



Nouvelles techniques couplées pour l'analyse d'arômes, parfums – Automatisation de la préparation d'échantillons

Bart Tienpont, Christophe Devos, Frank David and Pat
Sandra



RIC

Research Institute
for Chromatography

What do we expect?

- Volatility range **BROAD**
- Polarity range **WIDE**
- Complex samples **HIGHLY**
- Matrix interferences **OMG**



RIC

Research Institute
for Chromatography

Solutions in reach

- Sample preparation
 - Enrichment?
 - selective or generic?
- Separation: GC or GC-GC or GCxGC ?
- Detection: MS/ODP and... sensors?



RIC

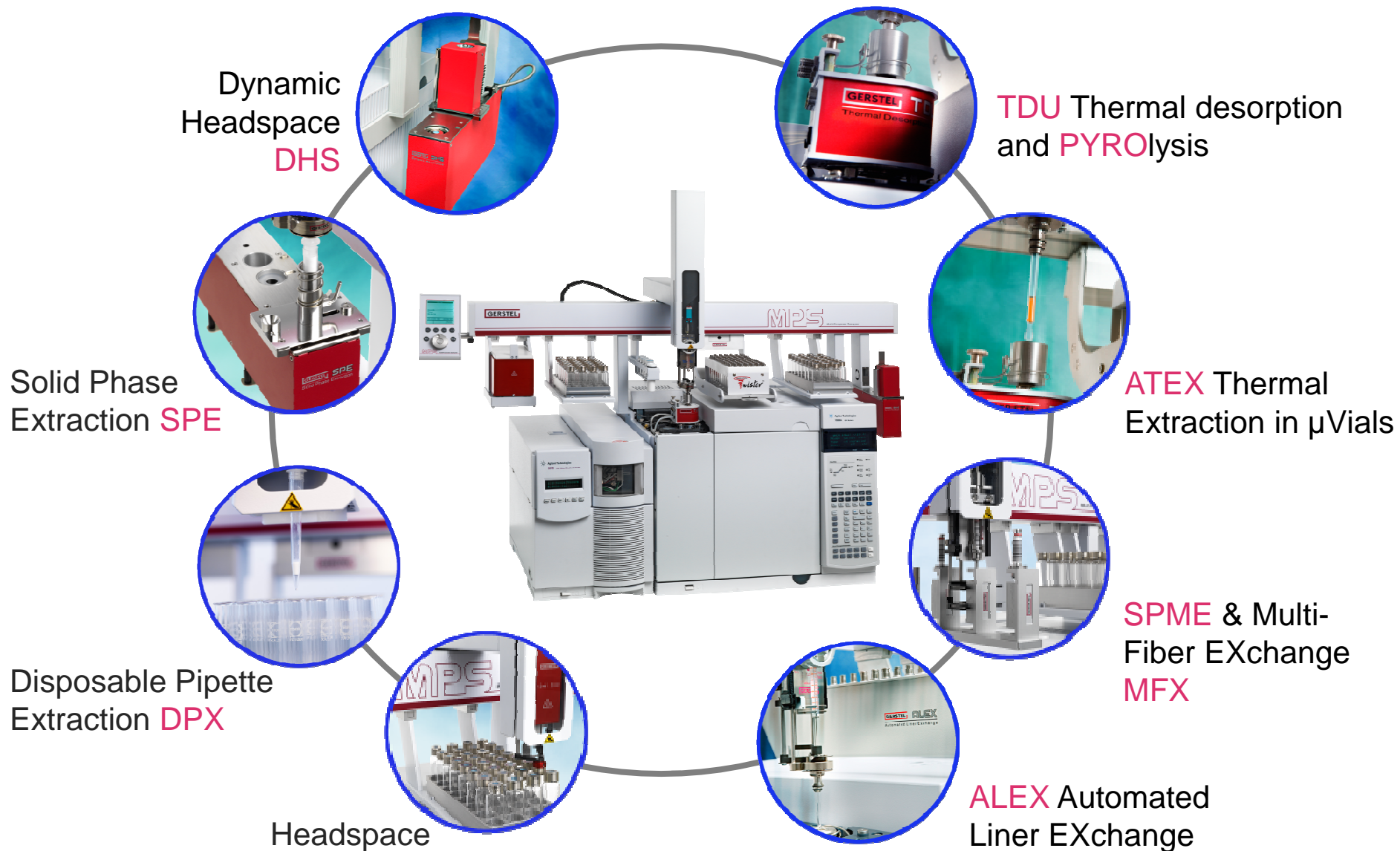
Research Institute
for Chromatography

1. Sample Preparation



RIC

Research Institute
for Chromatography



Personalized Solutions for Automated Sample Preparation



RIC

Research Institute
for Chromatography



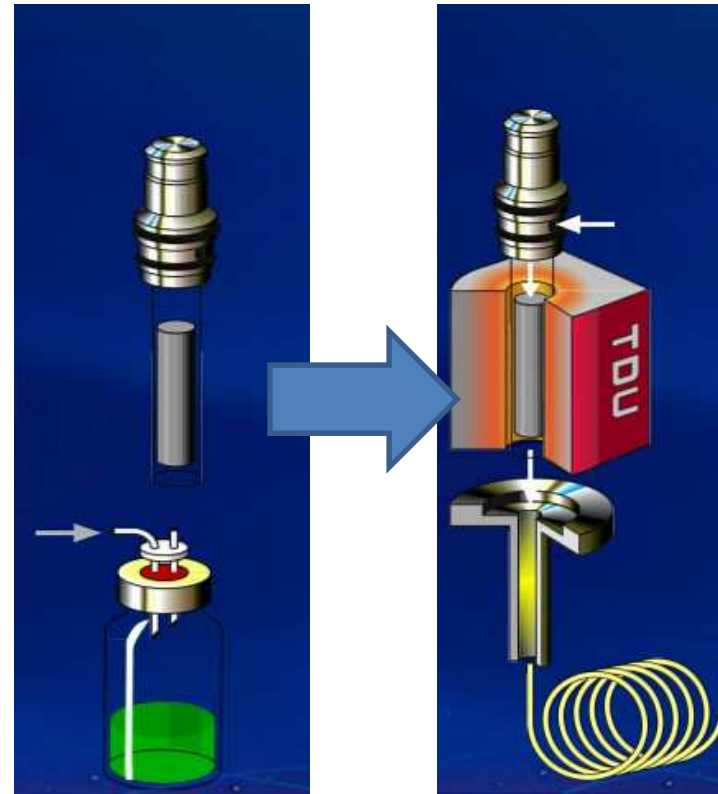
Personalized Solutions for Automated Sample Preparation



RIC

Research Institute
for Chromatography

Dynamic Headspace (DHS)

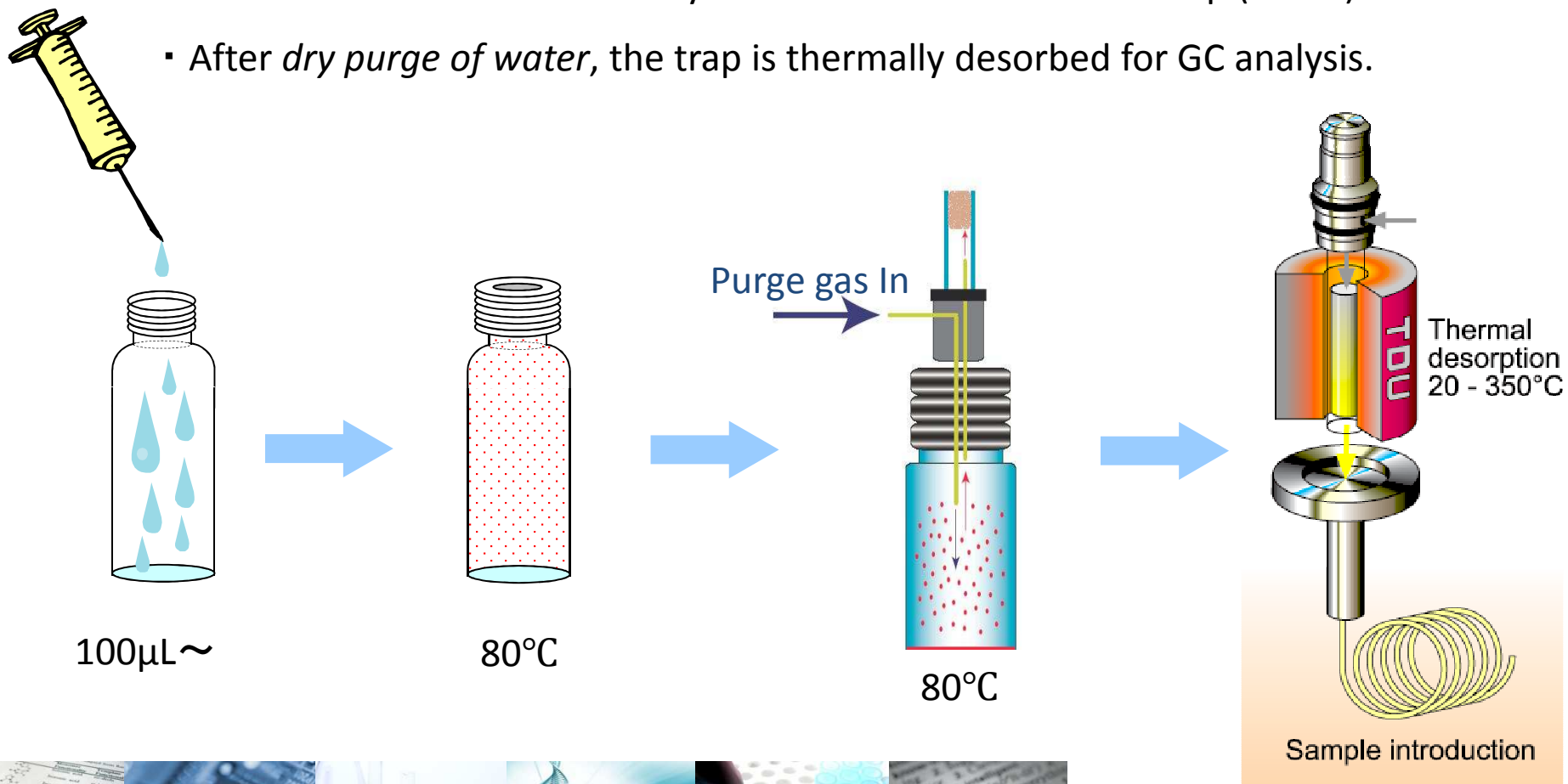


RIC

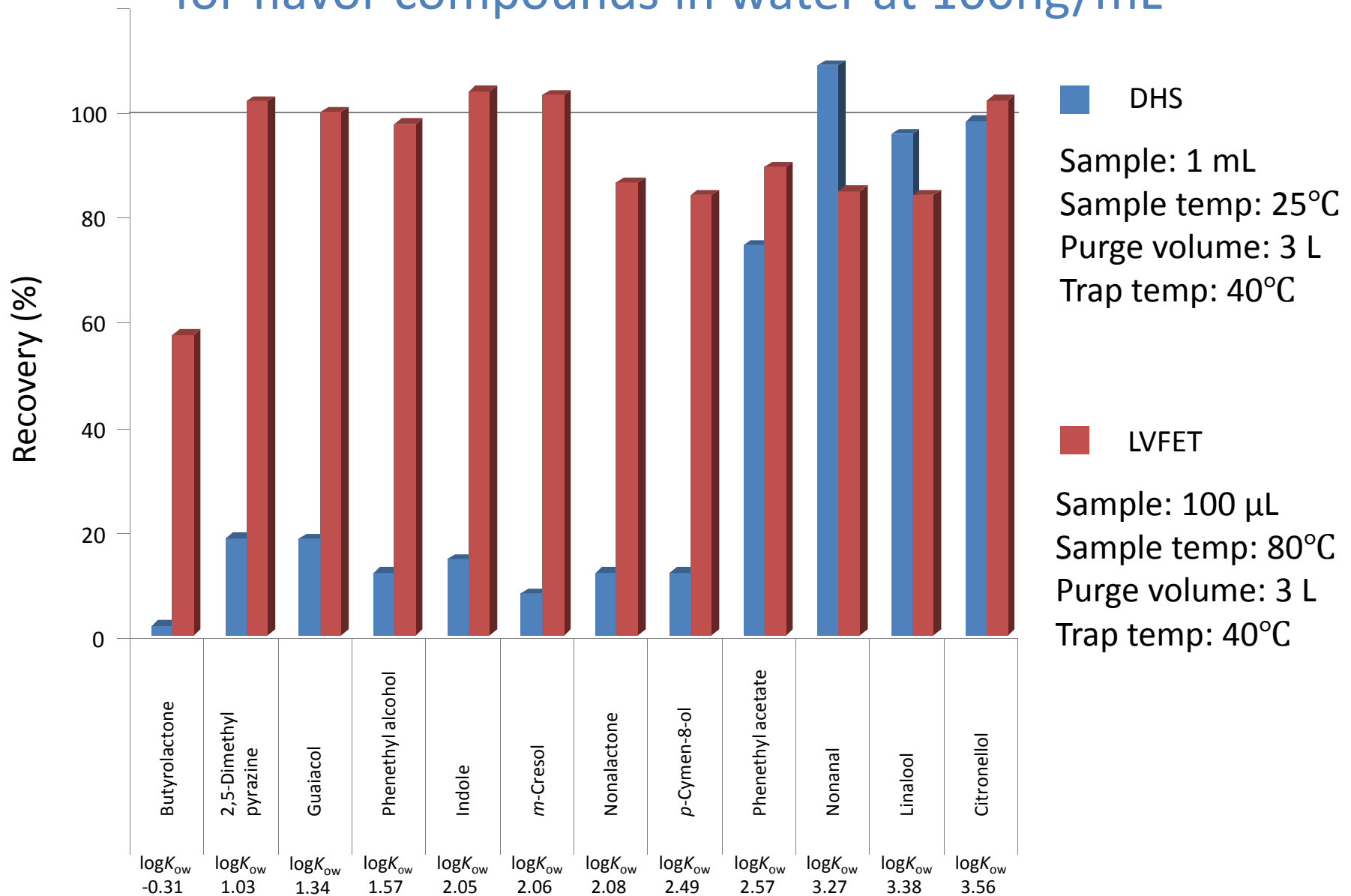
Research Institute
for Chromatography

Large Volume FEDHS (LV-FEDHS)

- **100 μL of aqueous sample !**
purged with inert gas at an elevated temperature (80°C) using DHS.
- Volatile and semi-volatile analytes are transferred into the trap (Tenax)
- After *dry purge of water*, the trap is thermally desorbed for GC analysis.



Comparison of recovery between DHS and FEDHS for flavor compounds in water at 100ng/mL



FEDHS of Flavors and Fragrances in Cosmetics

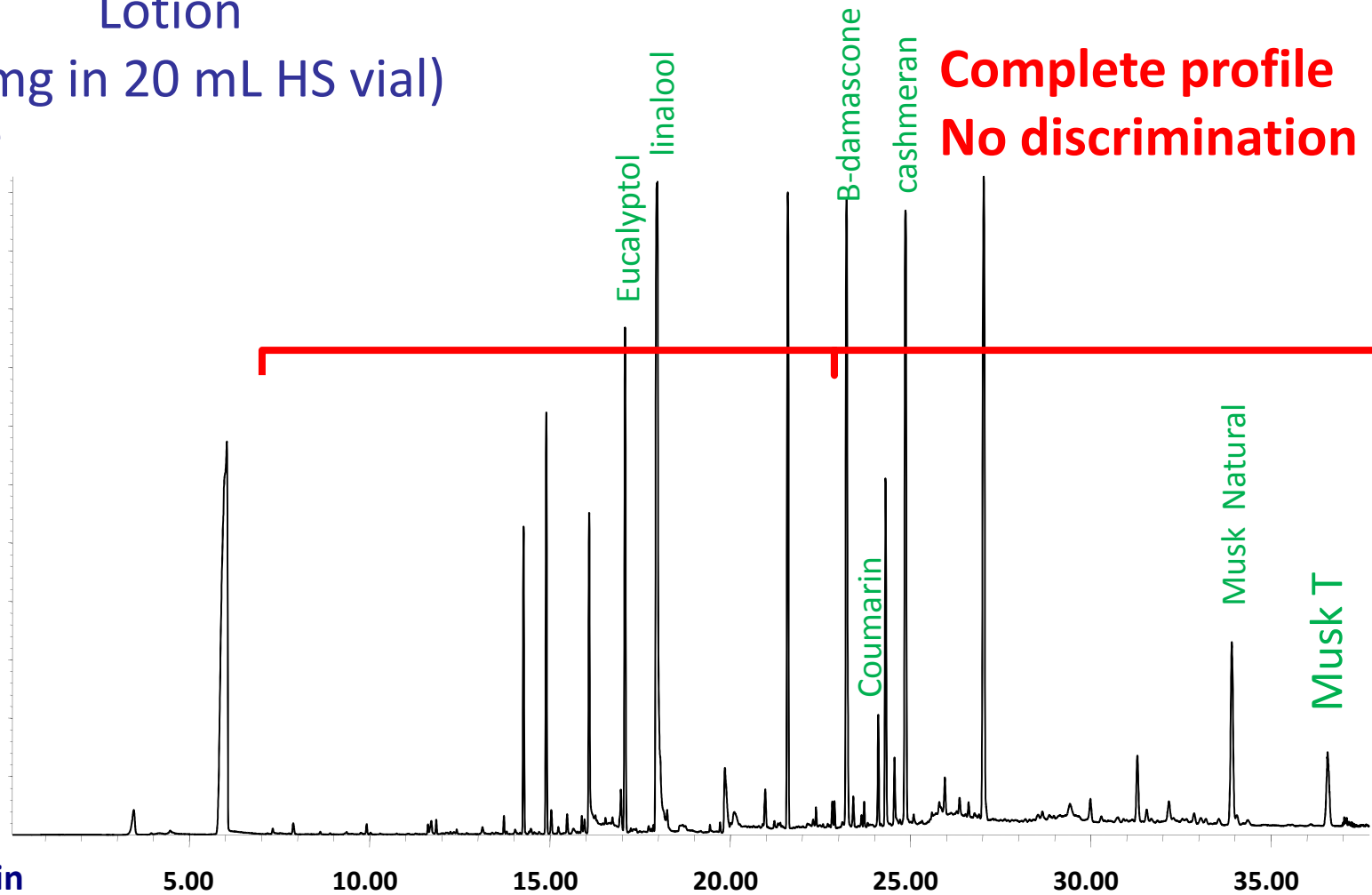
Lotion
(50 mg in 20 mL HS vial)

Abundance

5.5e+07
5e+07
4.5e+07
4e+07
3.5e+07
3e+07
2.5e+07
2e+07
1.5e+07
1e+07
5000000

Time, min

5.00 10.00 15.00 20.00 25.00 30.00 35.00



RIC

Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 1: Very Volatile Analytes



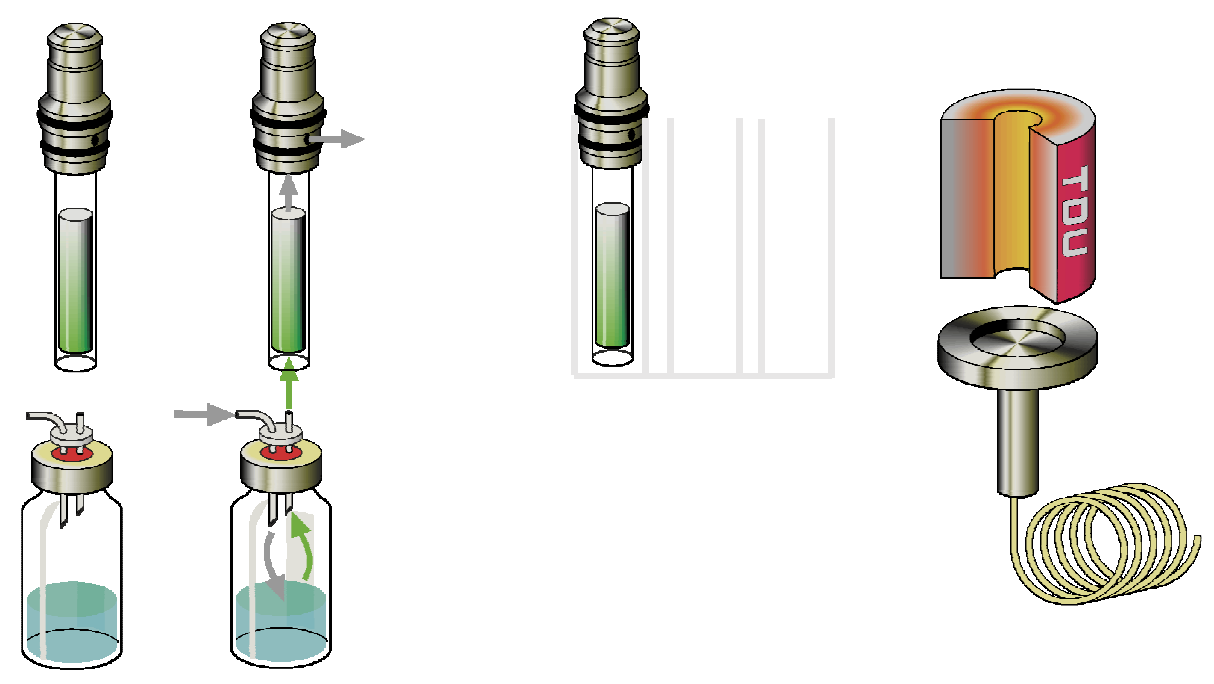
25°C - carbon-based adsorbent trap
150 mL @ 50 mL/min



RIC | Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace Method 1 for Volatile and Semi Volatile Analytes



25°C - carbon-based adsorbent trap
650 mL @ 100 mL/min

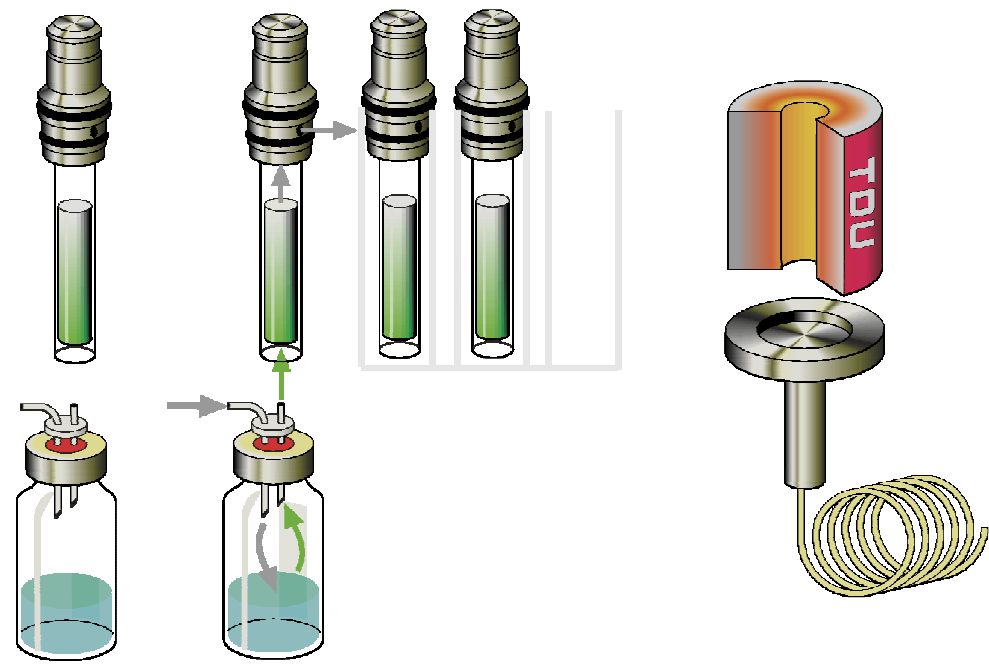


RIC | Research Institute for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 2: Volatile and hydrophilic analytes



80°C - Tenax TA trap
3L @ 100 mL/min

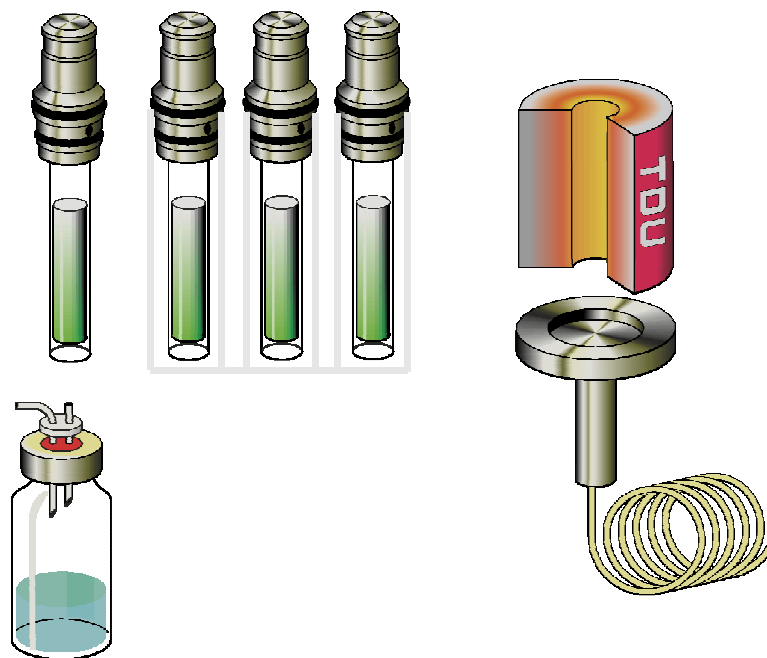


RIC | Research Institute for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 3: Volatile, non volatile and hydrophilic analytes

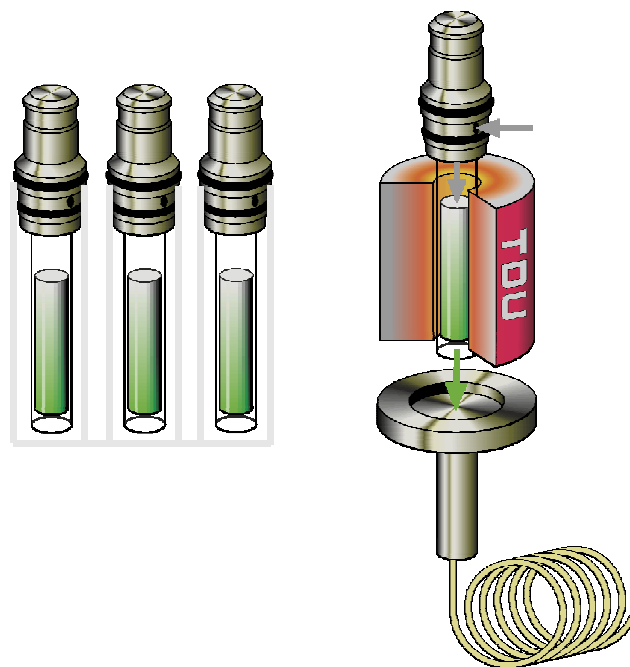


RIC | Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 4: TDU Multi Desorption

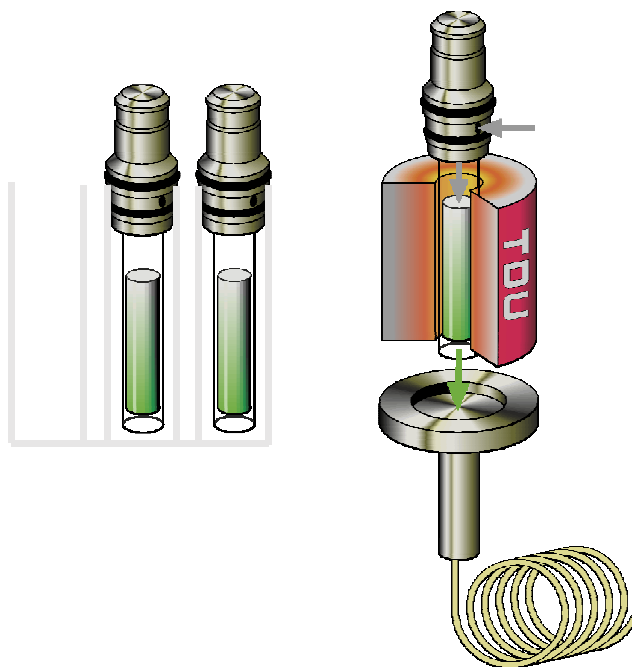


RIC | Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 4: TDU Multi Desorption

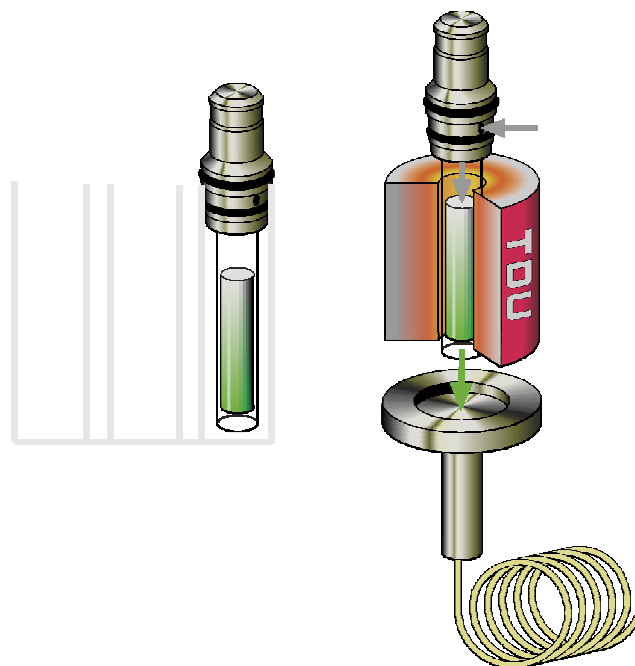


RIC | Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

Method 4: TDU Multi Desorption

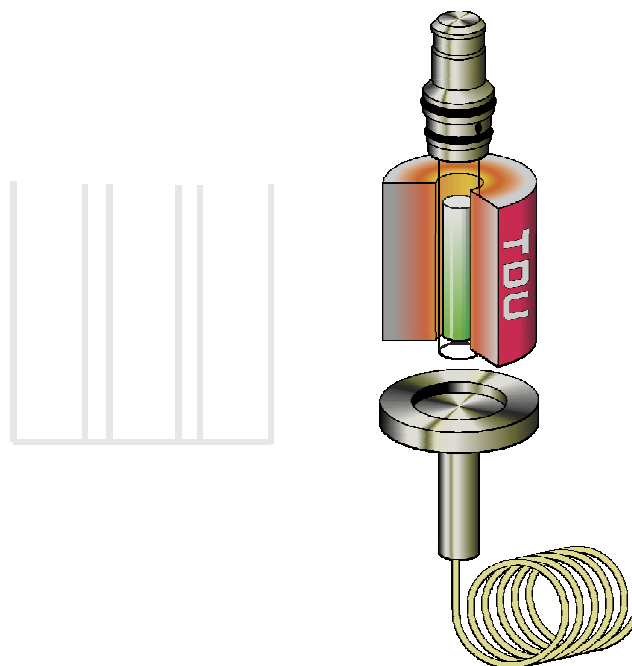


RIC | Research Institute
for Chromatography

Sequential Dynamic Headspace Sampling - Multi-Volatile Method (MVM) for Aroma Analysis of Beverages

Dynamic Headspace

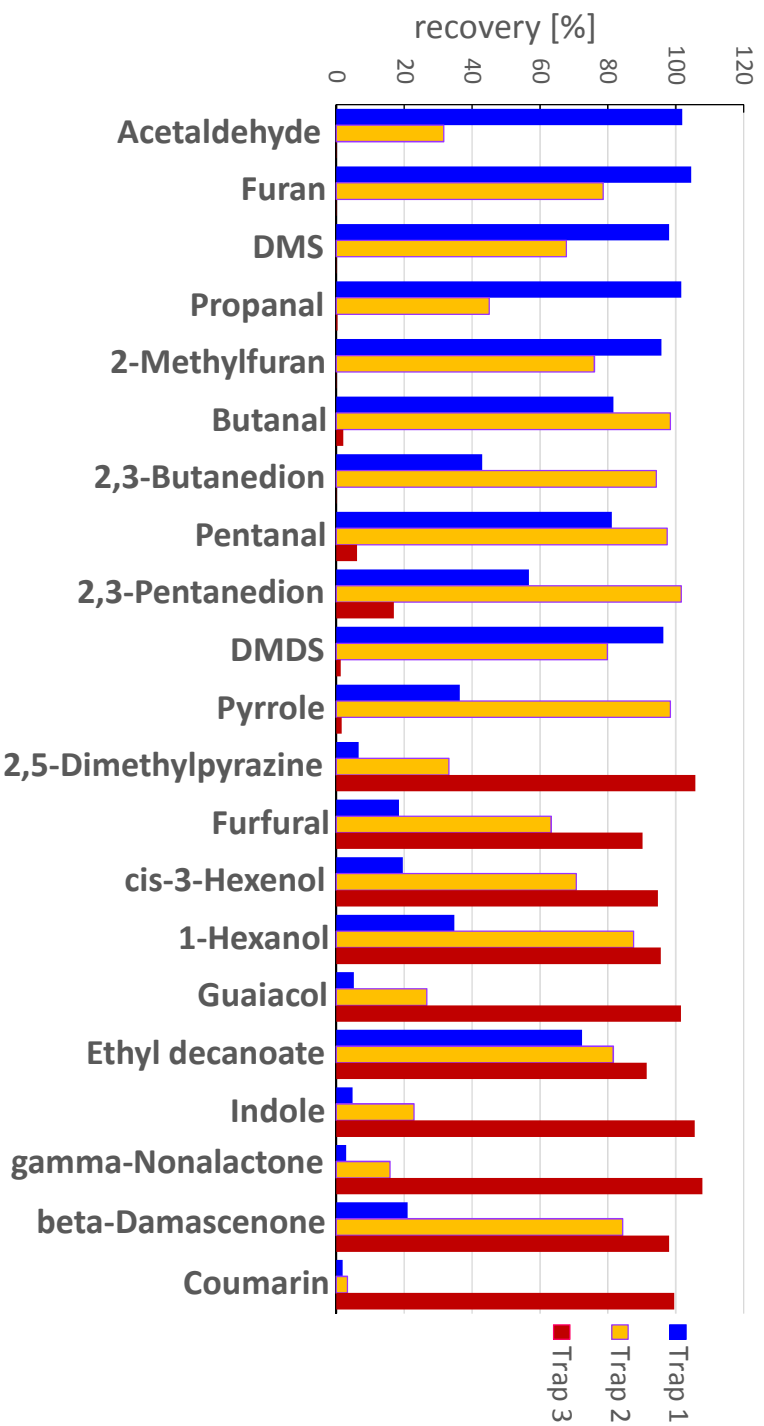
Method 4: TDU Multi Desorption



RIC

Research Institute
for Chromatography

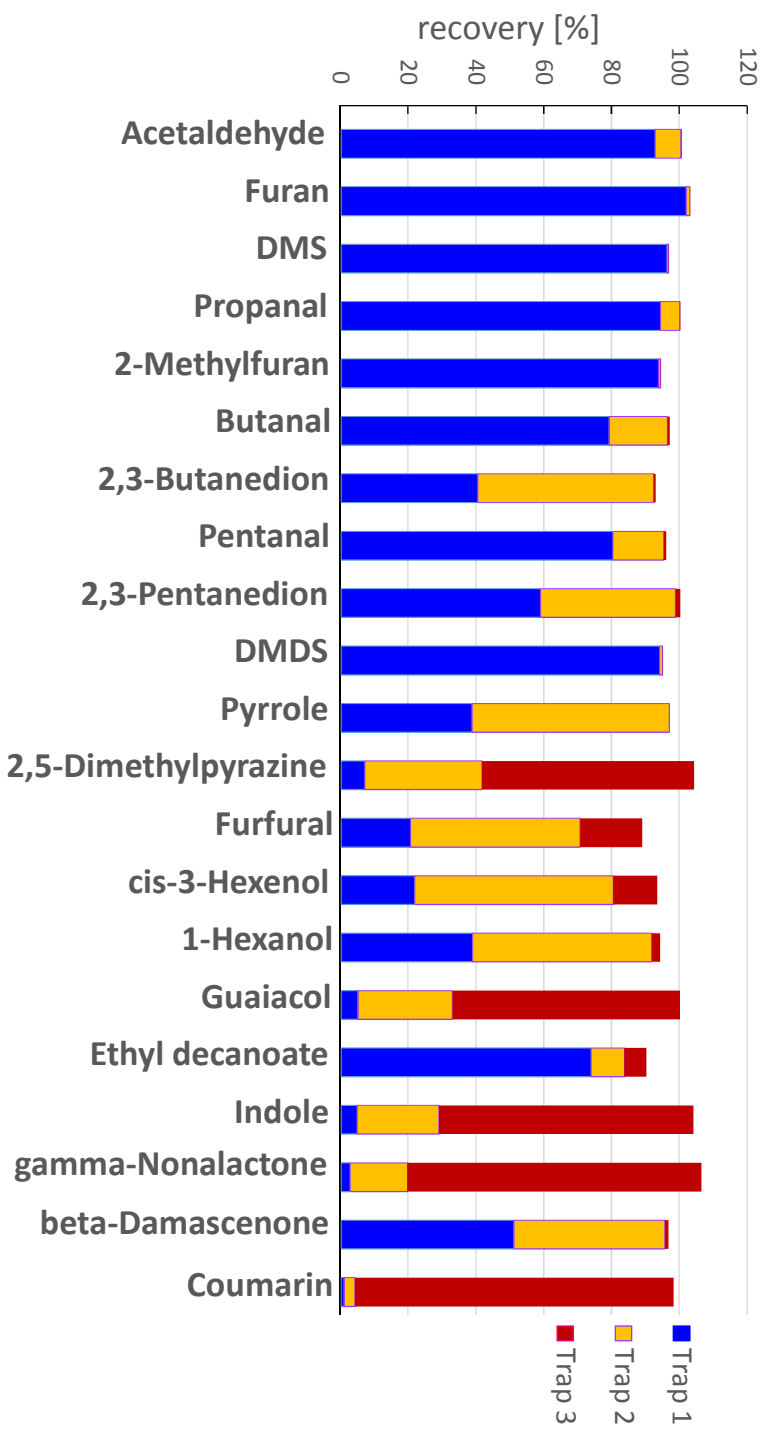
Brewed Coffee MVM Extraction



RIC

Research Institute
for Chromatography

Brewed Coffee MVM Extraction



RIC

Research Institute
for Chromatography

Wine Analysis using SBSE-GC-MS

Céline Franc, Frank David, Gilles de Revel, JCA



Multi-residue methods

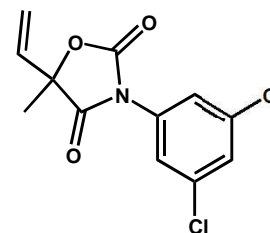
- 2007 : Off-flavours

IBMP, EP, EG, TCA, TeCA, PCA, TBA, Géosmine

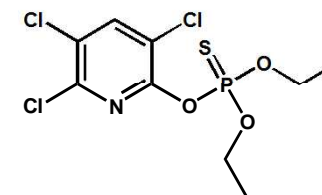
- 2010 : Markers of wine aroma (fruity)

C13-norisoprenoides and lactones

- 2010 : Pesticide residues



Vinclozoline



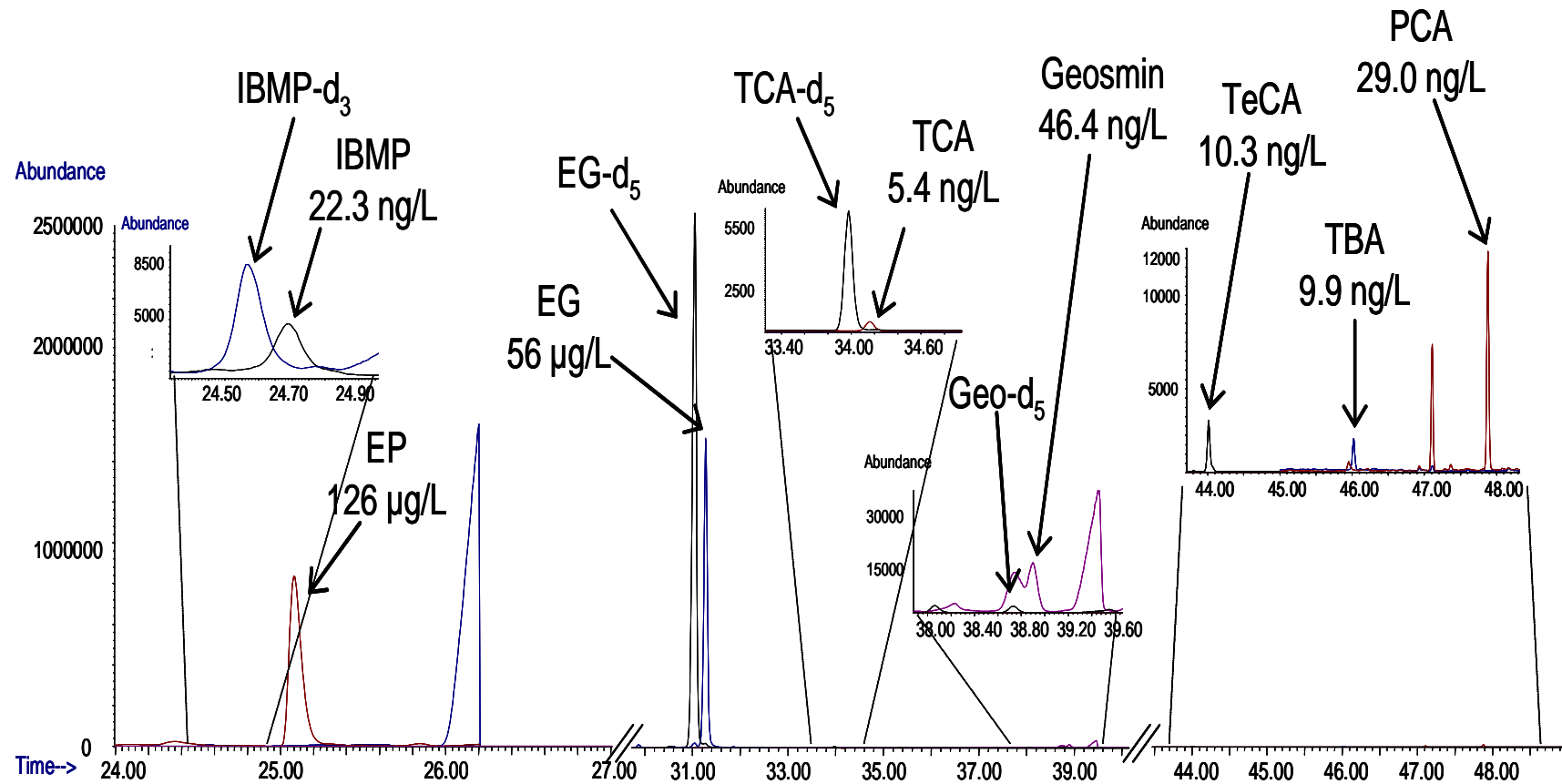
Chlorpyrifos



RIC | Research Institute
for Chromatography

Multiresidue Analysis of Wine Defects

Céline Franc, Frank David, Gilles de Revel, JCA



RIC | Research Institute
for Chromatography

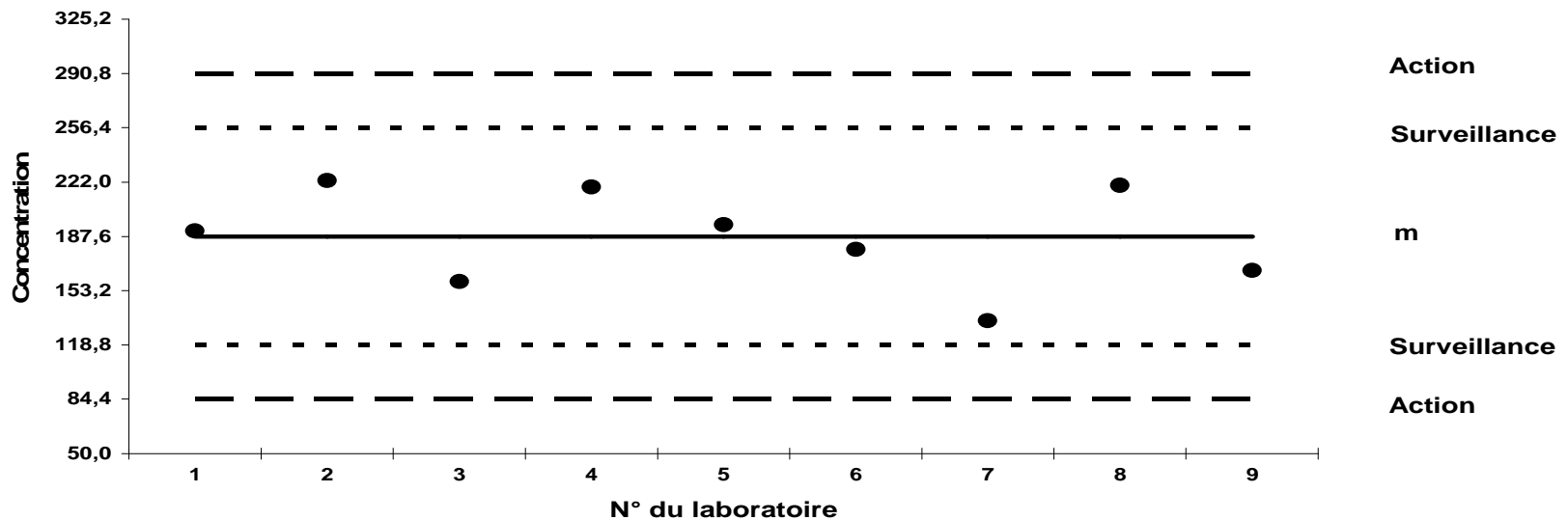
Application : Round-Robin test SBSE-GC-MS « off-flavors"»

2009 : inter-laboratory test, 8 compounds – 9 laboratoires

Volatile phenols

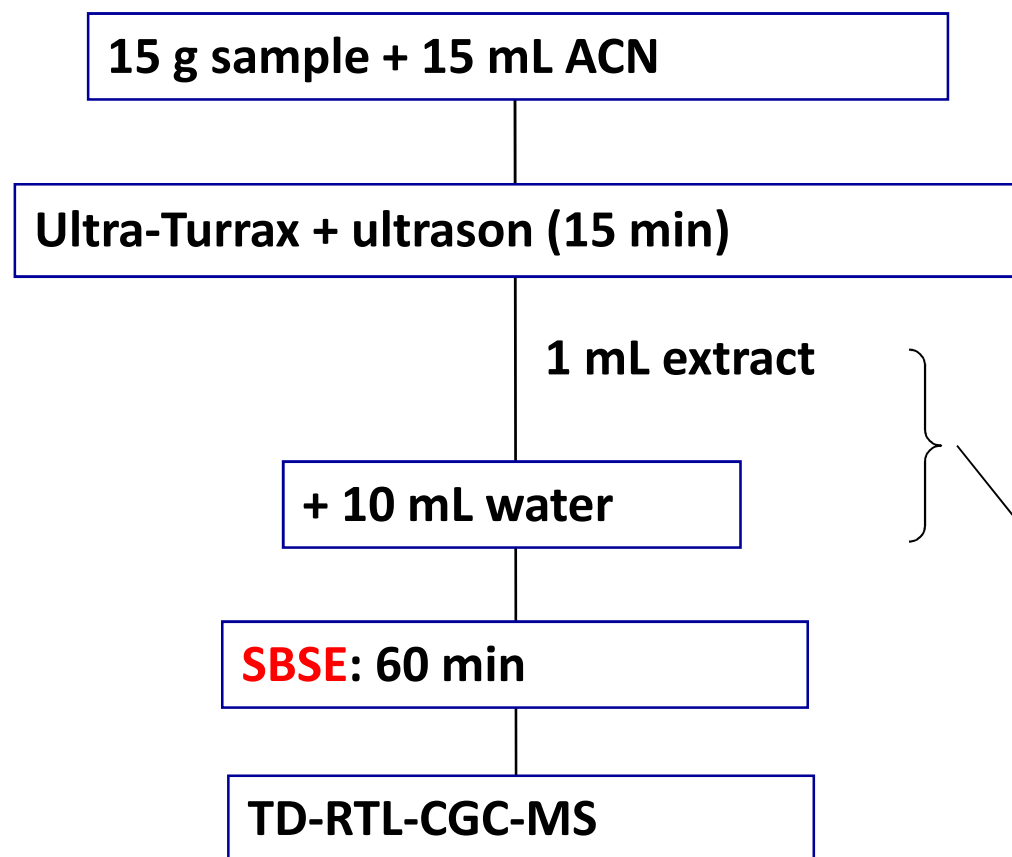
nombre de laboratoires retenus :	9		
valeur assignée m :	187,6		
écart-type s* :	34,4		
limites de surveillance :	118,8	;	256,4
limites d'action :	84,4	,	290,8

Répartition des résultats des laboratoires



RIC | Research Institute
for Chromatography

SBSE procedure for vegetables and fruit



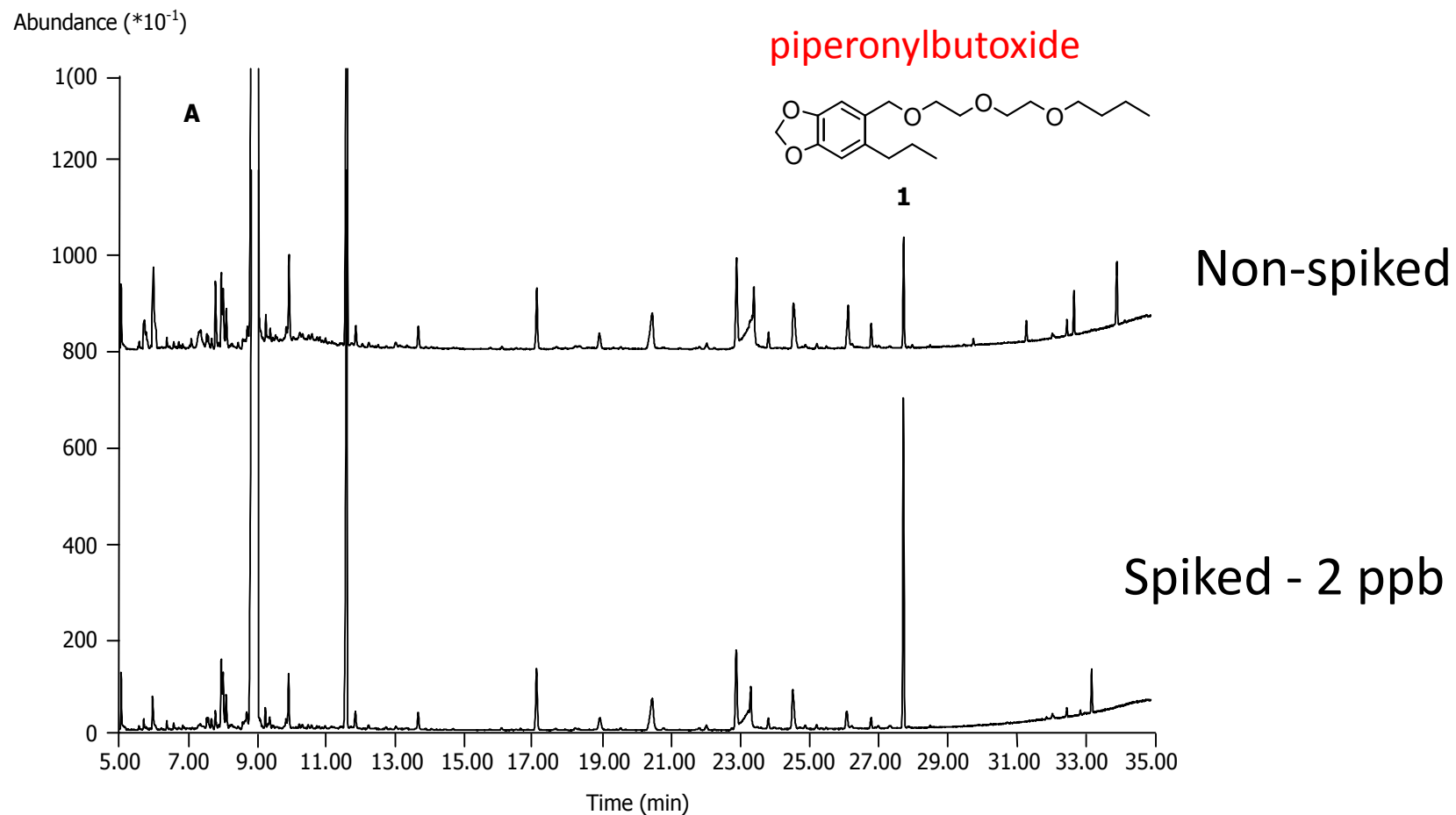
(=QuEChERS)

*P. Sandra, B. Tienpont and F. David
J. Chromatogr., 1000 (2003) 299-309*

**Modified ratio water/organic
Multi-Twister extraction**
N. Ochiai, K. Sasamoto, H. Kanda, T. Yamagami, F. David, B. Tienpont and P. Sandra
J. Sep. Sci 28 (2005) 1083-1092



Analysis of Baby Food by SBSE-TD-GC-MSD (mixed vegetables, rice, chicken)



QC on solid material

'Passive' Extraction

10 corks – leaching during 24h

in 500ml wine simulant (10 % Ethanol)



SBSE extraction for leachable haloanisoles:

- 100ml « extract (10 % Ethanol) »
- IS: TCA-d5 (10ng/l),
- 2 h extraction with 20 mm x 0.5 mm stir bar



RIC

Research Institute
for Chromatography

Polarity?

3 Strategies



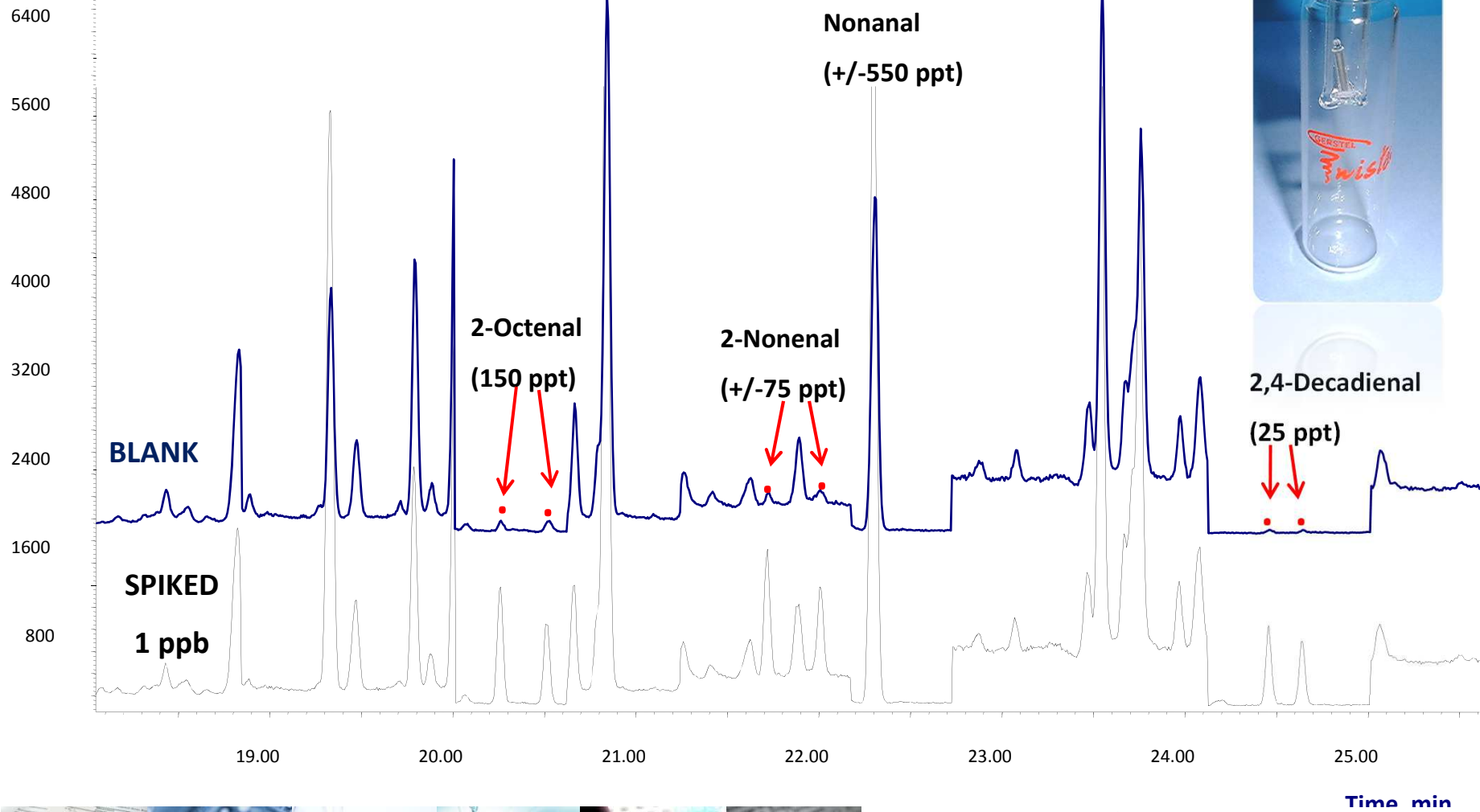
RIC

Research Institute
for Chromatography

Derivatization HSSE

perfluorobenzyl hydroxylamine (PFBHA) - BEER

Abundance

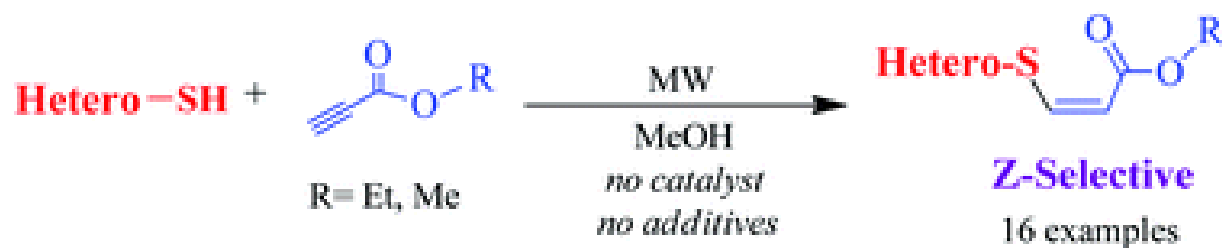


RIC

Research Institute
for Chromatography

Analysis of Thiols (wine, beer,...)

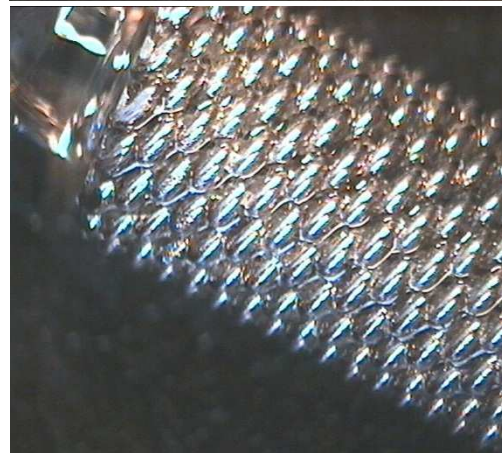
- Polyfunctional thiols [3-mercaptohexan-1-ol (3MH), 3-mercaptohexyl acetate (3MHA) and 4-mercapto-4-methylpentan-2-one (4MMP)] are important aroma compounds. Current methods lack specificity and sensitivity.
- Derivatization with alkyl propiolate (ethyl propiolate) can be performed in-situ and, combined with SBSE, high sensitivity and good selectivity are obtained for the detection of the thioacrylates.



- See: N. Ochiai et al.

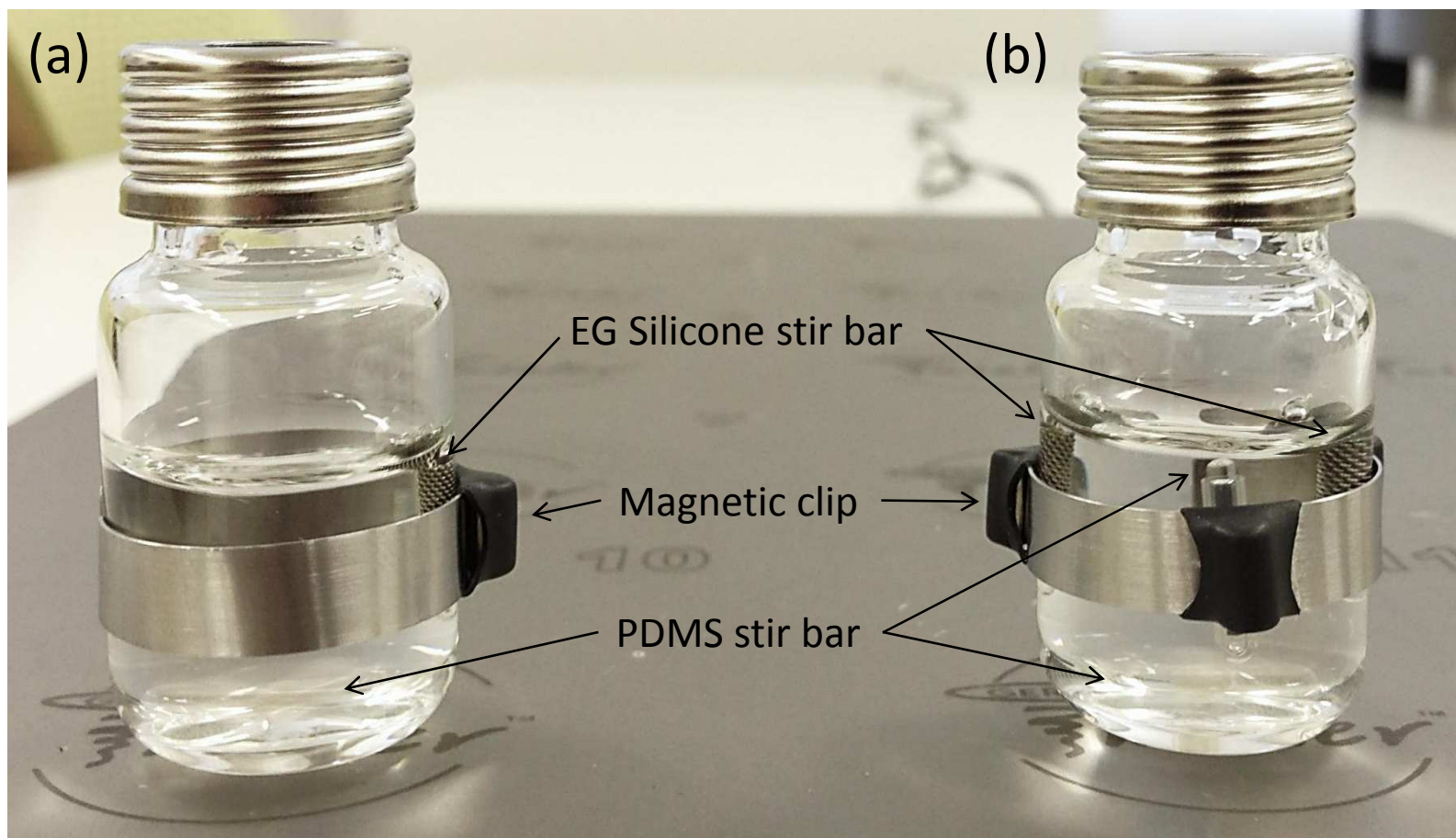


EG Silicon Twister



RIC | Research Institute
for Chromatography

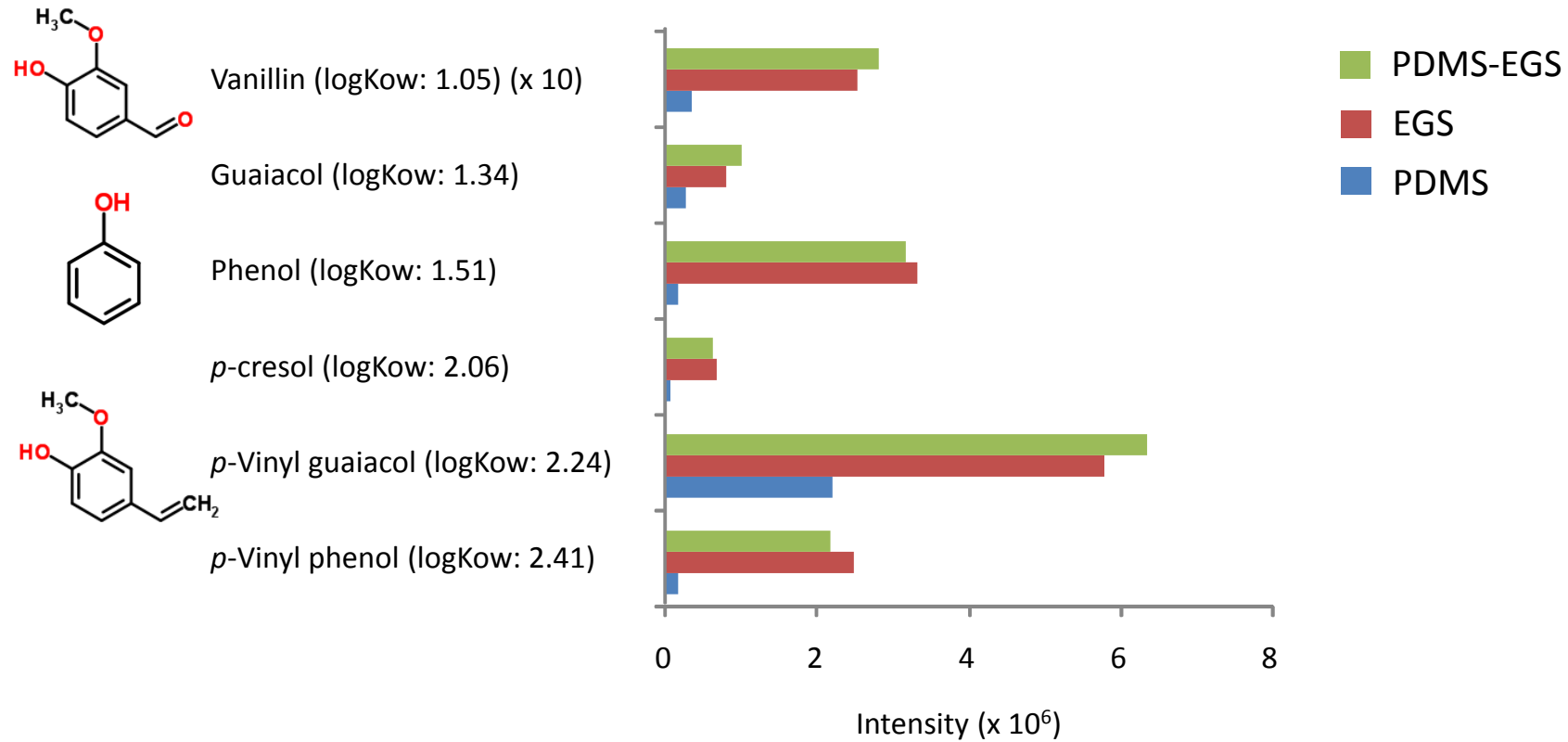
mSBSE using different coatings



RIC

Research Institute
for Chromatography

Comparison of recovery of phenolic compounds between EG Silicon, PDMS and Seq-SBSE

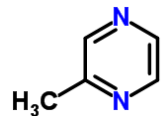


J.I. Cach et al, J. Pharm Biomed Anal 78-79 (2013) 255: Analysis of Bisphenols in Cosmetics



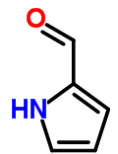
RIC | Research Institute
for Chromatography

Comparison of recovery of **nitrogen heterocyclic** compounds between EG Silicon, PDMS and Seq-SBSE



Methyl pyrazine (logKow: 0.49)

2-Acetyl pyrrole (logKow: 0.56)



2-Formyl pyrrole (logKow: 0.60)

Pyridine (logKow: 0.80)

2,5-Dimethyl pyrazine (logKow: 1.03)

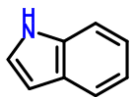


4,5-Dimethyl oxazole (logKow: 1.31) x10

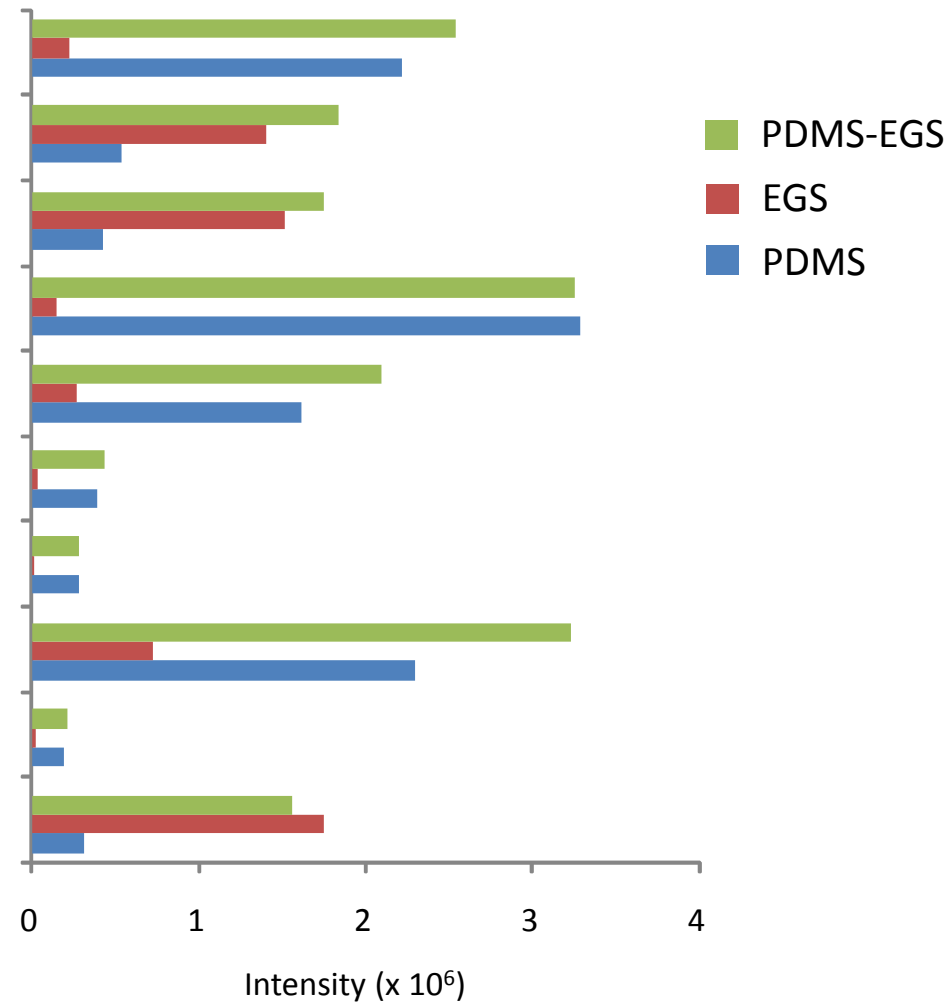
3,4-Dimethyl isoxazole (logKow: 1.31) x10

5-Methyl pyrazine (logKow: 1.53)

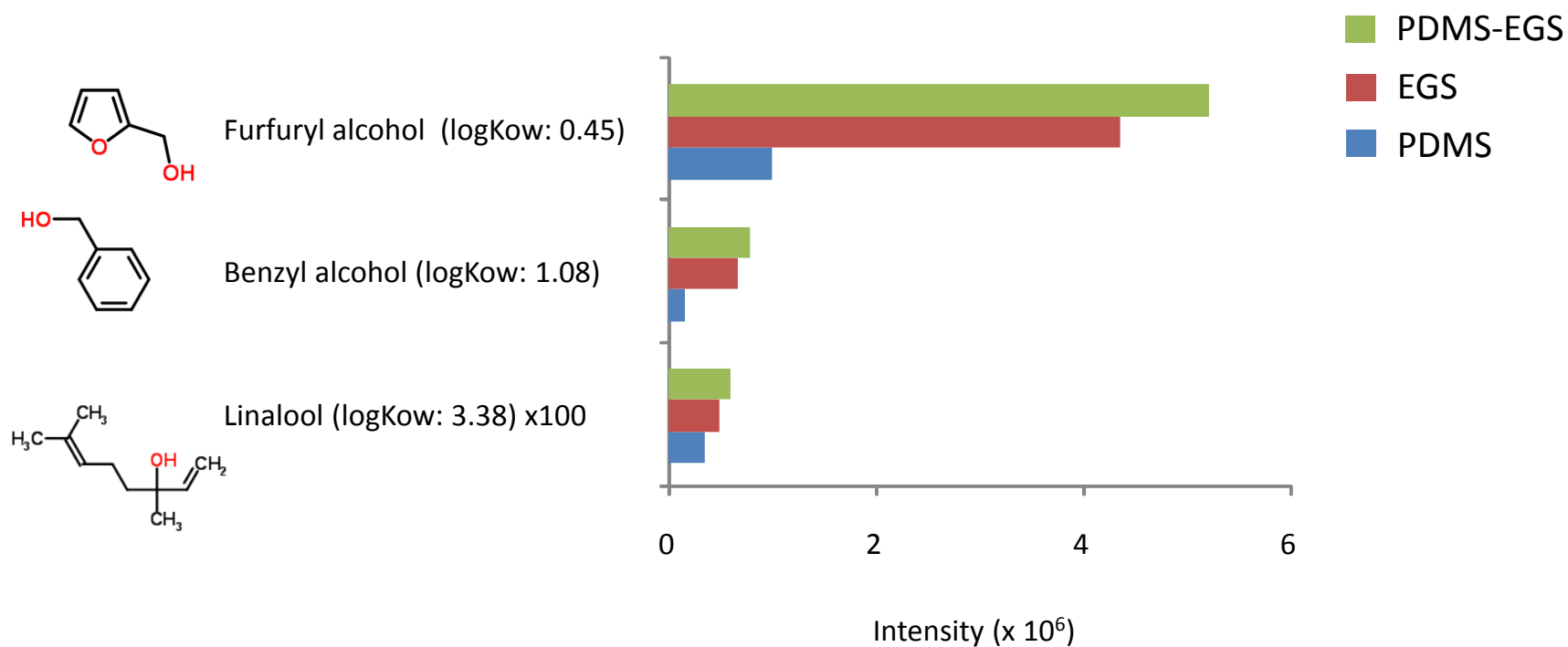
2,4,5-trimethyl oxazole (logKow: 1.86)



Indole (logKow: 2.05)



Comparison of recovery of **alcohols** between EG Silicon, PDMS and Seq-SBSE



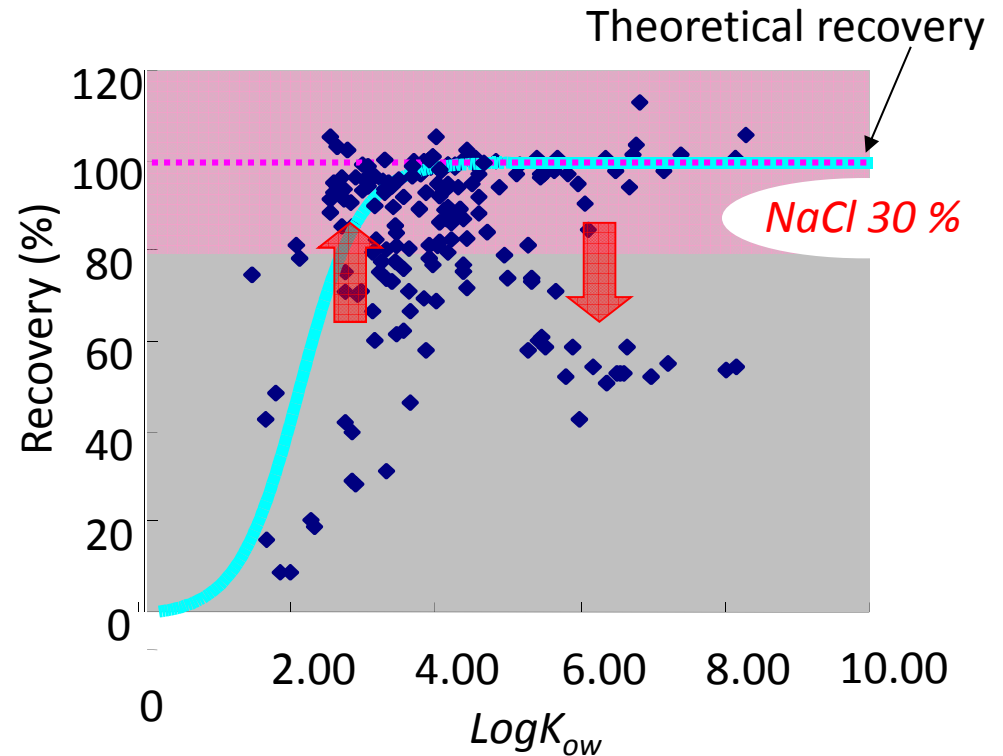
Sequential Stir Bar Sorptive Extraction

1. SBSE in pure water

Hydrophobic compounds
 $\text{LogKow} > 4$

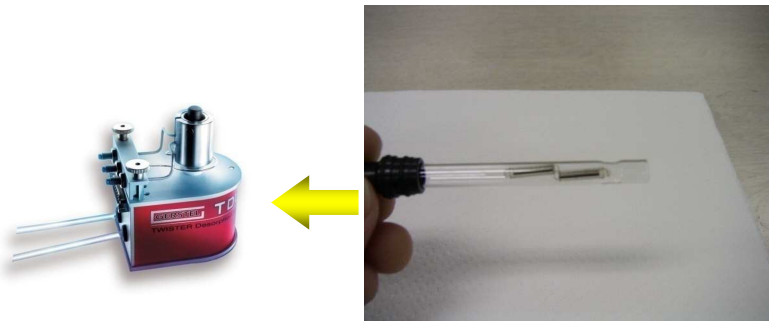
2. Add Salt to same sample - SBSE

Hydrophilic compounds
 $\text{LogKow} < 4$



Recovery of 88 pesticides by SBSE

N. Ochiai et al, J. Chromatography 1200 (2008) 72.



RIC | Research Institute
for Chromatography

2. Improve Analytical Performance



RIC

Research Institute
for Chromatography

Capillary Flow Technology (CFT)

(purged) 2 or 3-way
splitter

(purged) ultimate
union

Deans Switch

GCxGC flow
modulator

Quickswap

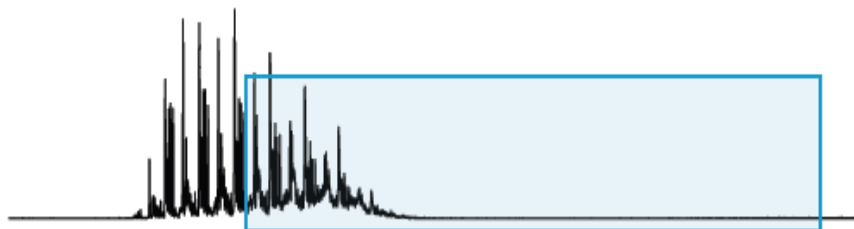


RIC

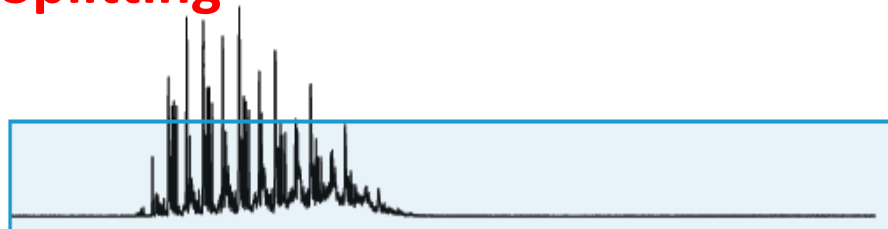
Research Institute
for Chromatography

Agilent CFT capabilities

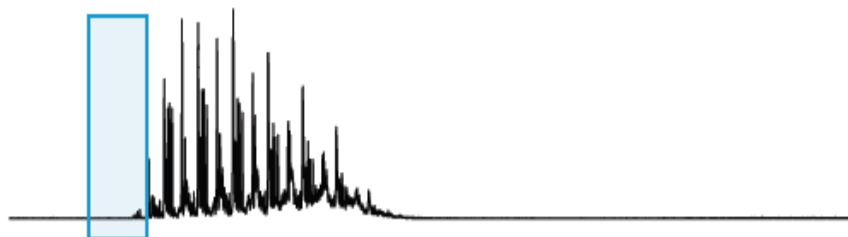
Back Flush



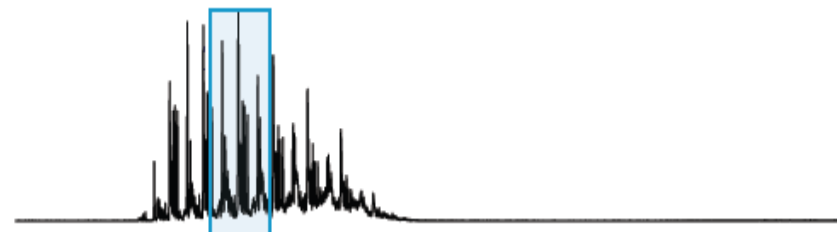
Splitting



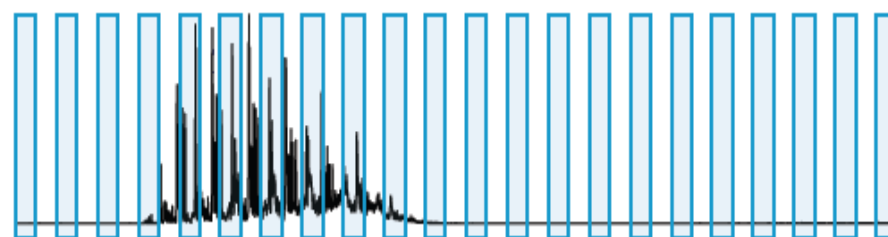
Solvent Bypass



Heart Cutting (Deans Switch)



GC x GC



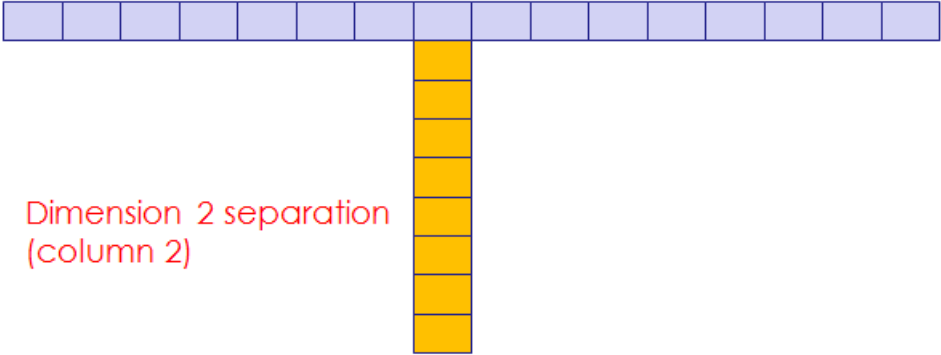
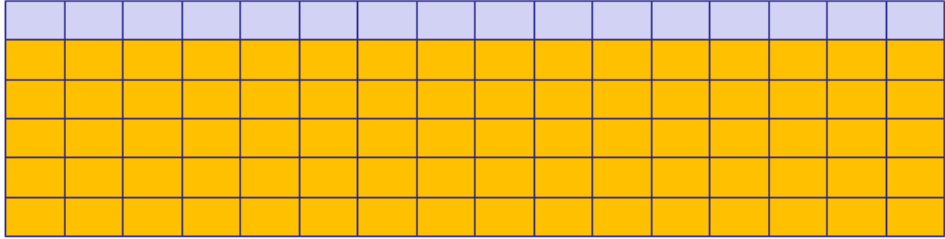
Specialties



RIC

Research Institute
for Chromatography

2D strategies – 4⁺ techniques

<p>Heart-cut</p> <p>Dimension 1 separation (column 1)</p>  <p>Dimension 2 separation (column 2)</p>	<p>GC-GC</p>
<p>Comprehensive</p> <p>Dimension 1 separation (column 1)</p>  <p>2D separation (column 2)</p>	<p>GCxGC</p>
	<p>LCxLC*</p>

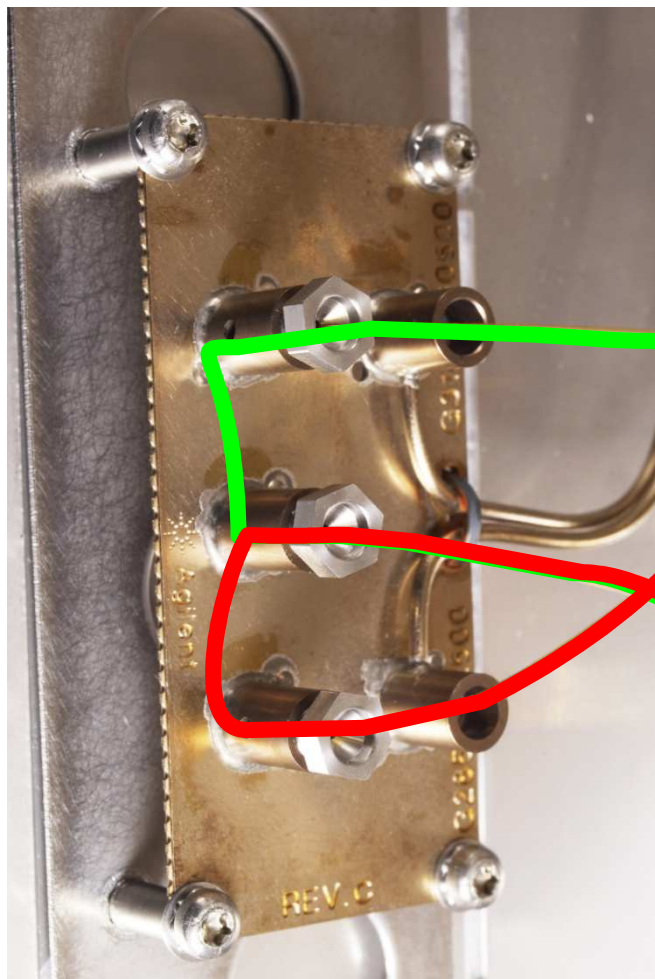
* SFC is an alternative



RIC

Research Institute
for Chromatography

Agilent Deans Switch = *flow selector*



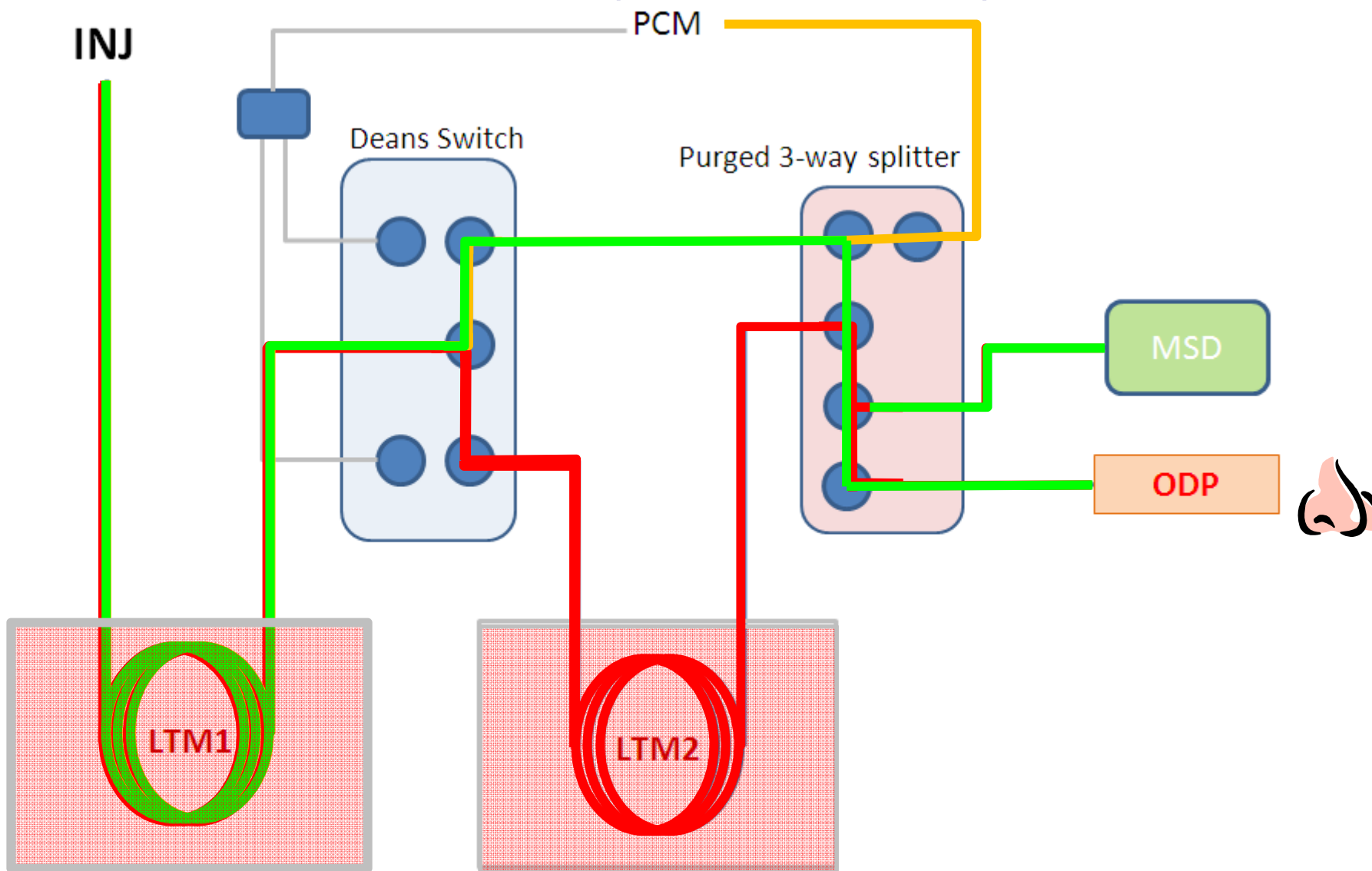
RIC

Research Institute
for Chromatography

Selectable 1D or 2D GCMS (RIC/ Gerstel KK)

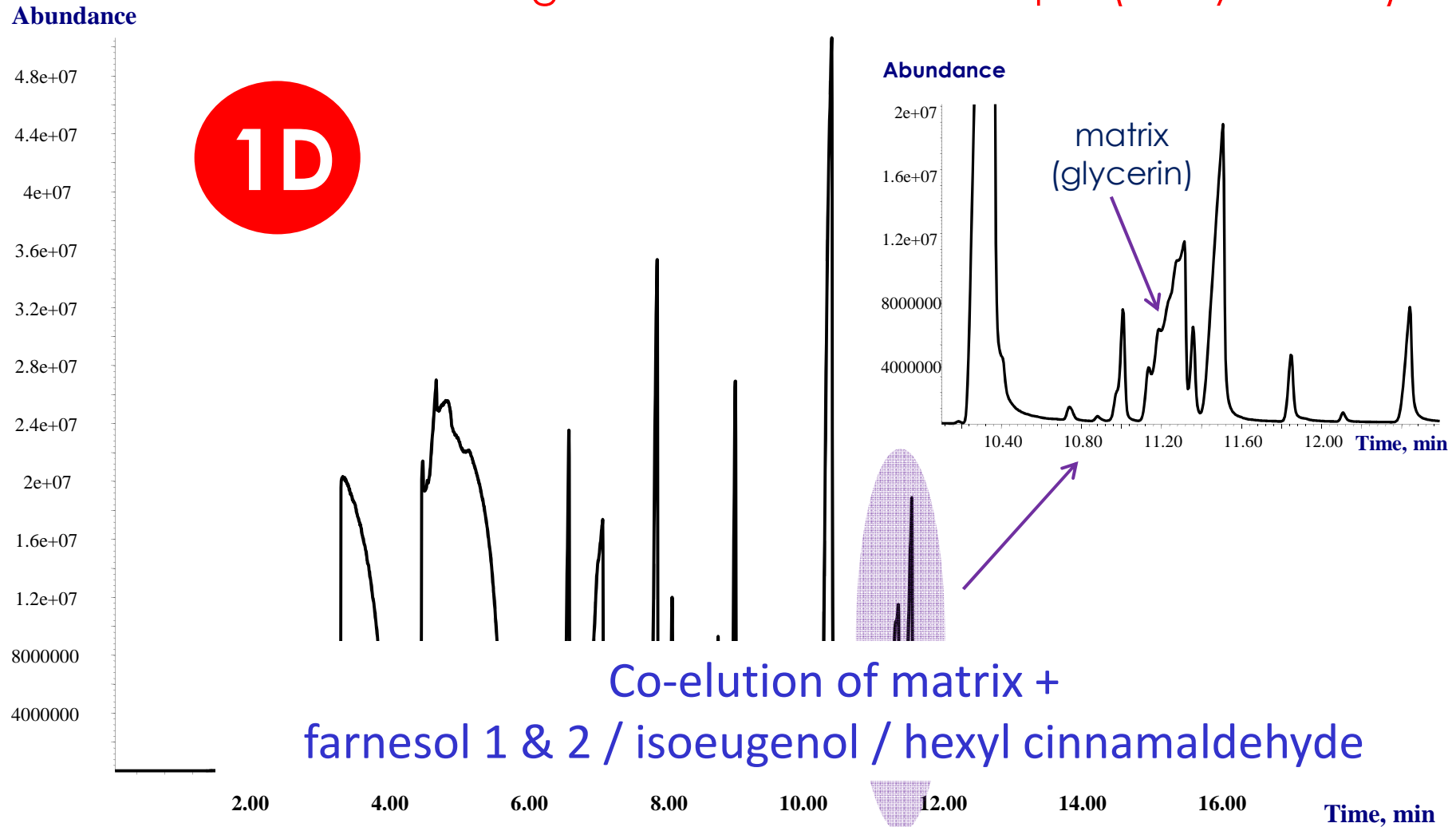


Selectable 1D or 2D GCMS (RIC/ Gerstel KK)



Selectable 1D/2D analysis of allergens in cosmetics

¹D chromatogram of a cosmetic sample (body cream)

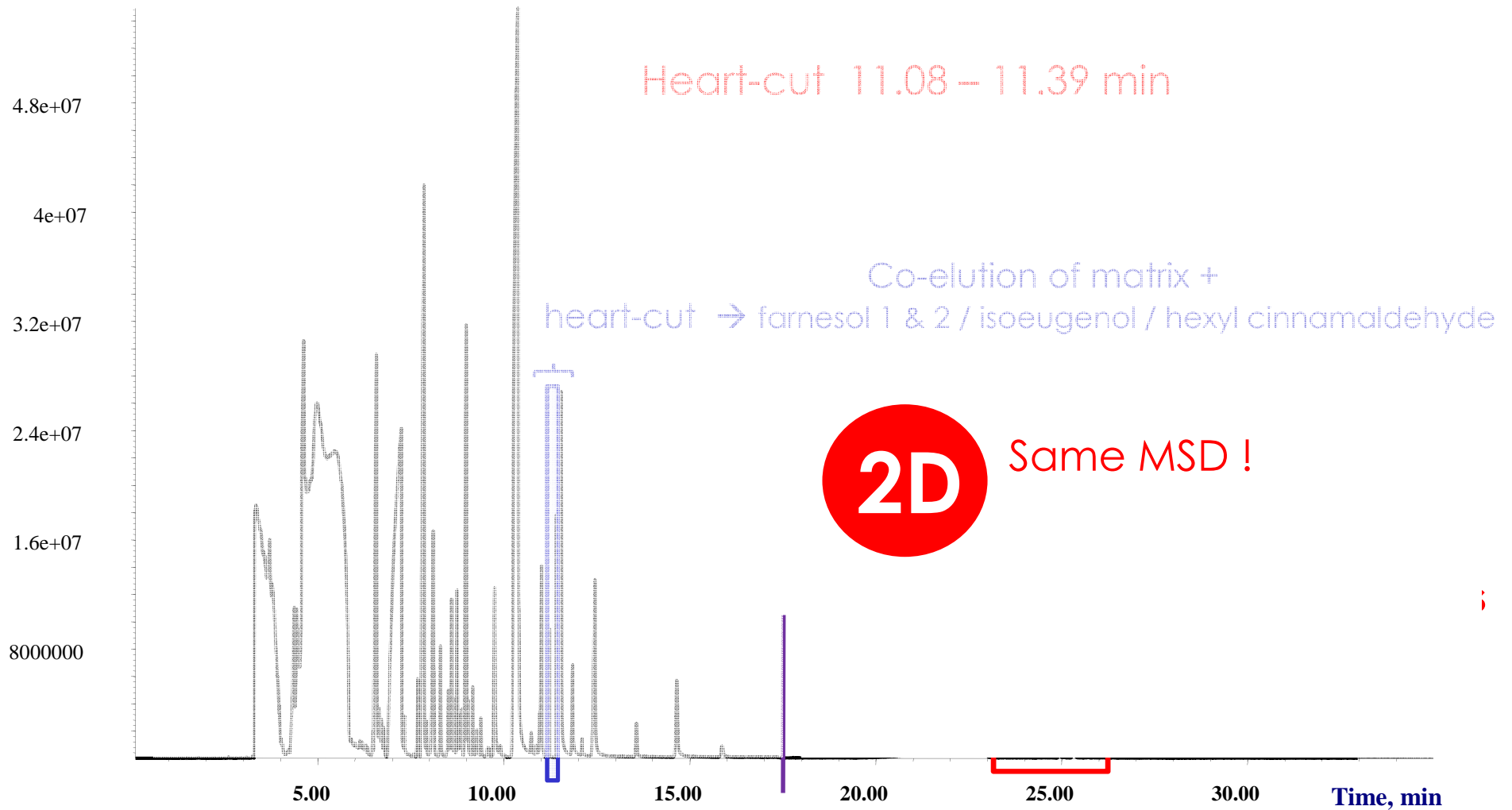


RIC

Research Institute
for Chromatography

Selectable 1D/2D analysis of allergens in cosmetics

Abundance

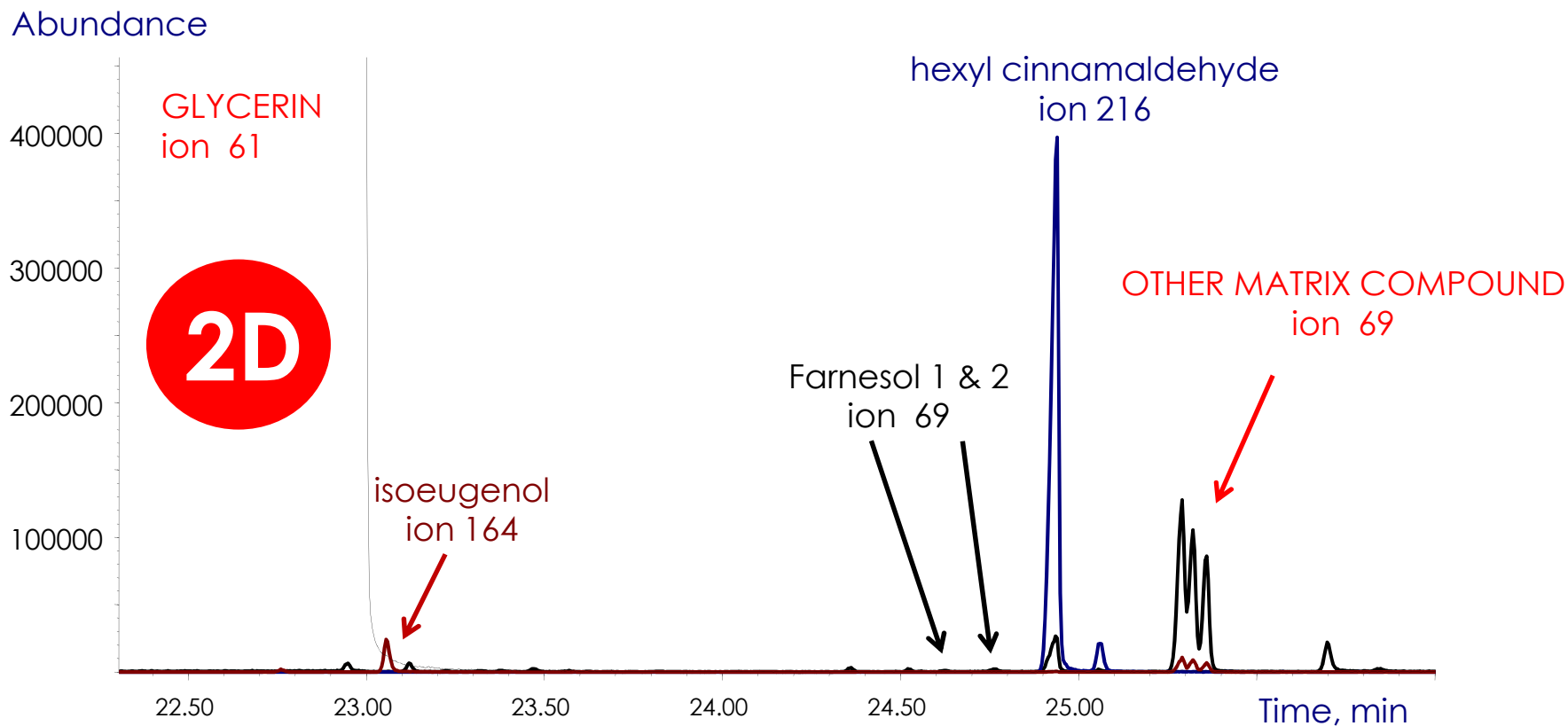


RIC

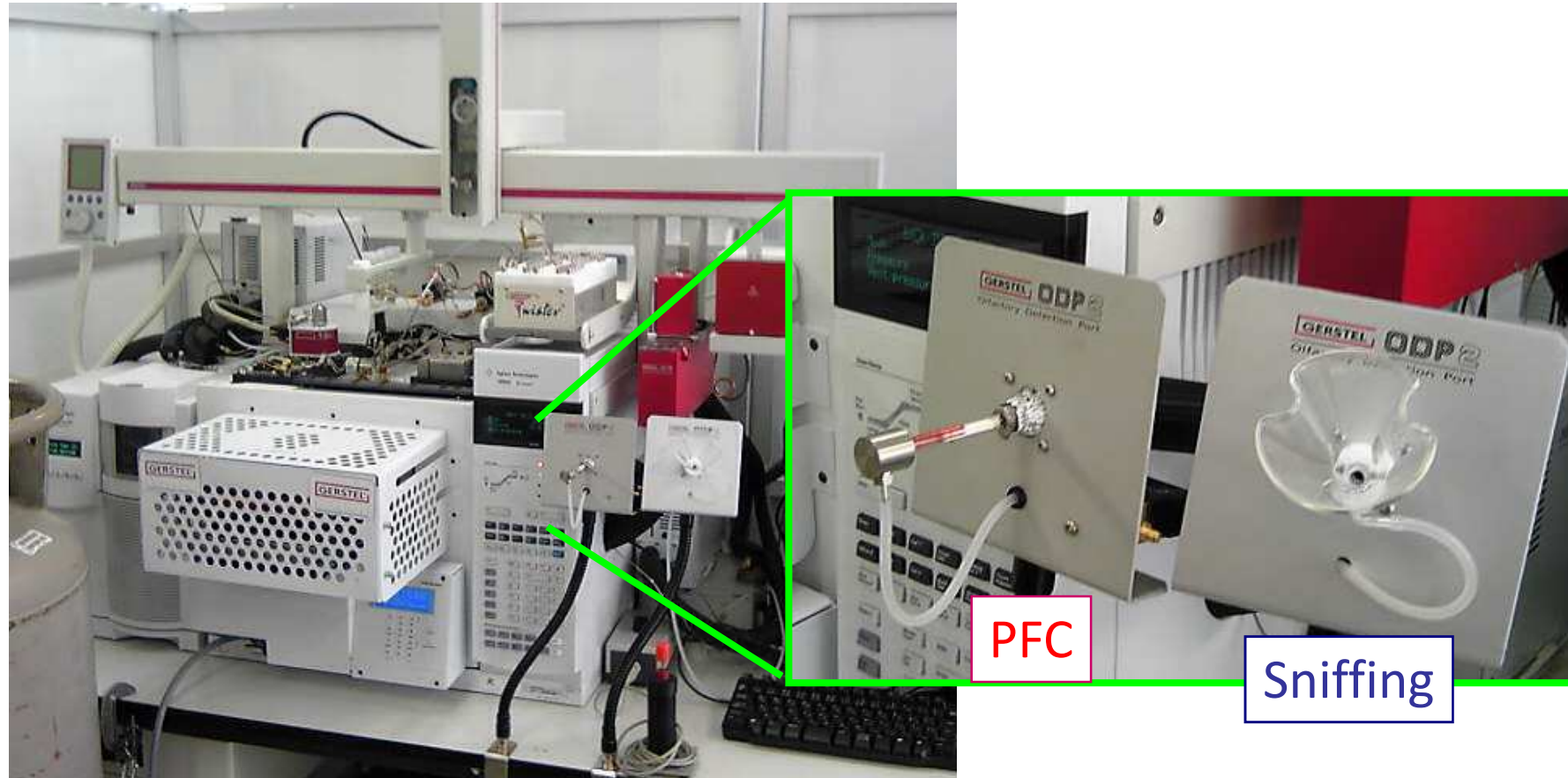
Research Institute
for Chromatography

Sel. 1D/2D analysis of allergens in cosmetics

EIC (m/z 164, 69, 216)



Selectable 1D/2D GC-O/MS with single PFC device



RIC

Research Institute
for Chromatography

Advanced Flavor and Fragrance Analysis using Two-dimensional (2D) GC-MS

Agro Aroma Office 2D



Flavor and Fragrance Compound Database for 2D GC-MS

Name	CAS No	RI	RI Diff	Column
3-methylthiopropanal (methional)	3268-49-3	1451	0	DBWAX
Methional	3268-49-3	1451	0	DBWAX
methional	3268-49-3	1451	0	DBWAX
methional	3268-49-3	1451	0	DBWAX

3-methylthiopropanal (methional)

CCSCC=O

- The searchable database contains RI information for a wide array of odor active compounds from many literature references.
- A cross search can be performed by using two different RIs acquired from both 1st and 2nd dimensional chromatograms, as well as a combined search with a single RI and mass spectrum.
- An RI search can be performed on conventional 1D data.
A cross search can be used for 2 RIs obtained on different columns.

- Selection of 1st and 2nd column
- RI acceptance window
- Elemental information (when it is necessary)

Library Search Report

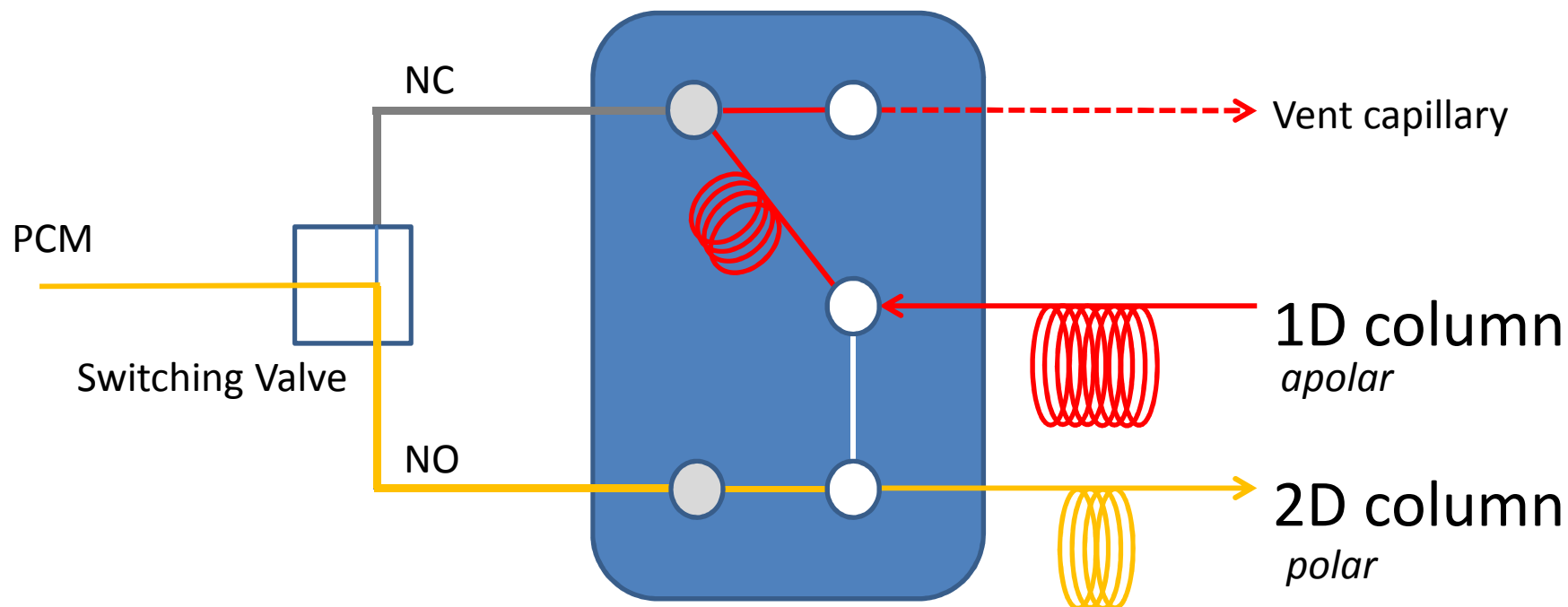
DataFile C:\msdchem\1\data\DEMO_Aroma Office\DBWAX_Floralys5%.D									
PeakNo	RT	RI	Area	Area%	Hit	Name	CASNo	Entry	Library
1)	5.614	1179	84198258	0.355	99	D-Limonene	005989-27-5	15682	NIST11.L
					91	Limonene	000138-86-3	15667	NIST11.L
					76	Cyclohexene, 1-methyl-4-(1-methylethenyl)-, (S)-	005989-54-8	15879	NIST11.L
2)	11.239	1442	1.05E+08	0.442	83	7-Octen-2-ol, 2,6-dimethyl-	018479-58-8	28299	NIST11.L
					74	2-Octanol, 2-methyl-6-methylene-	018479-59-9	28320	NIST11.L
3)	12.814	1514	2.59E+08	1.093	97	1,6-Octadien-3-ol, 3,7-dimethyl-	000078-70-6	26774	NIST11.L
4)	16.448	1688	76859118	0.324	99	Azulene, 1,2,3,5,6,7,8,8a-octahydro-1,4-dimethyl-7-(1-methylethenyl)-, [1S-(1.alpha.)]	003691-11-0	64530	NIST11.L
					99	.alpha.-Bulnesene	1000374-19-9	64321	NIST11.L
					83	.alpha.-Guaiene	003691-12-1	64292	NIST11.L
5)	17.225	1727	5.52E+08	2.327	96	Citronellol	000106-22-9	28222	NIST11.L
					96	6-Octen-1-ol, 3,7-dimethyl-, (R)-	001117-61-9	28325	NIST11.L
					70	2-Octen-1-ol, 3,7-dimethyl-	040607-48-5	28296	NIST11.L
8)	17.6	1747	1.04E+08	0.44	80	Nopyl acetate	000128-51-8	67677	NIST11.L
					64	Bicyclo[4.1.0]hept-3-ene, 7,7-dimethyl-3-vinyl-	113003-13-7	22846	NIST11.L
7)	18.704	1803	3.77E+08	1.587	94	Geraniol	000106-24-1	26608	NIST11.L
					72	Geranyl vinyl ether	1000132-11-4	45595	NIST11.L
					72	2,6-Octadien-1-ol, 3,7-dimethyl-, formate, (E)-	000105-86-2	47027	NIST11.L
8)	19.52	1846	6.32E+08	2.663	94	Phenylethyl Alcohol	000060-12-8	9920	NIST11.L
					70	Spiro[2,4]hepta-4,6-diene	000765-46-8	2472	NIST11.L



RIC

Research Institute
for Chromatography

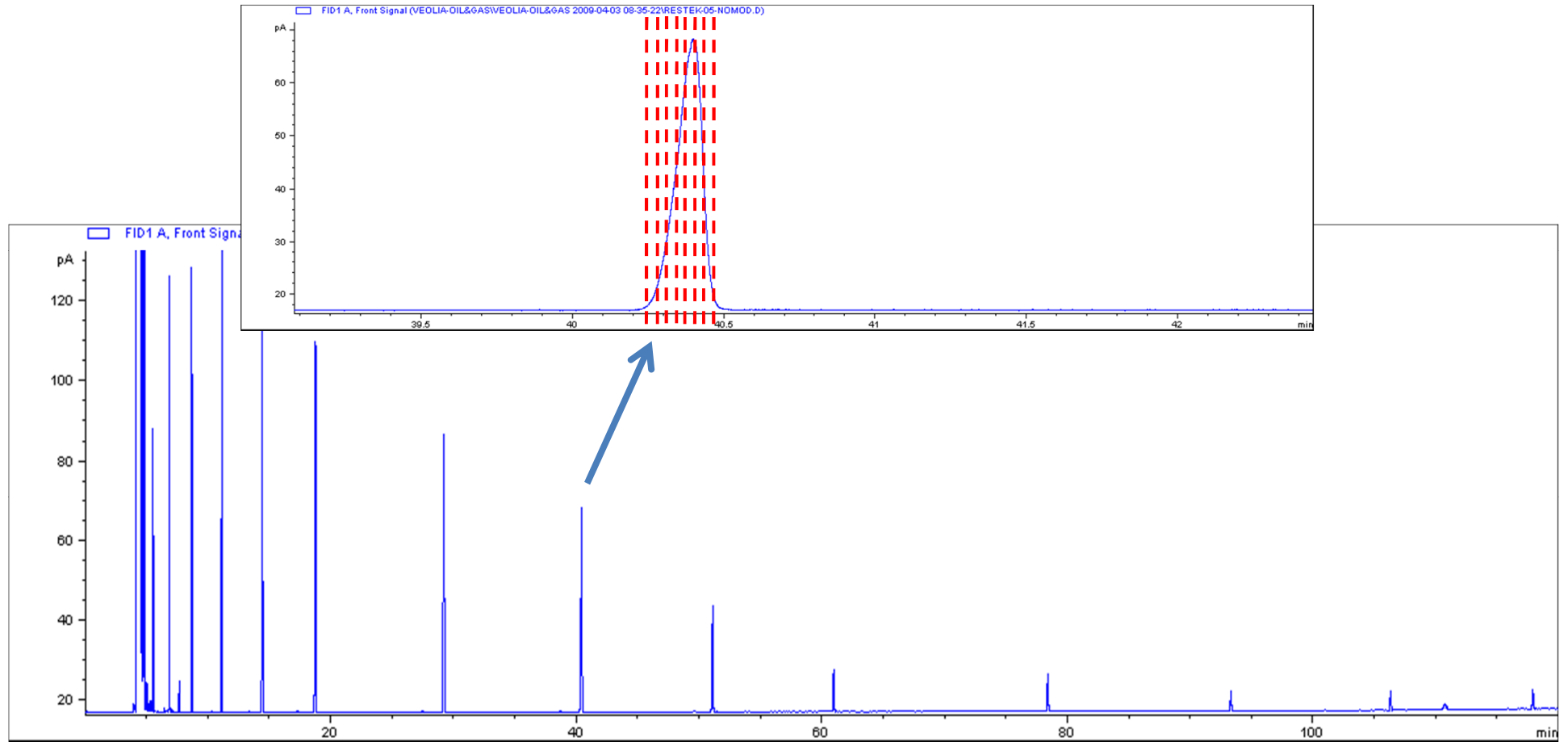
Comprehensive GCxGC Flow Modulator



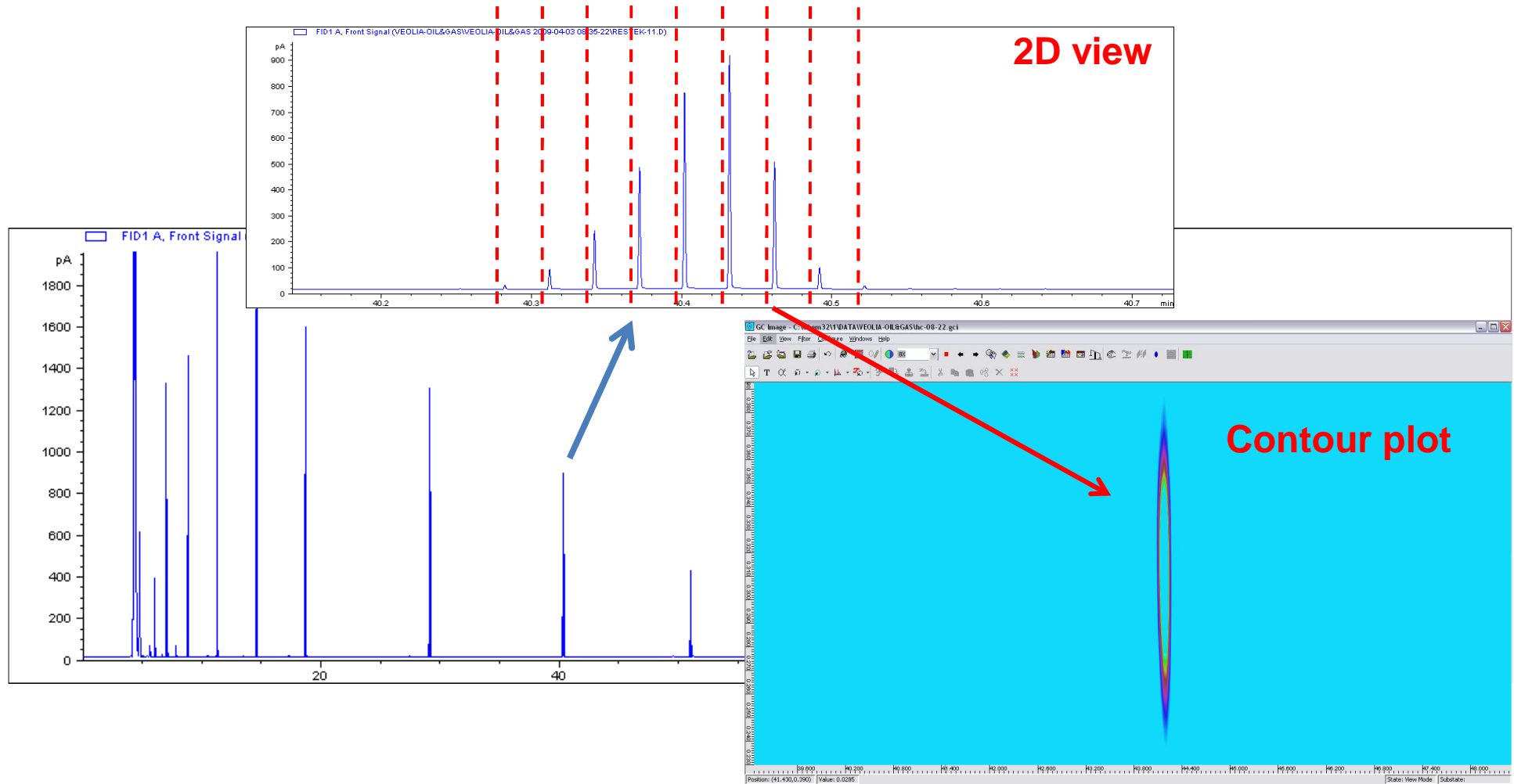
RIC

Research Institute
for Chromatography

1D analysis (ASTM HC mix)

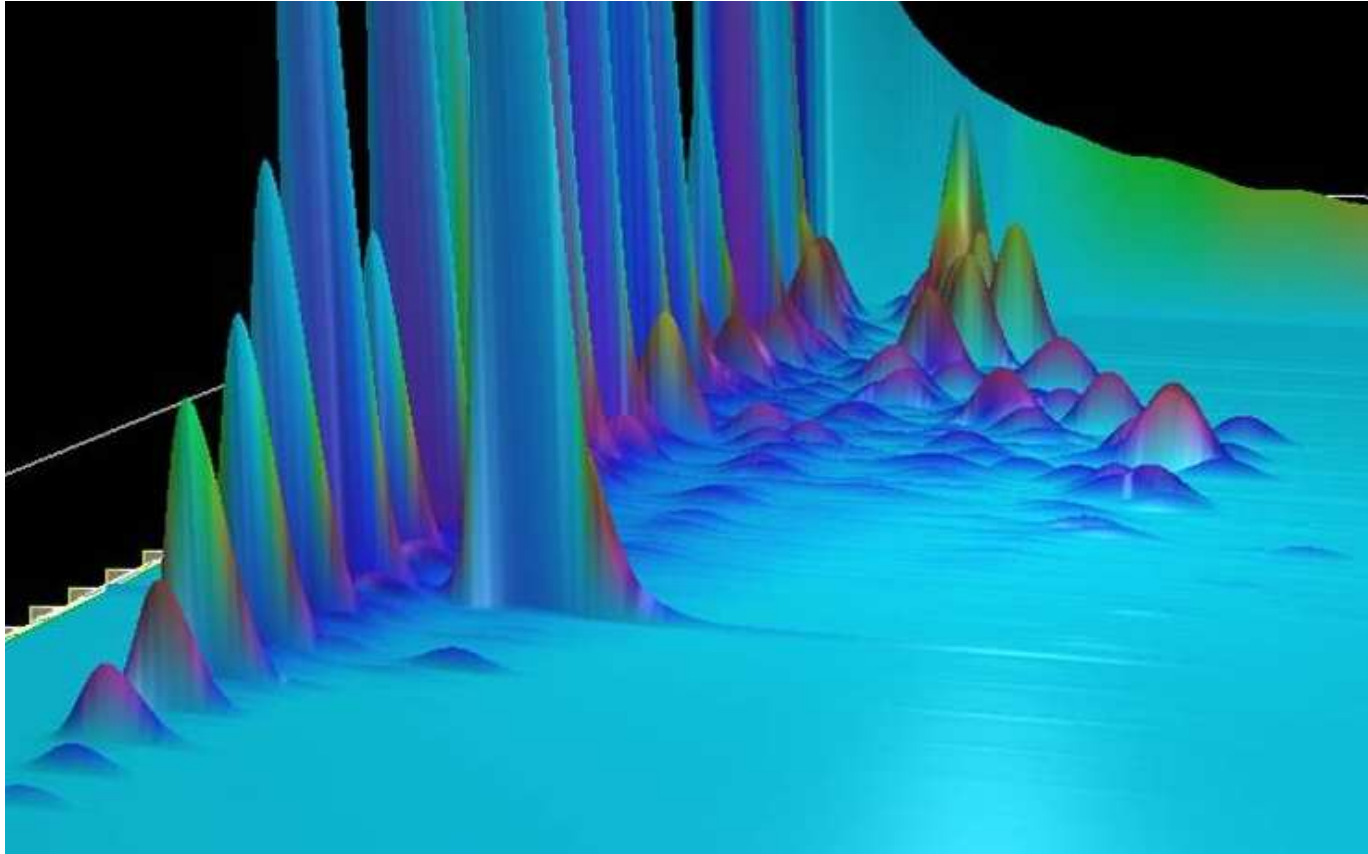


GCxGC modulated signal



Agilent G3486A CFI Modulator

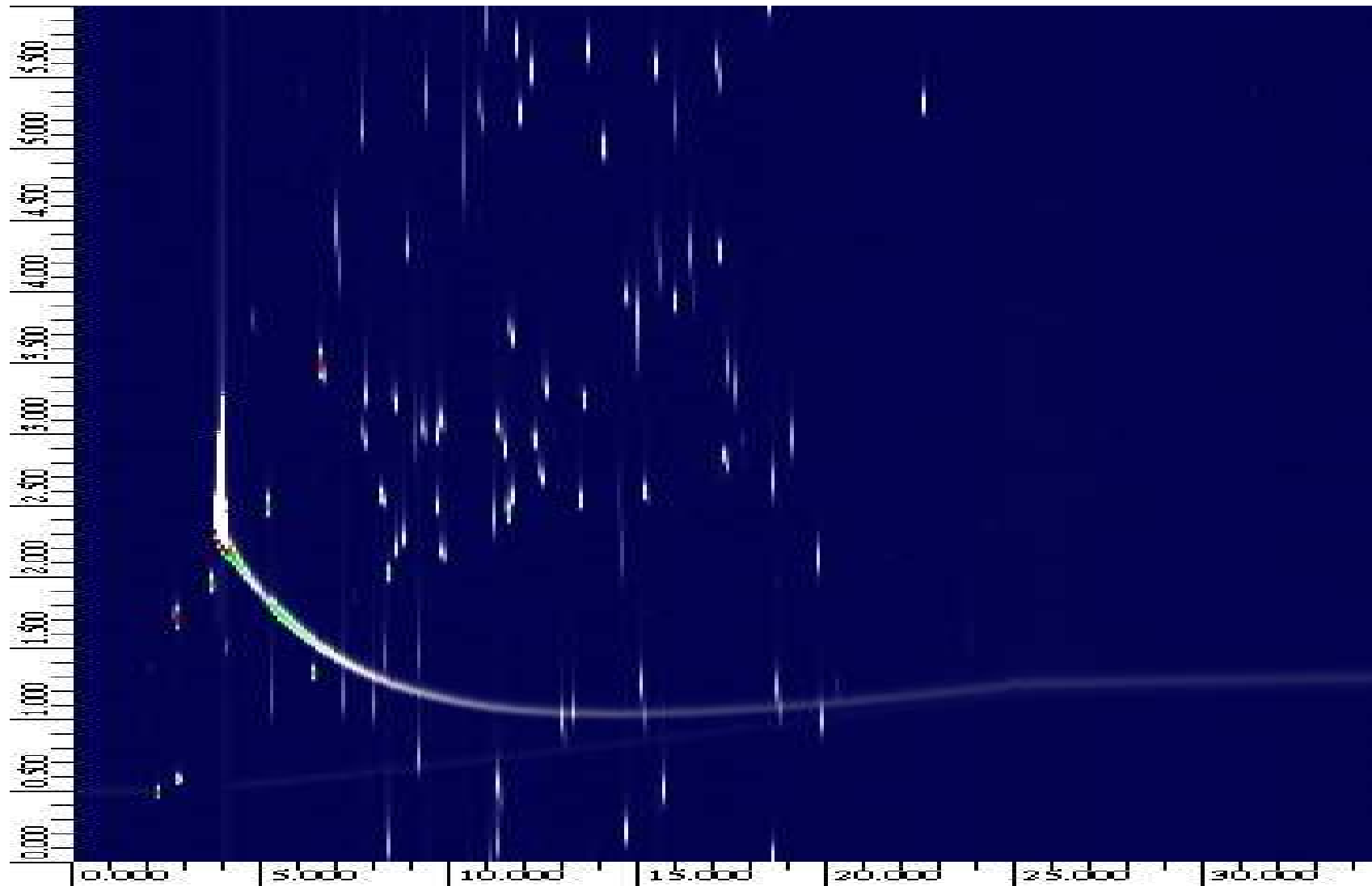
3D view



RIC

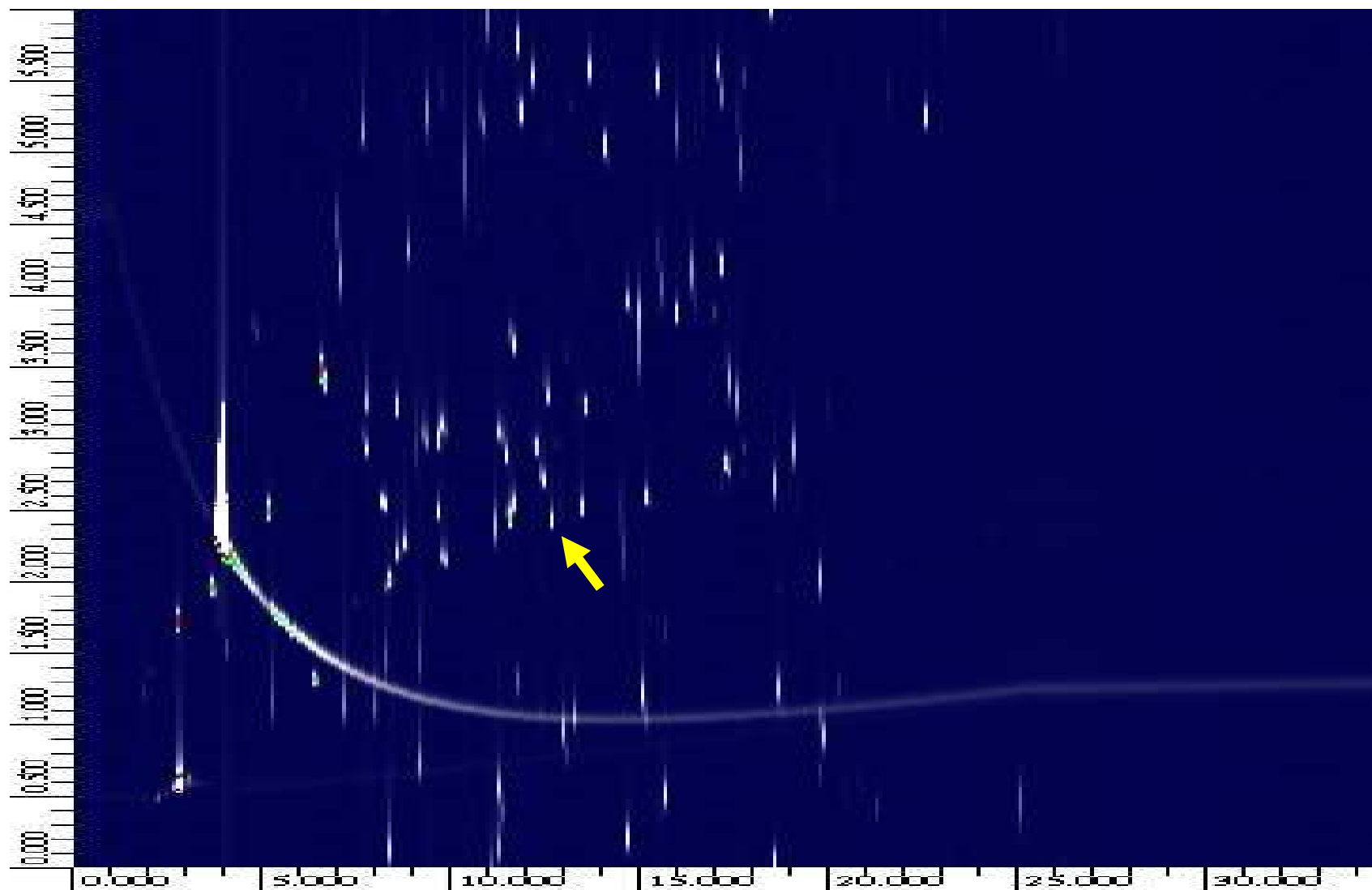
Research Institute
for Chromatography

GCxGC: Perfume - good quality



RIC | Research Institute
for Chromatography

GCxGC: Perfume with defect



RIC | Research Institute
for Chromatography

3. Detection

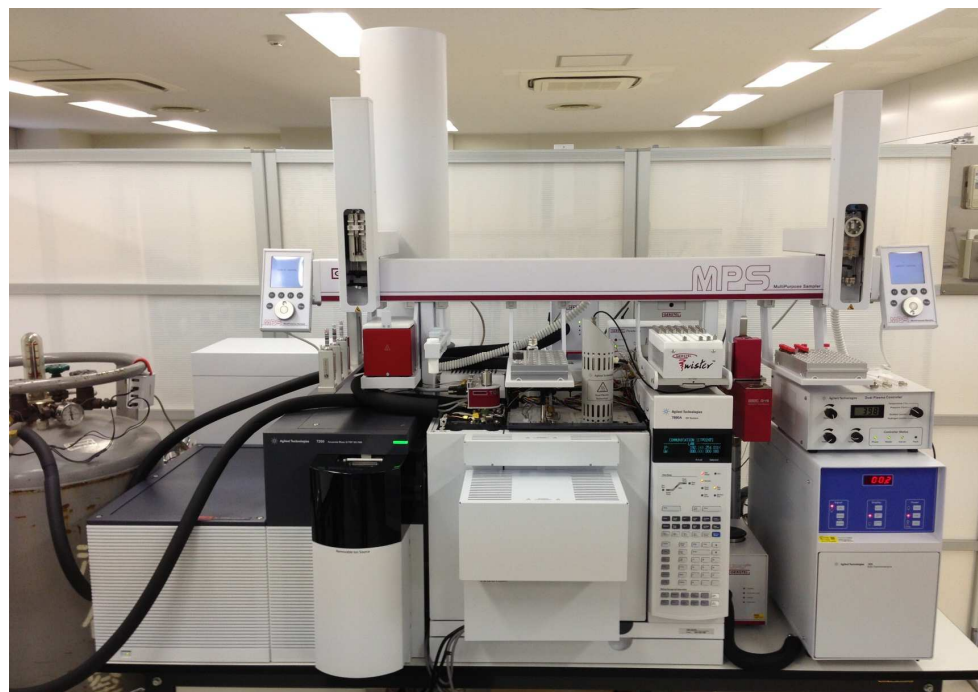


RIC

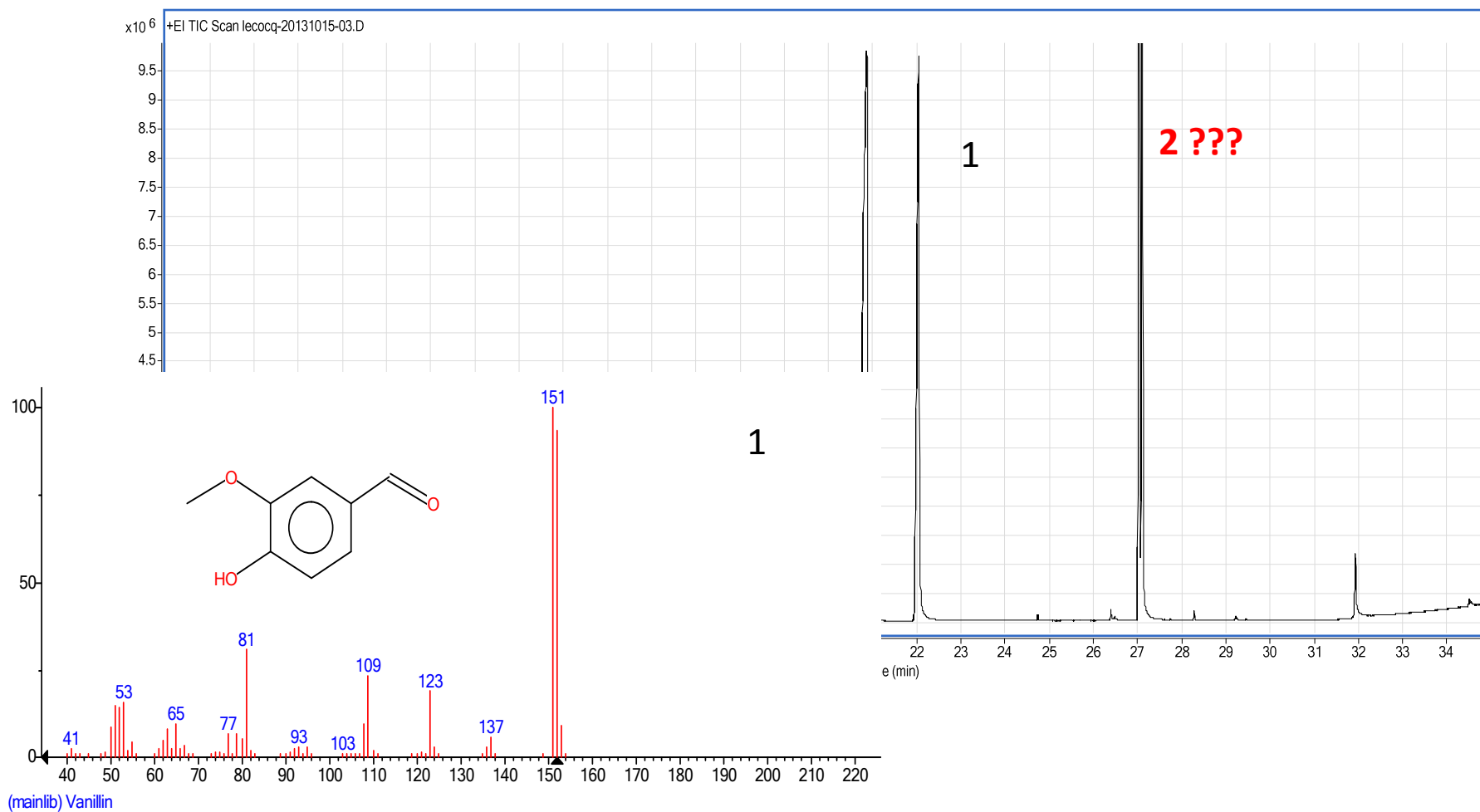
Research Institute
for Chromatography

Detection: Unleash the power of MS

- High sensitivity
- Accurate mass (TOF)
- EI, CI, APCI, SMB
- MS/MS



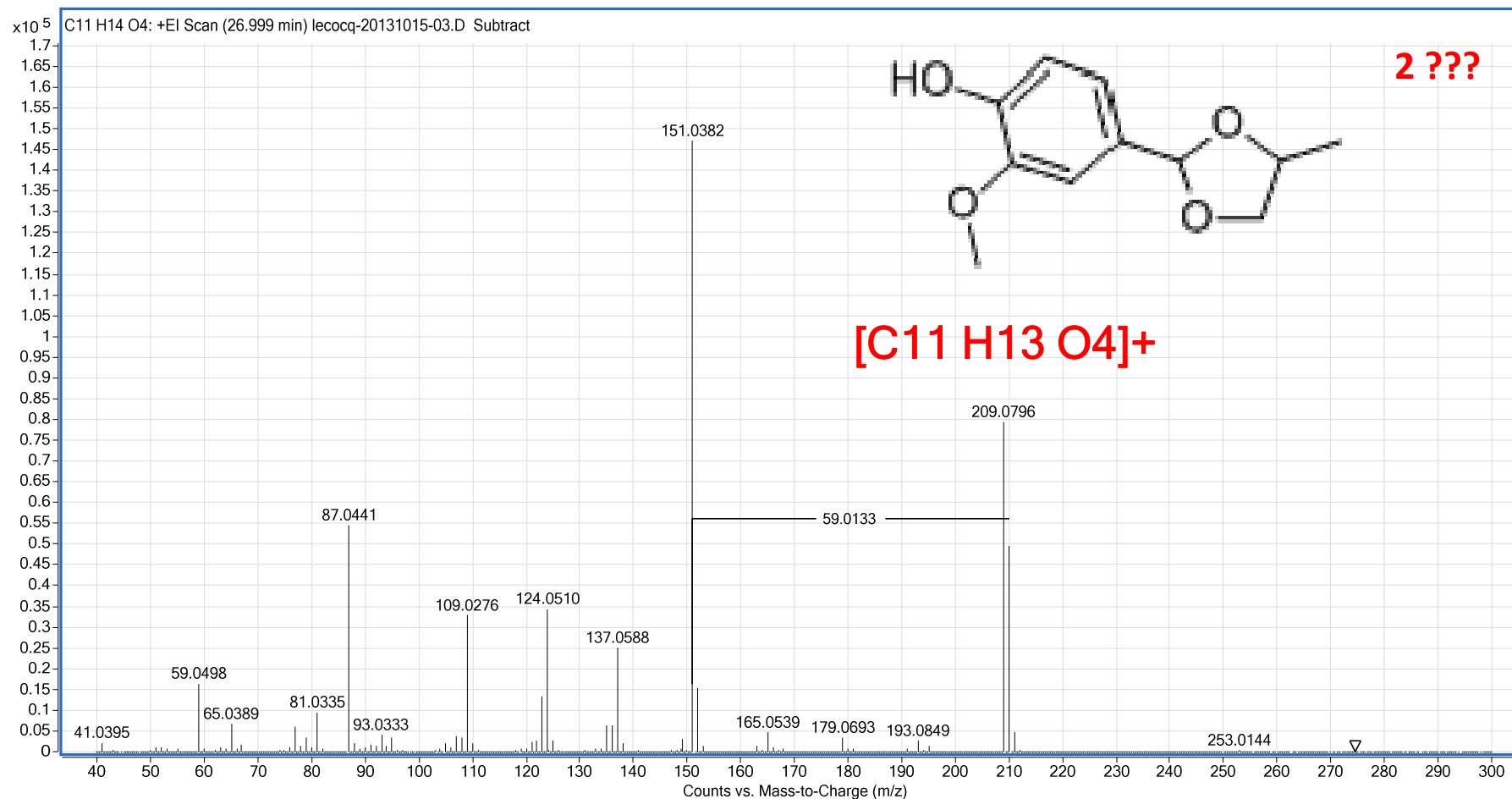
GC-Q-TOF: high resolution, high sensitivity



RIC

Research Institute
for Chromatography

GC-Q-TOF: high resolution, high sensitivity

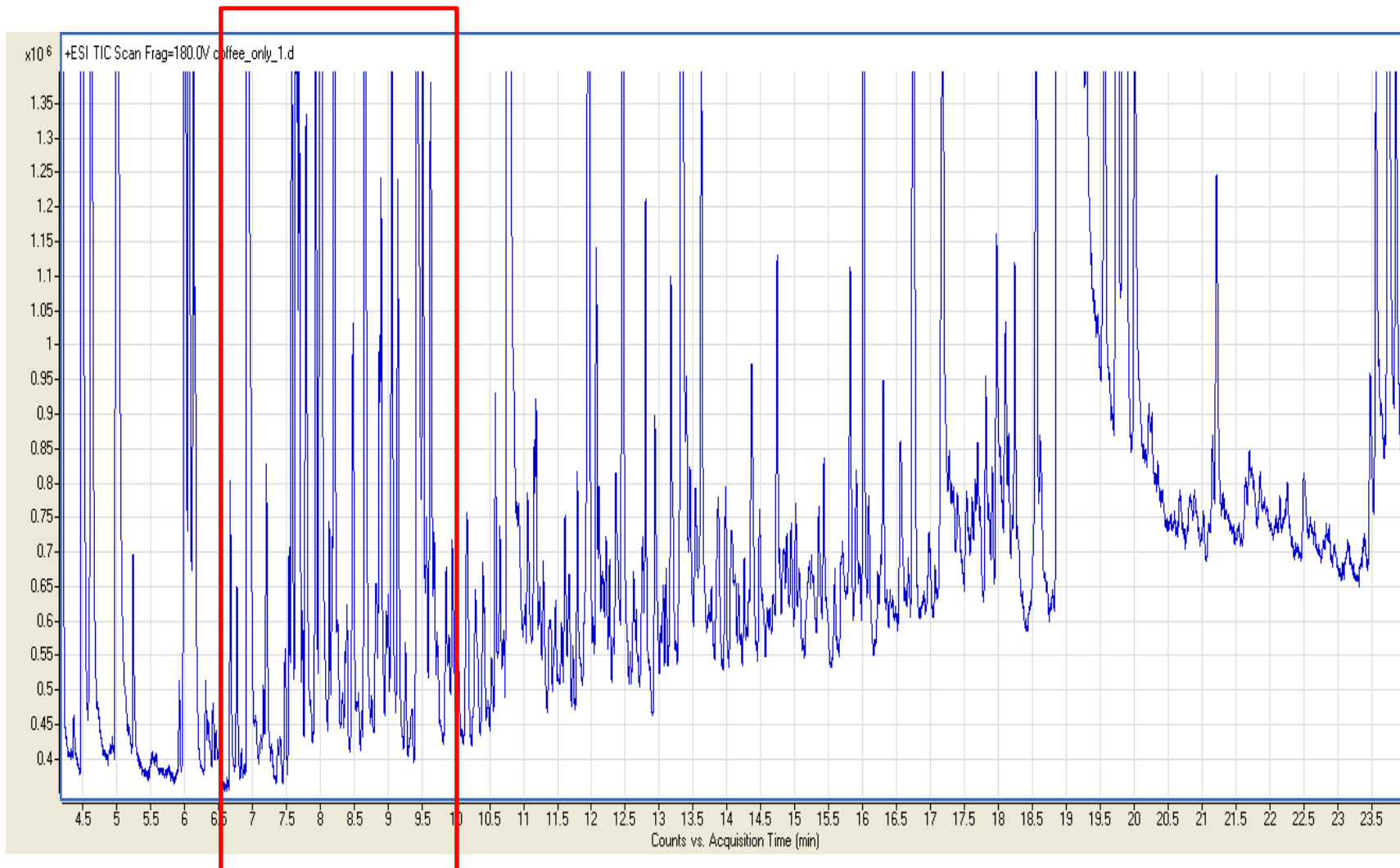


RIC

Research Institute
for Chromatography

GC-Q-TOF: high resolution, high sensitivity

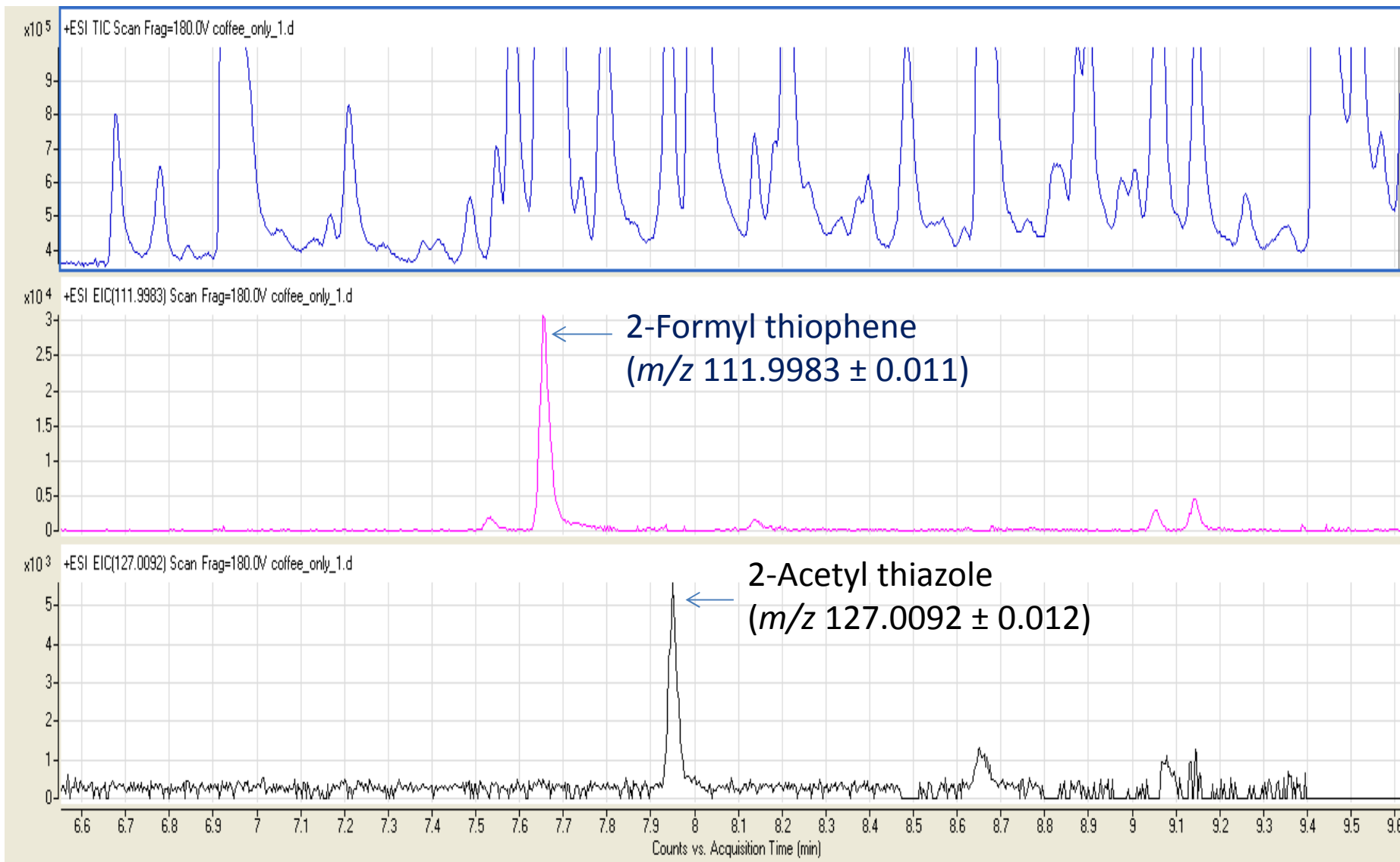
TIC of coffee extract (non-spike) (data from Gerstel KK)



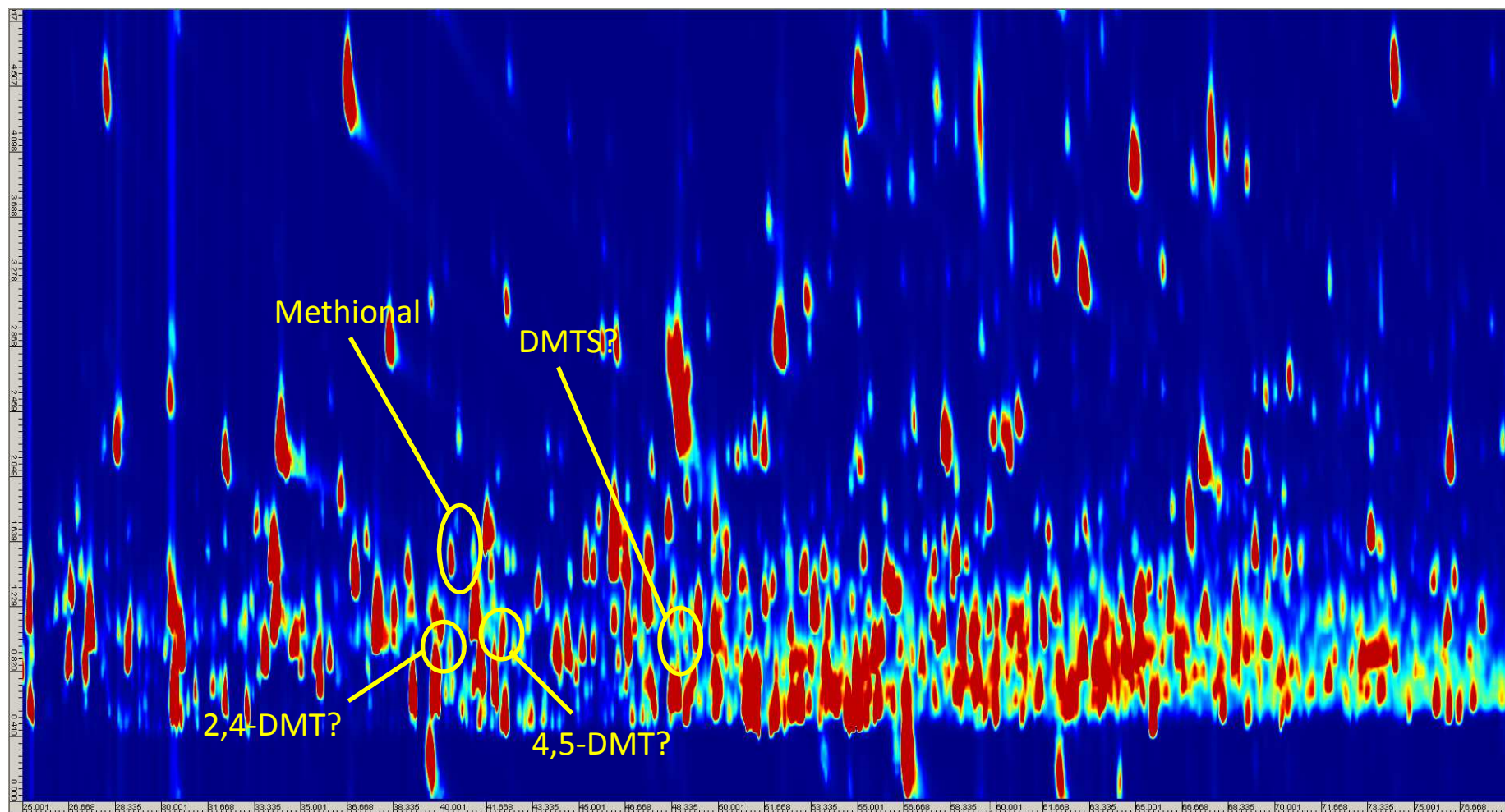
RIC

Research Institute
for Chromatography

TIC and mass chromatograms of coffee extract (non-spike)



GCxGC-QTOF-MS

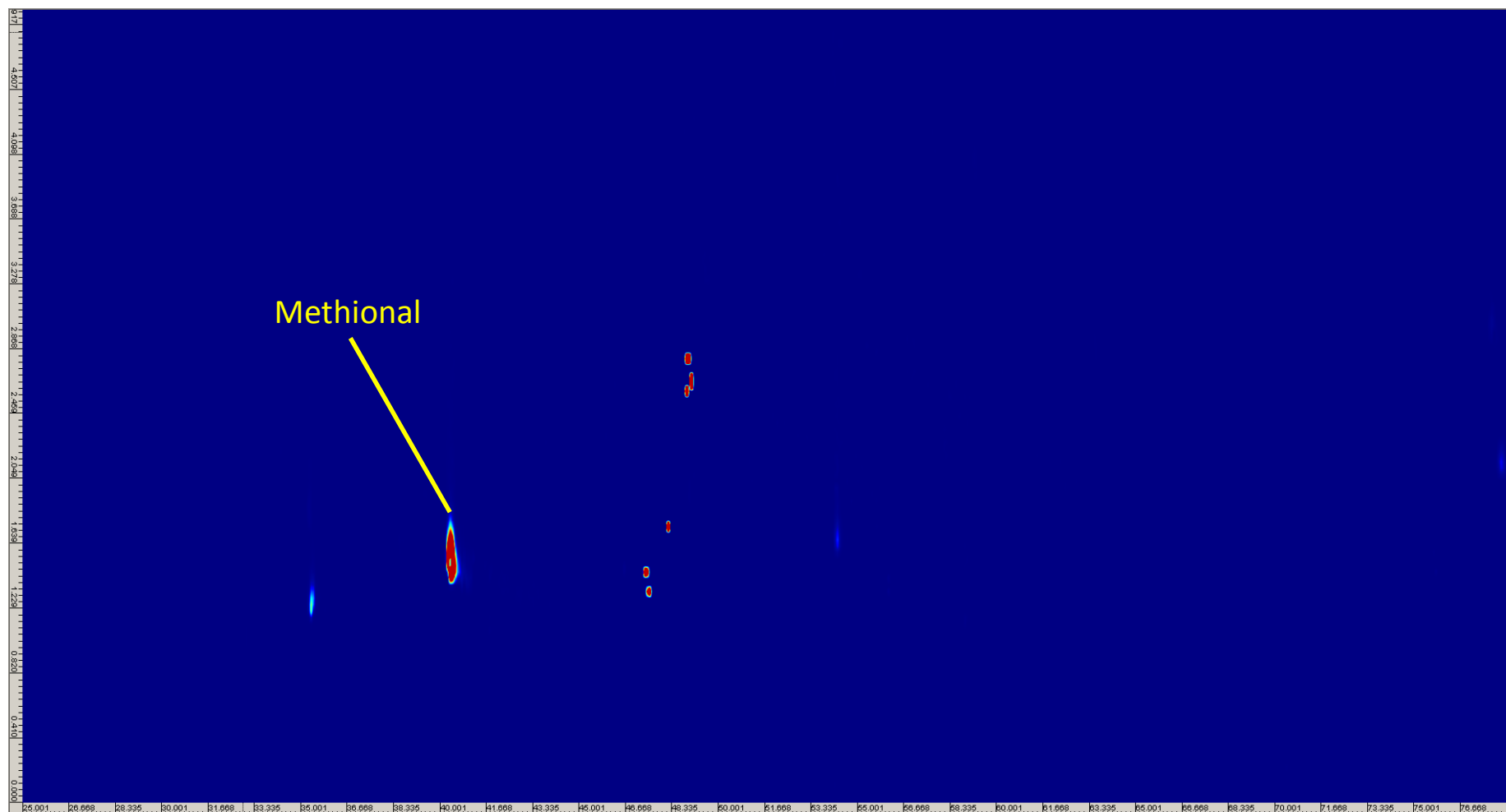


Tobacco smoke



RIC | Research Institute
for Chromatography

^2D mass chromatogram (m/z 104.0290 \pm 50 mDa)



RIC | Research Institute
for Chromatography

*Now we have: FEDHS-GC-GC-QTOF-MS and
FEDHS-GCxGC-QTOFMS*

What about Odor?



RIC

Research Institute
for Chromatography

SHS-E-nose

Objective approach

1 g, SHS @ 80°C (20 min)

Injection 2 mL

Alpha MOS Enose Fox 4000

Synthetic dry air: 150 mL/min

Acquisition time: 500 s

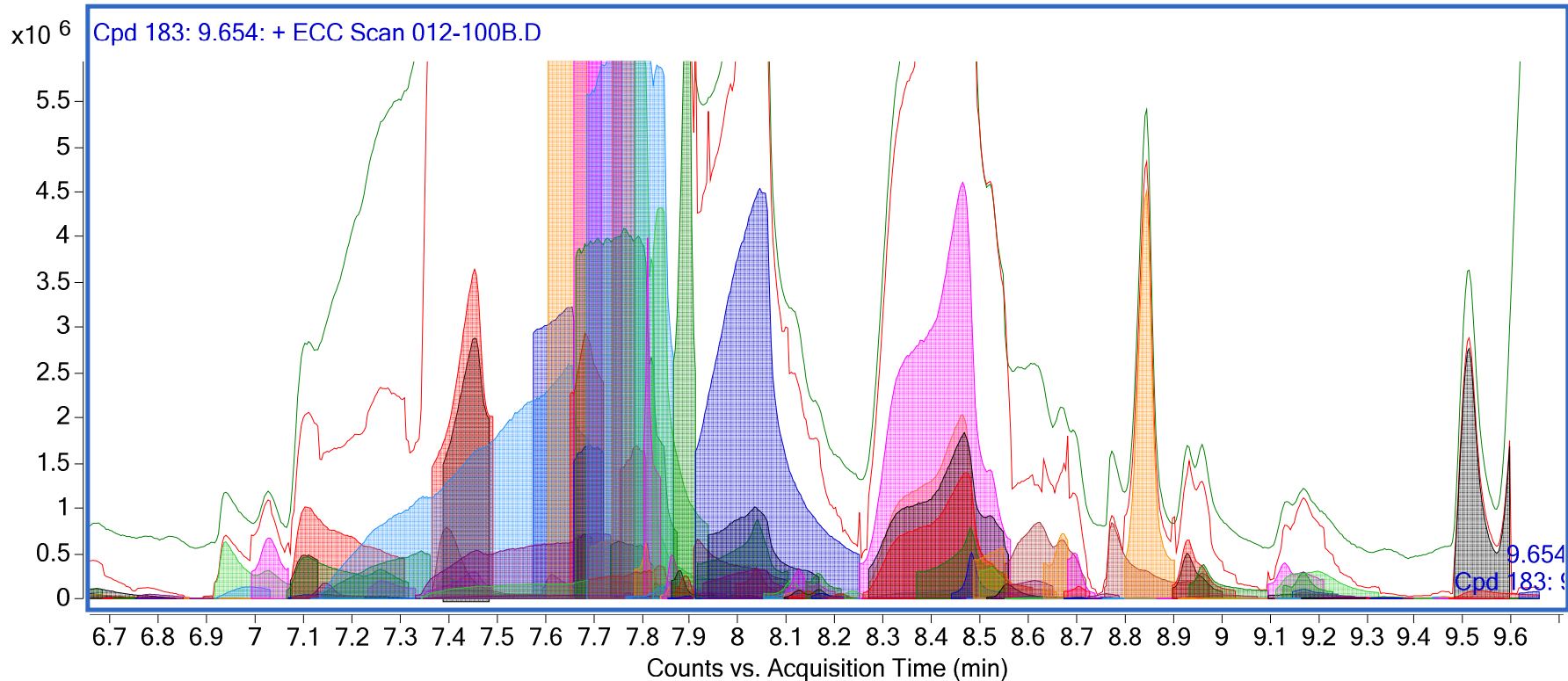
Acquisition period: 0.5 s



RIC

Research Institute
for Chromatography

GC-MS Deconvolution



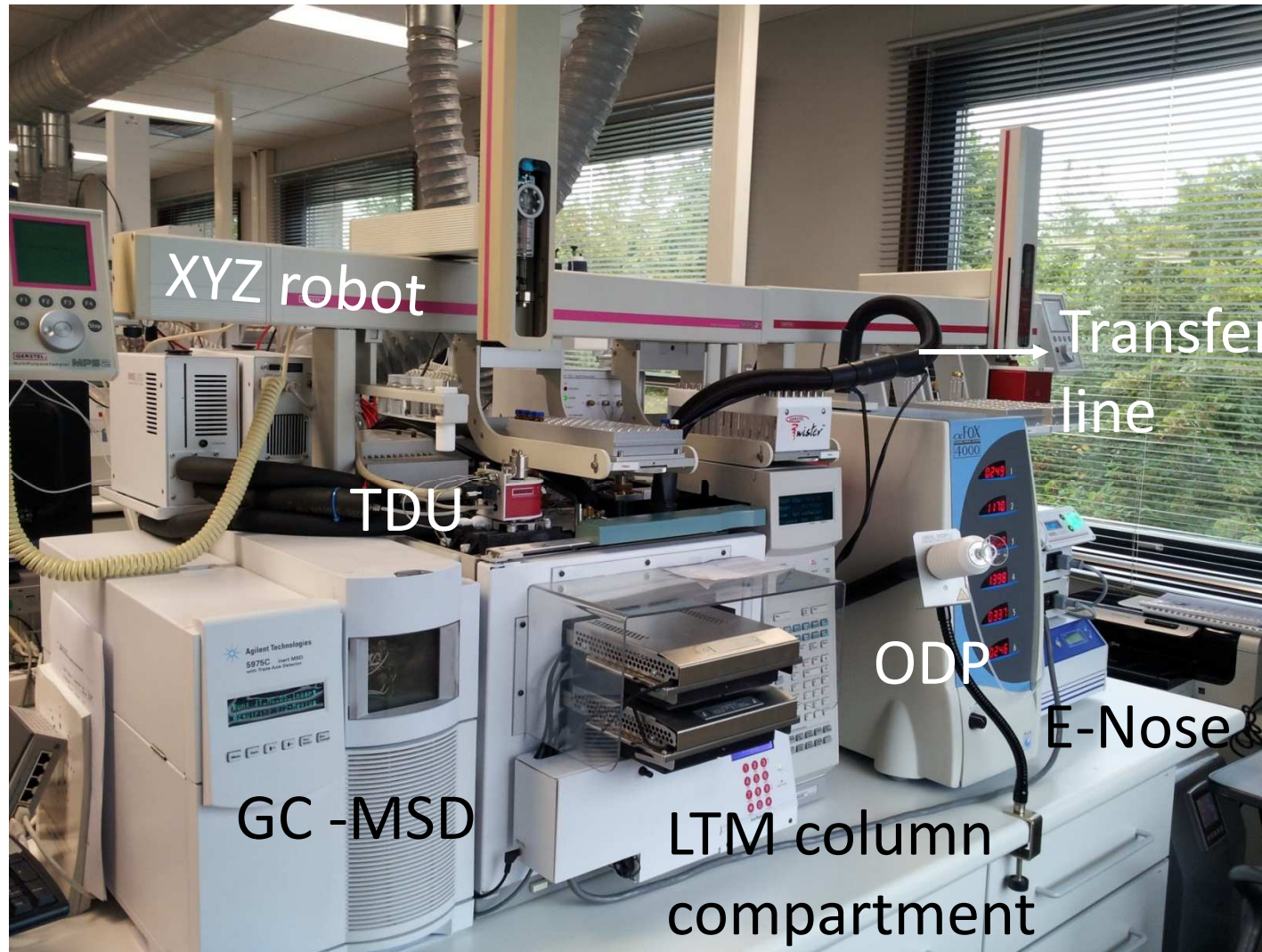
- 183 compounds in the 'odour' zone
- Concentration range 5 orders of magnitude



RIC

Research Institute
for Chromatography

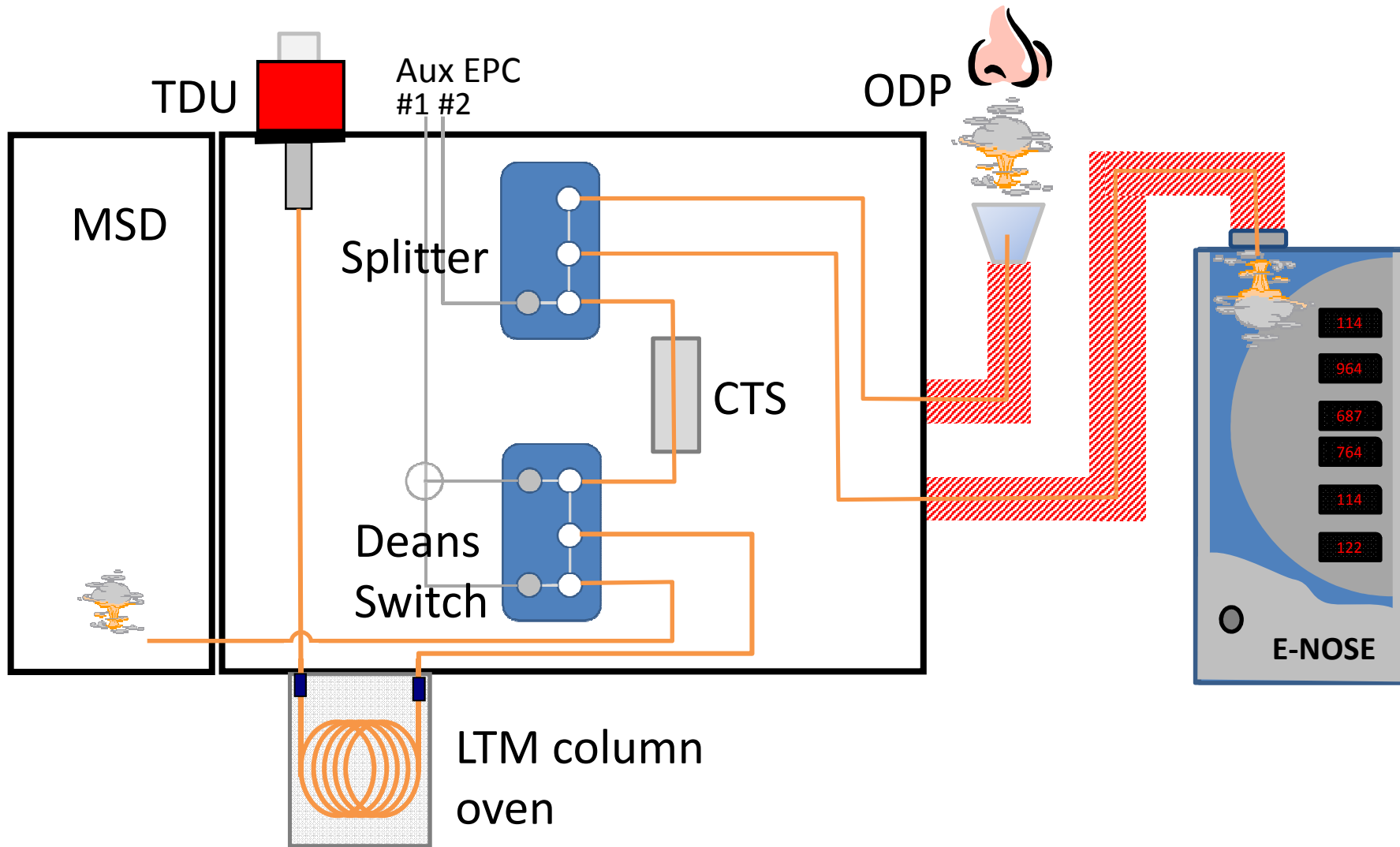
New GC-E-nose Configuration



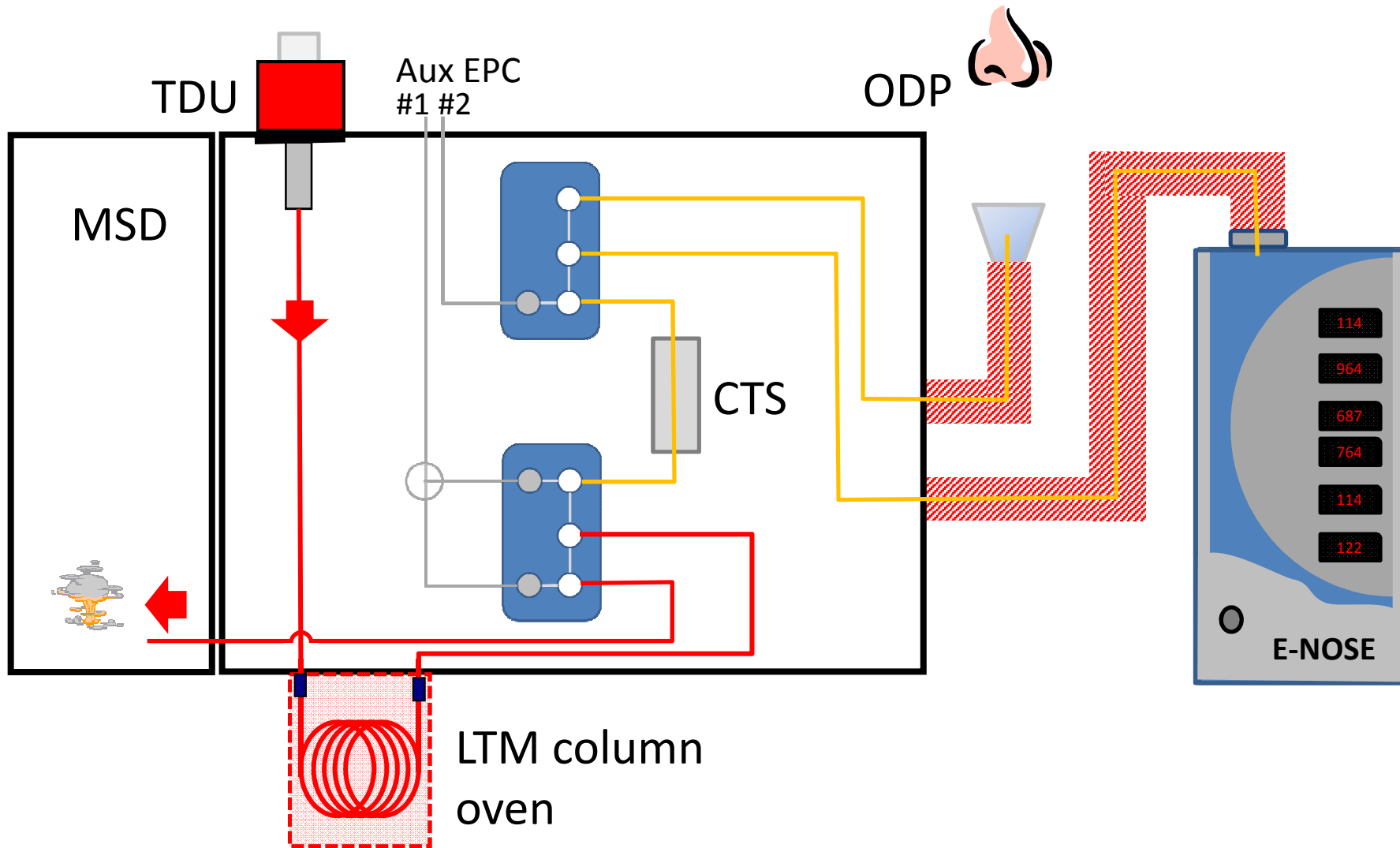
RIC

Research Institute
for Chromatography

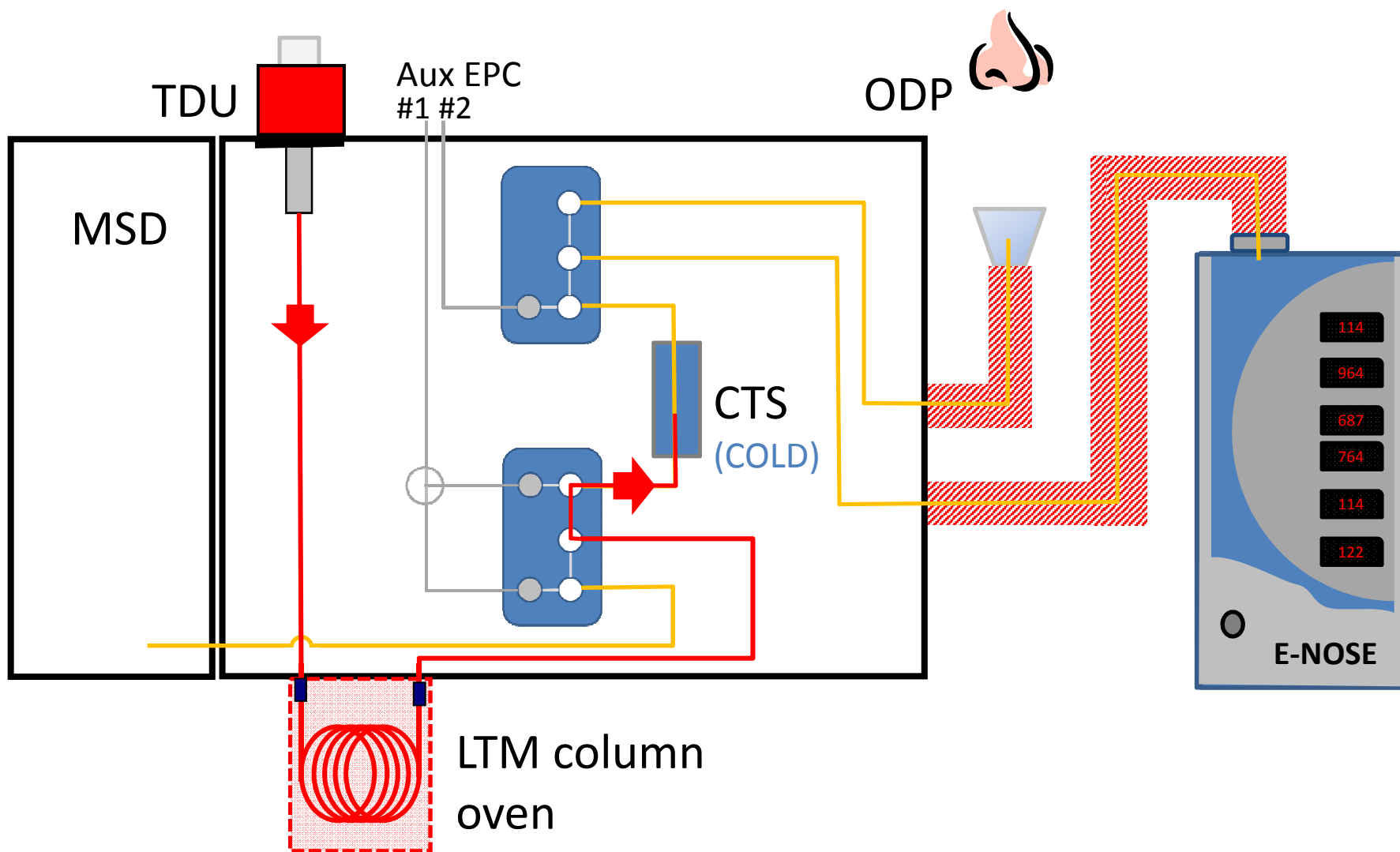
Configuration



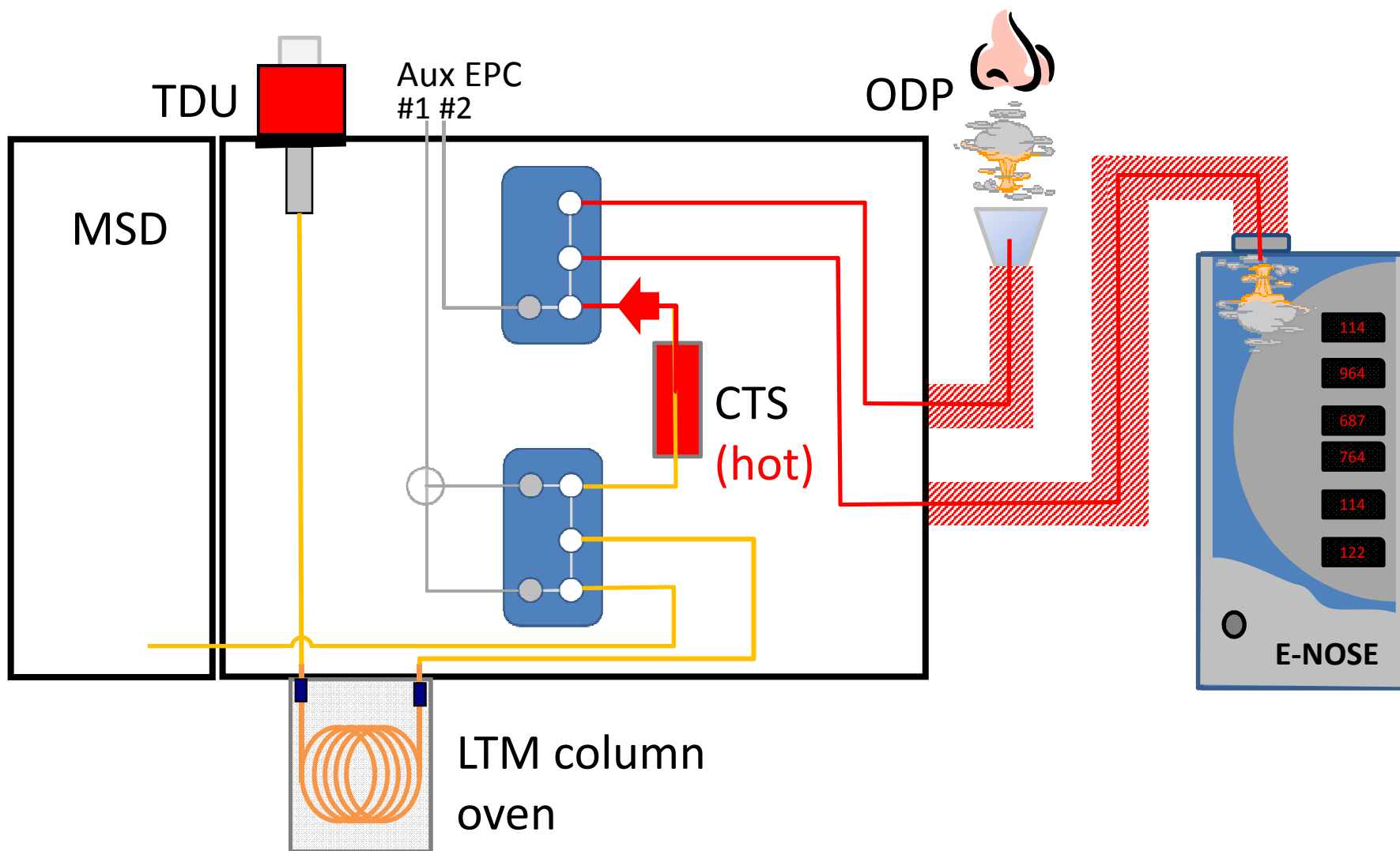
Step 1. GC-MS Analysis



Step 2. Collection of Fractions (heart-cut)

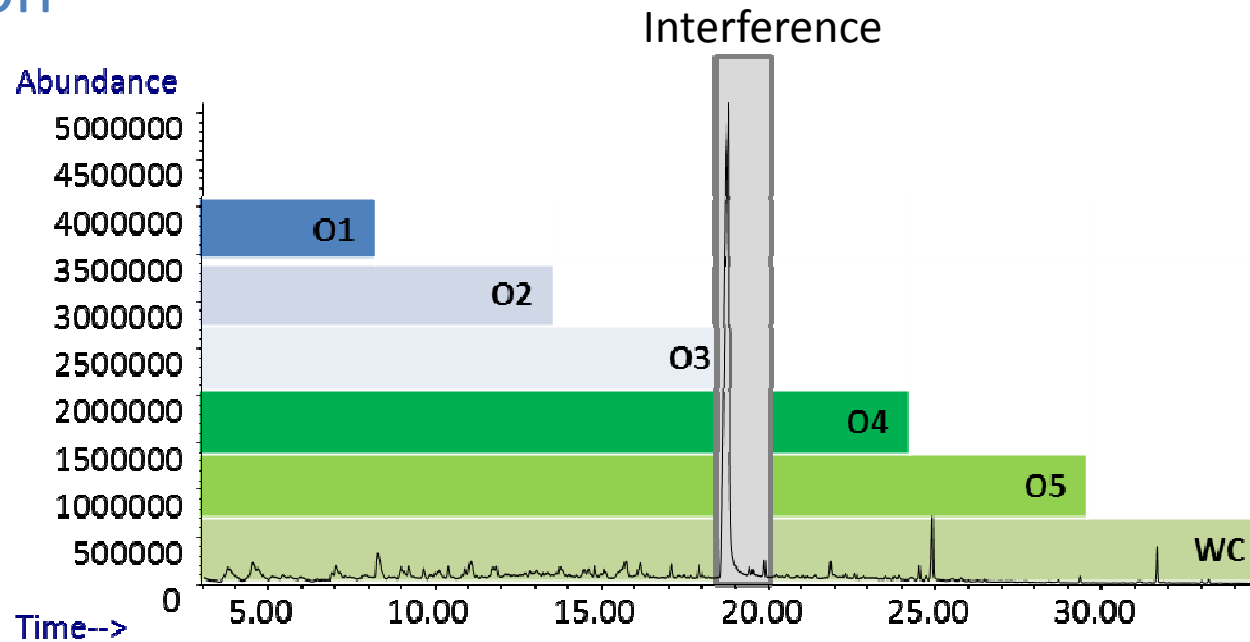


Step 3. Transfer of Fractions to Noses



2 Operation modes

1. Omission



2. Addition

- Perceptual interaction (heart-cut)
- Recombination



RIC

Research Institute
for Chromatography

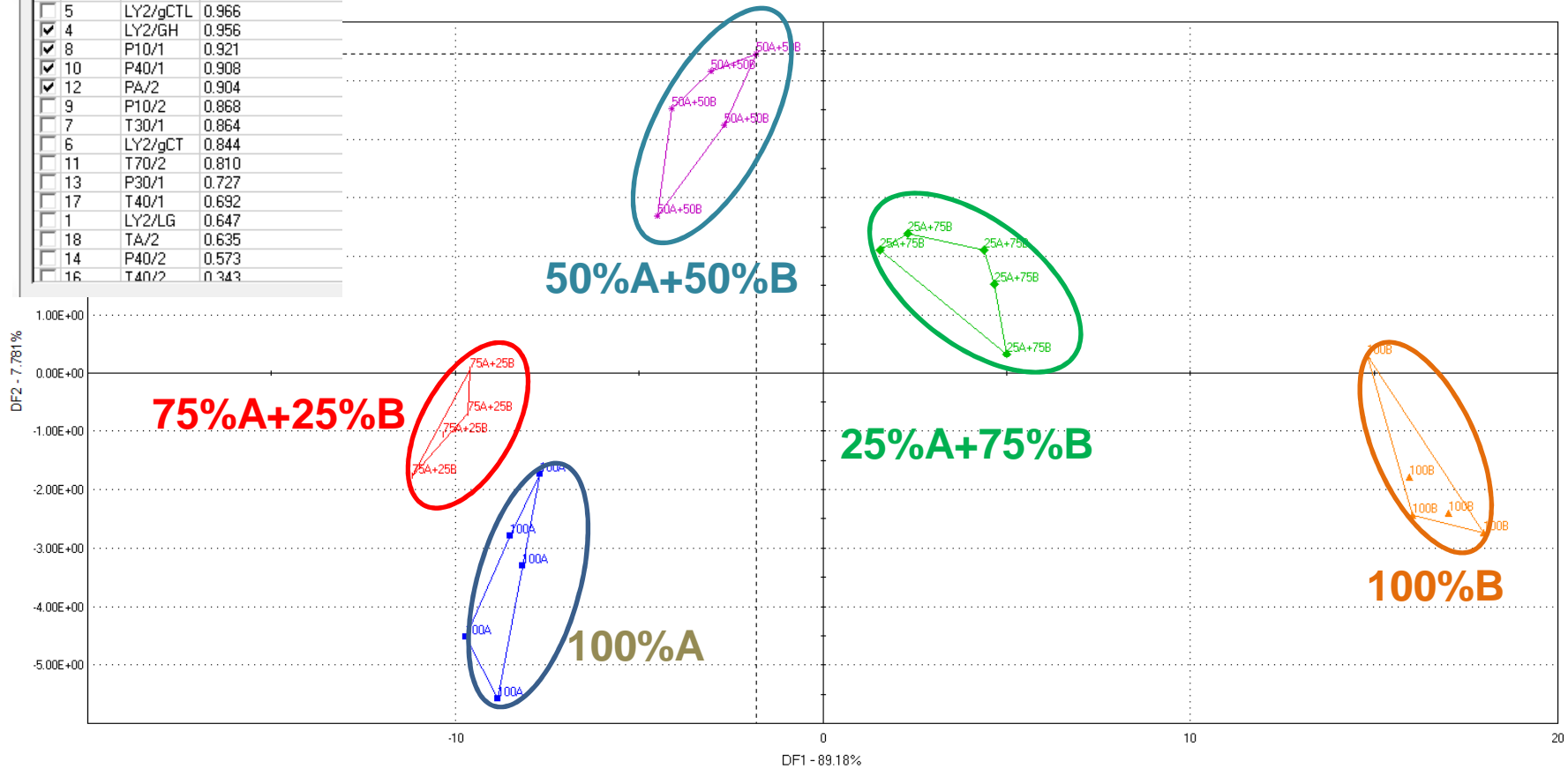
Discriminant Function Analysis (DFA) for Σ (HC)

6 Sensors

Sensor selection

Index	Sensors	Discrimination power	
<input checked="" type="checkbox"/>	2	LY2/G	0.970
<input checked="" type="checkbox"/>	3	LY2/AA	0.970
<input type="checkbox"/>	5	LY2/gCTL	0.966
<input checked="" type="checkbox"/>	4	LY2/GH	0.956
<input checked="" type="checkbox"/>	8	P10/1	0.921
<input checked="" type="checkbox"/>	10	P40/1	0.908
<input checked="" type="checkbox"/>	12	PA/2	0.904
<input type="checkbox"/>	9	P10/2	0.868
<input type="checkbox"/>	7	T30/1	0.864
<input type="checkbox"/>	6	LY2/gCT	0.844
<input type="checkbox"/>	11	T70/2	0.810
<input type="checkbox"/>	13	P30/1	0.727
<input type="checkbox"/>	17	T40/1	0.692
<input type="checkbox"/>	1	LY2/LG	0.647
<input type="checkbox"/>	18	TA/2	0.635
<input type="checkbox"/>	14	P40/2	0.573
<input type="checkbox"/>	16	T40/2	0.343

Validation Score = 96



... omics?

Discovering the difference in your data



RIC

Research Institute
for Chromatography

Mass Profiler Professional software

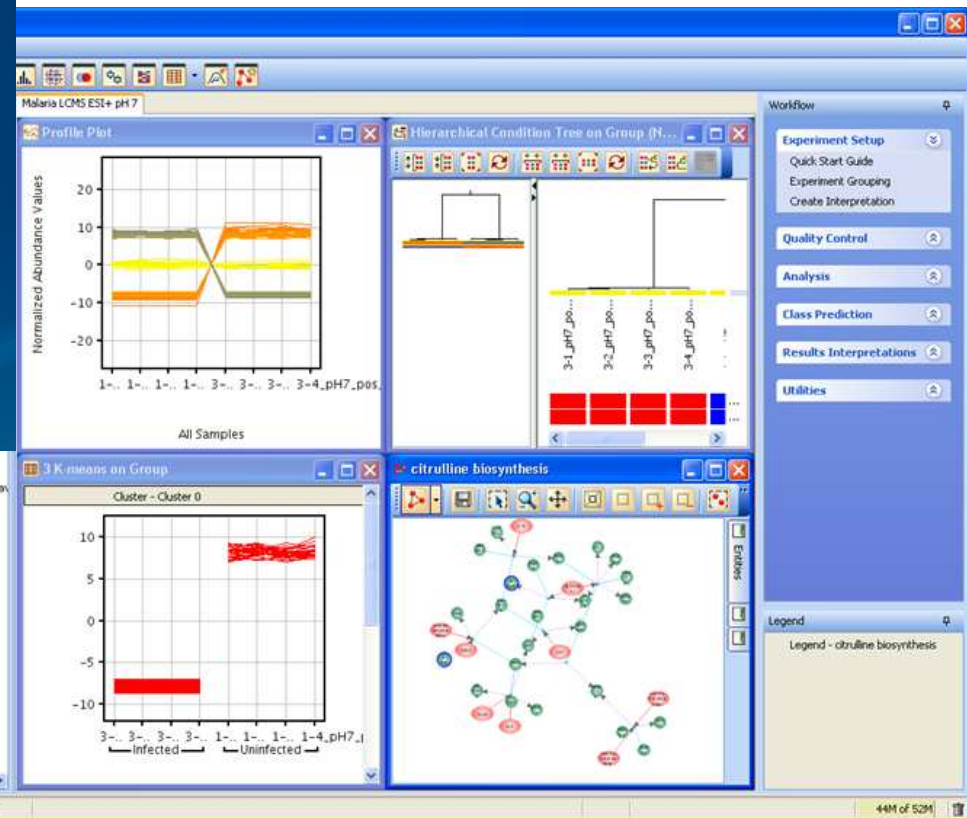
MassHunter
Mass Profiler
Professional
Software

MPP 12

Version B.12.00

© Agilent Technologies, Inc. 2012

www.agilent.com/chem



RIC

Research Institute
for Chromatography

Mass Profiler Professional software

- Analyze complex MS data (GC-MS, LC-MS, CE-MS, ICP-MS)
- Compare and (classify) sample groups
- Identify differences between samples
- *Untargeted*
- Application areas: proteomics, metabolomics, food safety, environmental, forensics, toxicology, ...

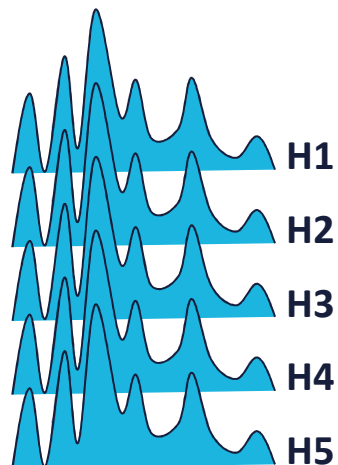


RIC

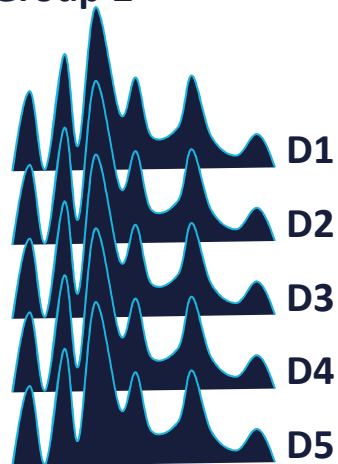
Research Institute
for Chromatography

Workflow: from chromatograms to potential markers

Group 1



Group 2



A diagram illustrating feature extraction for Group 1. It shows a hierarchy of peaks: H1, H2, H3, H4, and H5. Below each peak, there are smaller peaks representing metabolites. A table summarizes the data:

Metabolite	Value
Metab1	10000
Metab2	5000
Metab3	20000
Metab4	7500
Metab5	1000
Metab6	10000

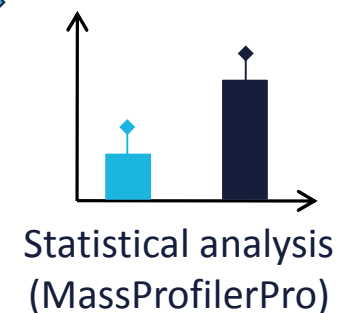
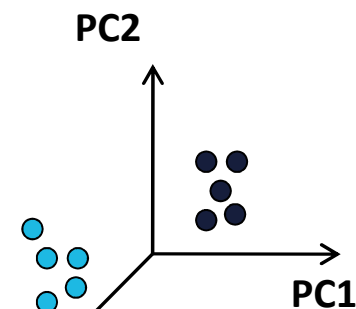
A diagram illustrating feature extraction for Group 2. It shows a hierarchy of peaks: D1, D2, D3, D4, and D5. Below each peak, there are smaller peaks representing metabolites. A table summarizes the data:

Metabolite	Value
Metab1	10000
Metab2	5000
Metab3	20000
Metab4	7500
Metab5	1000
Metab6	10000

A diagram illustrating alignment, normalization, and filtering. It shows a heatmap where the columns represent patients (Patient 1 to Patient 10) and the rows represent metabolites (Metab 1 to Metab 6). The cells are colored in shades of blue and dark blue, representing the intensity of each metabolite in each patient.

	Patient 1	Patient 2	Patient 3	Patient 4	Patient 5	Patient 6	Patient 7	Patient 8	Patient 9	Patient 10
Metab 1	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue	Dark Blue
Metab 2	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue
Metab 3	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue
Metab 4	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue
Metab 5	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue
Metab 6	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue	Light Blue

Alignment, normalization and filtering (MassProfilerPro)



Feature extraction (Mass Hunter, AMDIS, XCMS, MFE)

ANOVA, PCA, t-tests, volcano plots, hierarchical trees, SOMs, QT clustering, and SVMs for class prediction



RIC | Research Institute for Chromatography

Conclusions

- volatility range
- Polarity range
- Complex samples
- matrix interferences

DHS (LVDHS)

SBSE + variation

2D-GC

2D + MS



RIC

Research Institute
for Chromatography