


ESSENTIAL SEPARATIONS:

ANALYSIS OF AROMA OILS BY ONE-DIMENSIONAL AND TWO-DIMENSIONAL GAS CHROMATOGRAPHY AND MASS SPECTROMETRY

Multidimensional Chromatography Workshop
Los Angeles, CA January 2024

Robert (Chip) Cody



JEOL mass spectrometers used for this study



Q1600GC UltraQuad™ SQ-Zeta
GC-MS (Single Quadrupole)
Ionization used: EI, CI



TQ2000GC AccuTOF GC-Alpha
GC-HRTOFMS
GCxGC-HRTOFMS
Ionization used: EI, CI, FI



1. JMS-Q1600GC UltraQuad™ SQ-Zeta



High-precision Hyperbolic-shaped Quadrupole

- Superb ion transmission enhances high sensitivity
- Reduces ion-quenching when introducing a large volume of ions, allowing a wide dynamic range

Pre-filter

- Suppresses the degradation of resolution and sensitivity due to contamination of the quadrupole

Dual Filament

- Reduces instrument down-time during filament exchange

Draw-in Lens

- Reduces diffusion of ions from the quadrupole outlet and reduces chemical noises
- Improved sensitivity by re-acceleration of ions

Secondary Electron Multiplier Detector

- Detector is resistant to degradation in an atmospheric environment to maintain high performance over a long period of time

Split Flow Turbo Molecular Pump

- Quick evacuation after column replacement and maintenance work
- Use of mega bore, packed column is possible
- Compatible with the latest Low-Pressure GC (LPGC) column

No tools required
Easy maintenance!

High pumping
capacity!





2: GC-MS High-Resolution Time-of Flight Mass Spectrometer: AccuTOF™ GC-Alpha

- Simple ion optics
- High mass resolving power (>30,000)
- High mass accuracy (sub-mmu)
- High speed (50 Hz) -- compatible with GCxGC
- Wide mass range (m/z 4 up to m/z 6400)
- All ion sources and direct sample introduction methods
- Field Ionization (FI) and Field Desorption (FD) options
- Combination ion sources (EI/FI and EI/PI)
- Easy ion source exchanges with very fast pump-down
- The most powerful data analysis software available



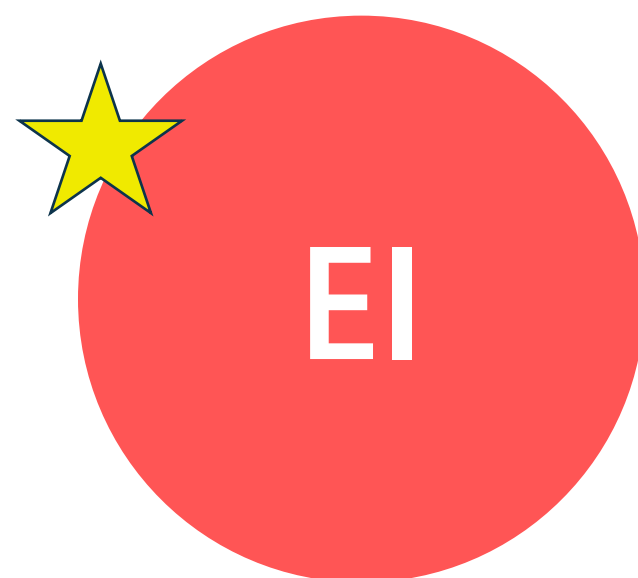
Dual-stage reflectron



Ionization Methods for GC-MS



Used in this talk



Electron Ionization

- Hard ionization
- Good for structural analysis using the generated fragment ions
- Able to change the ionization energy, typically use 70eV



Chemical Ionization

- Soft ionization
- Effective for compounds with higher proton affinities
- Negative-ion mode is available
- Reagent gas required

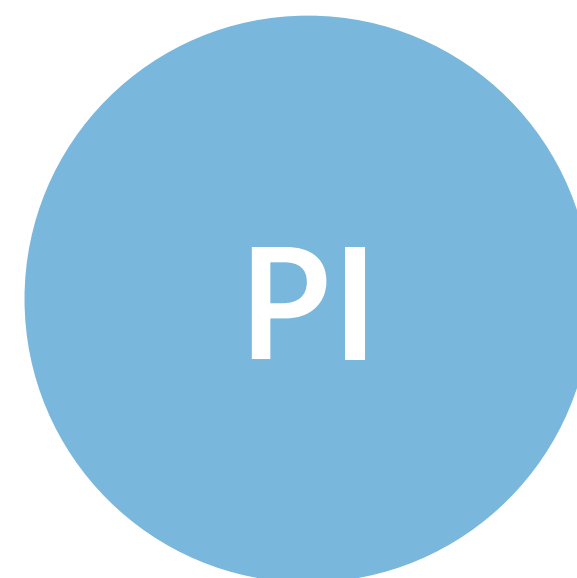
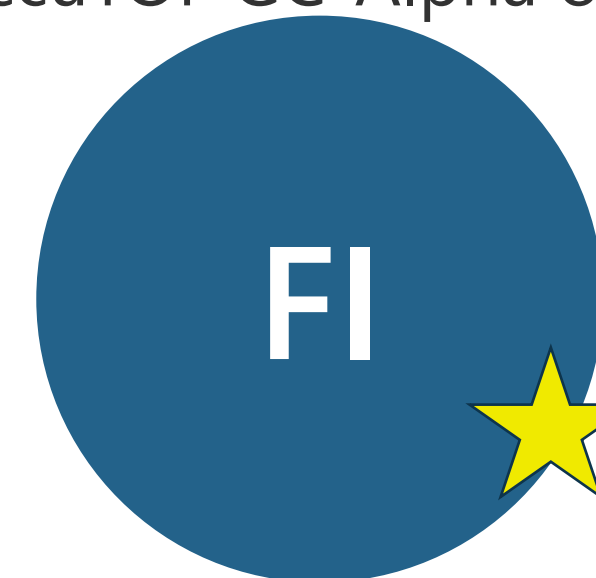


Photo Ionization

- Soft ionization
- Ionization energy is about 10.3eV (D2 lamp)
- Good for aromatic compounds
- No reagent gas required

AccuTOF GC-Alpha only



Field Ionization

- Softest ionization
- Energies imparted to the molecules are less than 1eV
- Effective even for lower polarity molecules
- No reagent gas required



Software used

- Data acquisition:
 - JEOL msPrimo (single quadrupole)
 - JEOL msAxel (AccuTOF GC-Alpha)
- GC-MS data analysis with deconvolution
 - msFineAnalysis iQ (single quadrupole)
 - msFineAnalysis AI (AccuTOF GC-Alpha)
- GC x GC data analysis
 - GC Image
 - AnalyzerPro XD (SpectralWorks Ltd.)
 - *GC-MS, GCxGC-MS & Direct analysis MS*

- Databases
 - NIST23 main and replicate libraries
 - Wiley main and replicate libraries
 - Wiley Food, Flavors, Fragrances, and Related Compounds
 - JEOL AI for compounds not in experimental databases
- NIST Hybrid Search



JEOL 1D GC data analysis with chromatographic deconvolution

msFineAnalysis iQ (Single quadrupole)

- Deconvolution
- Correlate EI with CI or PI
- Database search
- RI match
- Isotope match
- Variance analysis

msFineAnalysis AI (HRTOF)

- Deconvolution
- Correlate EI and soft ionization (EI, CI, PI, FI)
- Database search
- **Elemental composition**
 - **Molecular ion and fragment coverage**
- RI match
- Isotope match
- Variance analysis
- **Structure analysis with AI database**



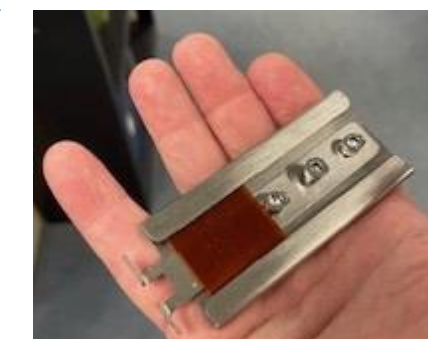
GC conditions

- GC-MS

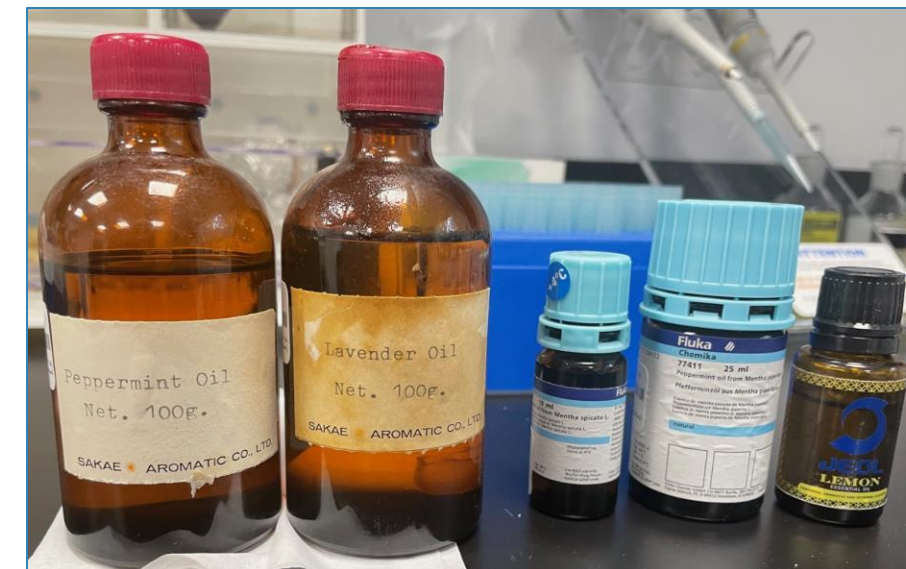
- Agilent 8890 GC
- 30m DB5-MS column (single quadrupole)
- GC x GC column set used in 1D GC mode for the AccuTOF GC-Alpha data
- Split injection: 200:1
- Oven:
 1. 50 °C (0 min)
 2. 5°C min⁻¹ to 220°C (0 min)
 3. 20°C to 300 °C
 4. Hold 8 minutes

- GCxGC-HRTOFMS

- ZB-5HT (30m), ZB35-HT (2m)
 - A portion of the ZB35-HT column was used as the modulator loop
- Zoex ZX-2 thermal modulator, 8 s modulation period
- InfoMass LLC Column holder
- Split injection (100:1)
- Oven:
 1. 60°C (0 min)
 2. 3°C min⁻¹ to 315°C (0 min)



Essential oil samples analyzed



1. Peppermint Oil, Sakae

- *Mentha piperita*
- >35 years old
- Stored at room temperature

2. Peppermint Oil, Fluka

- *Mentha piperita*
- < 5 years old
- Stored in refrigerator

3. Spearmint Oil, Fluka

- *Mentha spicata*
- < 5 years old
- Stored in refrigerator

4. Lavender Oil, Sakae

- >35 years old
- Stored at room temperature

5. Lemon Oil

- *Citrus limon*
- "JEOL" -- unidentified source

6. Orange Oil

- *Citrus sinensis*
- a.k.a. *Citrus × aurantium f. aurantium*

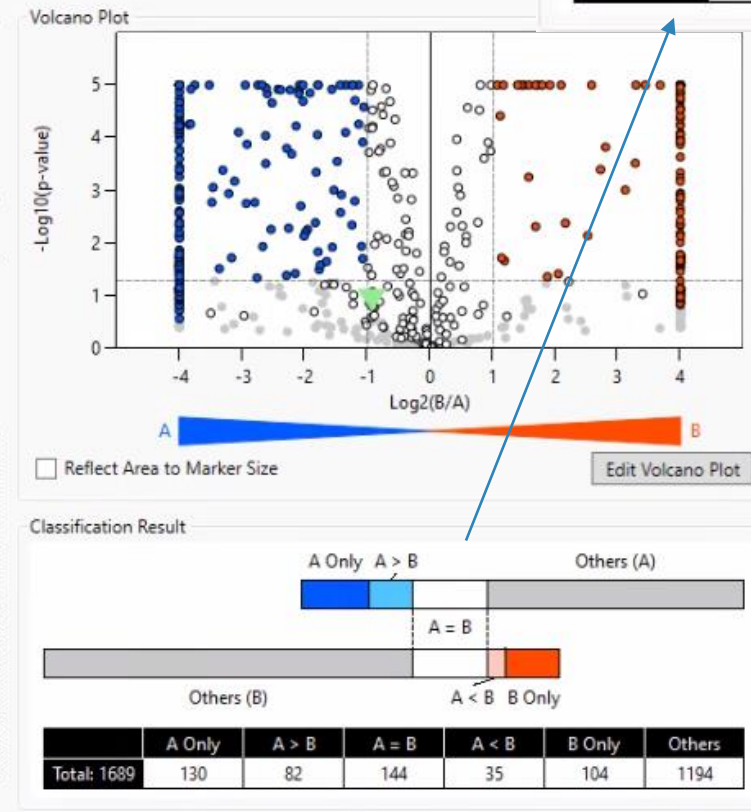
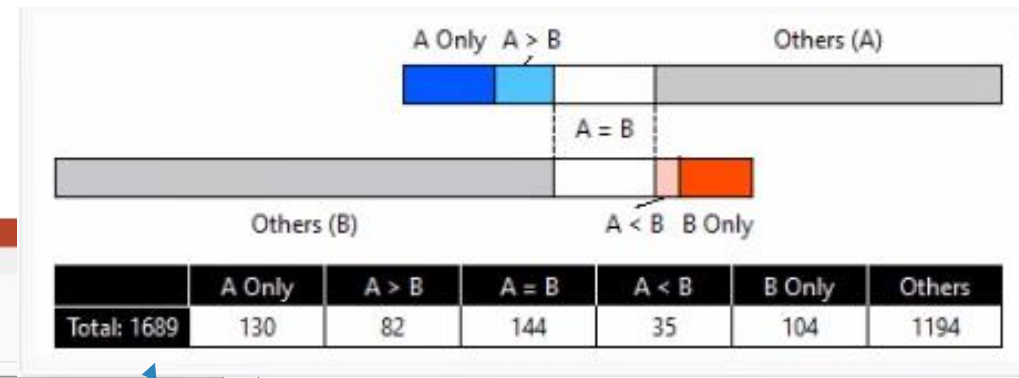
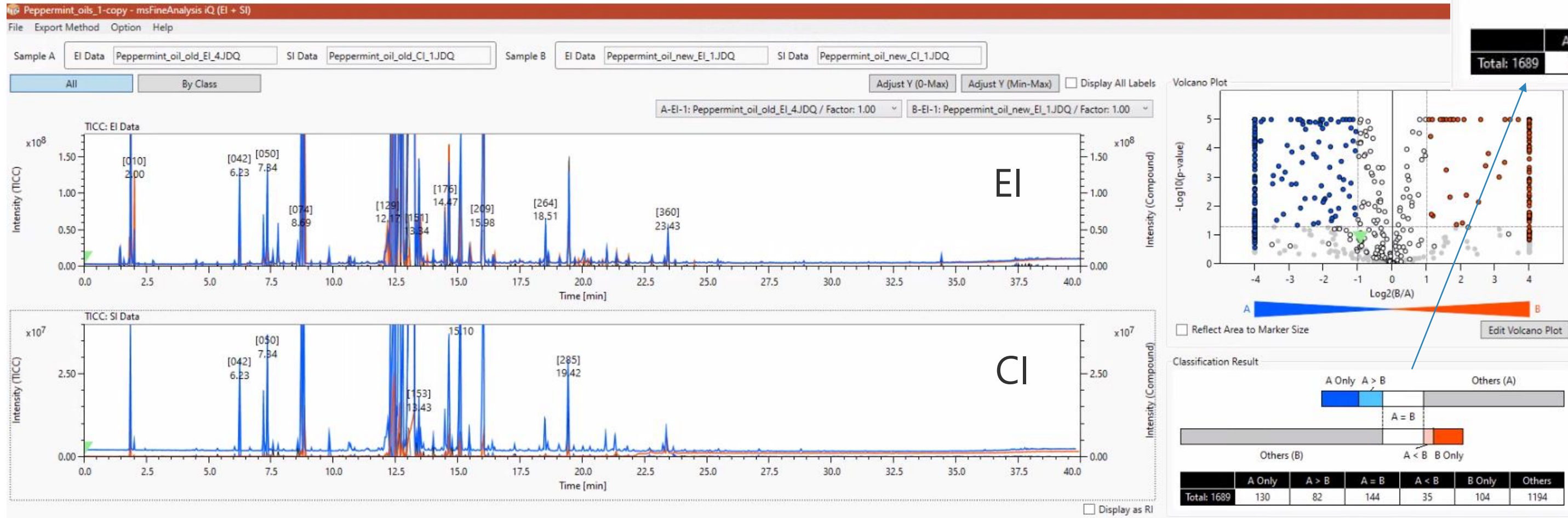
7. "Sweet Orange" Oil

- *Citrus sinensis*
- a.k.a. *Citrus × aurantium f. aurantium*

Difference analysis (1D GC and GC x GC)

- **Five EI replicates (Single Quad and HRTOF)**
- **One CI (Single Quad) or FI (HRTOF) measurement**

Single Quadrupole: GC-MS comparison of Peppermint Oils with EI and CI data, msFineAnalysis iQ software



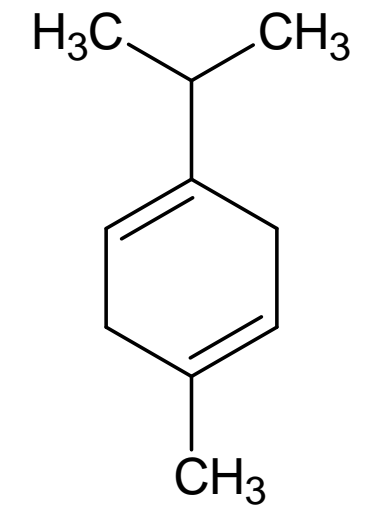
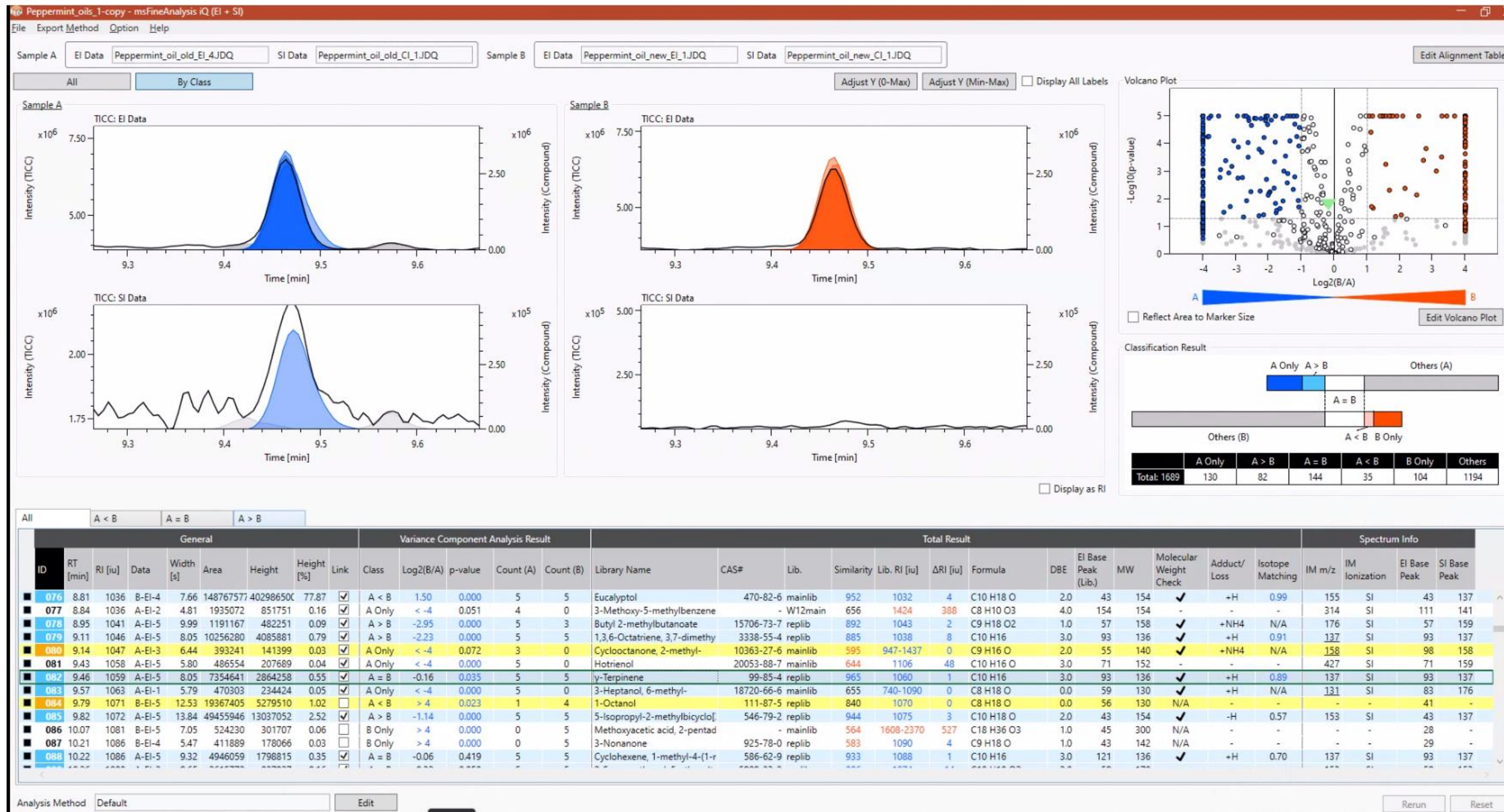
495 reproducible peaks in the 2 samples

- Deconvolution
- Correlate EI and CI
- Database search
- RI match
- Isotope match
- Variance analysis

General							Variance Component Analysis Result					Total Result										Spectrum Info								
ID	RT [min]	RI [u]	Data	Width [s]	Area	Height	Height [%]	Link	Class	Log2(B/A)	p-value	Count (A)	Count (B)	Library Name	CAS#	Lib.	Similarity	Lib. RI [u]	ΔRI [u]	Formula	DBE	EI Base Peak (Lib.)	MW	Molecular Weight Check	Adduct/Loss	Isotope Matching	IM m/z	IM Ionization	EI Base Peak	SI Base Peak
074	8.69	1031	A-EI-2	7.69	229411871	48934348	9.46	✓	A Only	< -4	0.000	5	0	1,4-Dimethyl-4-vinyl-1-cyclot	1743-61-9	W12main	703	1015	16	C10 H16	3.0	68	136	-	-	425	SI	68	103	
075	8.72	1032	A-EI-5	7.37	739333156	232214102	44.87	✓	A = B	-0.94	0.000	5	5	Limonene	138-86-3	W12main	821	1018	14	C10 H16	3.0	68	136	✓	+NH4 0.65	154	SI	39	81	
076	8.81	1036	B-EI-4	7.66	148767577	402986500	77.87	✓	A < B	1.50	0.000	5	5	Eucalyptol	470-82-6	mainlib	952	1032	4	C10 H18 O	2.0	43	154	✓	+H 0.99	155	SI	43	137	
077	8.84	1036	A-EI-2	4.81	1935072	851751	0.16	✓	A Only	< -4	0.051	4	0	3-Methoxy-5-methylbenzene	-	W12main	656	1424	388	C8 H10 O3	4.0	154	154	-	-	314	SI	111	141	
078	8.95	1041	A-EI-5	9.99	1191167	482251	0.09	✓	A > B	-2.95	0.000	5	3	Butyl 2-methylbutanoate	15706-73-7	replib	892	1043	2	C9 H18 O2	1.0	57	158	✓	+NH4 N/A	176	SI	57	159	
079	9.11	1046	A-EI-5	8.05	10256280	4085881	0.79	✓	A > B	-2.23	0.000	5	5	1,3,6-Octatriene, 3,7-dimethy	3338-55-4	replib	885	1038	8	C10 H16	3.0	93	136	✓	+H 0.91	137	SI	93	137	
080	9.14	1047	A-EI-3	6.44	393241	141399	0.03	✓	A Only	< -4	0.072	3	0	Cyclooctanone, 2-methyl-	10363-27-6	mainlib	395	947-1437	0	C9 H16 O	2.0	55	140	✓	+NH4 N/A	158	SI	98	158	
081	9.43	1058	A-EI-5	5.80	486554	207689	0.04	✓	A Only	< -4	0.000	5	0	Hotrienol	20053-88-7	mainlib	444	1106	48	C10 H16 O	3.0	71	152	-	-	427	SI	71	159	
082	9.46	1059	A-EI-5	8.05	7354641	2864258	0.55	✓	A = B	-0.16	0.035	5	5	γ-Terpinene	99-85-4	replib	965	1060	1	C10 H16	3.0	93	136	✓	+H 0.89	137	SI	93	137	
083	9.57	1063	A-EI-1	5.79	470303	234424	0.05	✓	A Only	< -4	0.000	5	0	3-Heptanol, 6-methyl-	18720-66-6	mainlib	655	740-1090	0	C8 H18 O	0.0	59	130	✓	+H N/A	131	SI	83	176	
084	9.79	1071	B-EI-5	12.53	19367405	5279510	1.02	✓	A < B	> 4	0.023	1	4	1-Octanol	111-87-5	replib	840	1070	0	C8 H18 O	0.0	56	130	N/A	-	-	-	41	-	
085	9.82	1072	A-EI-5	13.84	49455946	13037052	2.52	✓	A > B	-1.14	0.000	5	5	5-Isopropyl-2-methylbicyclo[546-79-2	replib	944	1075	3	C10 H18 O	2.0	43	154	✓	-H 0.57	153	SI	43	137	
086	10.07	1081	B-EI-5	7.05	524230	301707	0.06	✓	B Only	> 4	0.000	0	5	Methoxyacetic acid, 2-pentad	-	mainlib	564	1608-2370	527	C18 H36 O3	1.0	45	300	N/A	-	-	-	28	-	
087	10.21	1086	B-EI-4	5.47	411889	178066	0.03	✓	B Only	> 4	0.000	0	5	3-Nonanone	925-78-0	replib	583	1090	4	C9 H18 O	1.0	43	142	N/A	-	-	-	29	-	
088	10.22	1086	A-EI-5	9.32	4946059	1798815	0.35	✓	A = B	-0.06	0.419	5	5	Cyclohexene, 1-methyl-4-(1-r	586-62-9	replib	933	1088	1	C10 H16	3.0	121	136	✓	+H 0.70	137	SI	93	137	

Compare peppermint oil samples with msFineAnalysis iQ

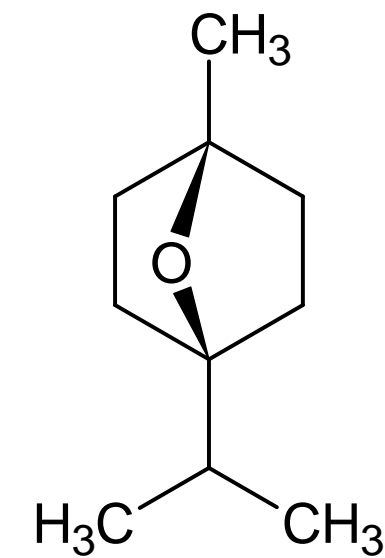
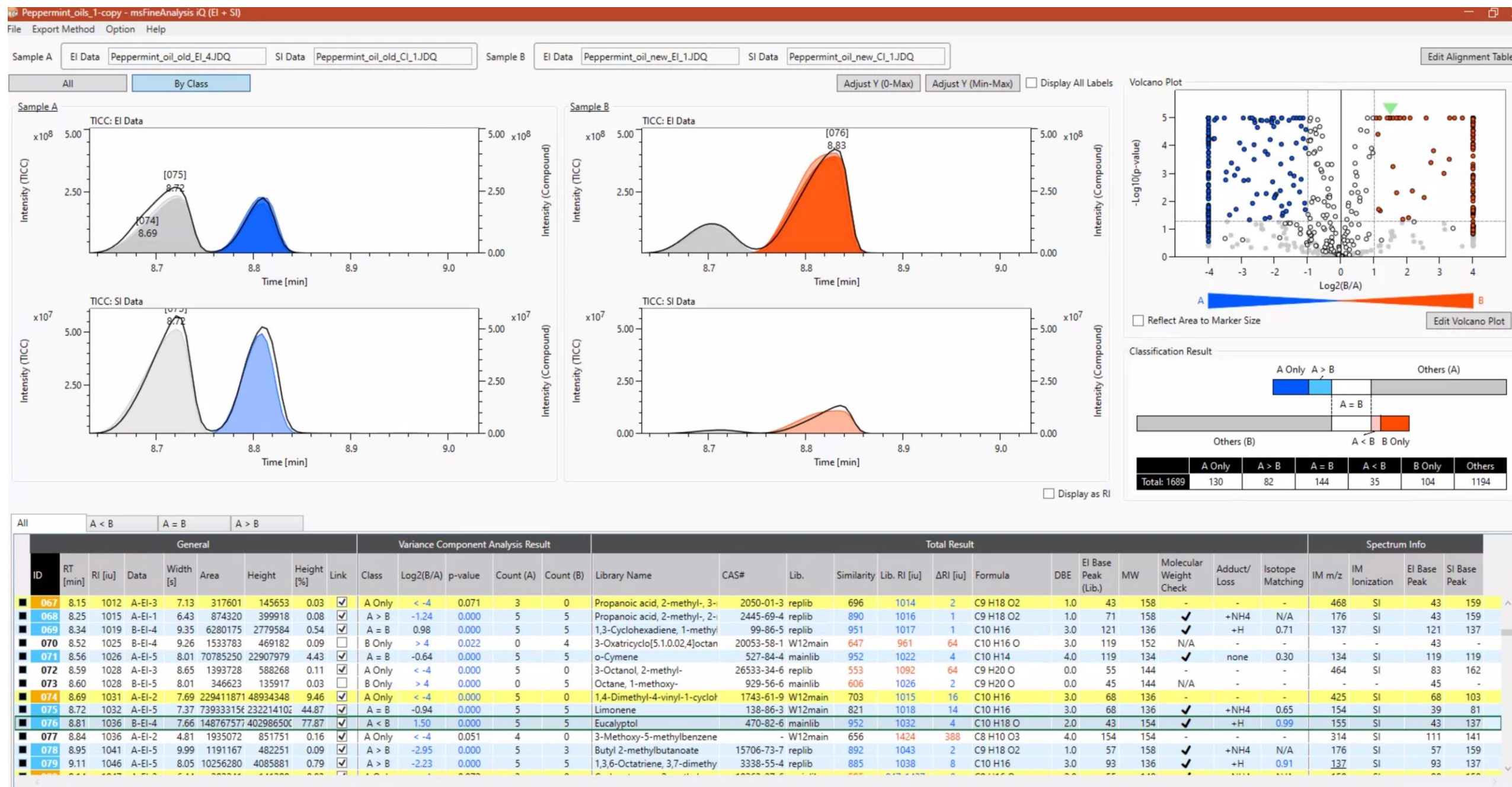
Single quadrupole: EI and CI data



γ -terpinene
Equal in both samples

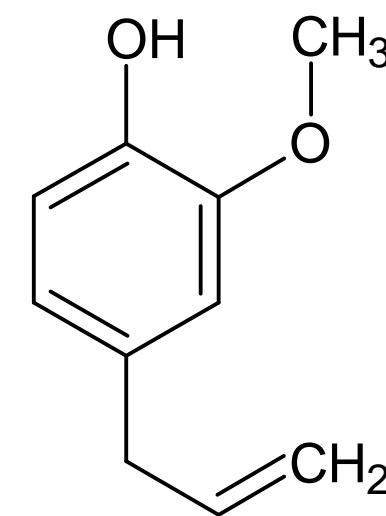
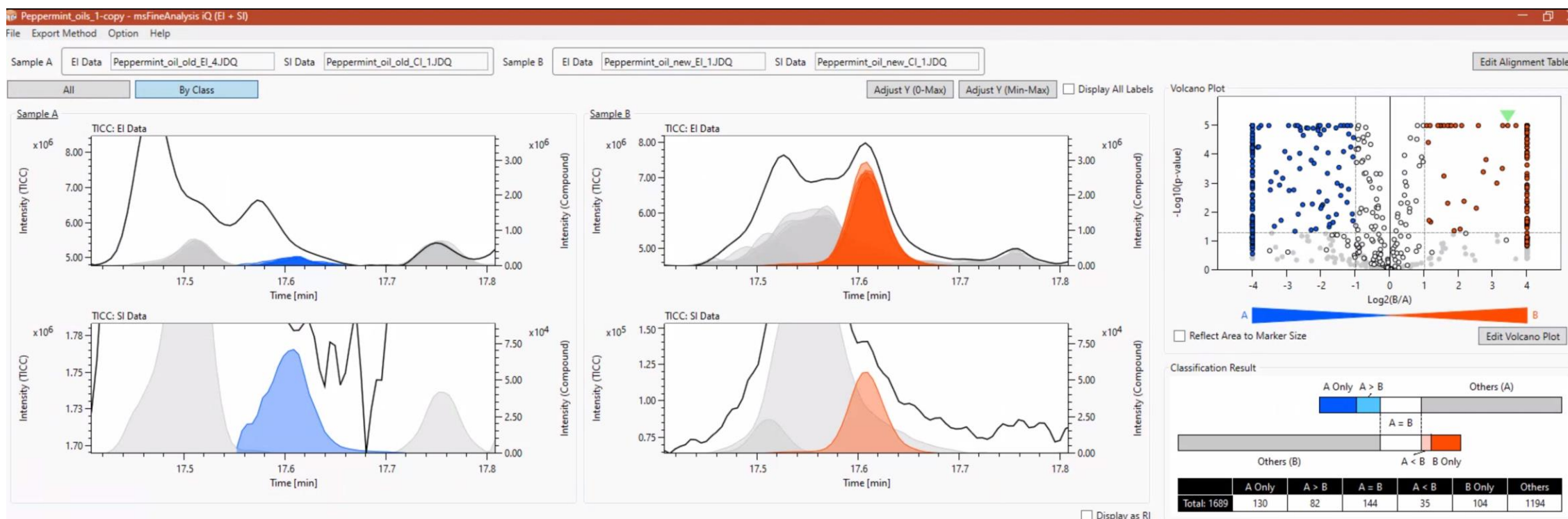
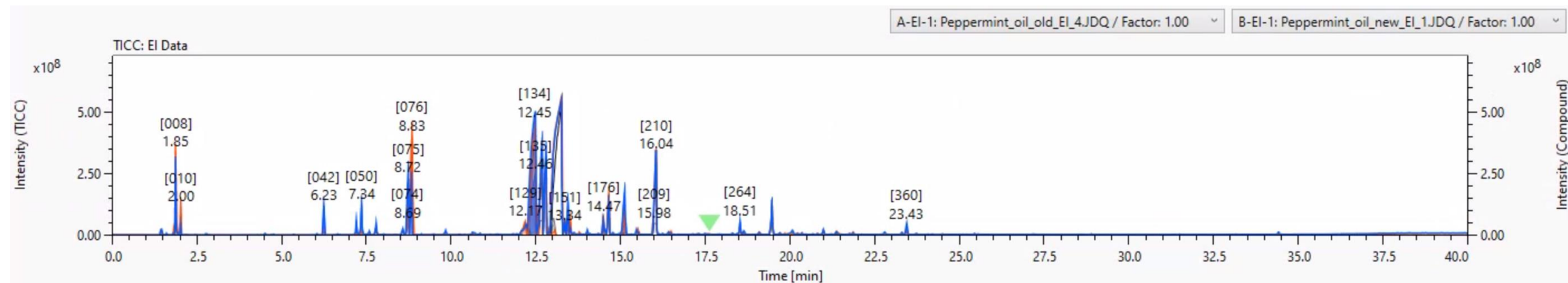
Compare peppermint oil samples with msFineAnalysis iQ

Single quadrupole: EI and CI data



Eucalyptol
More abundant in Fluka

Chromatographic deconvolution for small Eugenol peak



Eugenol
More abundant in Fluka
Height = 0.57%

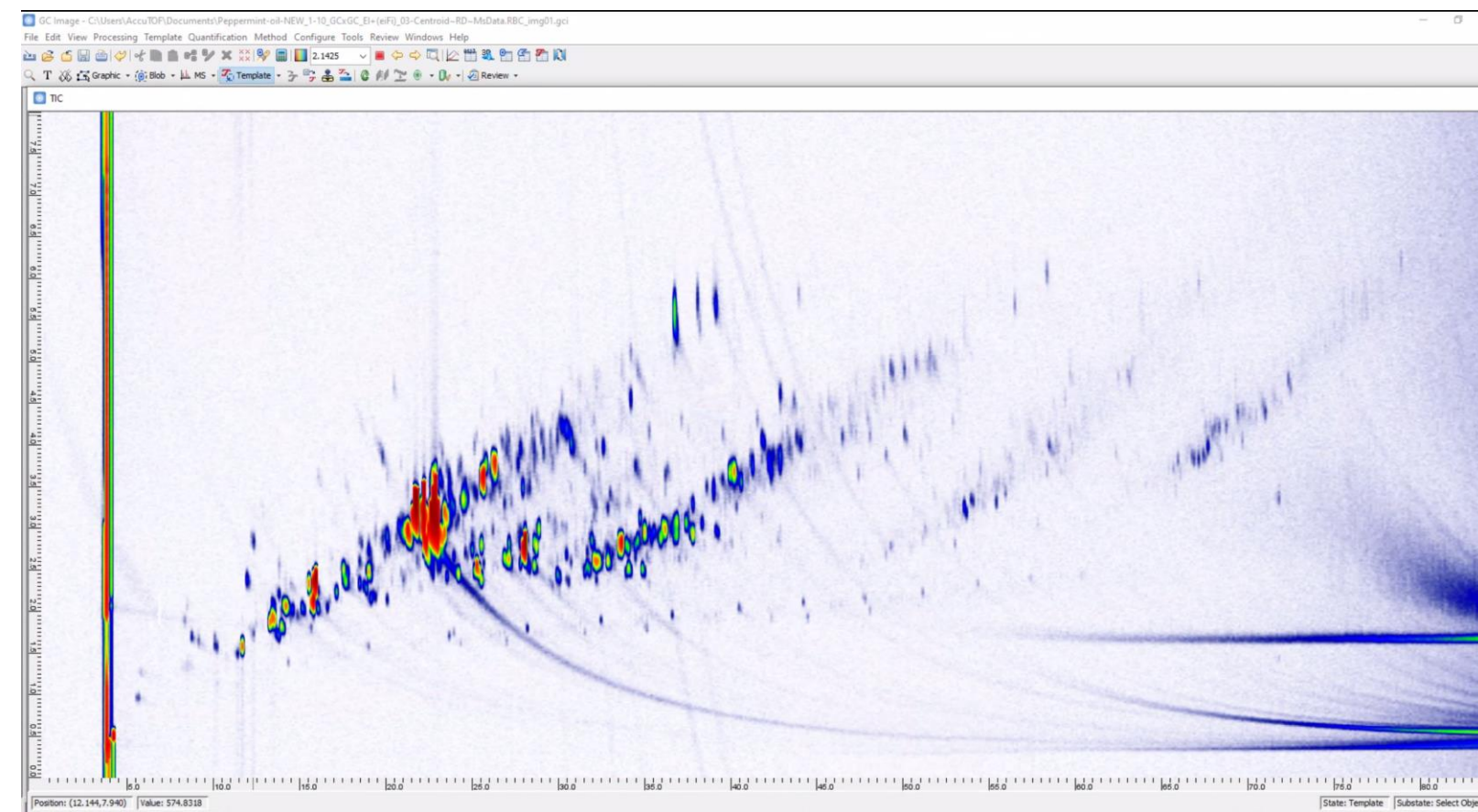
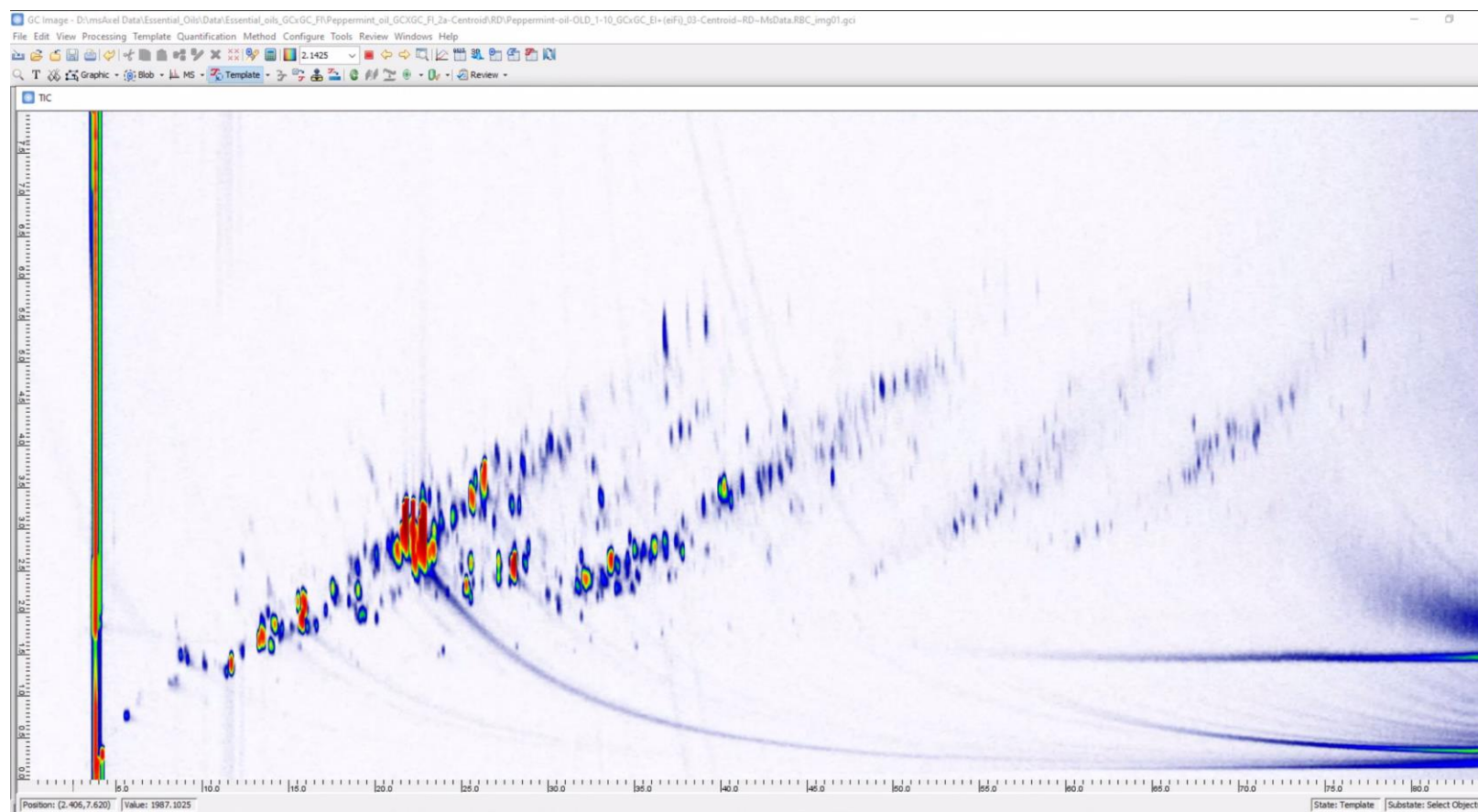
General										Variance Component Analysis Result					Total Result										Spectrum Info						
ID	RT [min]	RI [u]	Data	Width [s]	Area	Height	Height [%]	Link	Class	Log2(B/A)	p-value	Count (A)	Count (B)	Library Name	CAS#	Lib.	Similarity	Lib. RI [u]	ΔRI [u]	Formula	DBE	EI Base Peak (Lib.)	MW	Molecular Weight Check	Adduct/Loss	Isotope Matching	IM m/z	IM Ionization	EI Base Peak	SI Base Peak	
001	0.02	484	A-EI-5	5.13	1959112	852077	0.16	✓	A = B	-0.92	0.253	4	3	1H-Pyrrole-2-carbonitrile	9513-94-4	mainlib	546	689-1399	205	C5 H4 N2	5.0	92	92	-	-	-	509	SI	18	253	
002	1.41	610	A-EI-4	3.23	22802401	19509976	3.77	✓	A = B	-0.25	0.101	5	5	Nitrogen	7727-37-9	W12main	824	N/A	N/A	N2	2.0	28	28	-	-	-	513	SI	28	162	
003	1.44	613	A-EI-5	7.10	33243872	19396872	3.75	✓	A > B	-1.19	0.000	5	5	Water	7732-18-5	W12main	914	N/A	N/A	H2 O	0.0	18	18	-	-	-	476	SI	18	273	
004	1.47	615	A-EI-1	5.79	884766	631628	0.12	✓	A = B	0.33	0.259	5	5	Butane, 1-isocyano-	2769-64-4	replib	713	N/A	N/A	C5 H9 N	2.0	43	83	-	-	-	512	SI	29	234	
005	1.53	621	A-EI-2	2.91	519044	455011	0.09	✓	A Only	< -4	0.000	5	0	Acetic acid, ethoxyhydroxy-, e	49653-17-0	mainlib	679	642-1404	21	C6 H12 O4	1.0	29	148	-	-	-	475	SI	31	129	
006	1.58	625	A-EI-1	5.14	7192583	6354088	1.23	✓	A = B	-0.99	0.029	5	5	Acetone-oxime	-	W12main	795	652	27	C3 H7 N O	1.0	43	73	-	-	-	436	SI	43	309	
007	1.76	642	A-EI-1	5.79	3076091	2411445	0.47	✓	A > B	-3.46	0.001	5	4	Acetic acid	64-19-7	replib	910	610	32	C2 H4 O2	1.0	43	60	✓	none	0.95	60	EI	45	181	
008	1.85	650	B-EI-1	6.09	541481106	336917836	65.10	✓	A < B	1.19	0.022	5	5	Oxirane, 2-ethyl-2-methyl-	30095-63-7	mainlib	750	634	16	C5 H10 O	1.0	41	86	✓	none	0.95	86	EI	42	85	
009	1.97	660	A-EI-4	4.82	1854109	1502928	0.29	✓	A Only	< -4	0.075	3	0	1-Propanol, 2-methyl-	78-83-1	mainlib	928	624	36	C4 H10 O	0.0	43	74	✓	none	N/A	74	EI	43	217	
010	2.00	663	B-EI-1	3.53	133491926	115692976	22.36	✓	A = B	2.22	0.054	5	5	Cyclopentane, methyl-	96-37-7	replib	935	630	33	C6 H12	1.0	56	84	N/A	-	-	-	-	-	56	83
011	2.14	675	A-EI-1	5.79	1465304	1262437	0.24	✓	A > B	-3.76	0.000	5	3	Isovaleric aldehyde	590-86-3	FFNSC3	922	676	1	C5 H10 O	1.0	44	86	✓	+H	N/A	87	SI	43	87	
012	2.20	681	A-EI-2	3.53	1735661	1425185	0.28	✓	A Only	< -4	0.000	5	0	Butanal, 2-methyl-	96-17-3	mainlib	762	662	18	C5 H10 O	1.0	41	86	✓	+H	N/A	87	SI	29	87	
013	2.30	690	A-EI-2	5.46	624172	497365	0.10	✓	A Only	< -4	0.000	5	0	1-Penten-3-ol	616-25-1	W12main	884	671	19	C5 H10 O	1.0	57	86	-	-	-	458	SI	57	187	



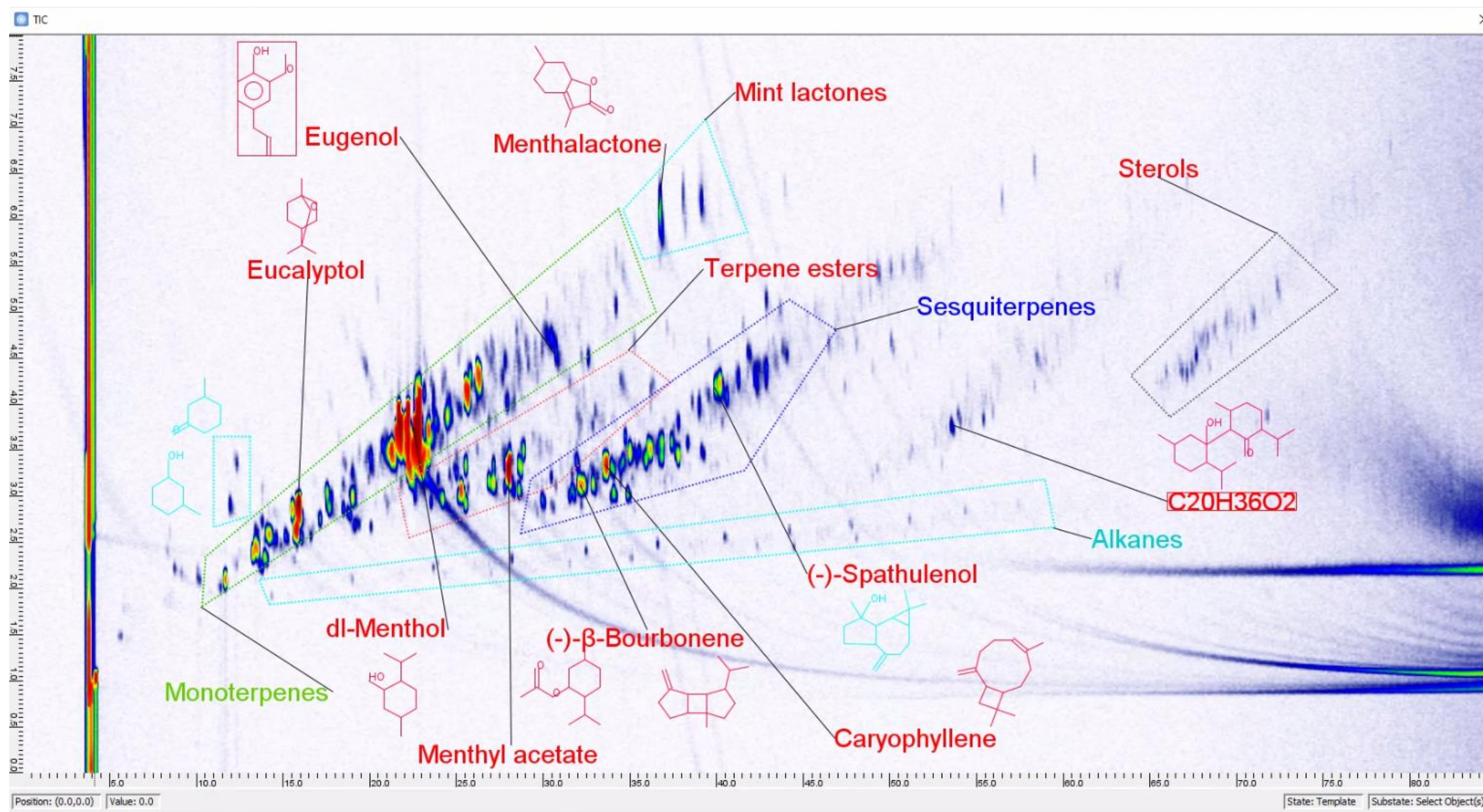
GC x GC-HRTOFMS chromatograms for the peppermint oils

Sakae (>35-year-old sample)

Fluka (Fresh sample)



Some details for the Fluka Peppermint Oil

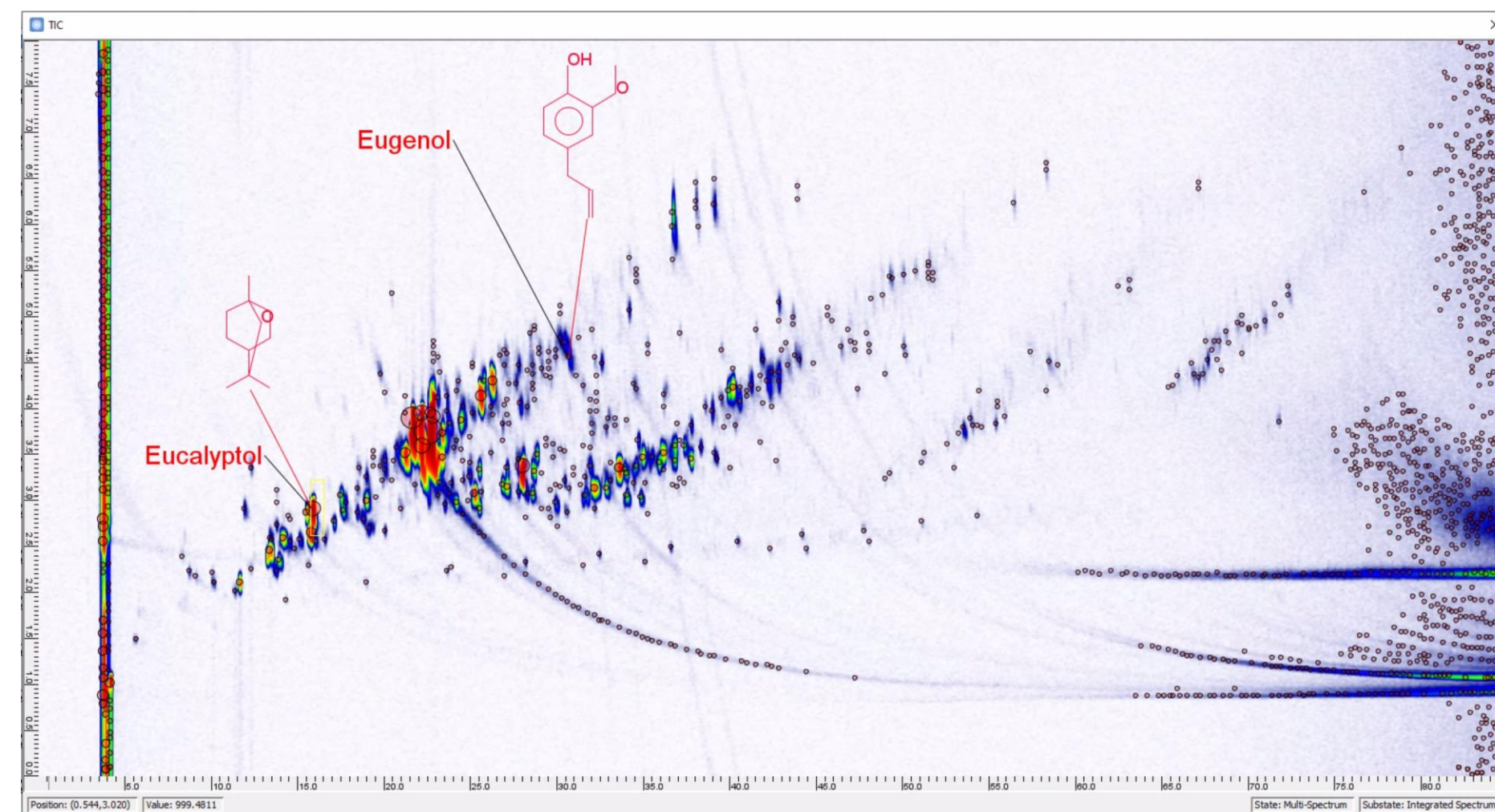
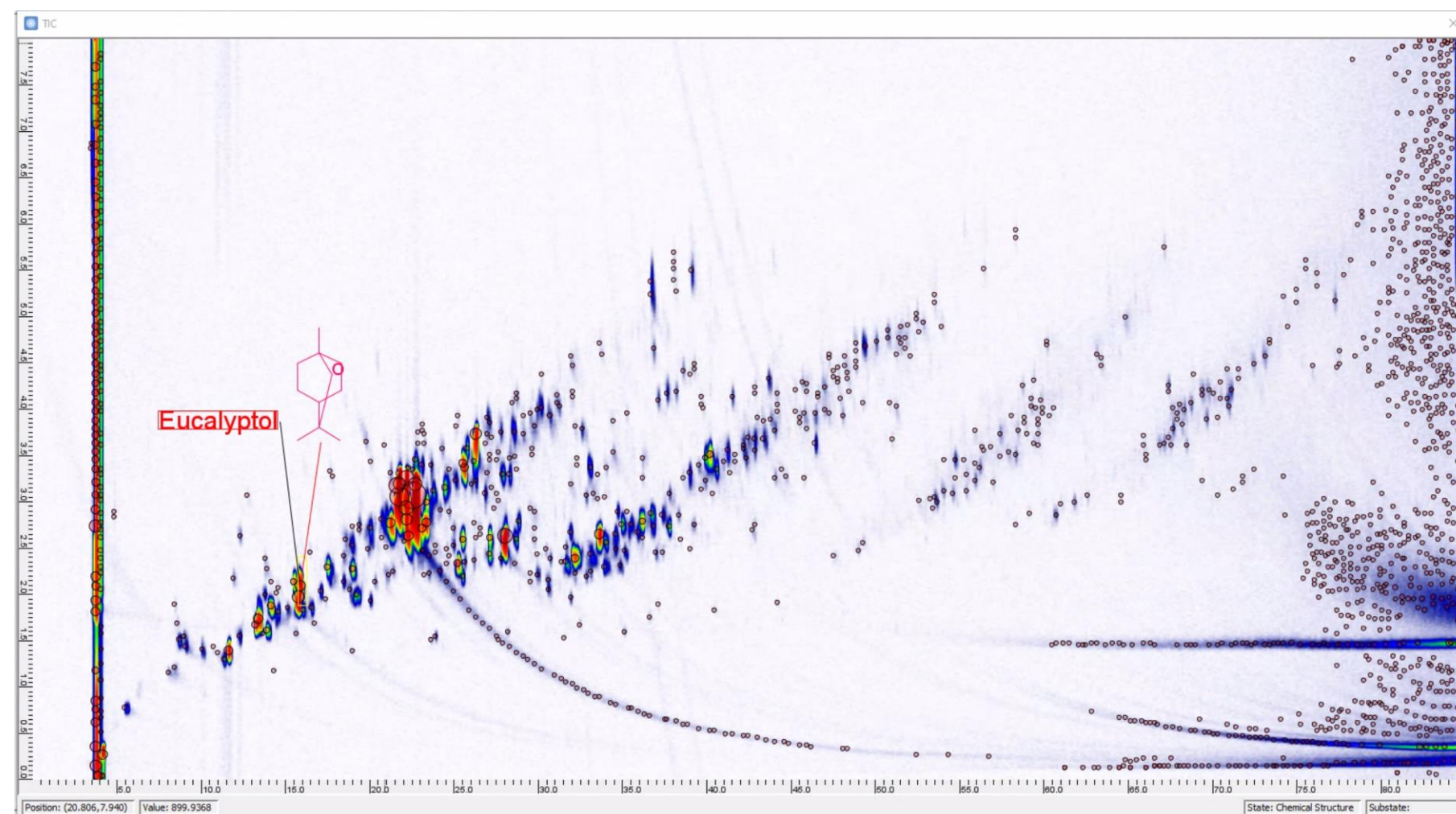




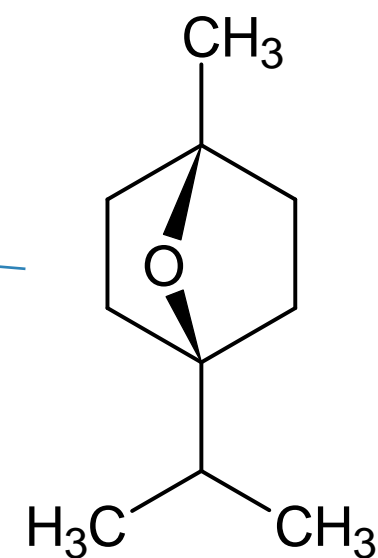
Eucalyptol and Eugenol blobs in Sakae and Fluka samples: Two compounds that were found to differ in the 1D GC-MS data

Sakae (>35-year-old sample)

Fluka (Fresh sample)

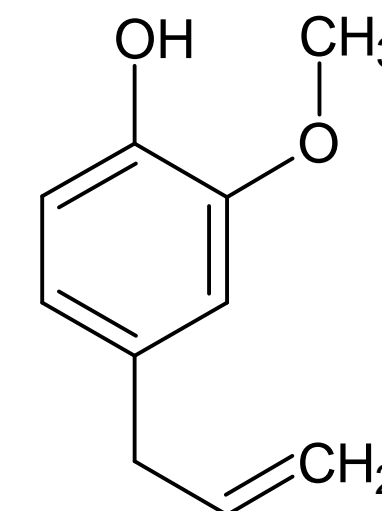
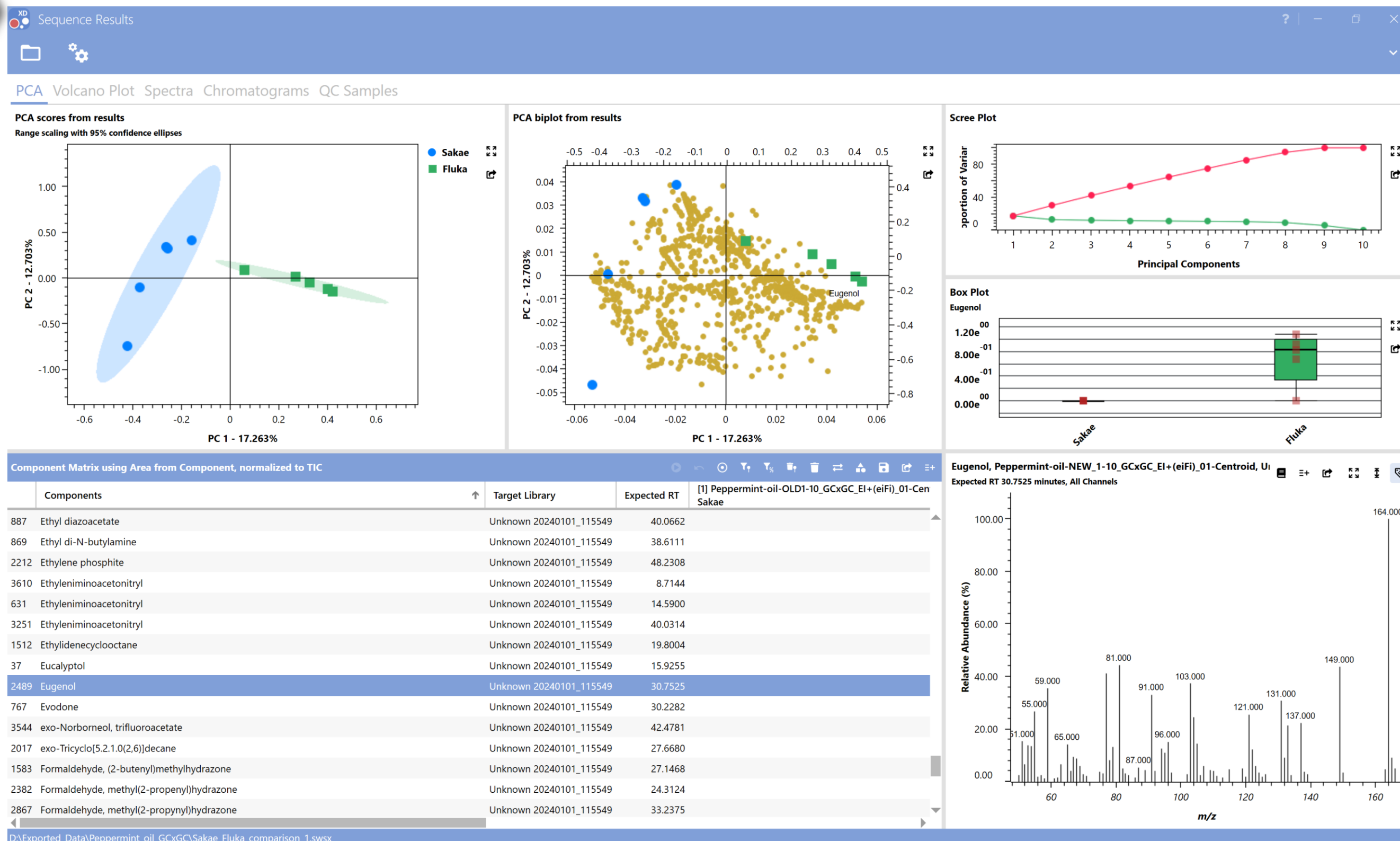


PCA and Eucalyptol entry for GC x GC-MS in AnalyzerPro XD



Eucalyptol
More abundant in Fluka

PCA and Eugenol entry for GC x GC-MS in AnalyzerPro XD



Eugenol
More abundant in Fluka
Height = 0.57%

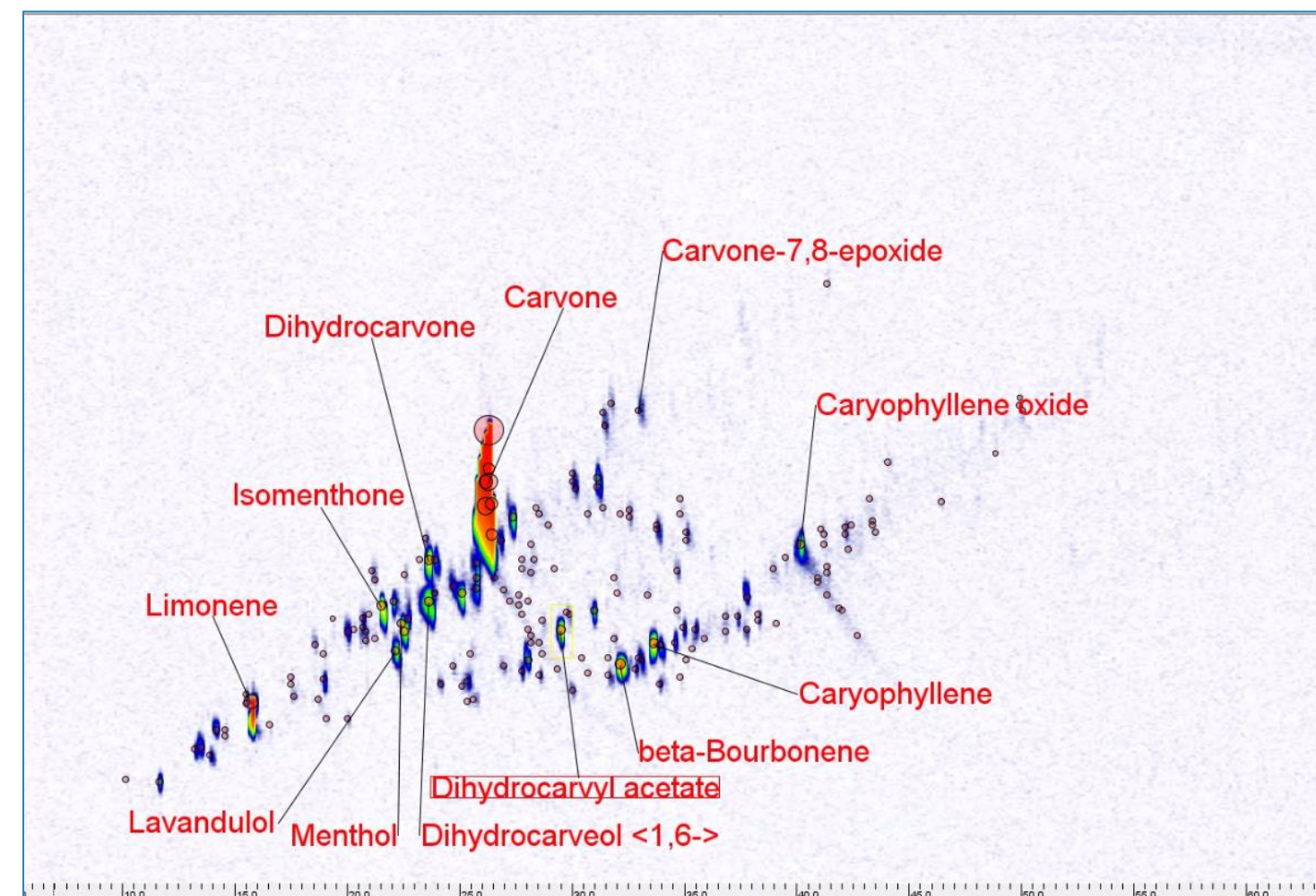
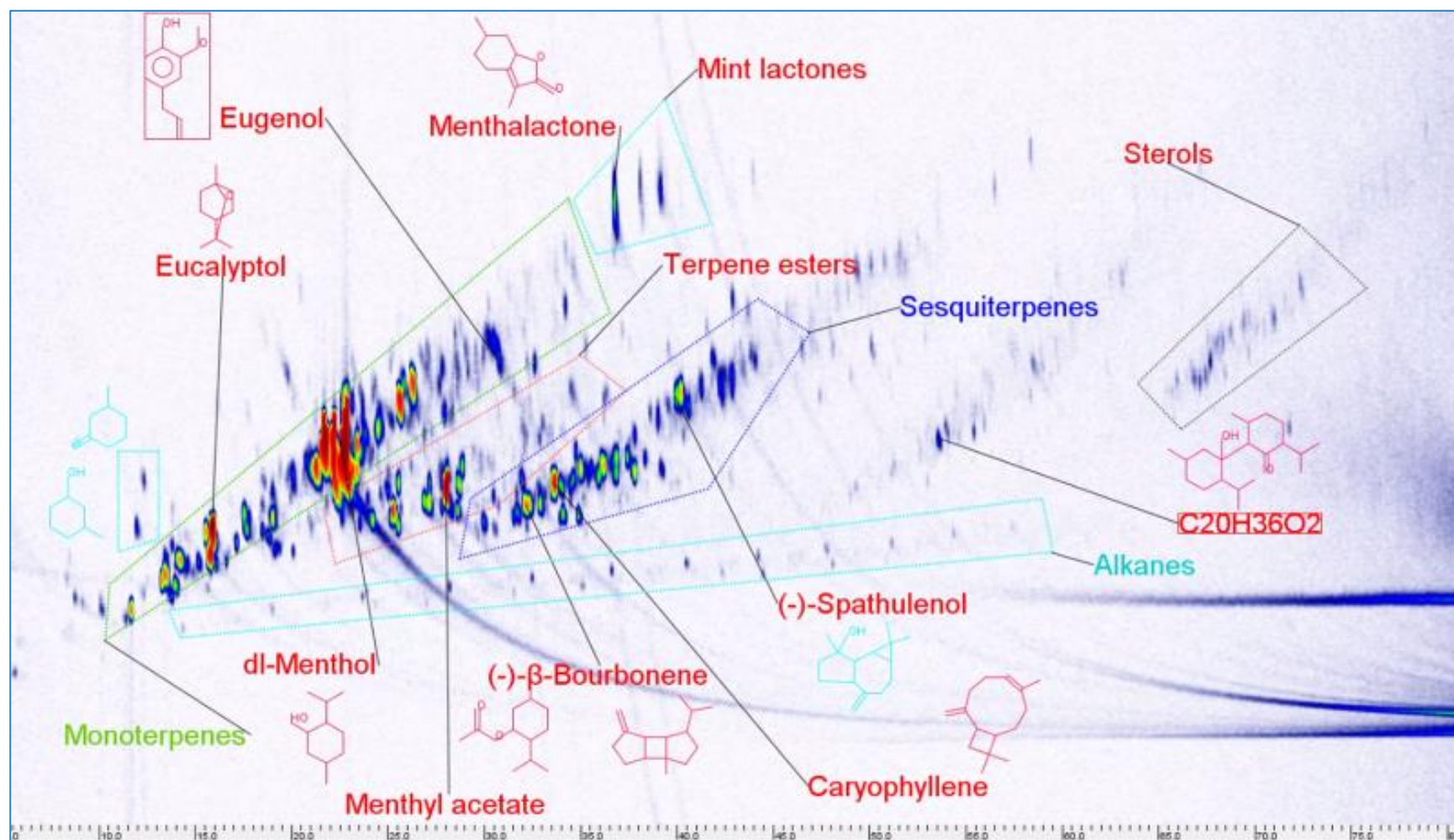
Peppermint Oil and Spearmint Oil: very different compositions!



Peppermint Oil

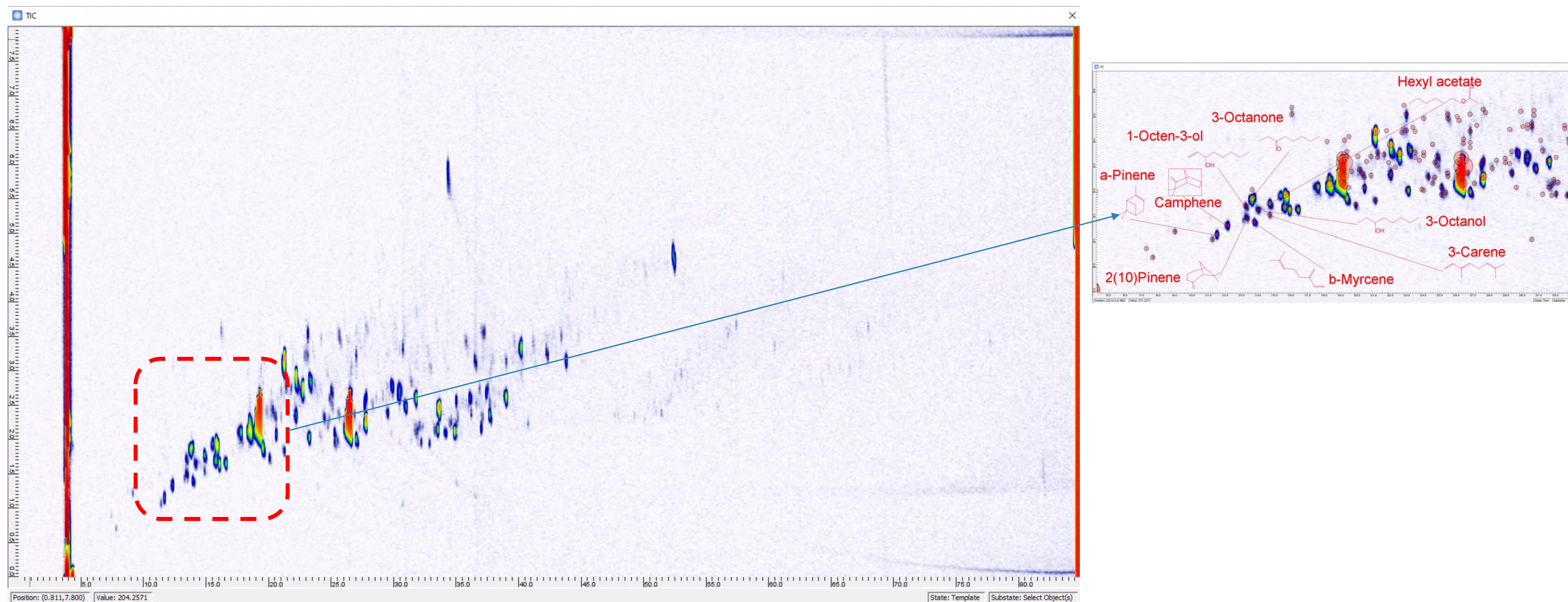


Spearmint Oil





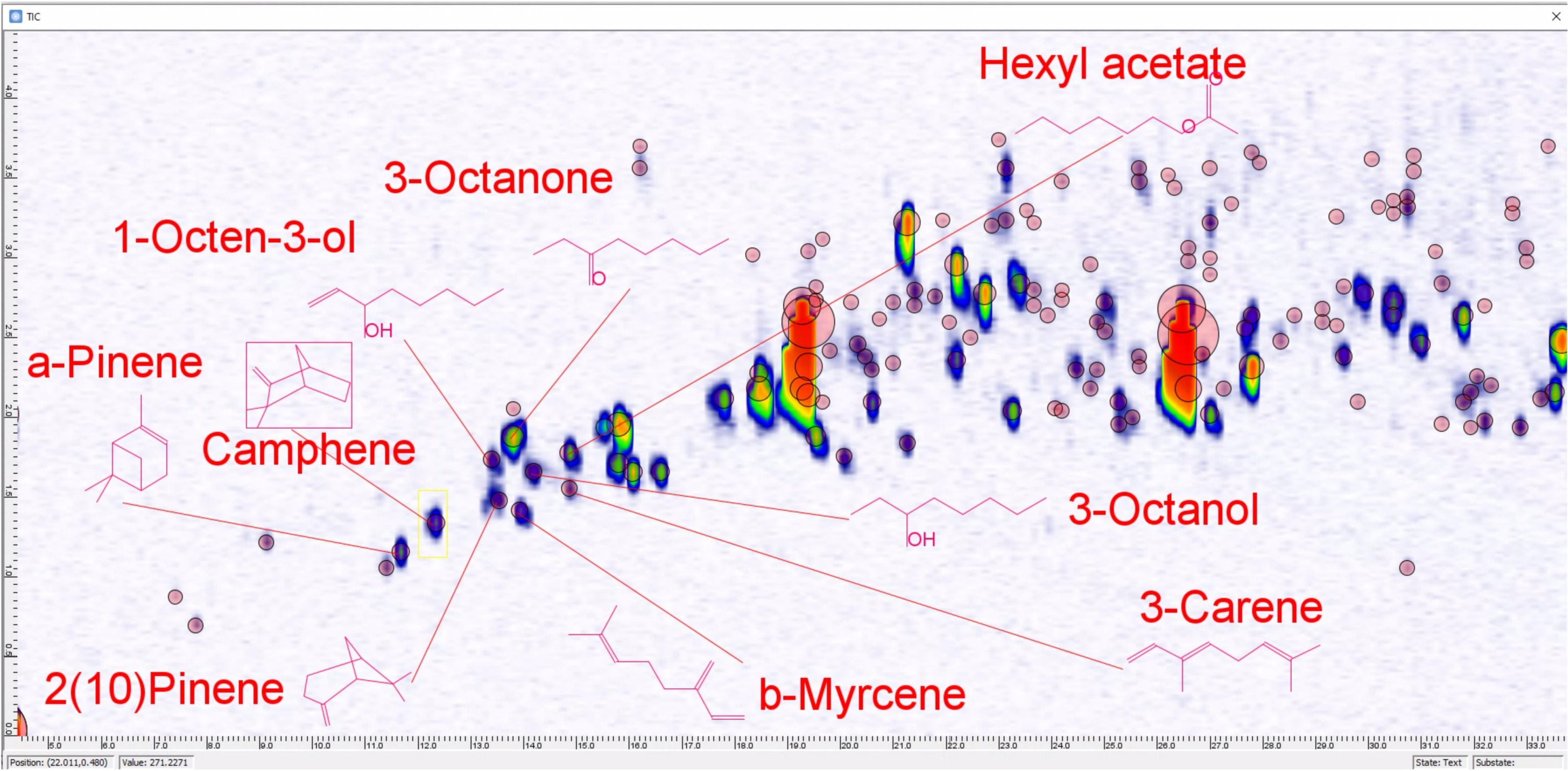
How does GC-MS (LR and HR) with deconvolution compare with GCxGC-HRTOFMS?



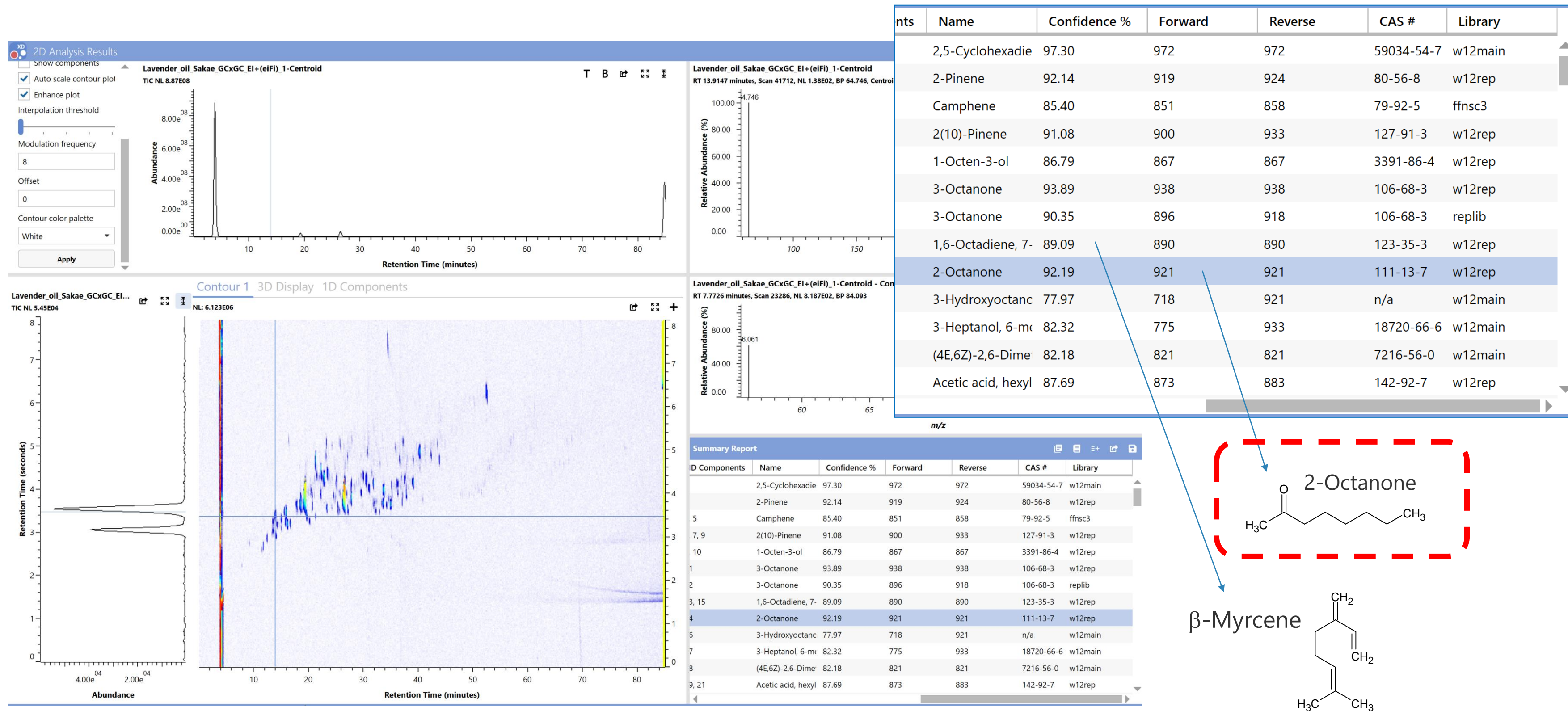
We'll use Lavender Oil as an example. Let's look at a small section of the chromatograms where several components elute closely

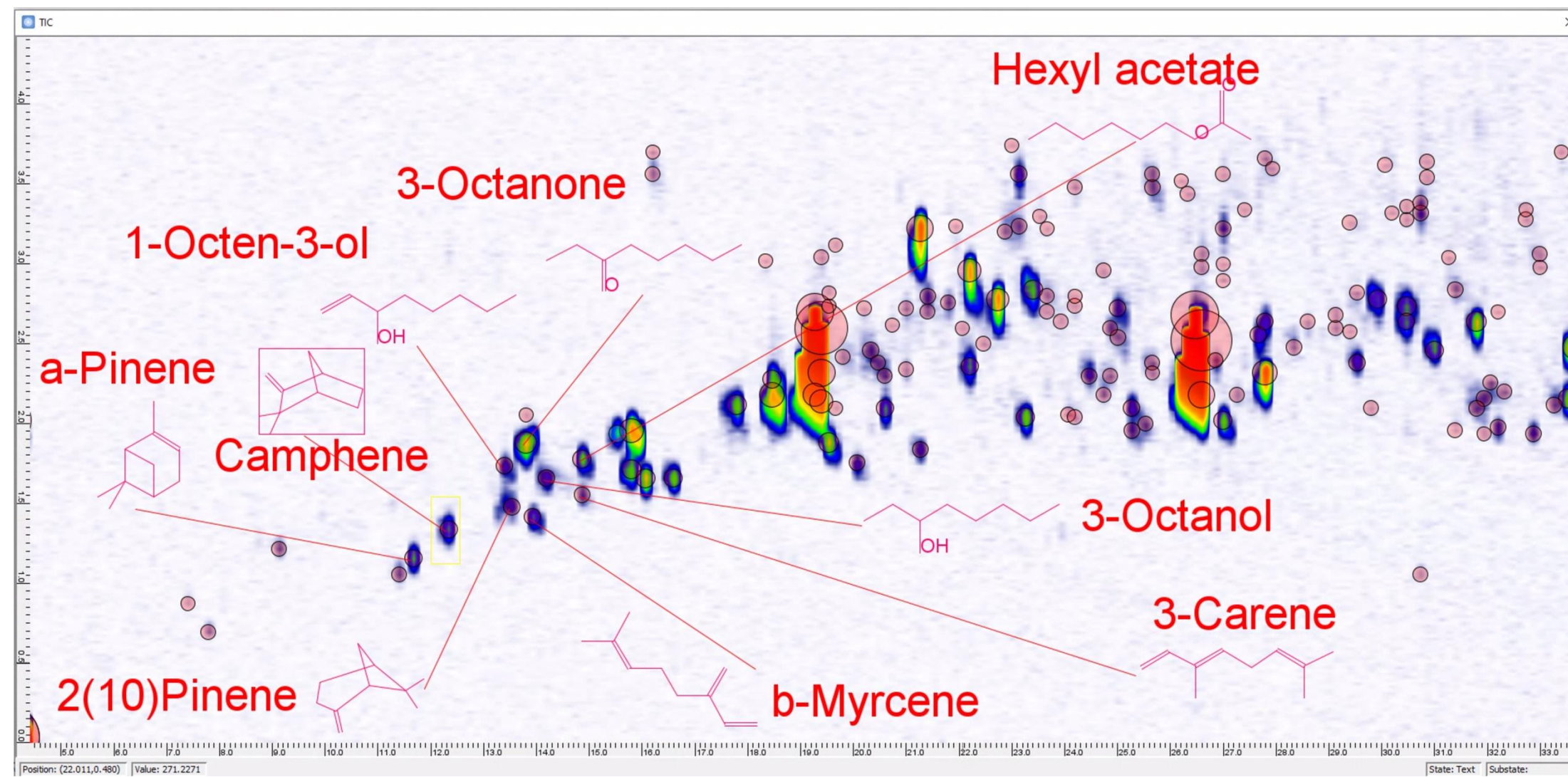


GC Image results for the selected region



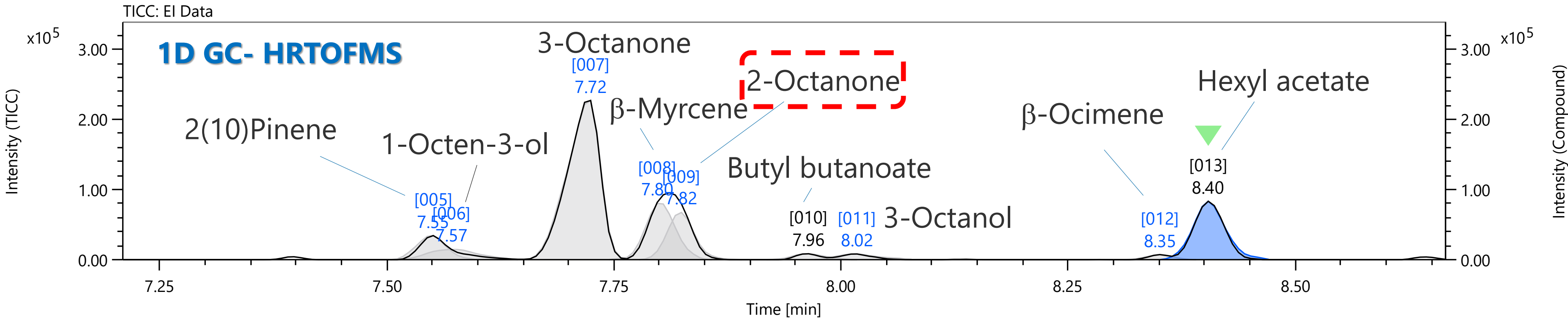
AnalyzerPro XD results for the selected region



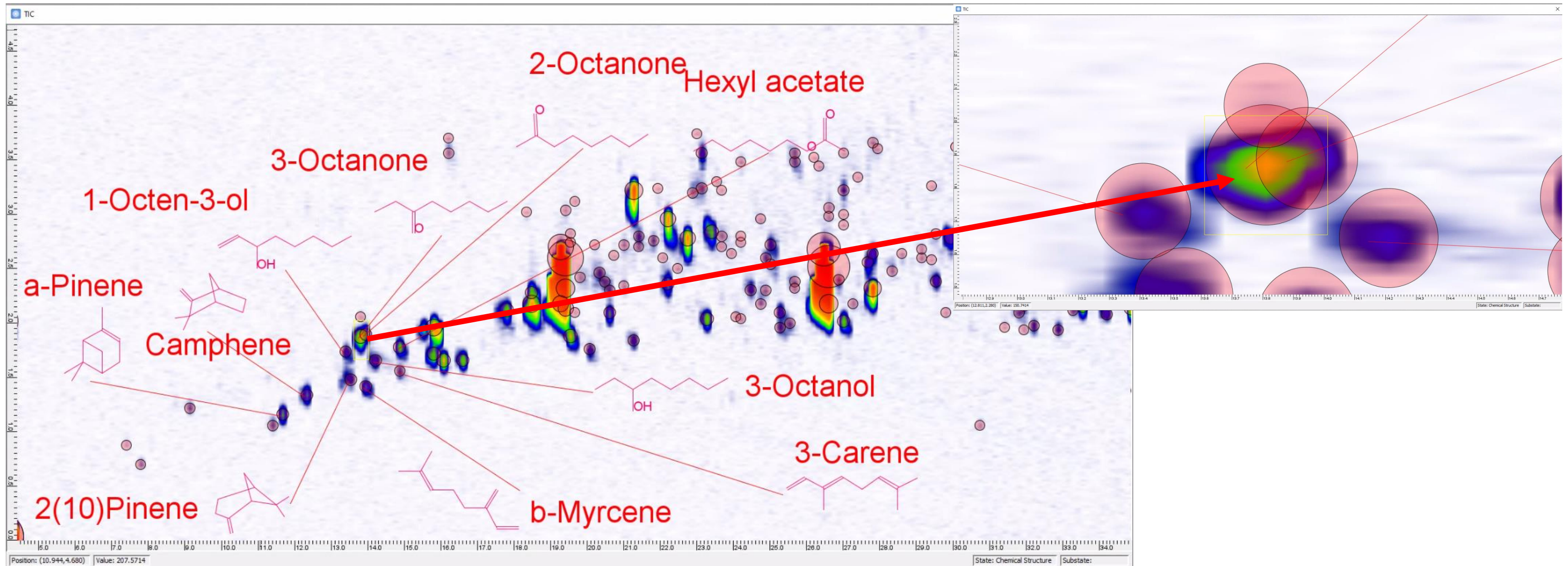


**GC x GC – HRTOFMS
+ GC Image**

2-Octanone not detected?



2-Octanone is detected in GC Image after blob deconvolution/unmixing



msFineAnalysis iQ results for GC-single quadrupole MS





Summary for a selected retention time region (lavender oil)

GCxGC/AnalyzerPro XD	GCxGC/GC Image	GC/Quadrupole	GC/HRTOF
2-Pinene (a-Pinene)	YES	YES	YES
Camphene	YES	YES	YES
2(10)-Pinene	YES	YES	YES
1-Octen-3-ol	YES	NO	YES
3-Octanone	YES	YES	YES
3-Octanone (not a separate blob)	YES	YES	YES
b-Myrcene	YES	NO	YES
2-Octanone	NO ¹	YES	NO
3-Hydroxyoctanoic Acid (Butyl Butanoate is Hit #6)	NO ²	NO	Butyl butanoate
3-Octanol	3-Octanol	NO	3-Octanol
b-Ocimene	YES	YES	YES
Acetic acid, hexyl ester	YES	YES	YES

12 Compounds
in this RT range

¹ YES after interactive blob deconvolution/unmixing

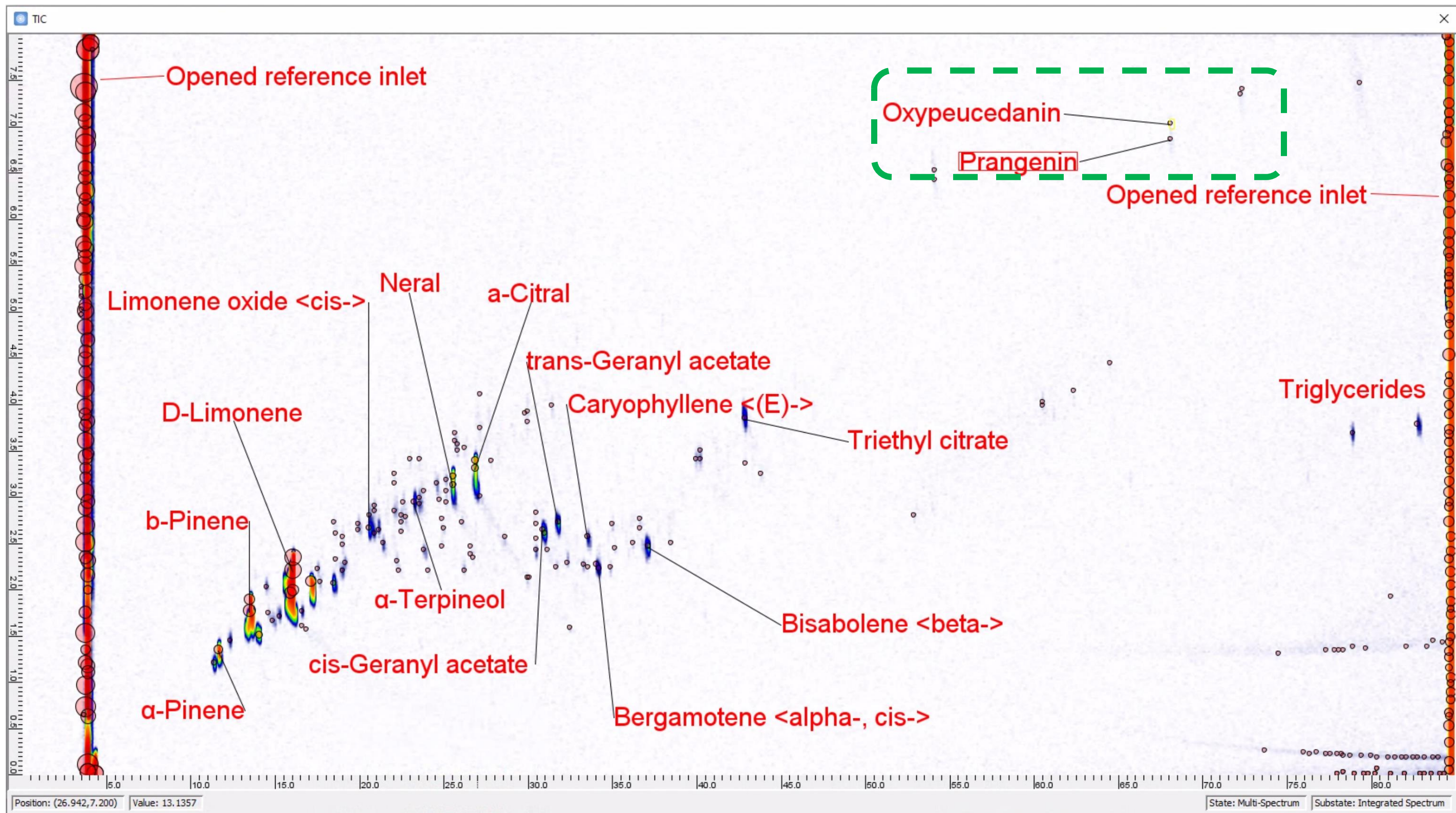
² 3rd hit after interactive blob deconvolution/unmixing

Both GCxGC data analysis programs examined were equally effective in detecting blobs and assigning compounds, although there were slight differences in the detection and ID of minor compounds.

- AnalyzerPro XD is easy to use for sample comparisons and uses a 2D deconvolution algorithm by default.
- GC Image has excellent graphics and has powerful tools for general GC x GC analysis
- For 1D GC-MS, msFineAnalysis iQ and AI use chromatographic deconvolution and integrate all available data



GCxGC-HRTOFMS of lemon oil. Let's look at the circled region.

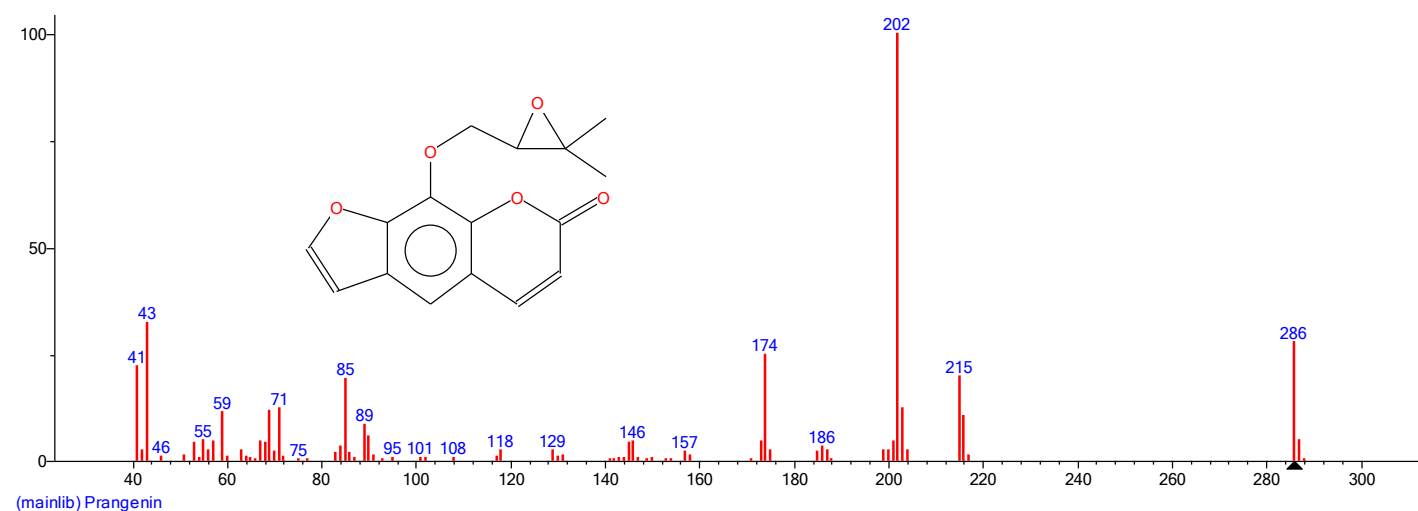




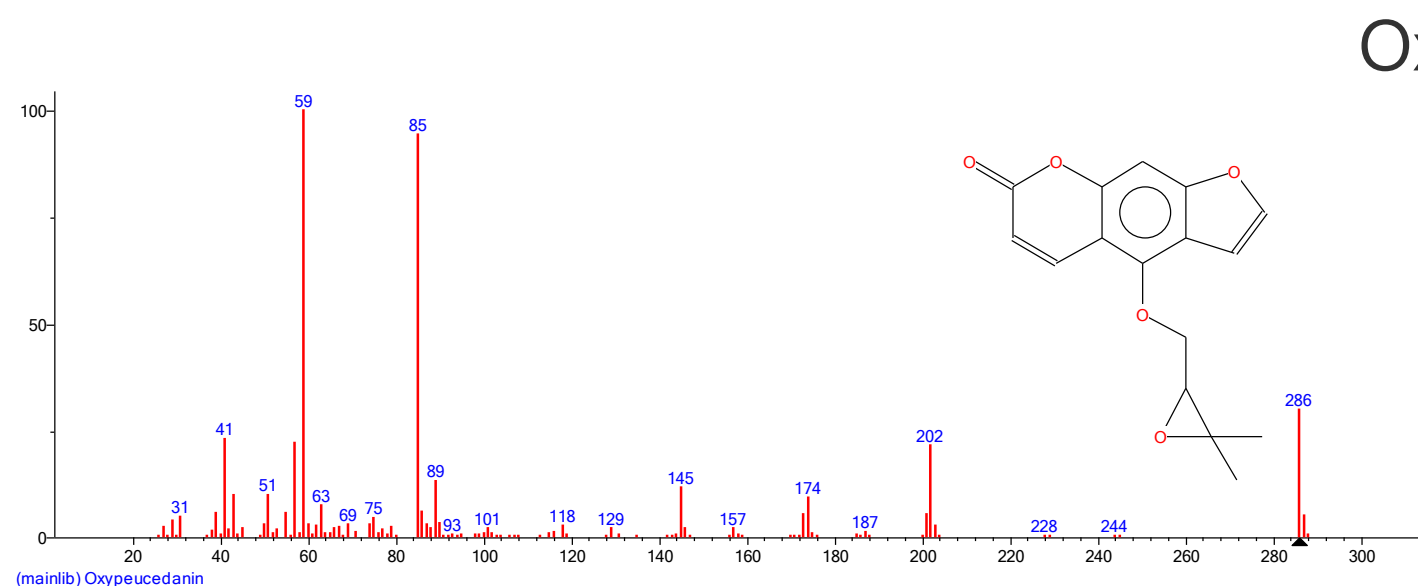
These furanocoumarins in lemon oil have an identical 1D retention index

C00002490	737-52-0	Oxypeucedanin	C ₁₆ H ₁₄ O ₅	286.08412356
C00037261	2880-49-1	Heraclenin	C ₁₆ H ₁₄ O ₅	286.08412356

http://www.knapsackfamily.com/knapsack_core/result.php?sname=organism&word=Citrus%20limon

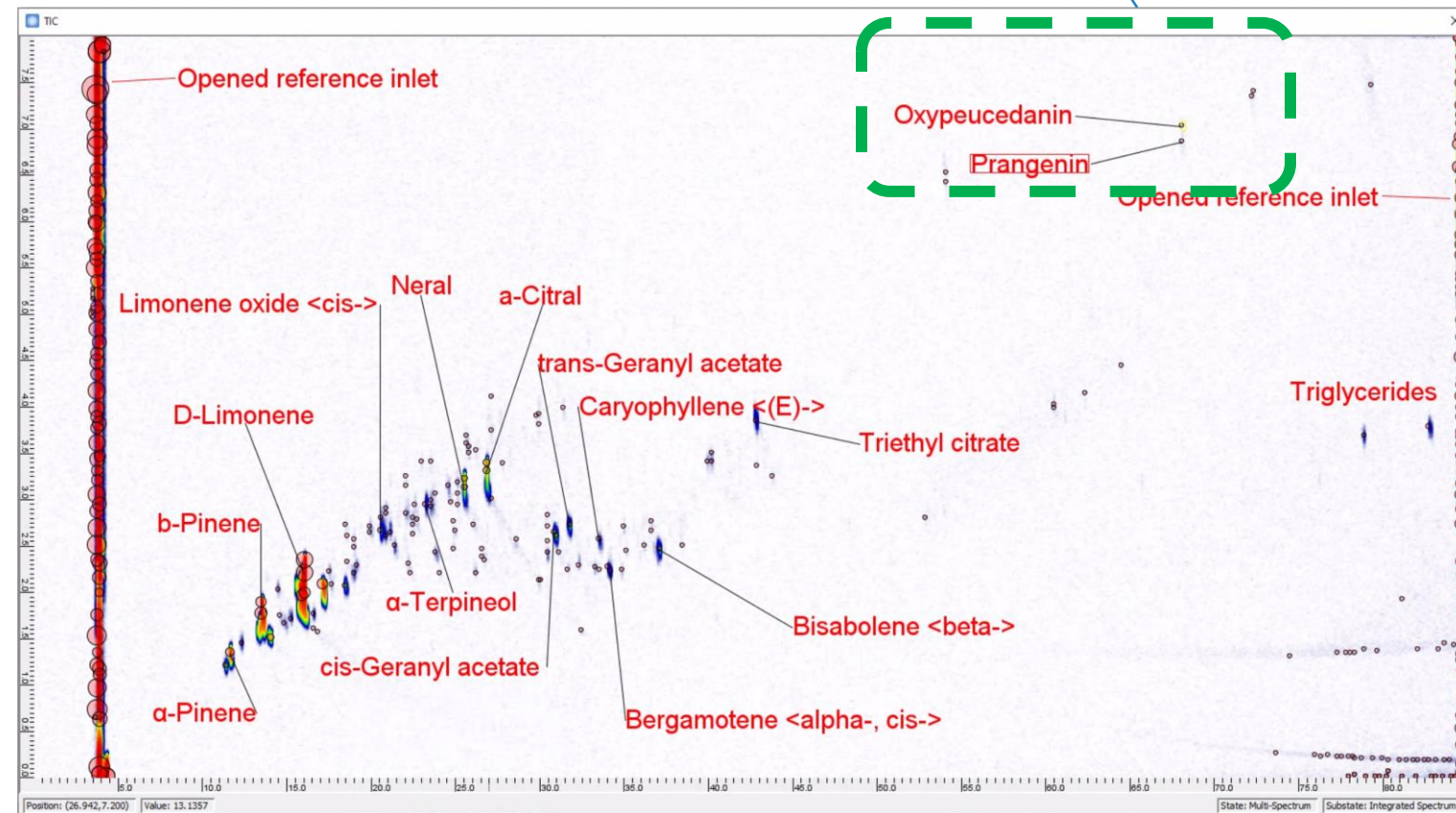
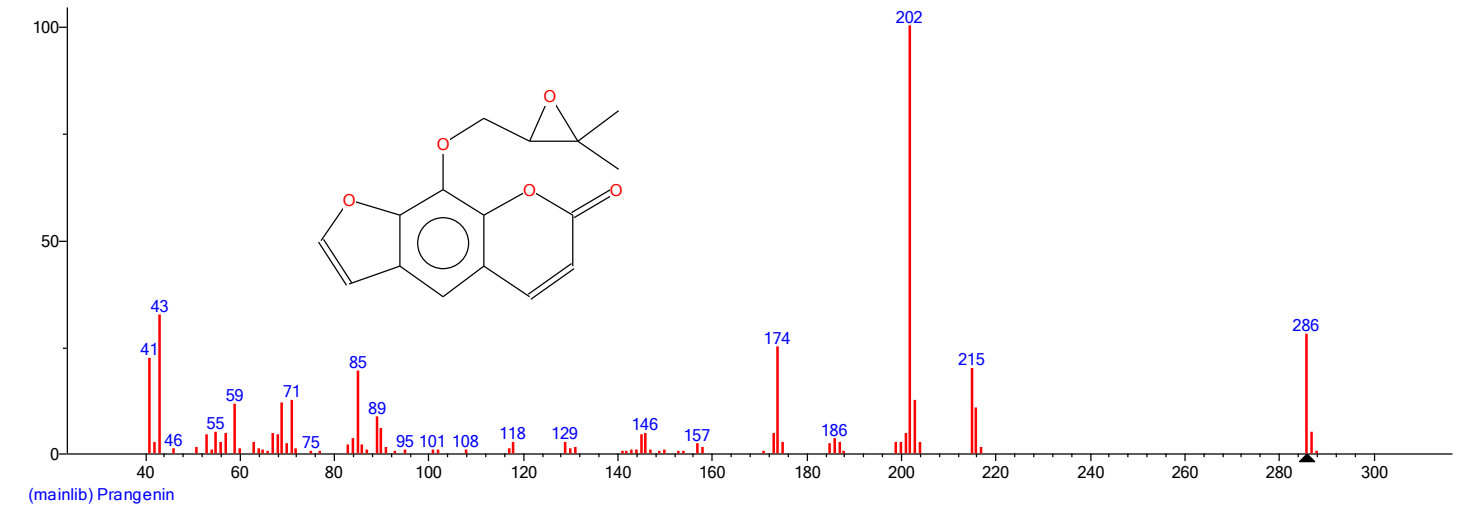


Calculated RI for semi-nonpolar column is
2324 IU for both compounds

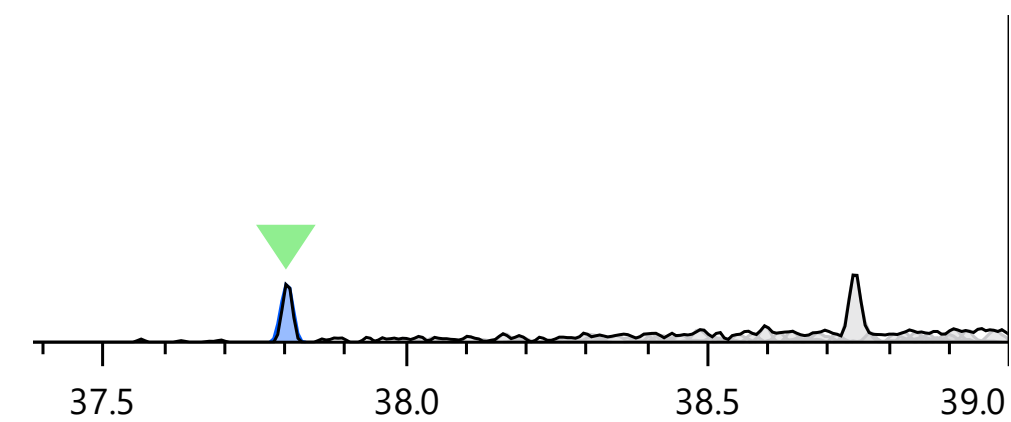




1D GC-MS does not see both compounds

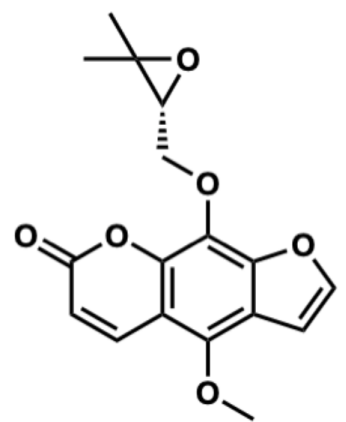


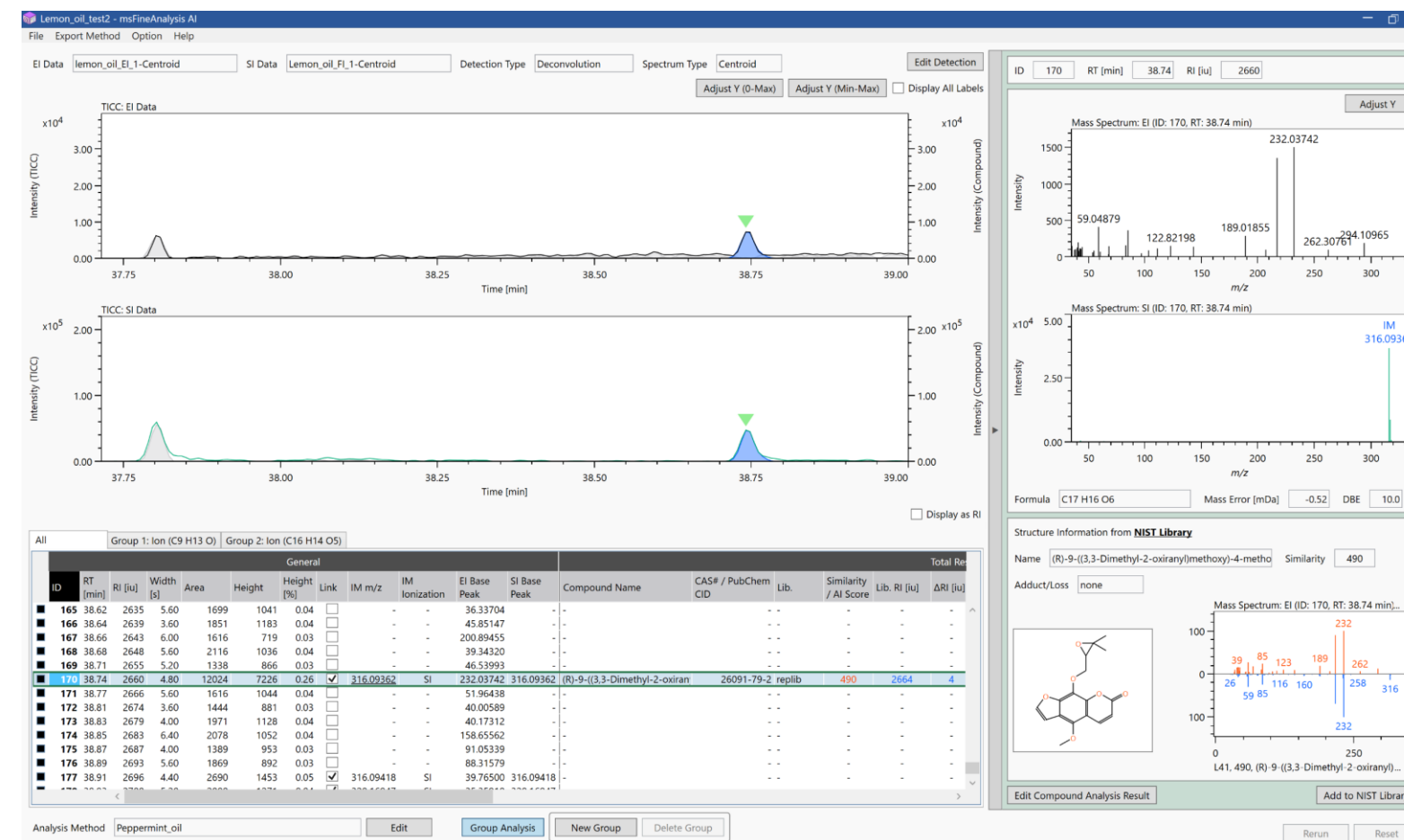
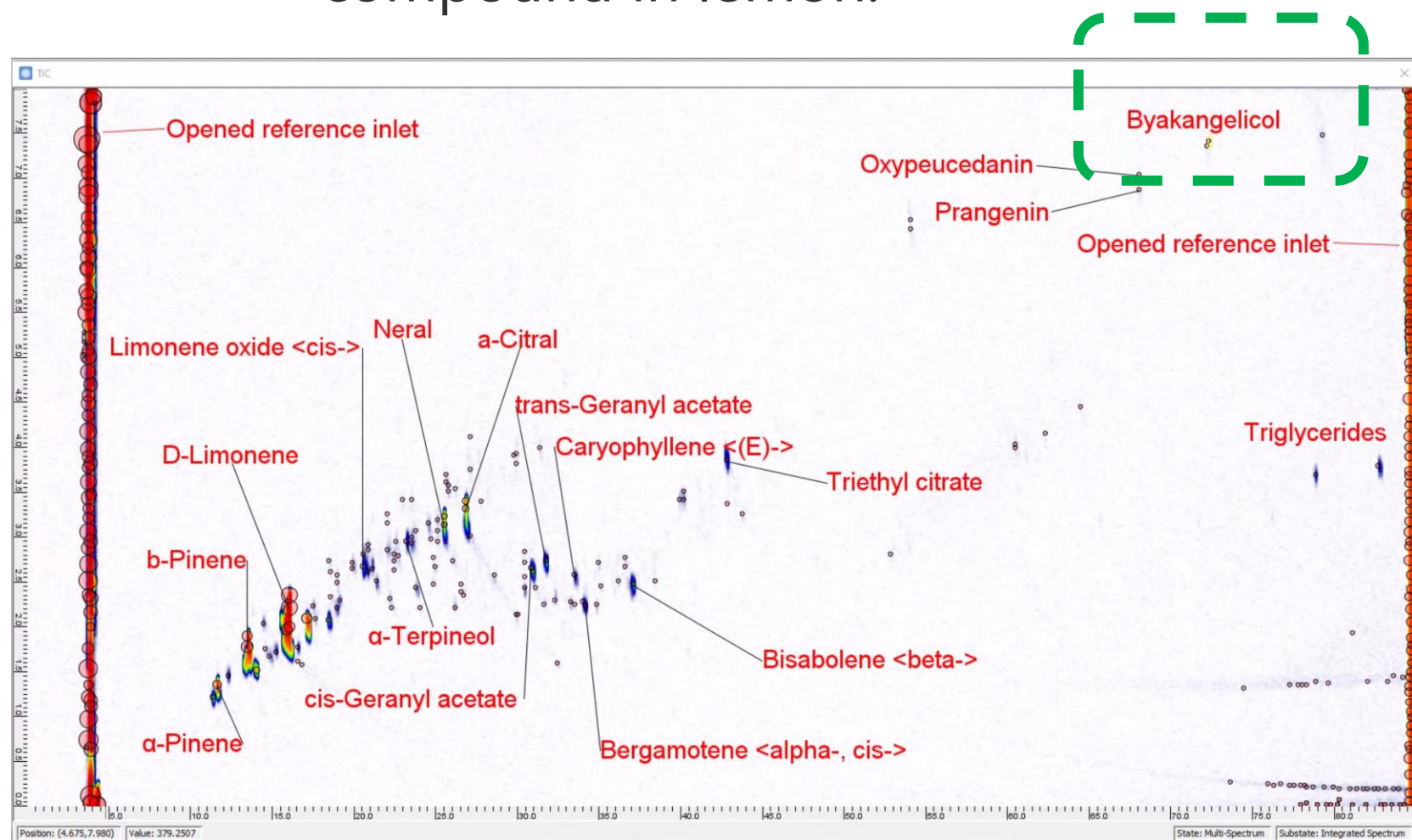
1D GC-HRTOFMS only detects one of these compounds



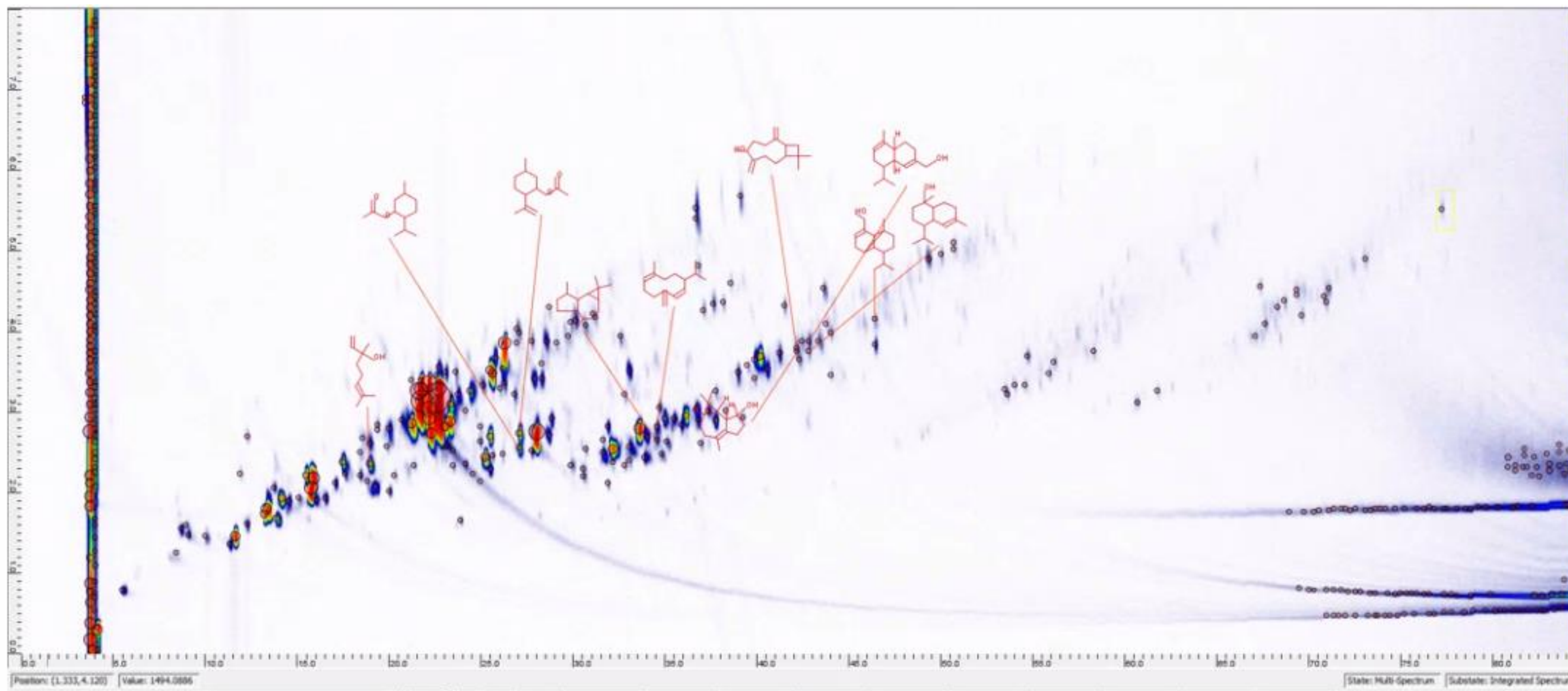


- GCxGC-HRTOFMS library search gave the correct ID as the first match.
- 1D GC-HRTOFMS gave the correct composition from the FI data. Needed to choose among DB hits.
- KNapSack search supports the presence of this furanocoumarin compound in lemon.

Metabolite Information				Structural formula
Name	Byakangelicol			 <p>zoom in</p>
Formula	C17H16O6			
Mw	316.09468824			
CAS RN	26091-79-2			
C_ID	C00053053			
InChIKey				
InChICode				
SMILES				
Start Substs in Alk. Biosynthesis (Prediction)				
Organism	Kingdom	Family	Species	
	Plantae	Rutaceae	Citrus lemon	Ref.
	Plantae	Rutaceae	Citrus limon (W)	Ref.



What 1D GC-MS does NOT detect in the Sakae peppermint oil



Structures for some compounds in peppermint oil that coelute with 1D GC.
GC x GC separates these nicely.



Summary

GCxGC

- More complex hardware setup, but stable once setup and optimized
- Longer run times – *one late-eluting compound detected in 1D GC for peppermint oil was missed in GC x GC data. (Data omitted – time constraints)*
- Most complete separation: a few compounds detected that were not detected by 1D GC. Clearly, this will be more evident for highly complex mixtures such as petrochemicals!
- Fast blob detection by GC Image and fast 2D deconvolution by AnalyzerPro XD
- Fast database search by both programs

1D GC

- Shorter run times and simpler hardware configuration
- Single quadrupole missed a few minor lavender compounds that were detected by the other methods
- *However, the high dynamic range of the Q1600 single quadrupole MS easily revealed trace differences between orange oils. (Data omitted – time constraints)*
- Chromatographic deconvolution on the HRTOF was surprisingly effective, especially when soft ionization and accurate- m/z data is incorporated.
- Chromatographic deconvolution and integrated data analysis a bit slower than GC x GC data processing



Conclusions



- Each method has its own benefits and disadvantages
- Running a 1D separation first can help to decide on optimal 2D separation conditions
- The most complete analysis came from cross-examining data from multiple methods
- Context aids in data analysis! – use information from natural products databases like KNapSack and Lotus.
- Use all the available information that you have: EI database search, soft ionization, accurate- m/z for fragments and molecular species, retention index values
- Structure tools like NIST Hybrid Search and msFineAnalysis AI structure tab can help with compounds that are not in databases.
- The >30-year-old peppermint oil showed little evidence of oxidation or degradation when compared to the fresh oil.