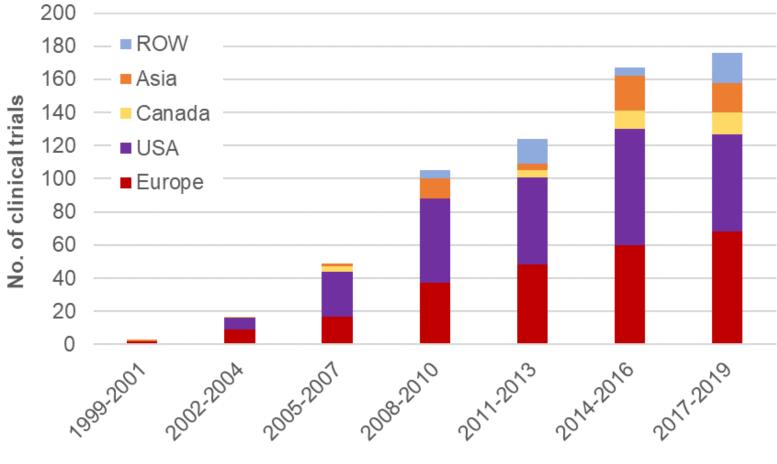
number of VOCs (all except Lung)

- Are these all 'real' biomarkers?
- The problem of lack of consensus





Breath VOC analysis in clinical research



Clinical trials involving breath analysis (based on start date)

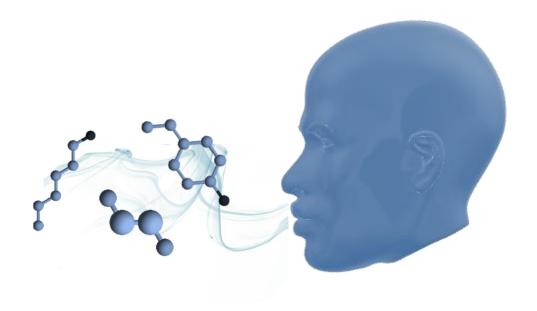
Clinical trial start date



Data source: clinicaltrials.gov (using search terms "breath" and "biomarker")

Analytical challenges in breath analysis

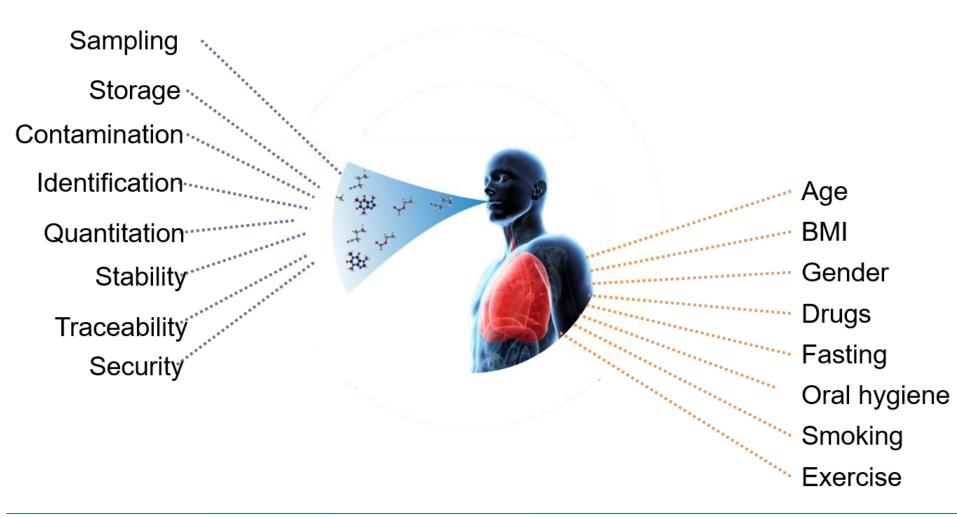
- Collection vessels:
 - Bags, tubes, Bio-VOC, ReCIVA
- Sampling alveolar air:
 - Low volumes, high humidity...
- High sample complexity:
 - Low concentration of volatiles (ppbv to sub-pptv)
 - Requires highly sensitive detection
- System reproducibility:
 - Sampling device
 - TD-GC(×GC)-MS
- Minimal chemical background of all devices and apparatus used





Breath VOCs: sources of variability

Application of breath analysis in clinical practice is limited





Why thermal desorption?

Breath VOC sampling, transport and concentration

Problem

- Collect breath from multiple clinical sites, transport and store samples
- 2. Breath is saturated with water vapour
- 3. Untargeted analysis, biomarkers present at trace levels
- 4. Clinical samples are invaluable
- 5. Empty tubes and samples transported between lab and clinic. Lengthy clinical studies.

Solution

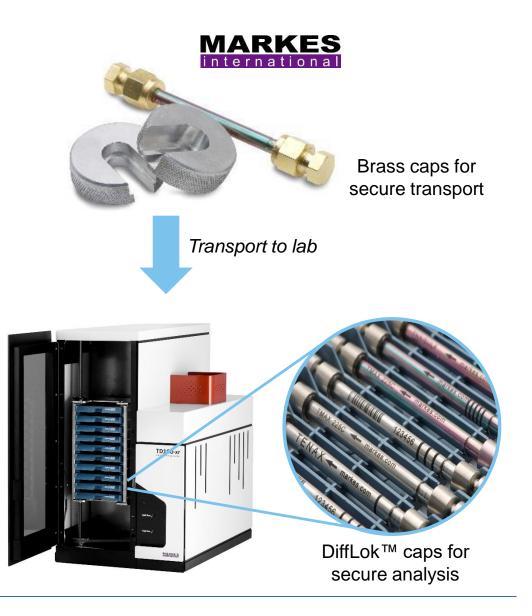
- TD tubes are compact, they can be sealed and conveniently transported
- Dry-purging of the tubes and focusing trap
- Cryogen-free trap focusing
- Re-collection
- Internal standard addition to tubes or focusing trap



Sample security

Breath analysis using thermal desorption (TD)

- Sorbent tubes are:
 - ✓ Sealed at point of use
 - ✓ Customised for breath analysis
 - Compatible with RFID tags for sample tracking and integrity
 - Compatible with fixed and variable-volume samplers





Standardisation

Type of QC samples in 'breathomics' studies and how to prepare them

Sample type	Description	Preparation
Extraction/process blank	What is not of biological origin/carryover	Run a blank of the whole system under the same conditions (e.g. empty tube)
System suitability sample	System check BEFORE precious biological samples	Load a clean TD tube with a standard mixture
Internal standards	Given sample complexity this is not applicable to all metabolites	Add ISTD to every sample. ISTD (or ISTDs) is similar to analytes but not overlapping
Within-laboratory inter-study sample/standard reference materials (SRMs)	Long-term reference material for use across research community	Standards are added to a clean tube under standardised conditions, typically by a third party/authority (e.g. National Physical Laboratory).

Abridged from: D. Broadhurst, R. Goodacre, S. N. Reinke, et al. 2018. "Guidelines and Considerations for the Use of System Suitability and Quality Control Samples in Mass Spectrometry Assays Applied in Untargeted Clinical Metabolomic Studies." *Metabolomics* 14 (6).



Standardisation

Type of standards in 'breathomics' studies and how to prepare them

- Automated standard addition to blank tubes prior to sampling
 - Intra-study 'pooled' sample
 - System check standard

- Automated standard addition to **sampled tubes** prior to analysis
 - Internal standard

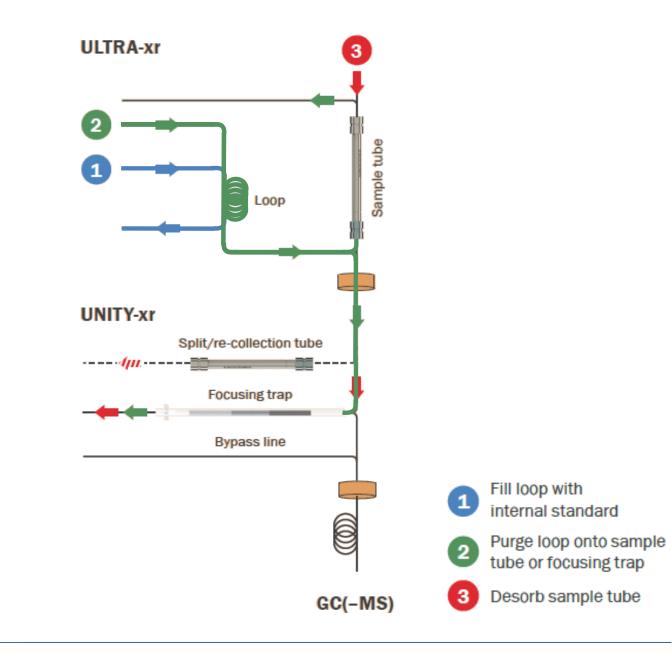
- Automated standard addition to focusing trap
 - System suitability sample



Automated standard addition

Gas standards

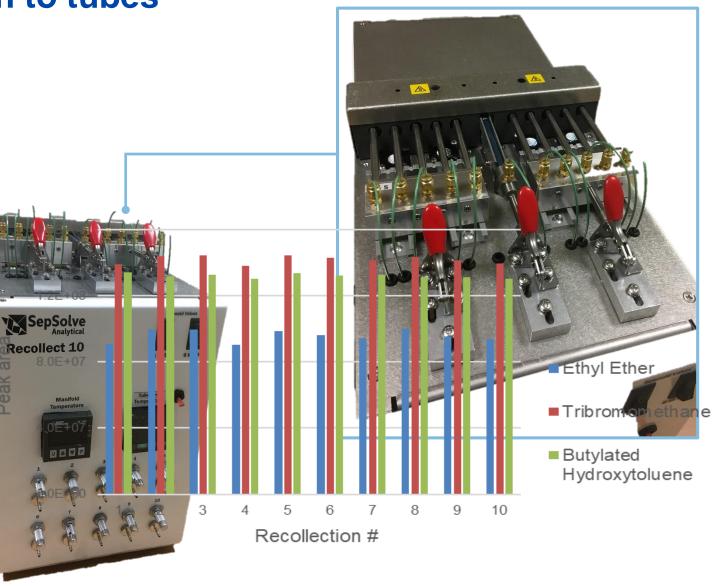
 Automated standard addition to tubes or focusing trap





Automated standard addition to tubes

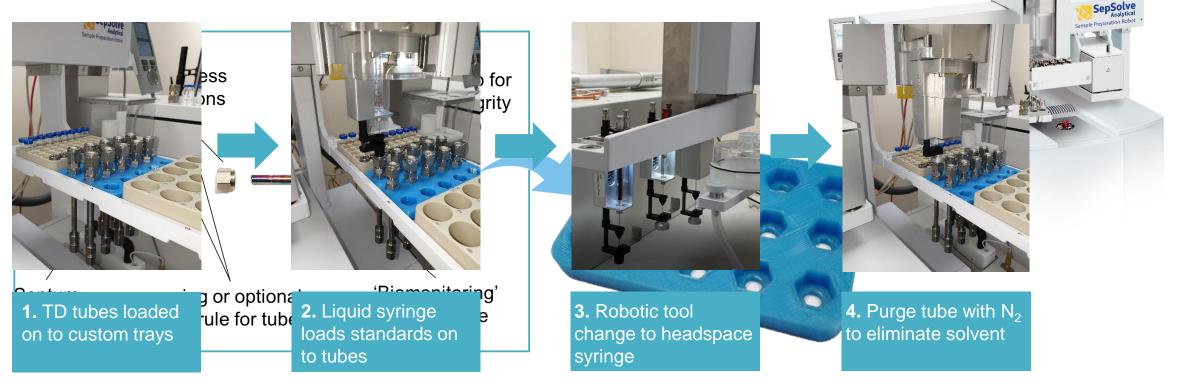
- Split high-loading samples between multiple tubes
- For replicate analysis and statistical comparison
- Stand-alone, and compatible with industry-standard TD tubes





Automated standard addition to tubes

Liquid standards



- Bespoke tray design to accommodate 15 thermal desorption tubes
- Workflow for fully automated spiking of sorbent tubes with liquid standards using a bespoke SPR

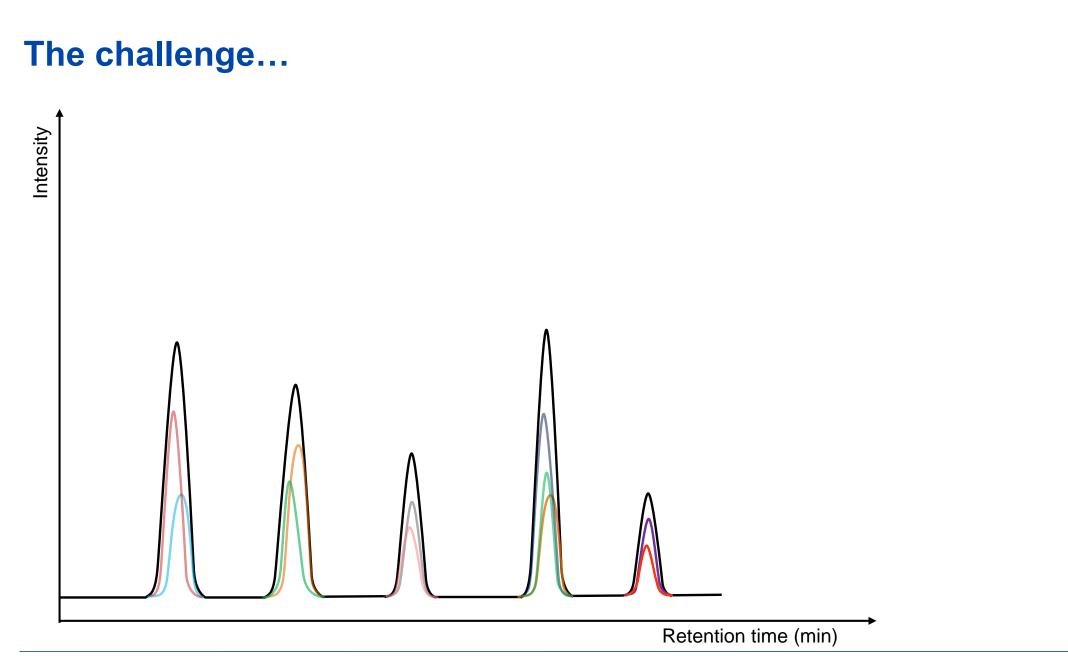


Increasing confidence in biomarker identification

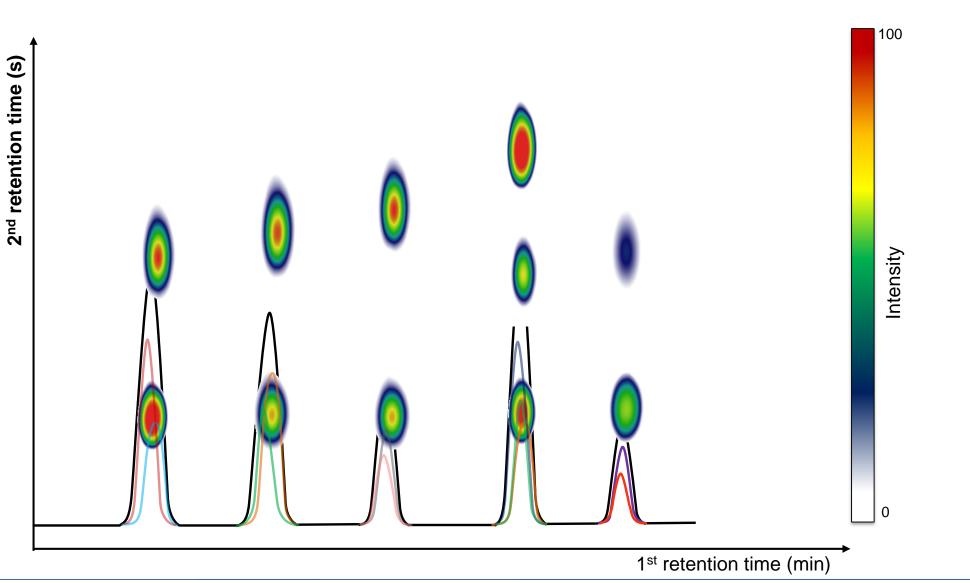
• Are you **<u>sure</u>** it's the compound you say it is?







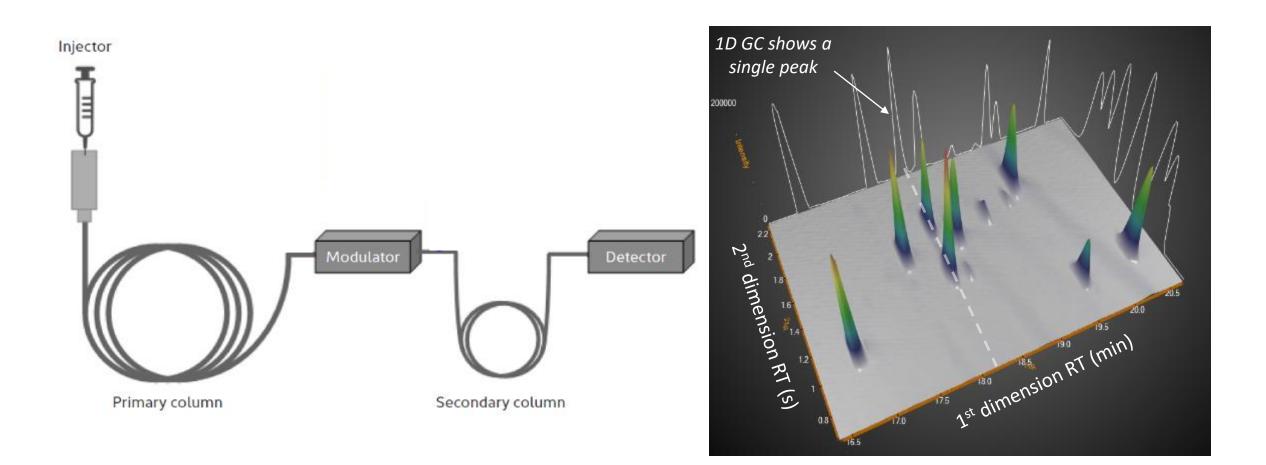






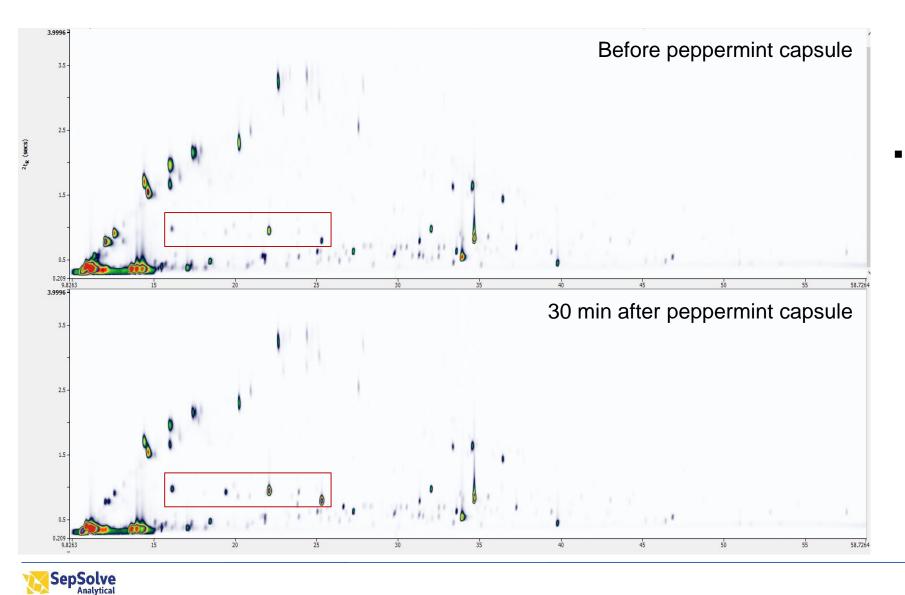
SepSolve Analytical

What is GC×GC?





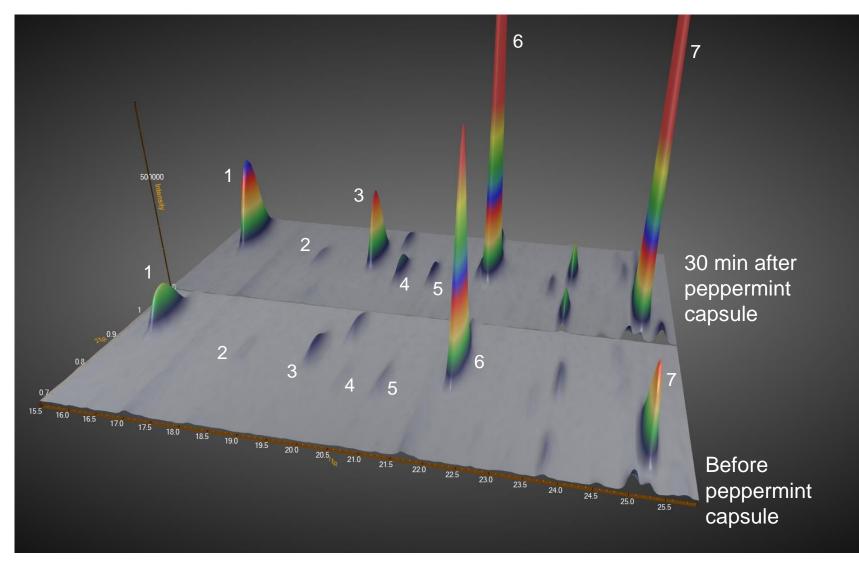
Improved separation by GC×GC





Breath VOCs were measured before & after consumption of a peppermint capsule

Improved separation by GC×GC



- α-Pinene
- 2 Camphene
- 3 β-Pinene
- 4 δ-3-Carene
- 5 α -Phellandrene
- 6 β-Mycene
- 7 Limonene
- 8 Eucalyptol



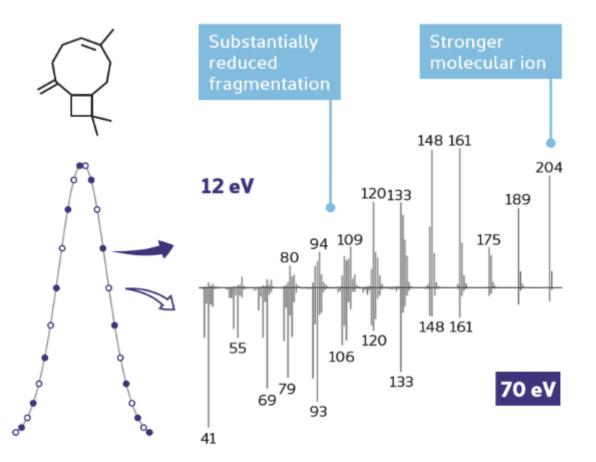
Increased confidence with Tandem Ionisation®

Acquire hard and soft EI simultaneously

Additional confirmation of analyte identity

No added analysis time

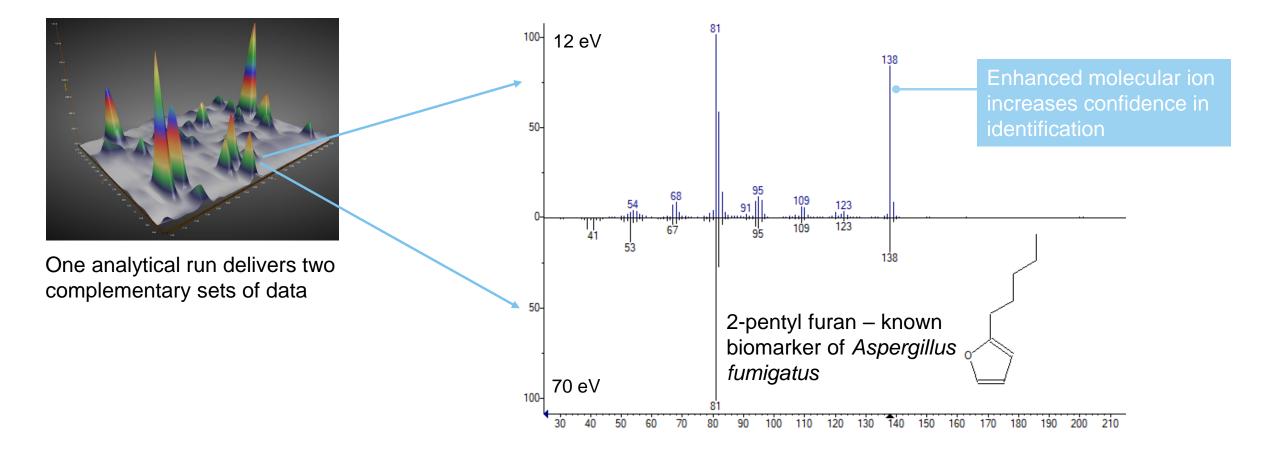
Patented technology exclusive to BenchTOF





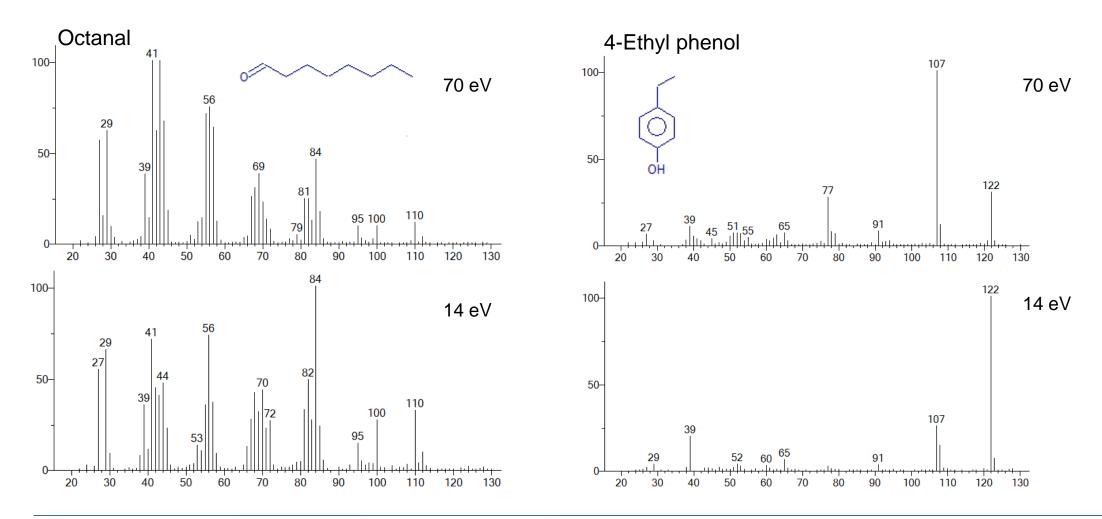
Delivering increased confidence in identification

Tandem Ionisation®





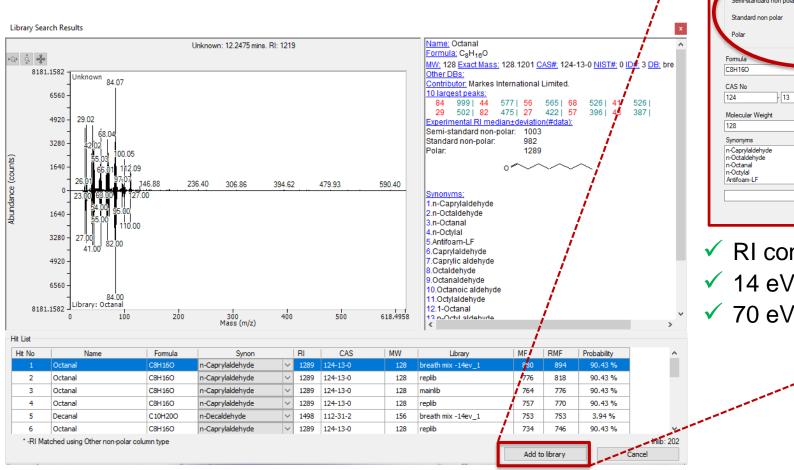
Other examples of TI for known breath biomarkers





Triple confirmation of identity

70 eV and 14 eV plus retention indices (RI)



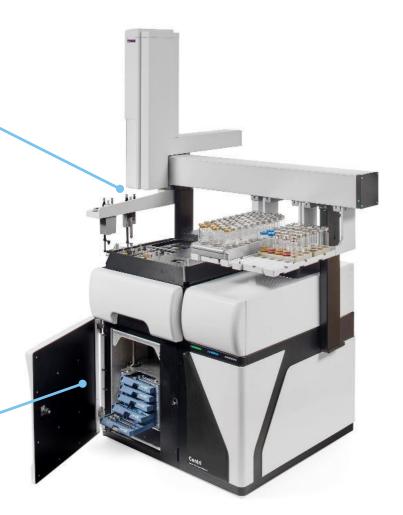
Compound Name	Compound ID		Target Libraries
Octanal	0	/	breath mix -14ev
Name in Library	Source Library Na	ame	breath mix -70ev
Octanal			demo
			If the second
Comment	Number of Mass-	Abundance Pairs	ri marker library
	217		test
Experimental RI median	Mass-Abundance		
Semi-standard non polar	Mass	Intensity ^	
Standard non polar 982	22	6.19884360456917	
	26.007	37.3339444366098	
Polar 💂 1289	27.007	607.345790438584	
	29.02 29.985	638.05824284304 103.12621633056	
Formula	30.873	14.9335777746439	
C8H16O	32,905	5.63531236779016	
	35.323	11.9750387815541	
CAS No	36.444	8.31208574249048	
124 · 13 · 0 <u>View CAS No. Calculation</u>	37	24.2318431814977	
Molecular Weight	39.031	360.800874347765	
128 Calculate from formula	40.001	76.4993653927514	
	41.016 42.023	631.295868001692 382.355944154562	
Synonyms n-Caprylaldehyde	43.022	330.651953180087	
n-Octaldehyde	43.998	485.341277675927	
n-Octanal n-Octylal	45.074	195.967987589903	
Antifoam-LF	46	16.7650542941757	
Add	47	19.3009448596813	
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RI confirmation			and the second sec
14 eV confirmation	n	and the second sec	
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70 eV confirmation	n 🦯		
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Beyond breath...

A single platform for the analysis of VOCs in breath and biological fluids

- Liquids, solids and semi-solids:
 - Urine, exhaled breath condensate, saliva, sputum as alternatives to breath
 - Skin, sweat: monitoring of physical or psychological stress, hydration state
 - Stool: GI infections, microbiome studies



• **Gases:** Analysis of breath biomarkers



What does Centri[®] do?



Headspace-trap (& classical headspace)



Tube-based thermal desorption





SPME-trap (& classical SPME)



HiSorb high-capacity sorptive extraction



Innovation at the heart of Centri

Why is a trap important?



Secure sample recollection



Improved chromatography



Selectivity



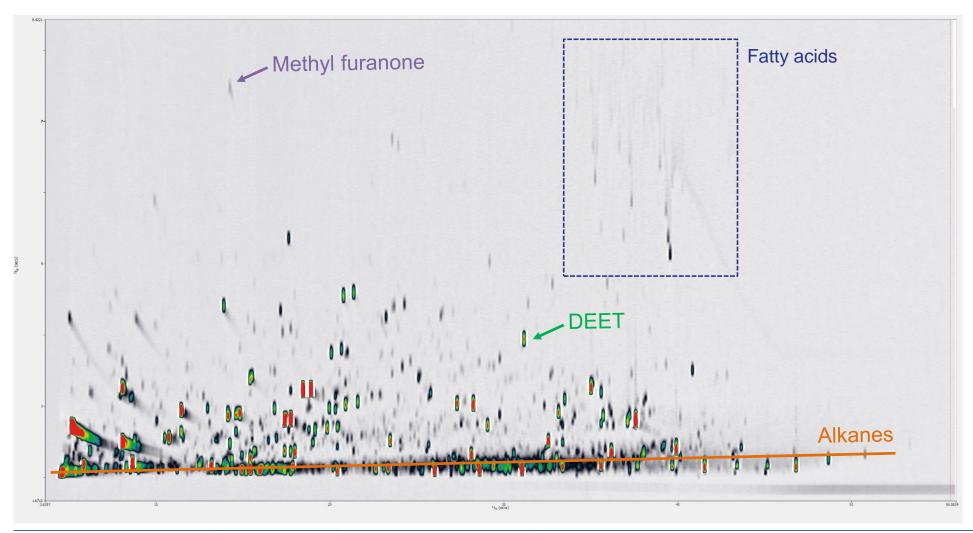
Sample enrichment





Beyond breath...

Thin Film-SPME-GC×GC-TOF MS of skin VOCs





Beyond breath...

HiSorb sorptive extraction

 Analysis of a range of biological samples (e.g. breath, urine, sputum...) using a single analytical platform

 HiSorb probes are ideal for high sensitivity immersive sampling of urine

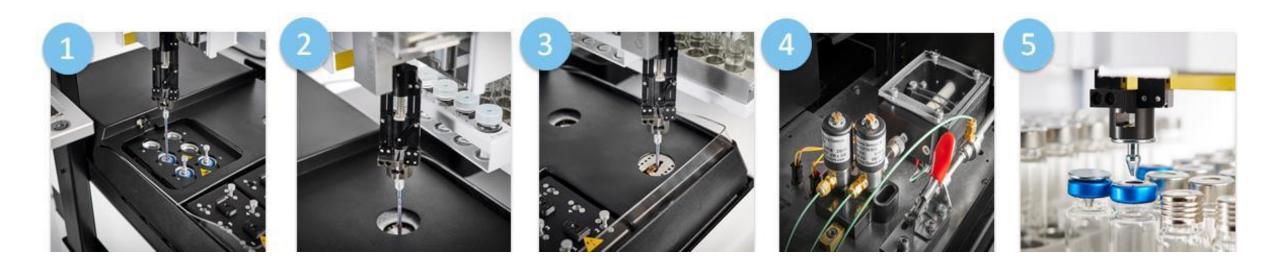
- log $K_{O/W} < 2$ = low recovery by PDMS e.g. Propanoic Acid $K_{O/W}$ = 0.33
- $2 < \log K_{O/W} < 4 \text{ partial extraction by PDMS}$ e.g. Octanal $K_{O/W} = 2.78$
- log K_{OW} > 4 excellent recovery by PDMS e.g. Decane K_{OW} = 5.25





HiSorb high-capacity sorptive extraction

Fully automated workflow

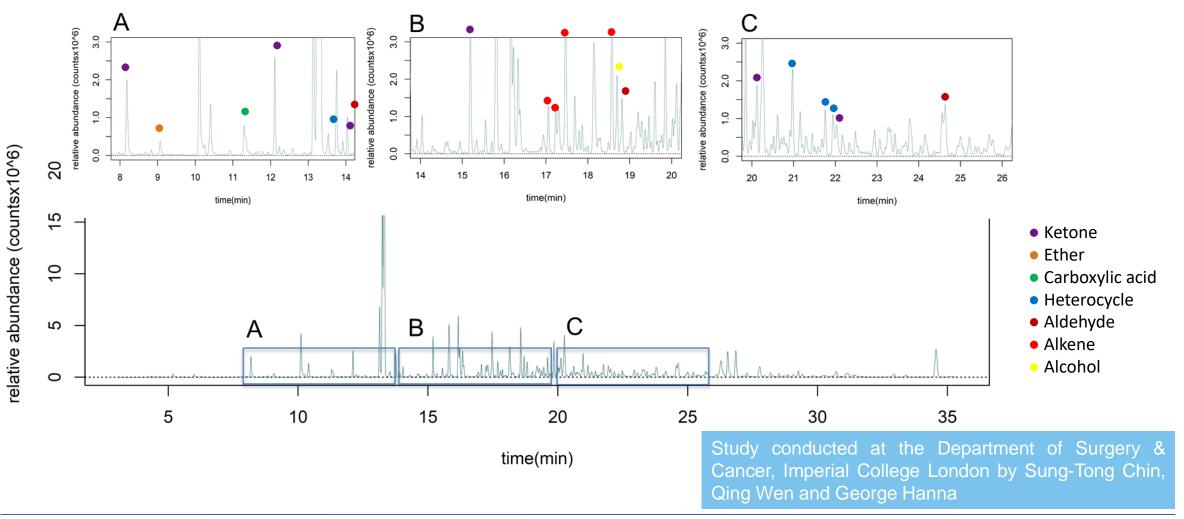


- Suitable for immersive sampling of liquids and headspace sampling of solids/liquids
- Urinary biomarkers have potential as a simple, inexpensive and non-invasive test for early cancer detection



Case study for HiSorb extraction

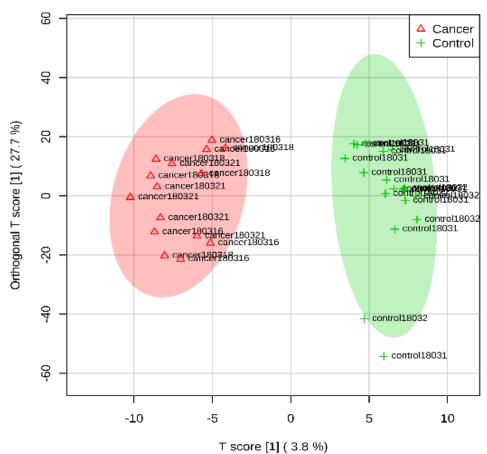
Urinary VOC analysis in oesophago-gastric cancer detection





Case study for HiSorb extraction

- Urinary VOC analysis in oesophago-gastric cancer detection
 - Good linearity R²= 0.991 0.994
 - Limits of quantitation: 30 70 ng/g
 - Urine is sampled from 50 cancer patients and 50 controls
 - 90% of correct discrimination between cancer and control
 - Applicable to large-scale clinical studies: convenient, low cost, rugged, sensitive



Scores Plot

Study conducted at the Department of Surgery & Cancer, Imperial College London by Sung-Tong Chin, Qing Wen and George Hanna



The workflow of breath analysis

TD-GC-MS is the "golden standard" technique



Automated thermal desorbers

Pre-concentration, automation, water management, leak testing...

Biofluids (urine, blood...), tissues, culture media

Automation & concentration

platform

$\mathbf{GC}{\times}\mathbf{GC}{\text{-}}\mathbf{TOF}\ \mathbf{MS}$

Biomarker 'discovery' platform



Summary

- TD-GC(×GC)-MS is the gold-standard technique for breath
- All metabolomics studies must ensure data quality
- When data are not properly QC checked, this leads to wrong data interpretations
- TD can make breath samples more amenable to QC sample preparation and QC checks
- Confident identification is possible with improved separation and/or additional checks (soft ionisation, retention indices...)







Thanks for listening! Any questions?

Contact SepSolve

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