

GCT-MS applied to the food and beverage analysis

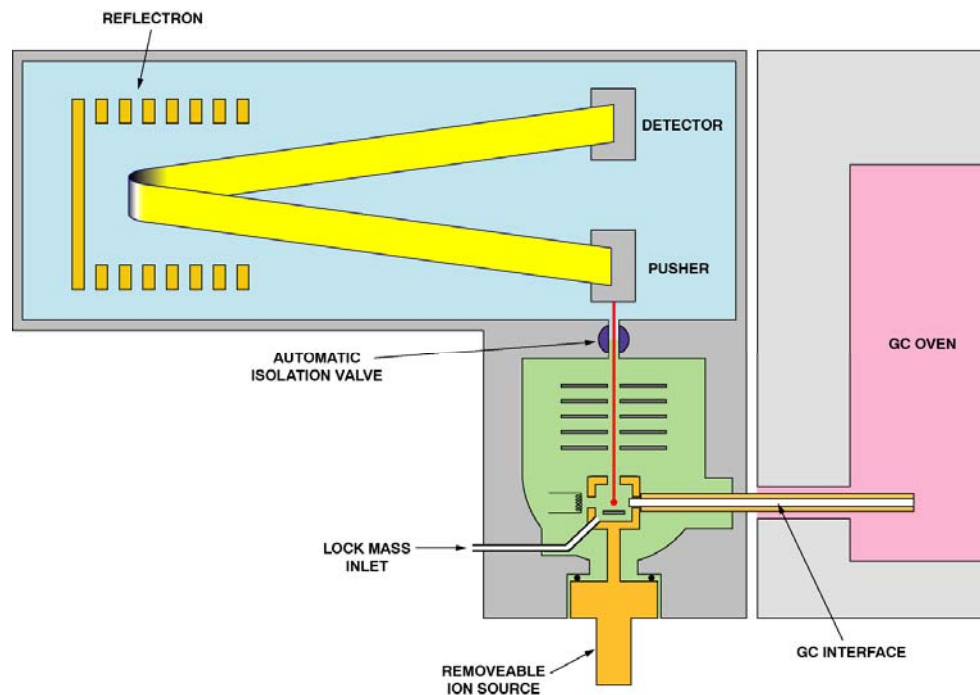
Universidad de Murcia, 17 Junio 2008

- Exact mass TOF
 - Selectivity / Mass accuracy
 - Dynamic Range Enhancement
- Screening in food: Analytical challenges
 - Targeted screening versus untargeted screening
 - Quantification of 'known' compounds
 - Deconvolution, identification
 - Detection of a non-targeted residue
- Flavour profiling: Beer analysis

GCT Premier - Features Orthogonal Acceleration TOF/MS

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- Plug in EI/CI ion volumes for rapid change over
- Resolution >7000 (Full width half maximum)
- Exact mass less than 5ppm
- 4 orders linear dynamic range
- Spectral acquisition rate of 20 spectra per second



GCT Premier - Benefits Orthogonal Acceleration TOF/MS

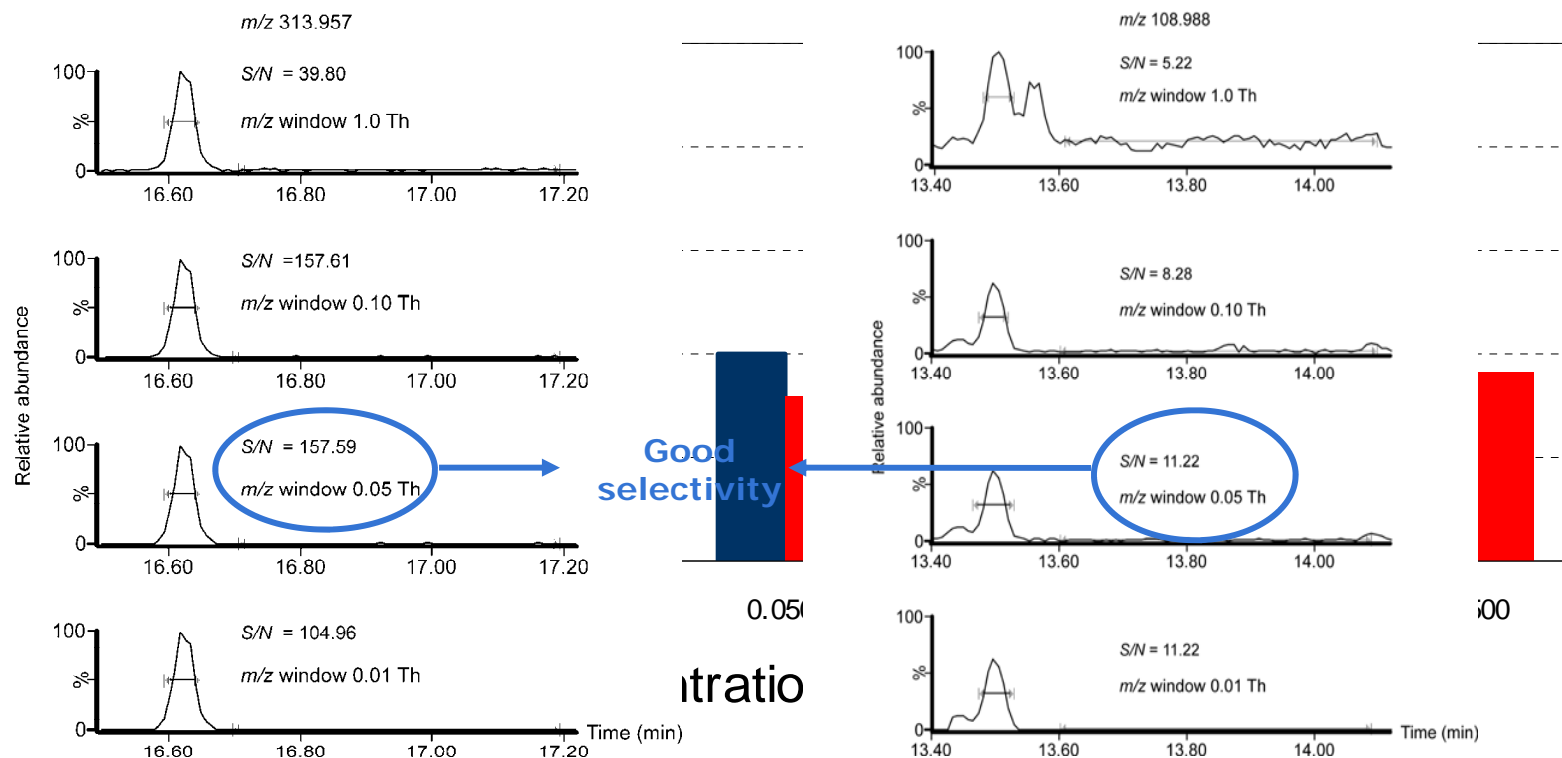
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- Rapid full spectral acquisition rates
 - Compatible with narrow GC chromatographic peaks
 - Produces Non-skewed spectra
 - Enables accurate chromatographic peak profiling and deconvolution
- Non scanning instrument with a high duty cycle
 - High full scan sensitivity
- Produces EI library searchable spectra
 - Allows compound identification using standard or user libraries
- Spectra recorded with accurate mass
 - Enables confirmation or elucidation of structure
- Elevated Resolution
 - Enhanced selectivity



Orthogonal Acceleration TOF High resolution = Selectivity

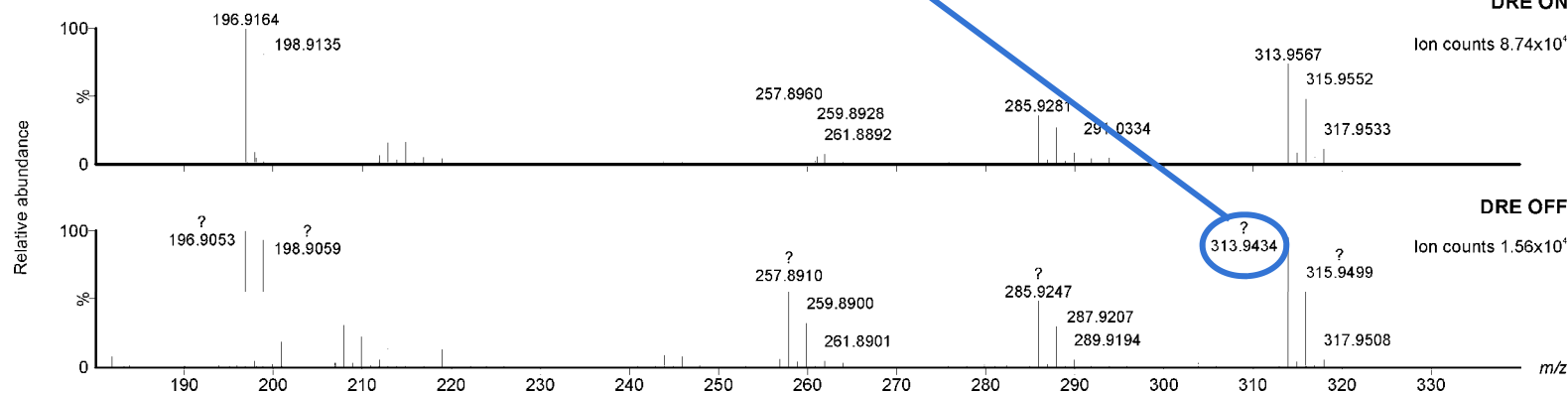
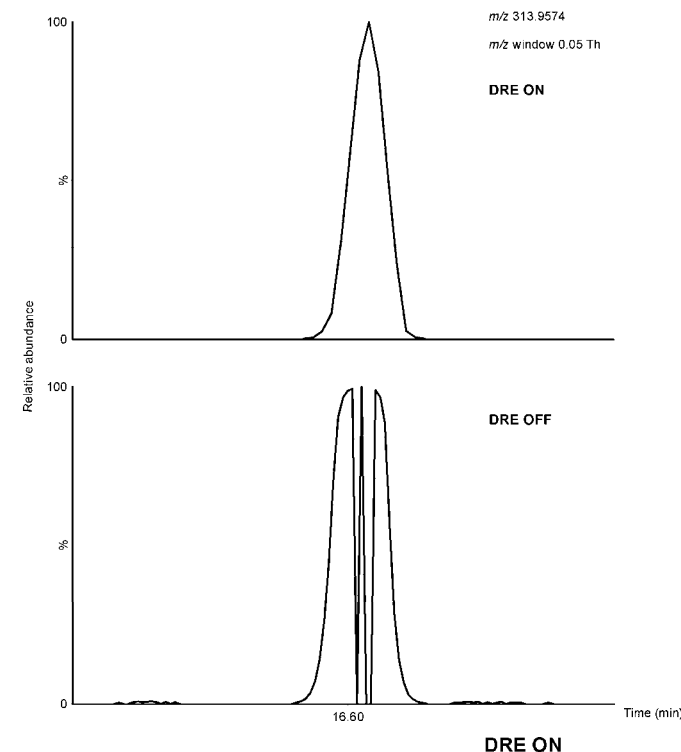
- Elevated mass accuracy allows for greater selectivity
 - Mass accuracy in fruit based baby food matrix matched standards
 - Chlorpyrifos (m/z 313.9574) and Fonofos (m/z 108.9877) in fruit based baby food
 - 0.05 Th is the routine mass window applied



Orthogonal Acceleration TOF DRE

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- DRE offers enhanced dynamic range
 - Chlorpyrifos in lettuce
 - Reduced saturation
 - With DRE off too many ions arrive at the detector simultaneously
 - Response falls outside mass window
- Enhanced mass accuracy
 - ? = Saturated peak, Mass error = 14 mDa



Screening in food: analytical challenges

- Definition as “an examination, usually methodically, in order to make a separation into different groups”
- In chemistry we refer to a rapid check of a sample for a large number of analytes, most of which will result in a negative result.
- Identification and monitoring of residues in different commodities.
 - Food safety
 - Pesticides, veterinary drugs, banned additives...
 - Environmental
 - Drugs of abuse, endocrine disruptors...
 - Impurity control

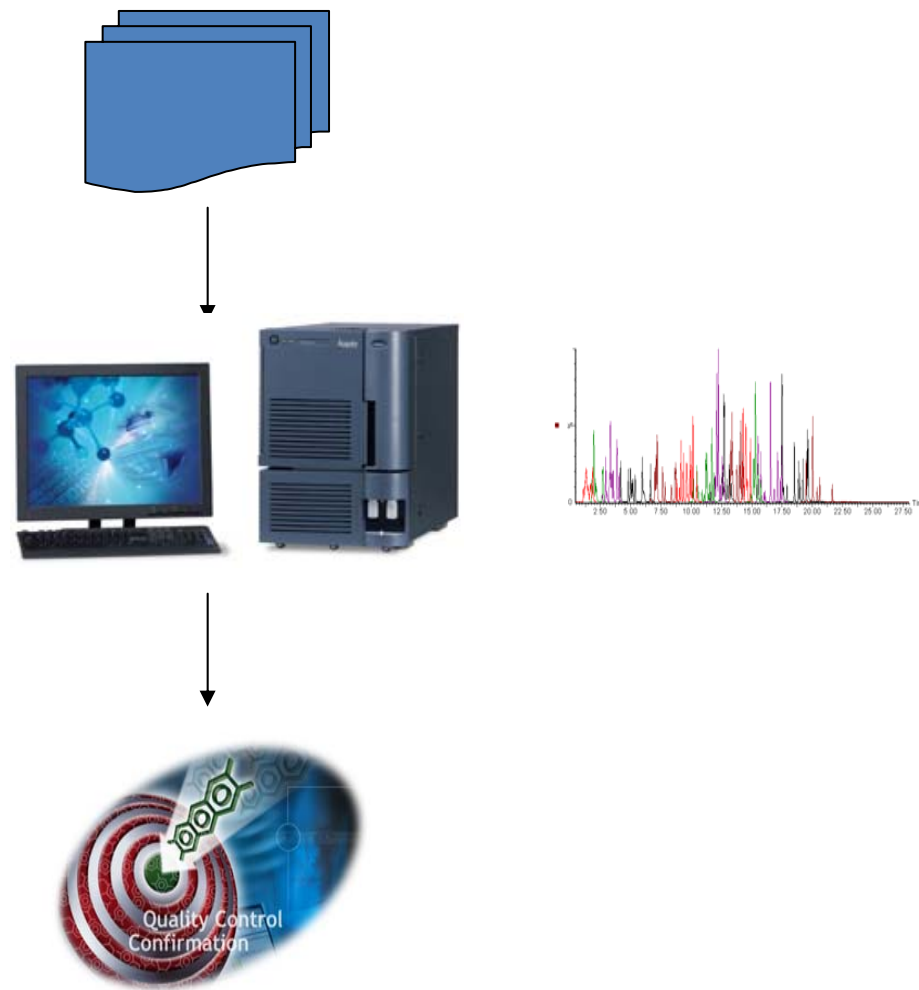
- **Screening**: to (rapid) detect the presence of a compound
 - Sample positive for an analyte? → **YES** or **NO**
 - Typically a second analysis is required to:
 - Correctly quantify or
 - Confirm the identity of the compounds detected
- **Confirmation**: to confirm the identity of the compound detected
- **Quantification**: to give information (accurate) on the analyte concentration in sample
- **Elucidation**: to discover (elucidate) the identity of a compound detected that is not a target analyte
 - Non-target analysis.

- **Challenge #1: Pesticide residue methods must cover a very wide range of analytes and matrices**
- Numerous chemically and structurally diverse analytes exist
 - Approximately 1000 pesticides registered worldwide
- Wide variety of matrices (fruits, vegetables, herbs, spices) need to be monitored
- Over 17,000 EU community MRLs have been established for about 140 pesticides.

- **Challenge #2**: High demands are placed on analytical method performance to meet legislative requirements

- Methods must be...
 - **Sensitive**, low default reporting limit of 0.01mg/kg
 - **Selective**, reduce or eliminate matrix interferences
 - **Multiresidue**, multiple targets in a single run
 - **Rugged**, complex samples with reduced or no sample clean-up
 - But also **generic** (ie. Having broad scope)...
 - 'Food scares' and changes in legislation require updating of methods and re-analysis of samples

- First the analytes to be screened are selected
- Data acquisition
 - ACQUITY – TQD
 - ACQUITY – QP XE
 - GC-MS/MS
- Data processing
 - TargetLynx



Strengths

- High sensitivity
 - MRM in triple quadrupoles
- High selectivity
 - Providing the MRM transitions are specific
- High dynamic range
 - Quantitation “gold standard”
- Legislation compliant
 - One trace for quantitation and one for confirmation

But...

- Sensitivity dependent on the number of analytes
 - Duty cycle vs. # transitions
- Not capable of reprocess historical data
 - New target needed, the analysis needs to be repeated.

Post-target screening

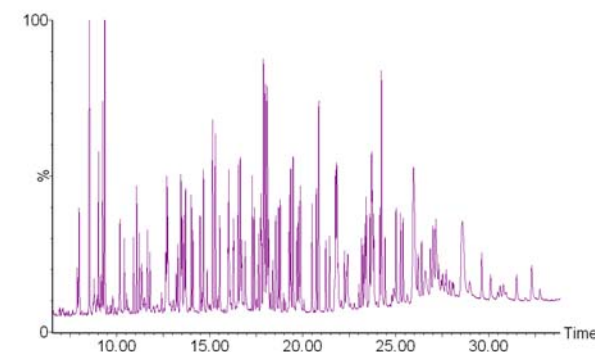
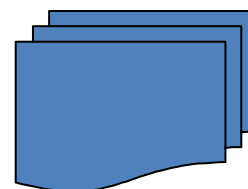
1. Data are acquired

- LCT Premier
- GCT Premier



2. Analytes to be screened are decided

- templates



3. Data processing

- TargetLynx



Strengths

- High sensitivity in full scan
 - Duty cycle not affected by # of analytes to screen
- High selectivity
 - Exact mass
 - Possibly retention time
- Dynamic range
 - DRE feature
- Capable of reprocessing historical data
 - Data are acquired, is just a matter of processing

But...

- Sensitivity less than most tandem quadrupoles in MRM mode
- Dynamic range less to the one of triple quadrupoles
- Legislation does not recognise exact mass as a proof of identity
 - Presumptive positives need to be submitted for confirmatory analysis be repeated.

Analytical Challenges

- New approaches with ToF/MS

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- Time-of-Flight MS (with both GC and LC inlets) is currently creating much interest for pesticide residue analysis.
- ToF/MS can support development of more generic screening methodologies
 - Can detect both target and non-target compounds using ion ratios and/or spectral matching to increase confidence
 - Very large numbers of residues may be determined in a single analysis
 - Presumptive positives can be submitted for confirmatory analysis by MS/MS if desired
 - Automated software processing
 - Ability to re-process historical data

System Solution

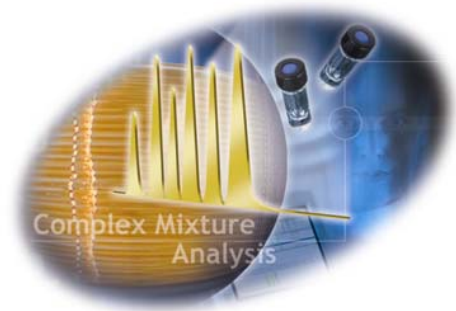
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GCT Premier



LCT Premier XE



ChromaLynx
Peak detection, deconvolution and library searching



TargetLynx
Advanced quantitation and automated QC

Application Example 1

Use of GC-ToF/MS for screening of pesticides residue in fruits, vegetables and baby foods

Sample Preparation and GC-TOF parameters

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- Extraction of the baby food, lettuce and pear samples
 - Modified QuEChERS extraction



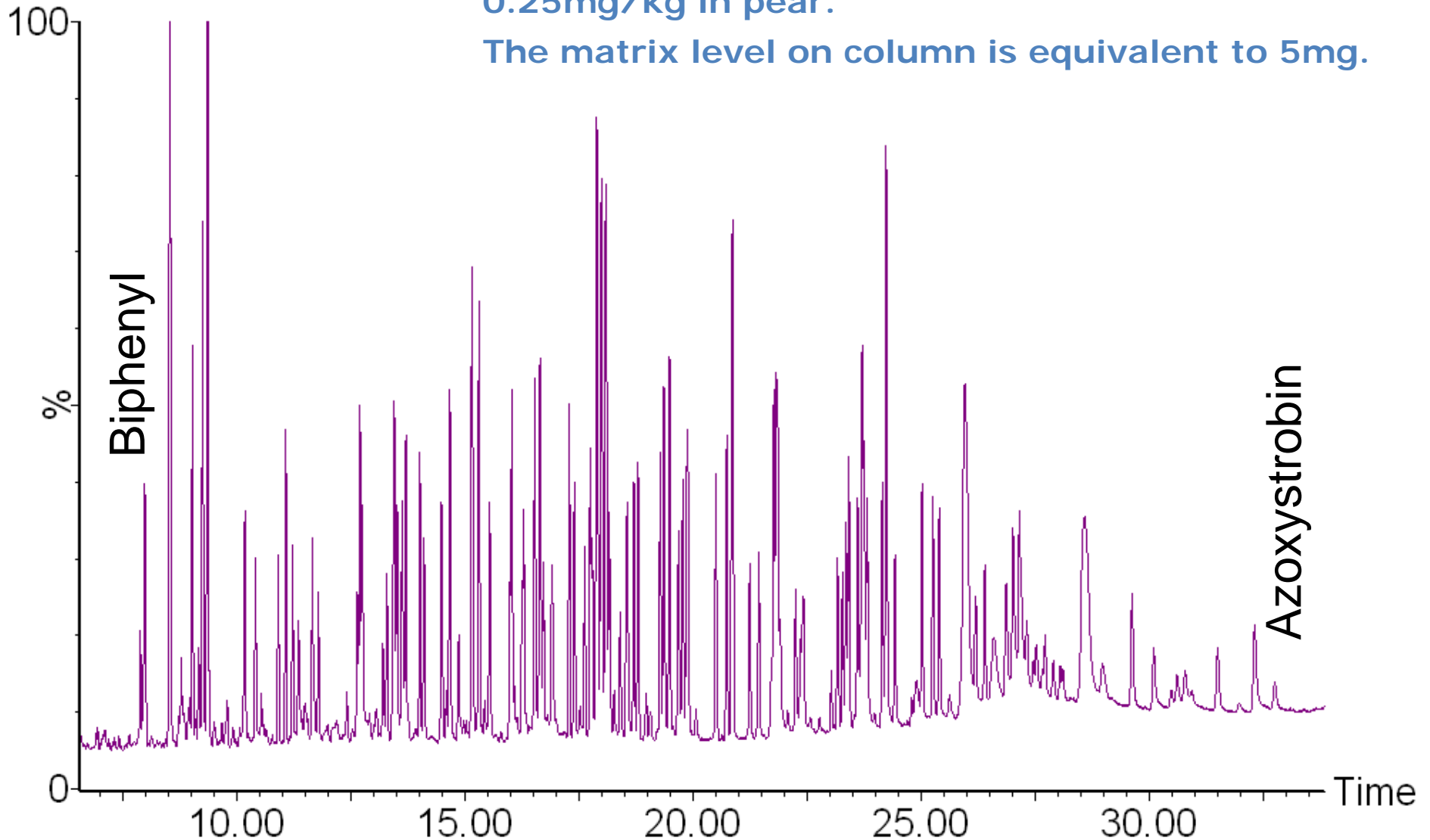
- Electron Impact (EI+) with a mass range of m/z 50 → 500
- Acquisition Speed = 4 spectra / s
- Dynamic Range Enhancement (DRE) On
- 4GHz TDC
- Pusher interval 35 μ s
 - 28571 raw spectra s^{-1}
- RXi-5ms GC column
 - 5 μ L PTV injection
 - Simple oven temperature ramp

Total Ion Chromatogram (TIC)

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The residue concentrations are equivalent to
0.25mg/kg in pear.

The matrix level on column is equivalent to 5mg.



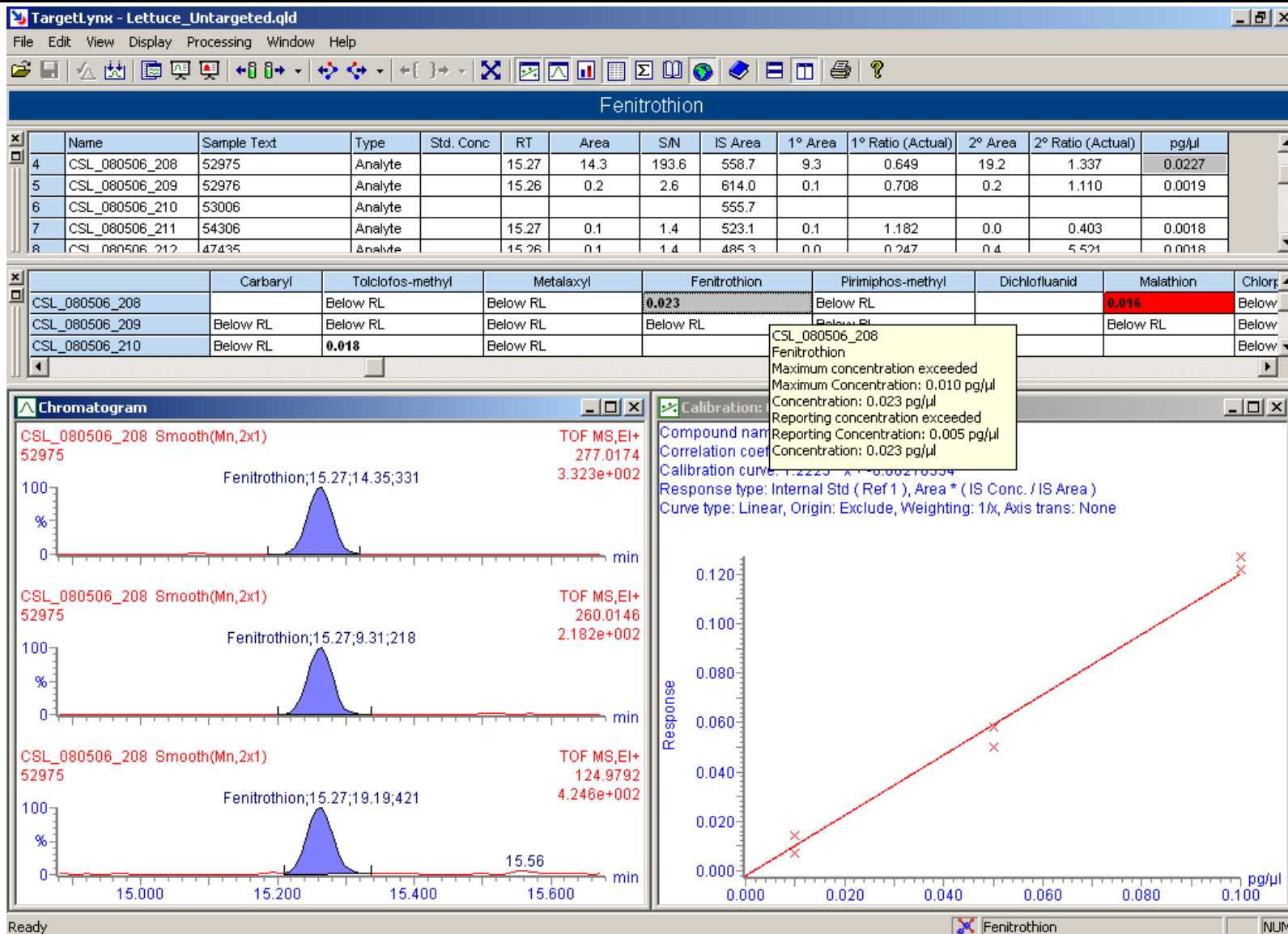
Targeted Screening GC-TOF/MS and TargetLynx

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- Compound list is known
- Detection is based on extracting exact mass chromatograms with 0.02 Da windows
 - Any ion can be chosen for detection and/or quantification
 - Number of residues/ions can be increased without loss in sensitivity
 - *Post-acquisition addition* of target compounds is possible
- Screening is based on one or more exact mass chromatograms
 - If a peak is found within a defined time window the target compound is detected
 - Ion ratios can be used to increase confidence in detection
 - Quantitation against standards allows presumptive positives to be identified
- TargetLynx application manager
 - Highlights samples with concentrations above the reporting level
 - Highlights samples that do not pass quality control criteria

Incurred Residue in Lettuce Fenitrothion – 0.023mg/kg

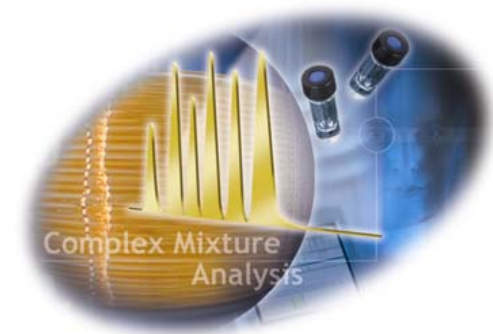
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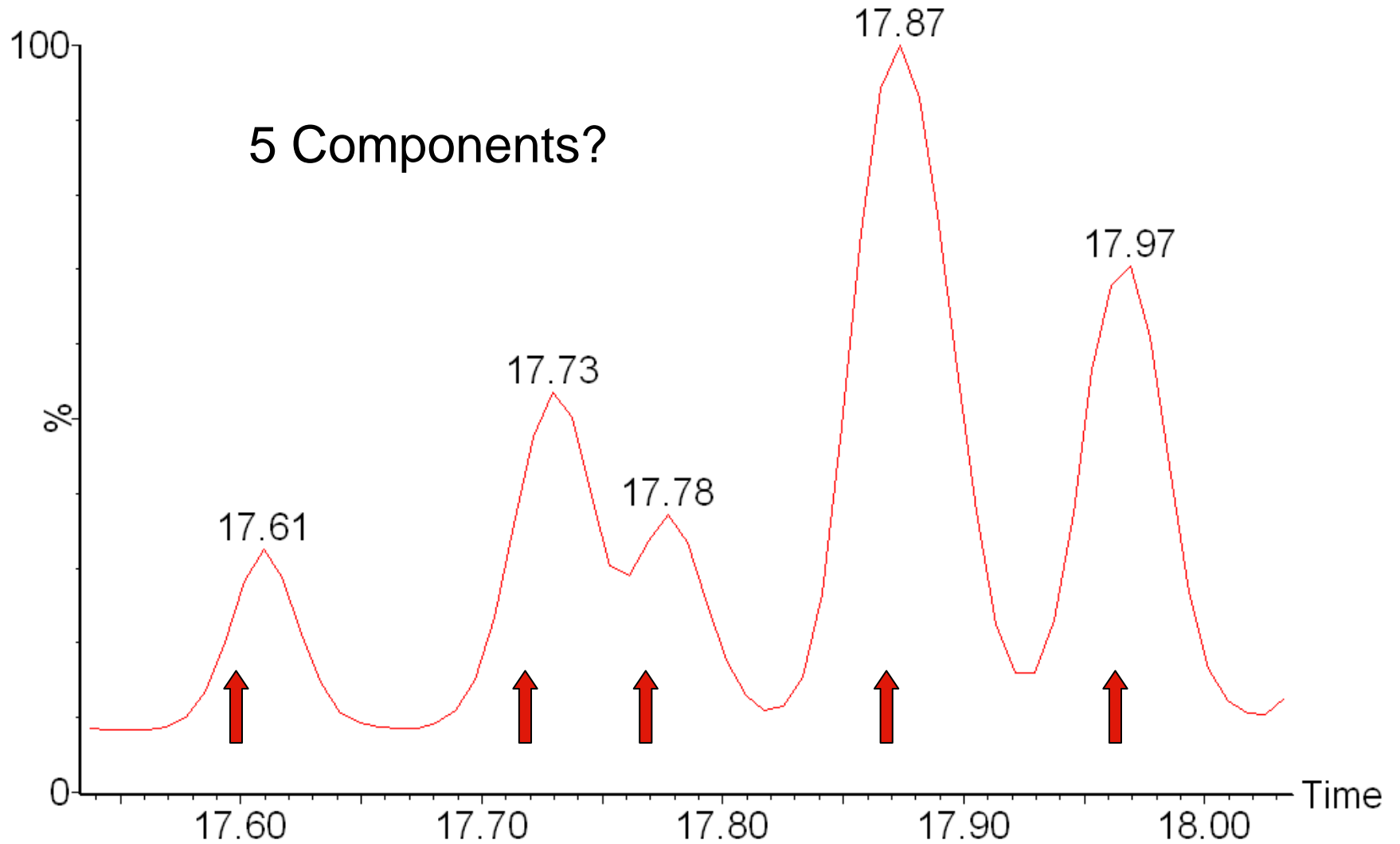
Non-Targeted Screening GC-TOF/MS and ChromaLynx

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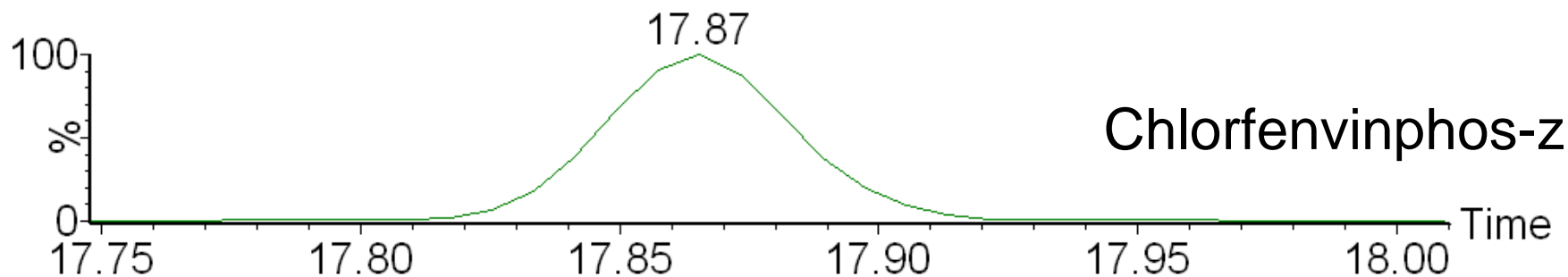
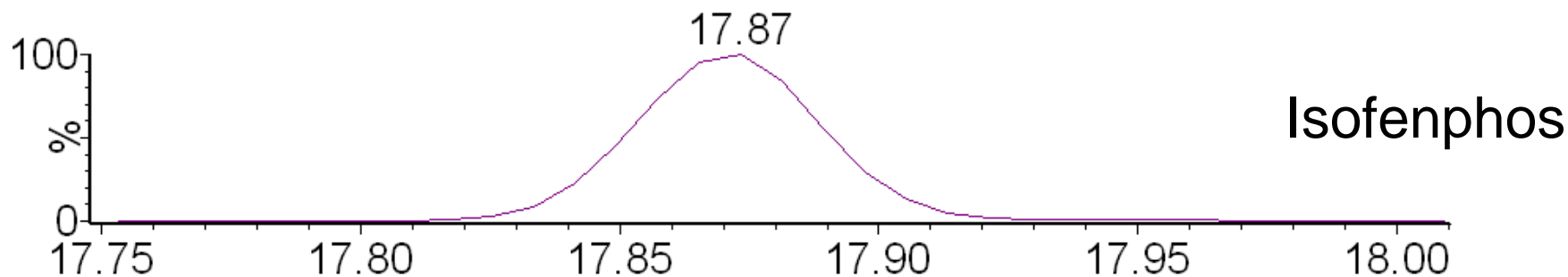
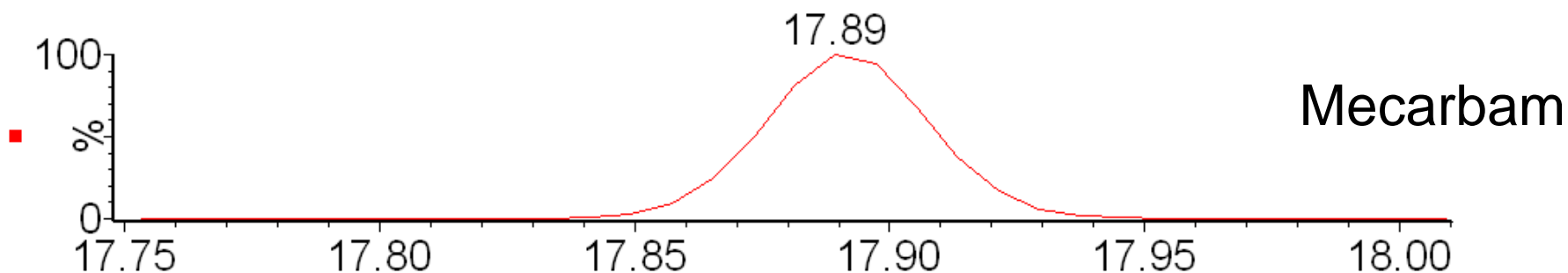
- Compound list is unknown
- ChromaLynx application manager
 - Automatic peak detection and deconvolution
 - Automatic production of “clean” component spectra
 - Standard EI spectra are generated by the GCT source
 - Automatic identification via library searching
 - Commercial or user generated libraries may be used
 - Exact mass scoring provides additional confirmation of identity
 - Exact mass spectra can facilitate structural elucidation if component is not in the library



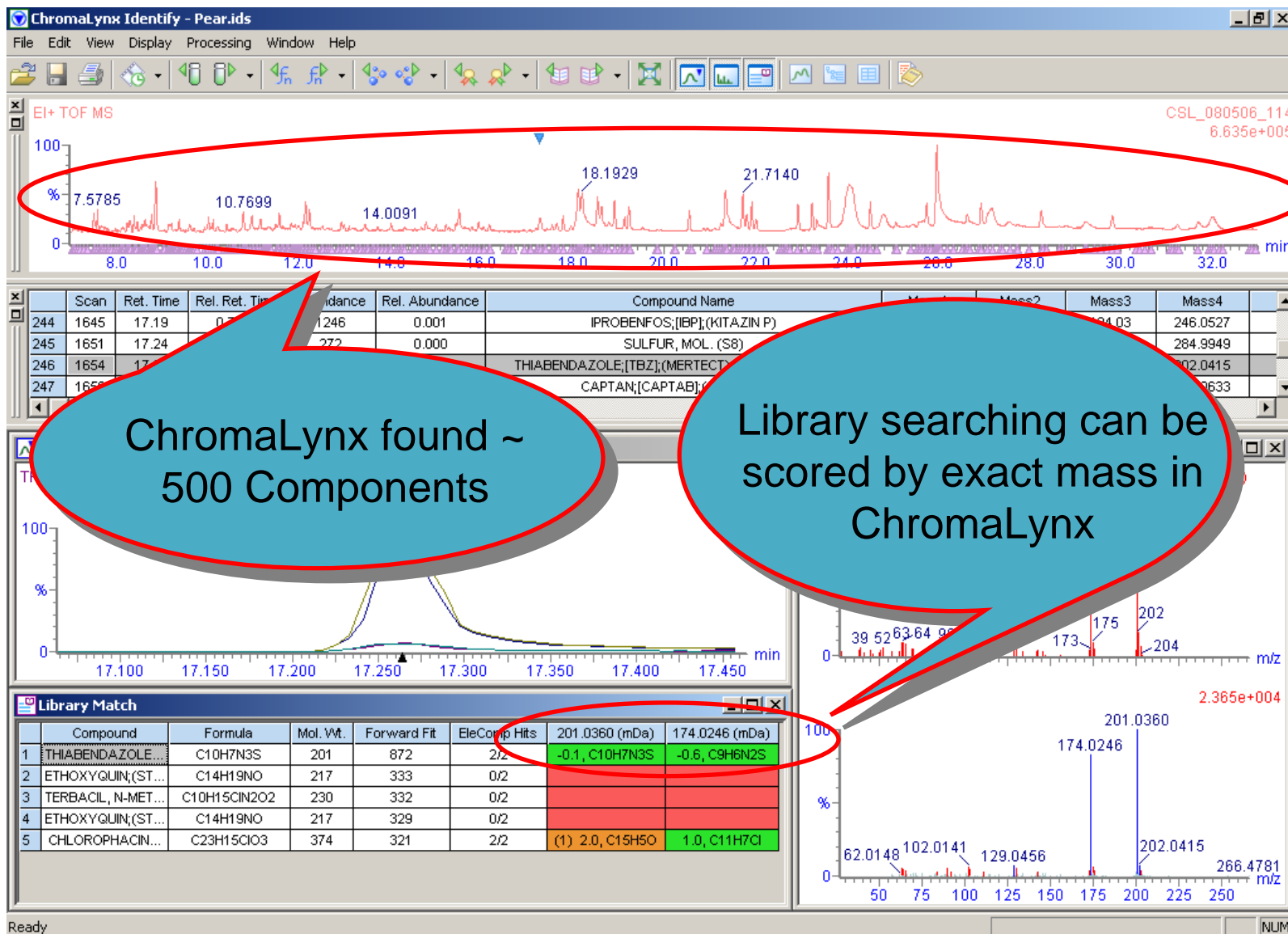
Deconvolution



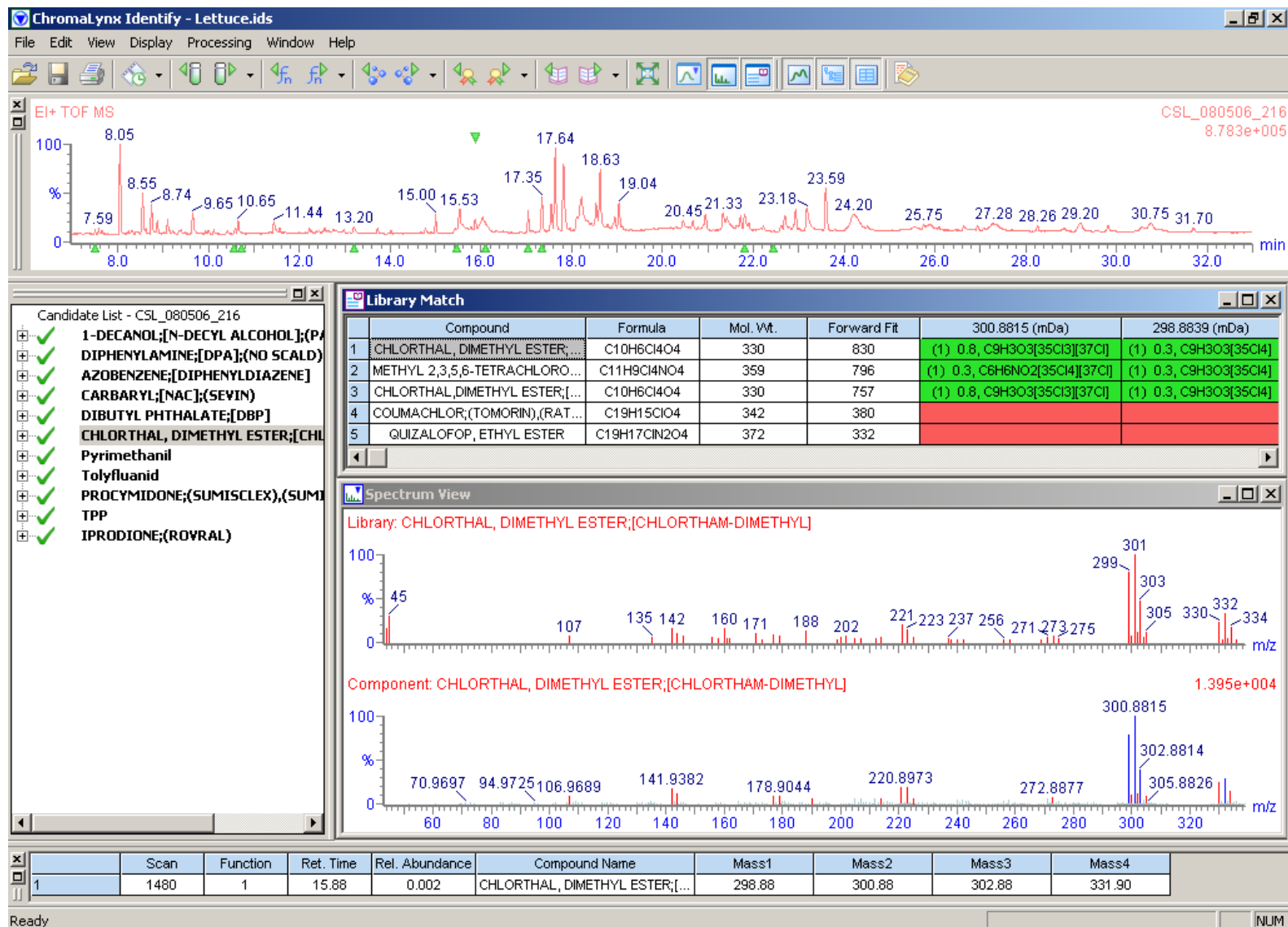
Extracted Mass Chromatograms 3 Components



Pear Extract - ChromaLynx



Example of Untargeted Screening DCPA in Lettuce



Application Example 2

Profiling of different beers

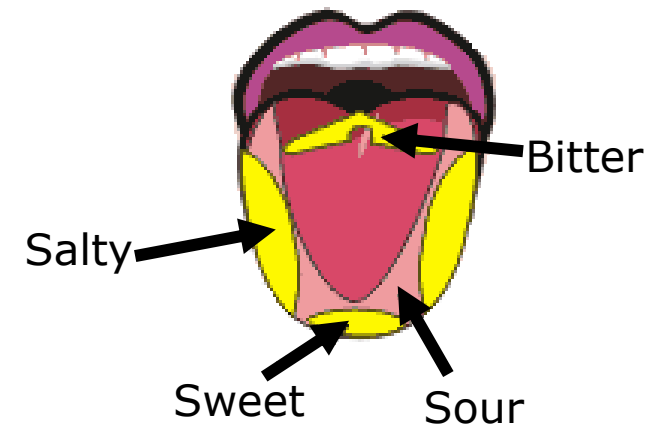
Importance for the Food Industry

Challenges for Analytical Chemists

- Some of the challenges for scientists in the food industry
 - Identifying **process changes** that may lead to a change in quality
 - Detecting **adulteration** in any ingredient
 - Identifying the **geographical origin** of raw materials
- Challenges for scientists in regulatory agencies
 - Detecting economic fraud due to product substitution and adulteration, as well as health risks from possible contamination
- These QC issues are traditionally assessed by experts
 - Intensive training to be able to determine a product's quality

Taste

- Occurs on the tongue
- Detects 4 primary tastes
- Mouthfeel (e.g. viscosity, smoothness)
- Pungency



Smell

- Occurs in the nose
- Detects 32 primary tastes e.g. vanilla, smoky, cereals, fruity, sulphury, floral

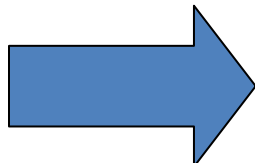
Appearance

- Colour
- But sometimes altered by the addition of caramel

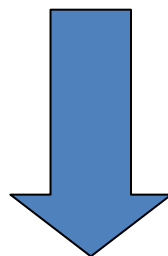
**Experts can distinguish
between different brands &
can often identify from which
geographical location they
come from but can we do this
by MS?**

Work Flow

Collect samples

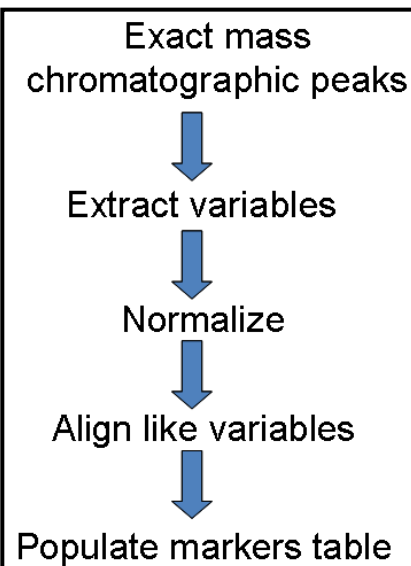


Analyse samples by UPLC/MS and GC/MS



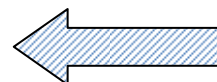
Multivariate Analysis

Process Data



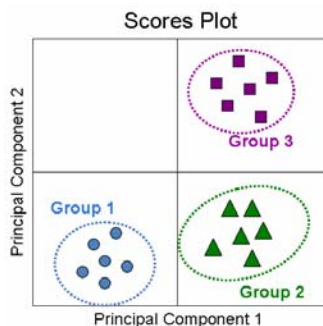
GC/MS = NIST
UPLC/MS = online database searching

Identify Results



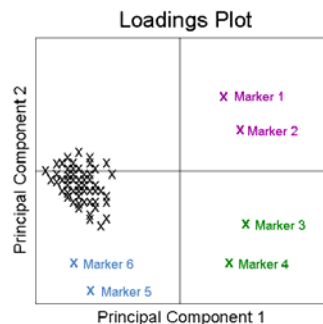
Statistical Analysis Scores and Loadings Plot

- PCA is an unsupervised technique that enables elucidation of main sources of variability in datasets, detects clustering formation and enables identification of 'markers' (t_R , m/z pairs)
- Scores plot



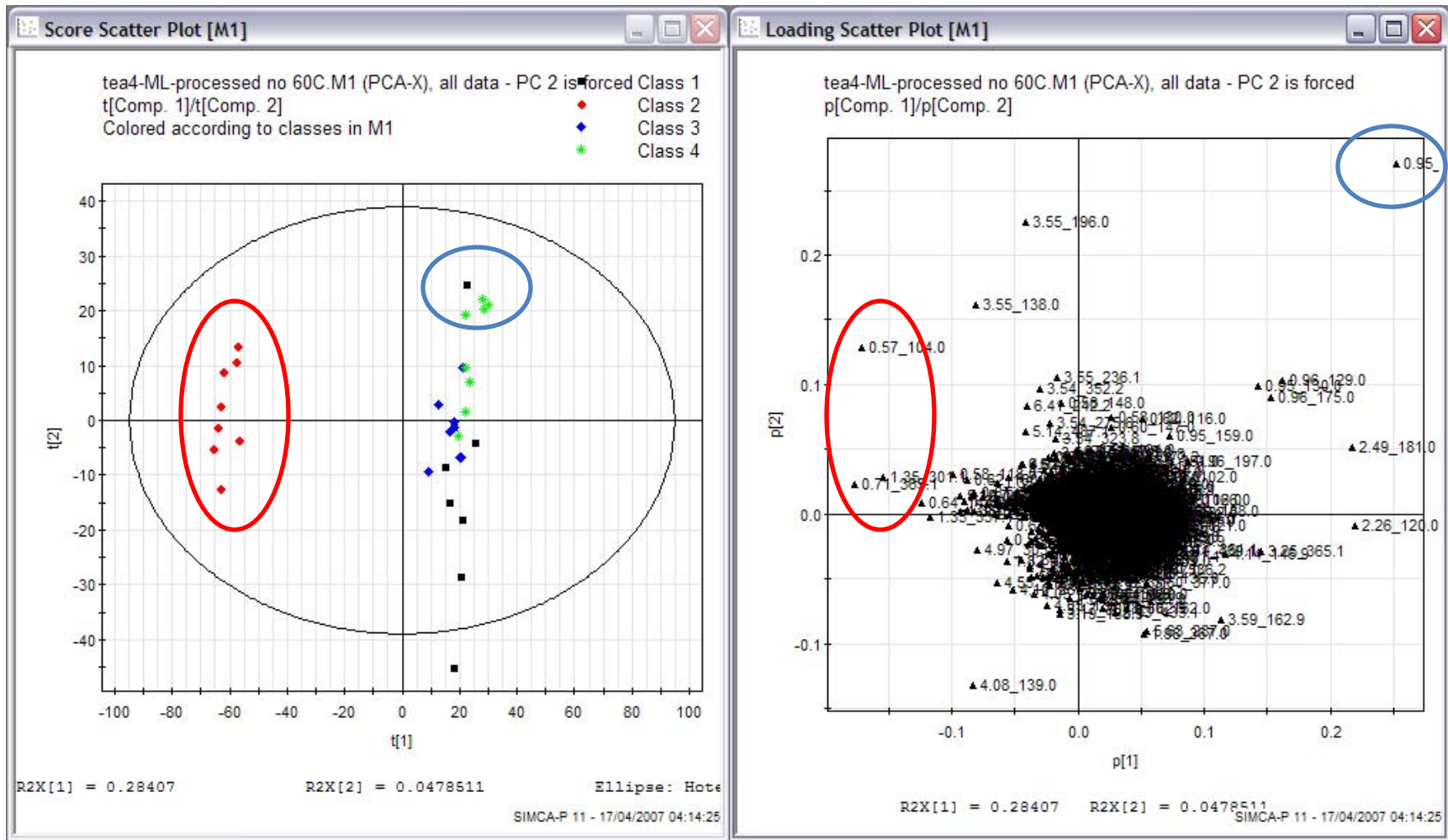
Two principal components make up a plane. When points are projected onto the plane similarities between objects are described

- Loadings plot



Describes the variables relationships and can interpret the scores plot by telling which variables are responsible for similarities

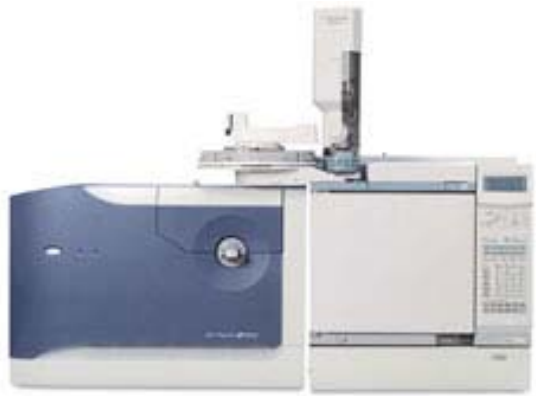
Scores and Loadings plot



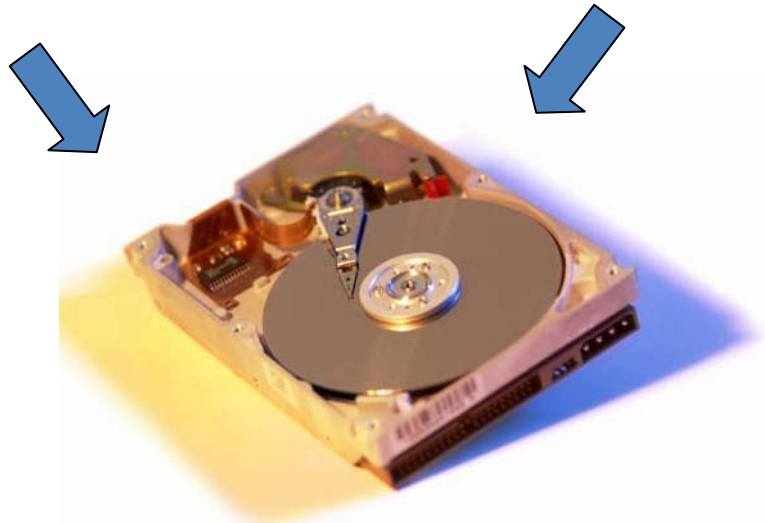
Data Acquisition

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GC/MS



LC/MS



Aims of the experiment

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- Instrument: GCT
- Ten samples:
 - Nine commercially available beers
 - One home-made beer
- Mix of dark and light beers, from different geographical locations

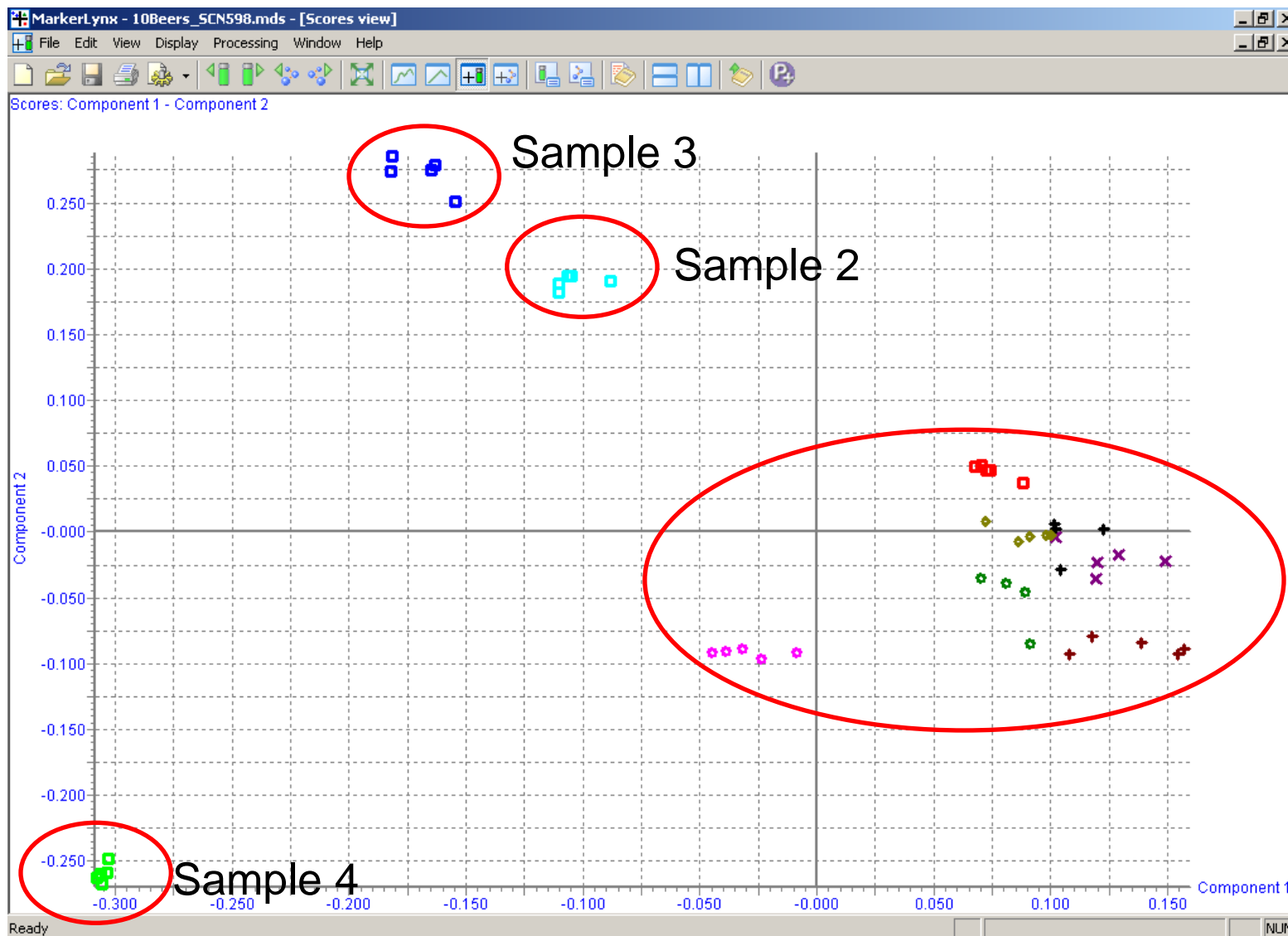


Questions:

- Do any of the beers have markers that identify them from the other beers
- How similar are the brands to one another
- Is there anything else that we can learn about the beers

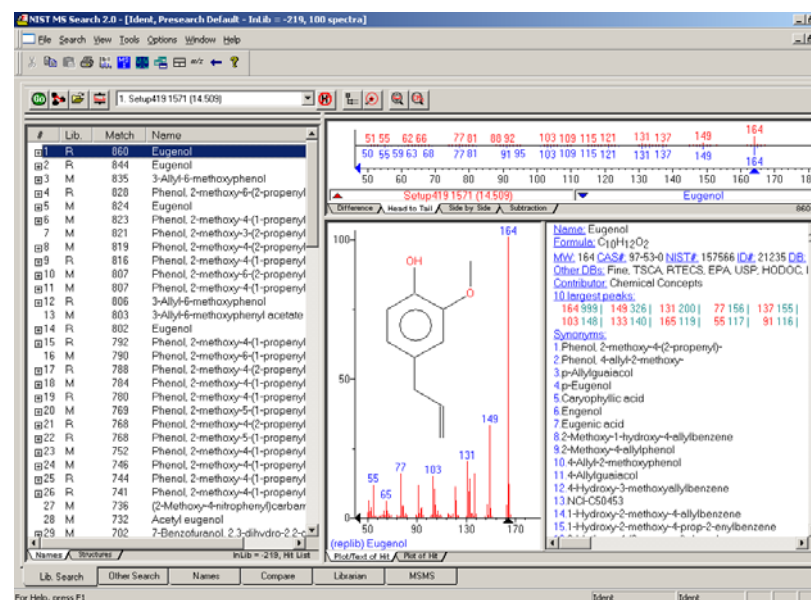
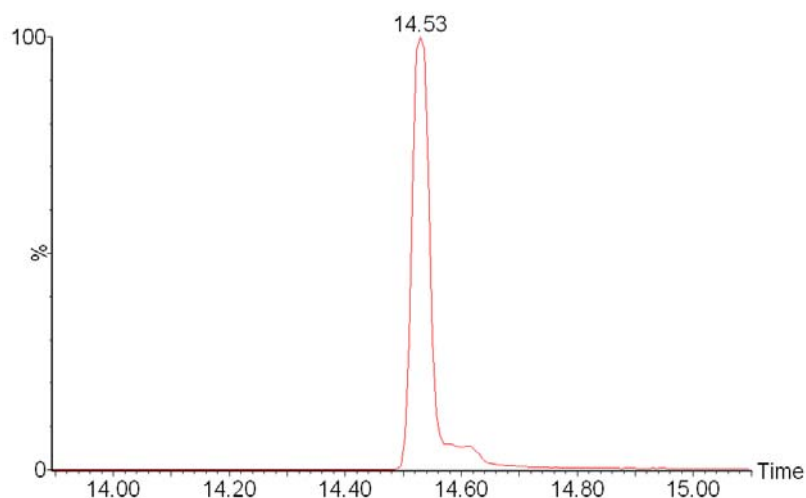
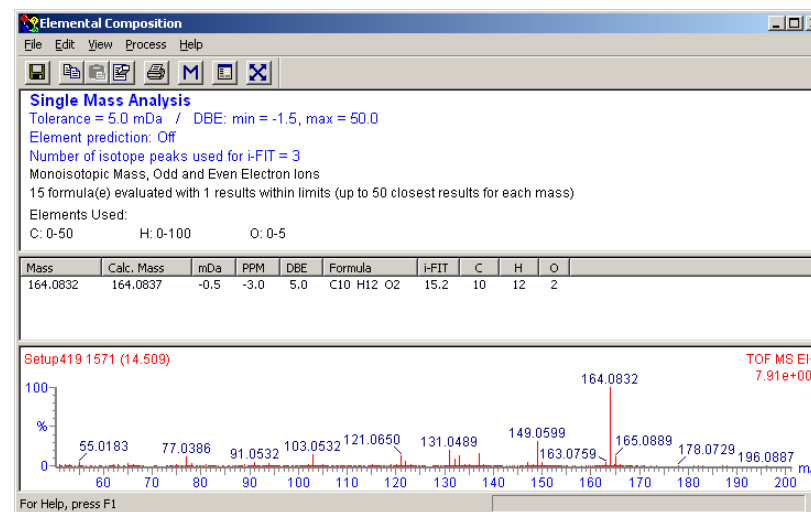
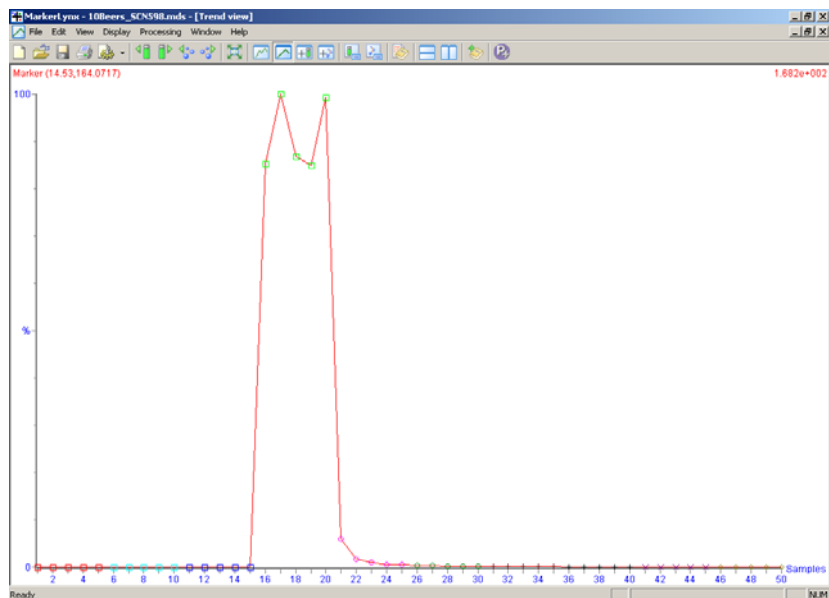
Scores Plot View: 10 Beers

3 Outliers



Library Search: 10 Beers

$R_t = 14.53$, $m/z = 164.0717 \equiv$ Eugenol



10 Beers Marker Identification

RT	<i>m/z</i>	Sample		Name	Formula
14.53	164.0717	4	Decreasing significance ↓	Eugenol	$C_{10}H_{12}O_2$
10.64	93.0674	2, 3		3-Carene	$C_{10}H_{16}$
17.34	85.0231	3		Persicol	$C_{11}H_{20}O_2$
9.86	70.0744	3		Isoamyl butylate	$C_9H_{18}O_2$
16.03	85.0259	3		Decanolactone	$C_{10}H_{18}O_2$
12.25	93.0689	2, 3, 4		Terpinyl acetate	$C_{12}H_{20}O_2$
15.13	178.0976	4		Methyl eugenol	$C_{11}H_{14}O_2$
9.39	93.0691	4		Limonene	$C_{10}H_{16}$
8.71	71.0478	3		Butyl butylate	$C_8H_{16}O_2$
8.02	106.0404	3		Benzaldehyde	C_7H_6O

- GCT is a powerful, versatile technology for a wide range of applications

- ToF are by principle an unbiased technology
 - Open data
 - Post-target analysis
 - Require POWERFUL SOFTWARE tools to mine the data

- GCT can be successfully used for
 - Screening of compounds
 - Profiling of commodities - NIST libraries facilitate the identification of markers