



# Tracing the origin of Lambrusco sparkling wines profiling the “aroma” by GC-QTOF

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# Overview



Introduction to  
Food Profiling



Authenticity



Food Degradation



Process Development



# Introduction to Food Profiling



## Standard Food Testing Applications with Mass Spec:

- Contaminant Identification and Quantitation
  - Pesticides
  - Mycotoxins
  - Marine Toxins
  - Veterinary Drugs
  - Adulterants (Melamine)
  - And others

*... but what are we missing?*

# Introduction to Food Profiling

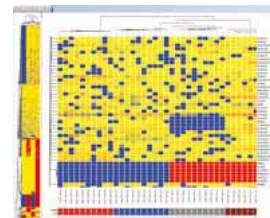


The questions contaminant testing won't answer:

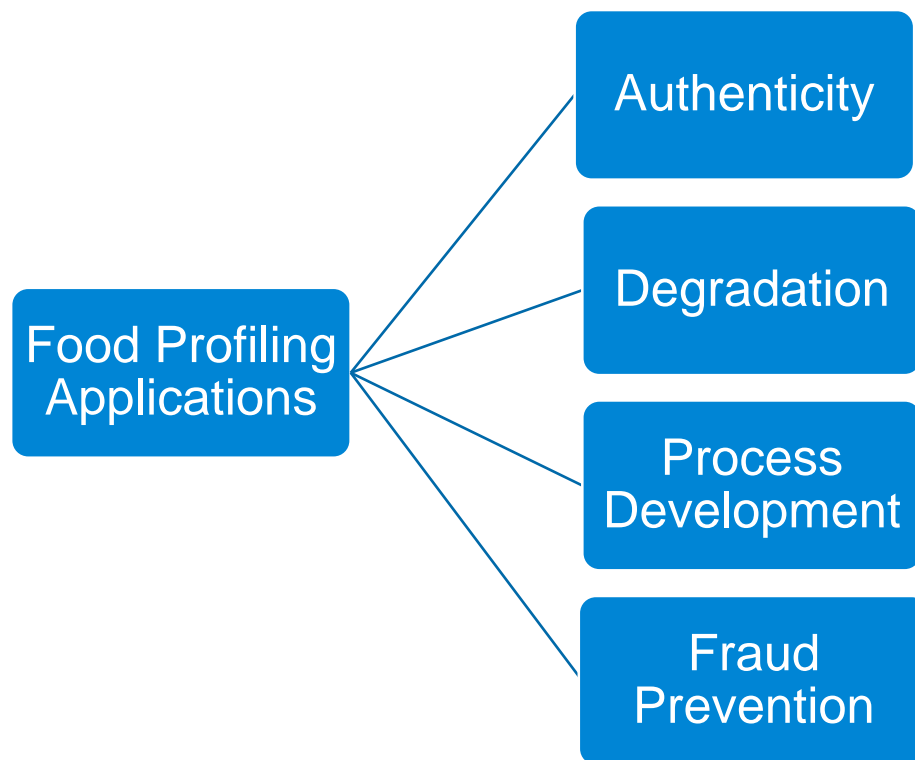
- Is this wine cabernet or pinot noir?
- Is this wine label true?
- Is this wine a real DOC wine?
- How did changing my growing or fermentation process affect my food product?

To answer these questions, we need  
**Food Profiling**

# Agilent Food Profiling Solution



## Q-TOFs and Mass Profiler Professional (MPP)



Food Profiling (def.) - *The processing of mass spectral data from two or more food samples in order to highlight the compounds which have a statistically different concentration.*



# The Agilent Profiling: Chemometric Workflow

Separate &  
Detect

Feature  
finding  
quantitate

Alignment &  
statistics

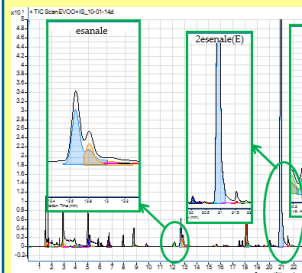
Identify

Pathways

GCMS



GC-QTOF

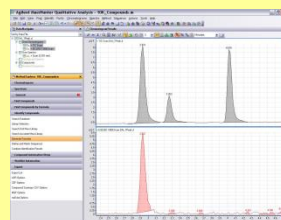


Deconvolution

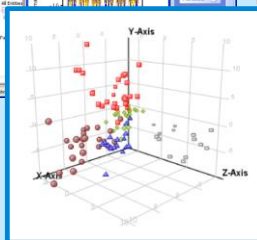
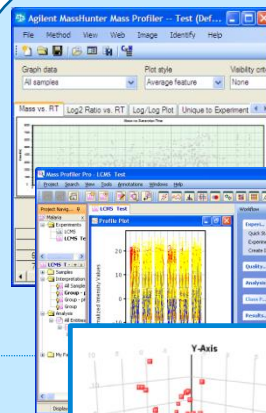
LCMS



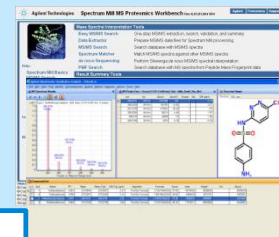
LC-QTOF



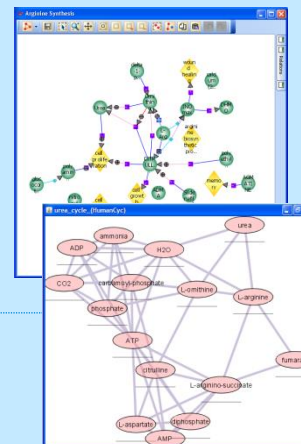
Molecular Feature  
Extraction



Mass Profiler  
Professional



ID Browser



Pathway Architect  
Cytoscape

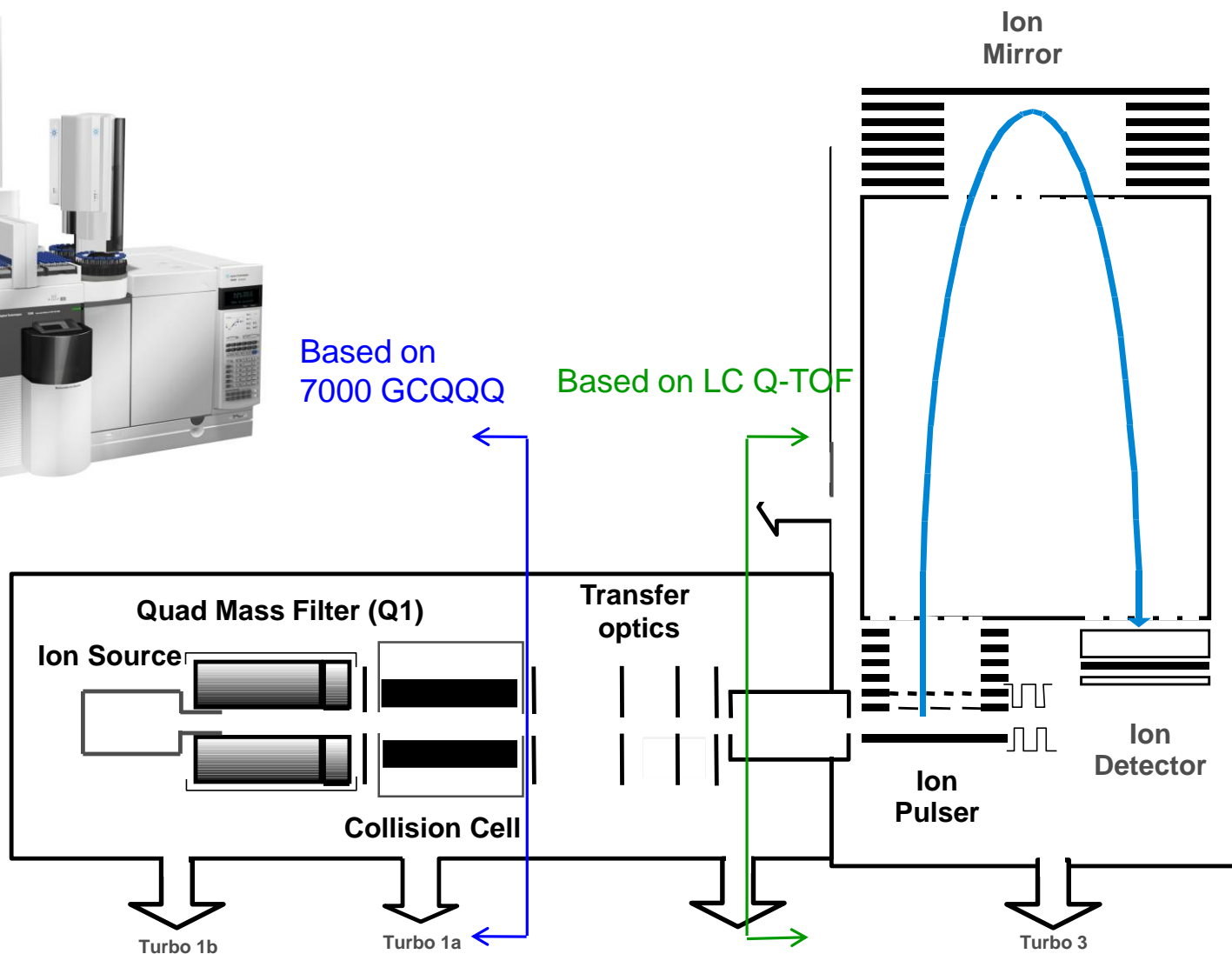


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# Agilent 7200 GC-QTOF



# What is the GC-QTOF



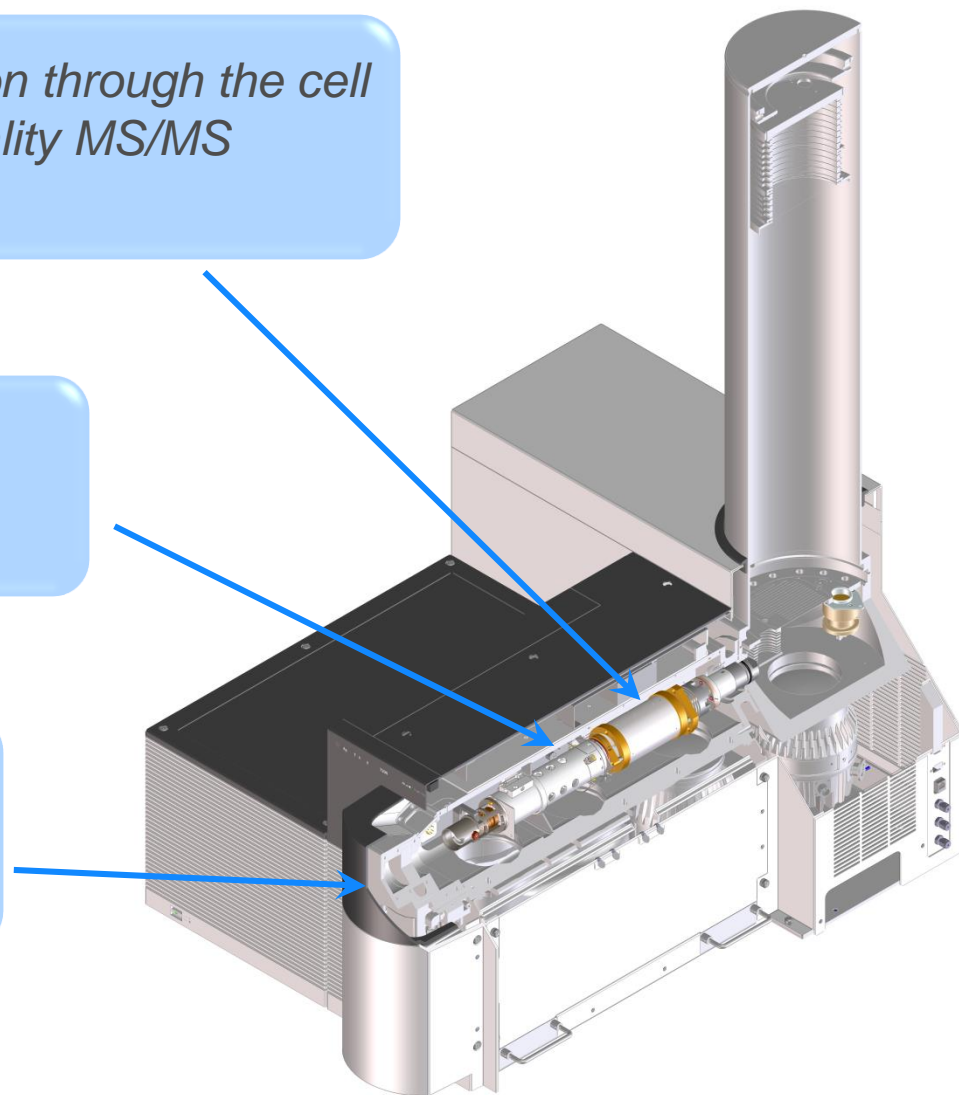


# 7200 GC/Q-TOF

**Hexapole collision cell** accelerates ion through the cell to enable faster generation of high-quality MS/MS spectra without cross-talk

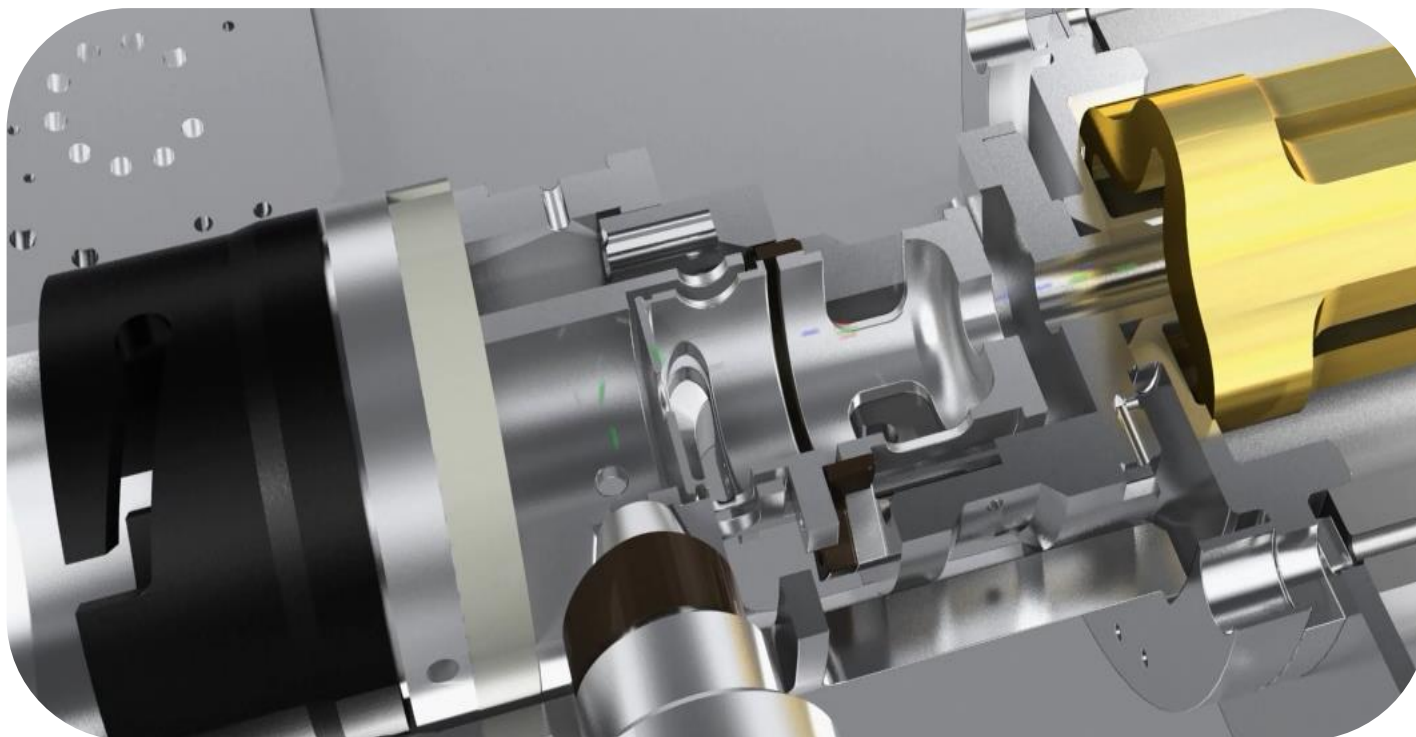
**Hot, quartz monolithic quadrupole analyzer** identical to the 7000 Quadrupole MS/MS

**Removable Ion Source** includes repeller, ion volume, extraction lens and dual filaments



# New Removable Ion Source

*includes repeller, ion volume, extraction lens and dual filaments*



# The Project “Lambrusco”

“New Analytical Methodologies for Geographical and Varietal Traceability of Oenological Products”



## Red sparkling wine

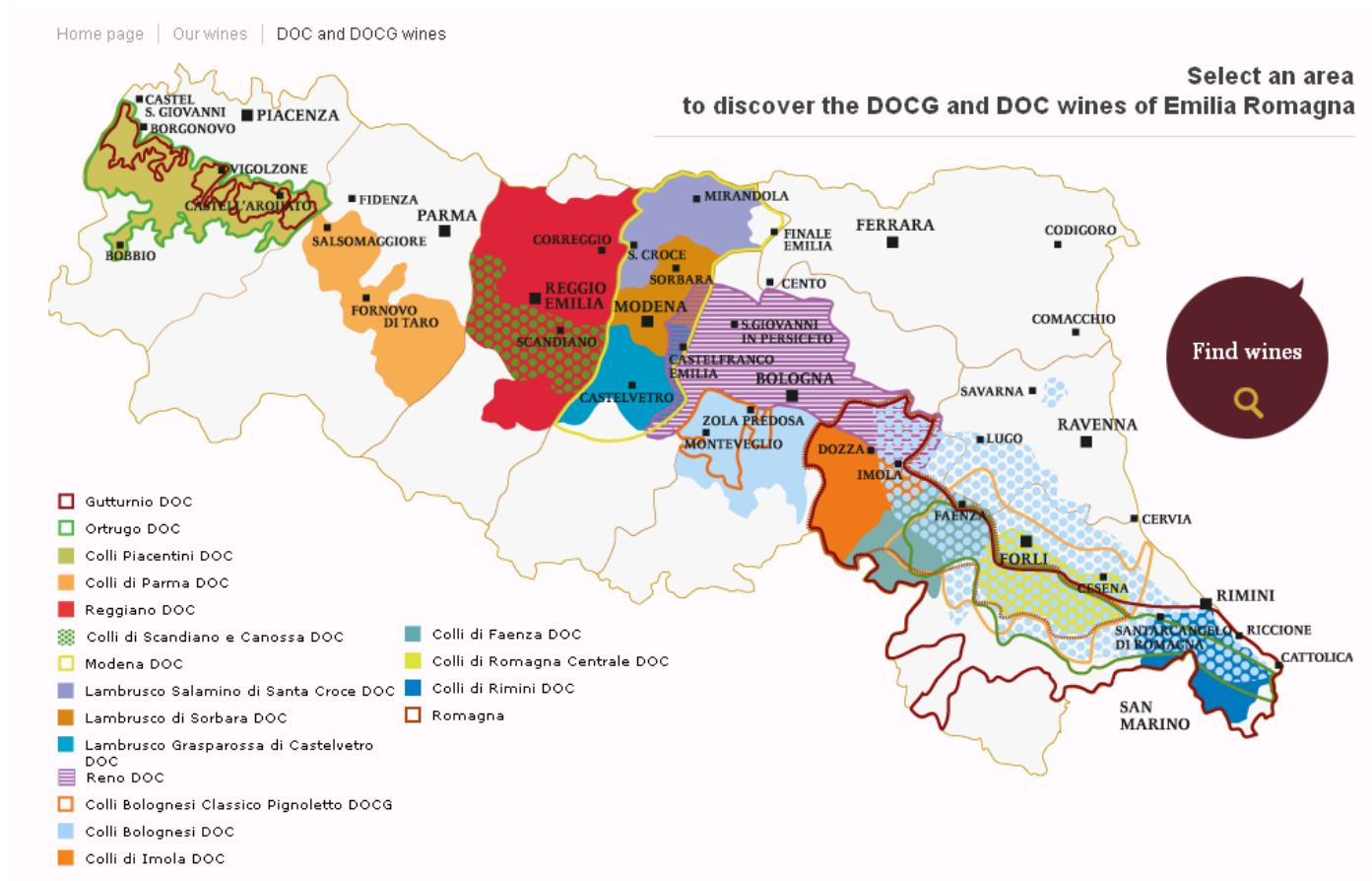
In Modena 4 DOC areas:

- Lambrusco Salamino di Santa Croce
- Lambrusco di Sorbara
- Lambrusco di Modena
- Lambrusco Grasparossa di Castelvetro

# Lambrusco Project

New Analytical Methodologies for Geographical and Varietal Traceability of Oenological Products

## Emilia-Romagna



Agilent Technologies

2nd MS Wine Day - Susegana

2017

# GC-QTOF Analysis

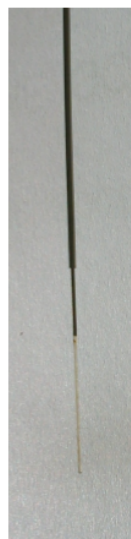
## Solid Phase Microextraction SPME

### Sample preparation

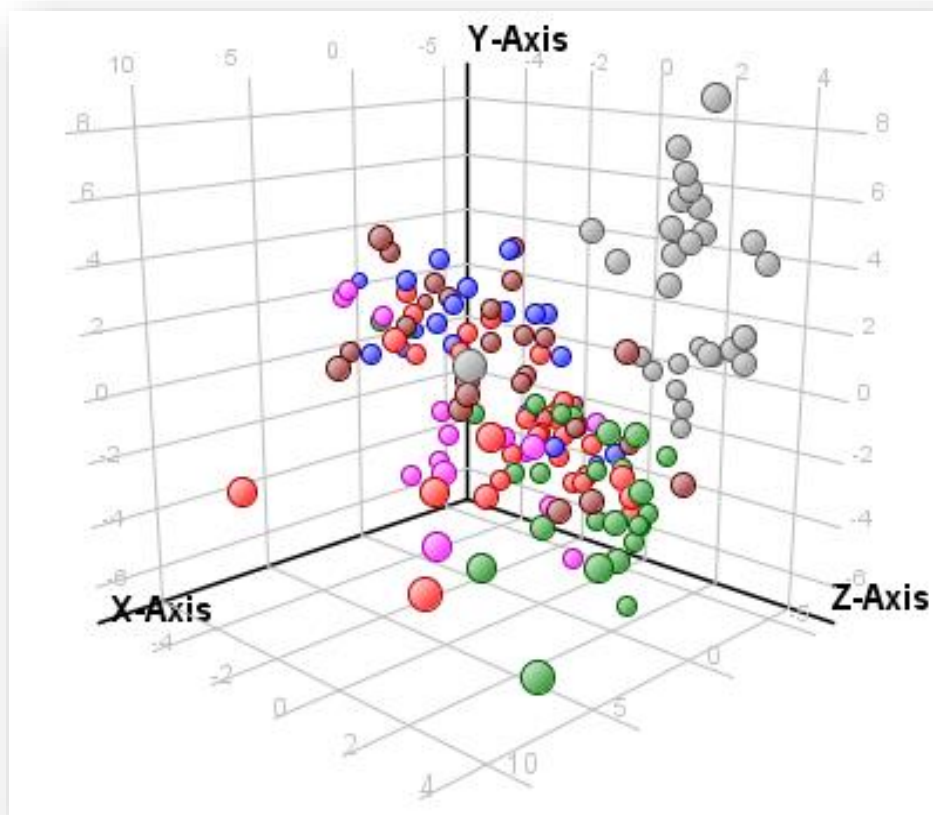
- ultrasonic bath to remove CO<sub>2</sub>
- Internal standard: ethyl hexanoate d11
- Quality control QC: mix of all samples

### Solid phase microextraction SPME

- 10 mL wine + 3 g NaCl in 20mL vial,
- incubation 10 min @ 35°C
- extraction 40 min @ 35°C
- DVB/CAR/PDMS 2cm fiber
- Agilent autosampler (like CTC combiPAL)
- Desorption 260°C in GC inlet 2 min splitless, narrow liner 0.75mm id
- Fibre cleaning 10min @ 265°C needle heater



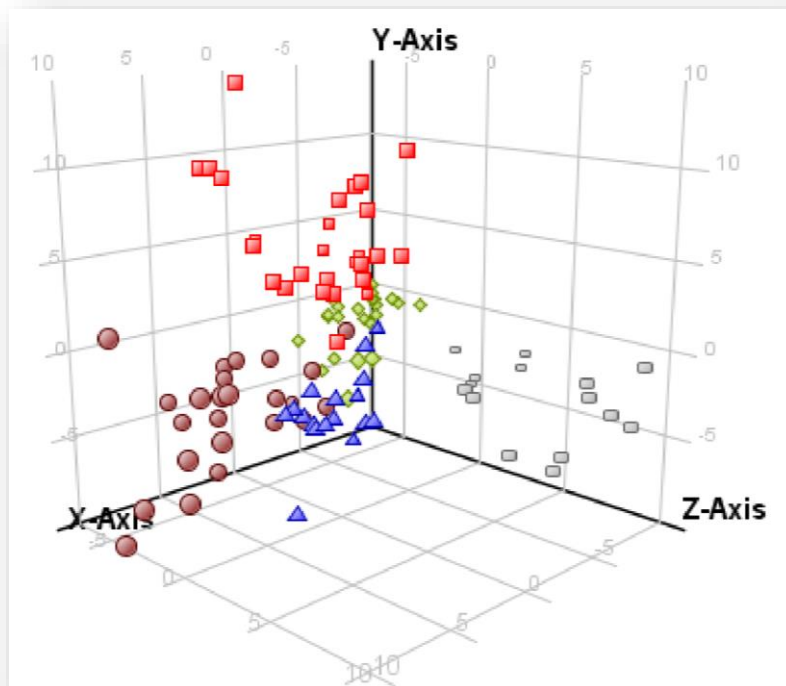
# Mass Profiler Professional MPP



PCA: initial evaluation

# Mass Profiler Professional MPP

Classification – PLS model



- Grasparossa
- ◆ Sorbara
- ▲ Salamino
- Modena
- outside

Overall model accuracy: 97%



# Identification of Compounds

Use accurate mass → formula calculator

**Formula Calculator**

Submit formulas to NIST MS SEARCH

Allowed Species Limits Scoring

Mass and charge

Mass or m/z: 171.1741

Charge: 1

Charge carrier

Positive ions: H Negative ions: H

MS ion electron state: allow both even and odd

Elements and limits

Element	Minimum	Maximum
C	1	30
H	0	62
O	0	5
N	0	3
S	0	2
Cl	0	3
[13C]	0	1

Formula (M)	Ion Formula	Mass (MFG)	m/z (Calc)	Diff (ppm)
C11 H22 O	C11 H23 O	170.1671	171.1743	1.42
C9 H20 N3	C9 H21 N3	170.1657	171.173	-6.47

No hits

46 hits  
And now?





# Identification of Compounds

## Chemical Ionization (CI):

- to define the Molecular Ion;

## Electron Impact spectra (EI):

- Identification of fragments formulas, library search;

Using Accurate Ms/Ms data, 200 compounds are ready for identification.

Actual Results: 31 compounds identified (mf>60%) matching expected RT and RI

limonene	ethyl 2-methyl butyrate
methylglyoxal	2-methylbutyl acetate
nonanal	ethyl hexanoate
p-cymene	ethyl heptanoate
undecanal	ethyl isobutyrate
α-methylbutyric acid	ethyl myristate
1-hexanol	ethyl nonanoate
1-nonanol	ethyl octanoate
E,E-2,4-hexadienal	ethyl phenylacetate
2-nonanone	ethyl trans-4-decenoate
2-pentanone	butyl benzoate
3-ethoxy-1-propanol	isobornyl acetate
4-ethylphenol	ethyl heptanoate
benzyl alcohol	ethyl sorbate*
decanal	
epiglobulol	
γ-octalactone	

\* Degradation product from sorbic acid, wine additive, found in only 1 sample



# Molecular Structure Correlator Software (MSC)

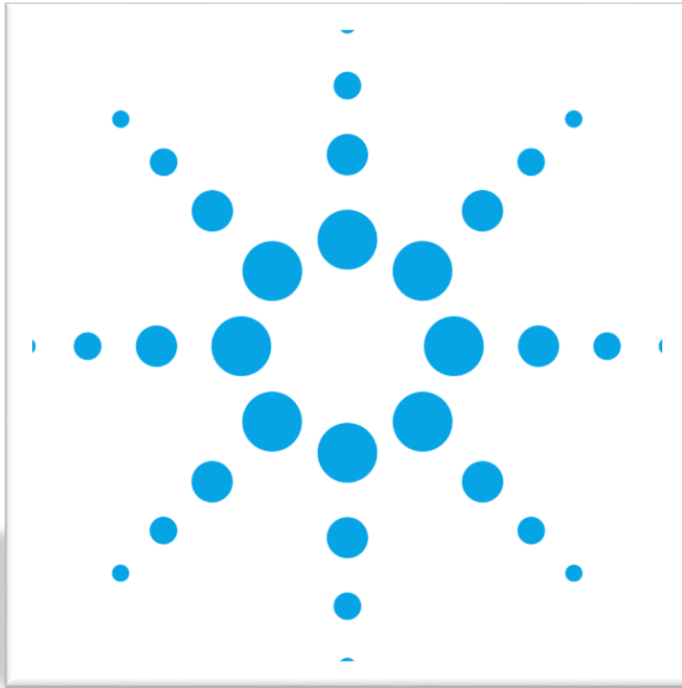
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# Conclusions

- GCQTOF suitable for food profiling applications
- Volatile Wine fraction as biomarker
- Smooth statistical data processing needed
- Identification by Ci+EI data, Libraries and MSC



# Thank you for attention



## Acknowledgments:



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