



**INSTITUTE OF  
CHEMICAL TECHNOLOGY PRAGUE**

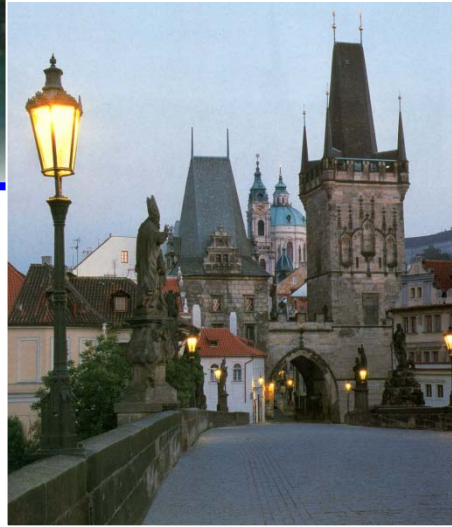
# **Direct analysis in real time and high resolution mass spectrometry: a new tool for food authenticity**

**JANA HAJŠLOVA**

**on behalf of her Ph.D. students & young scientists**



**17.5. 2013, Palermo, Italy**



Prague, Czech Republic, EU



# ICT Metrological and Testing laboratory

- research and education,
- collaboration with industry,
- collaboration with other research/testing laboratories (private, official),
- international co-operation within EU research projects aimed at detection techniques,
- expertise/testing, consultation
- laboratory audits,
- laboratory training and demonstration



# International projects



# Overview

- Introduction: food authenticity & fraud
- Metabolomics: what is its role in food analysis?
- Instrumental platforms for metabolomics
- DART ion source coupled with HRMS
- Case studies based on metabolomics
  - wine authenticity
  - oxidized fats detection



# FOOD AUTHENTICITY AND FRAUD

- ➔ Important food quality parameter
- ➔ Most of valued food commodities are subject to fraud
- ➔ Substitution or extension by cheaper product / ingredient / raw material
- ➔ Geographic origin
- ➔ Use of undeclared technology / processing

**ECONOMIC ASPECTS**

**CONSUMERS' DECEPTION**

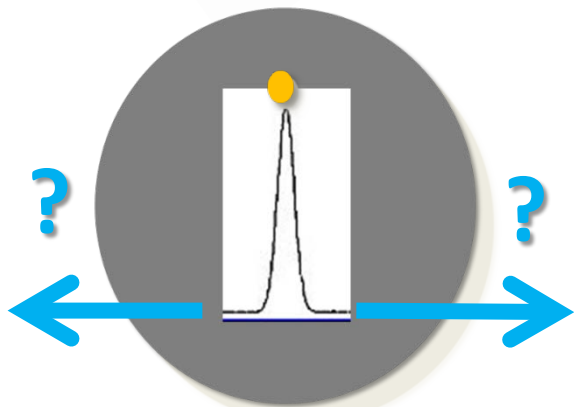
**THREAT TO CONSUMERS' HEALTH**



# Authentication strategies: where to go....?

'Classic' approach

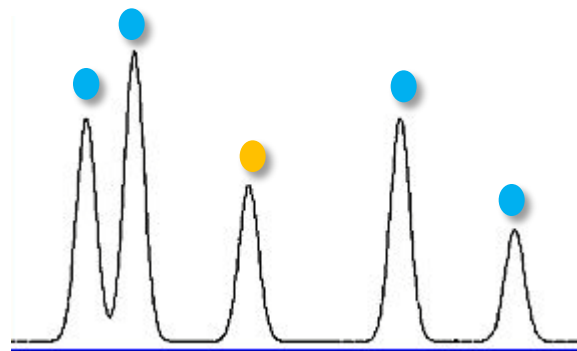
Analysis of one or few target markers



New strategy:  
non-target screening / profiling

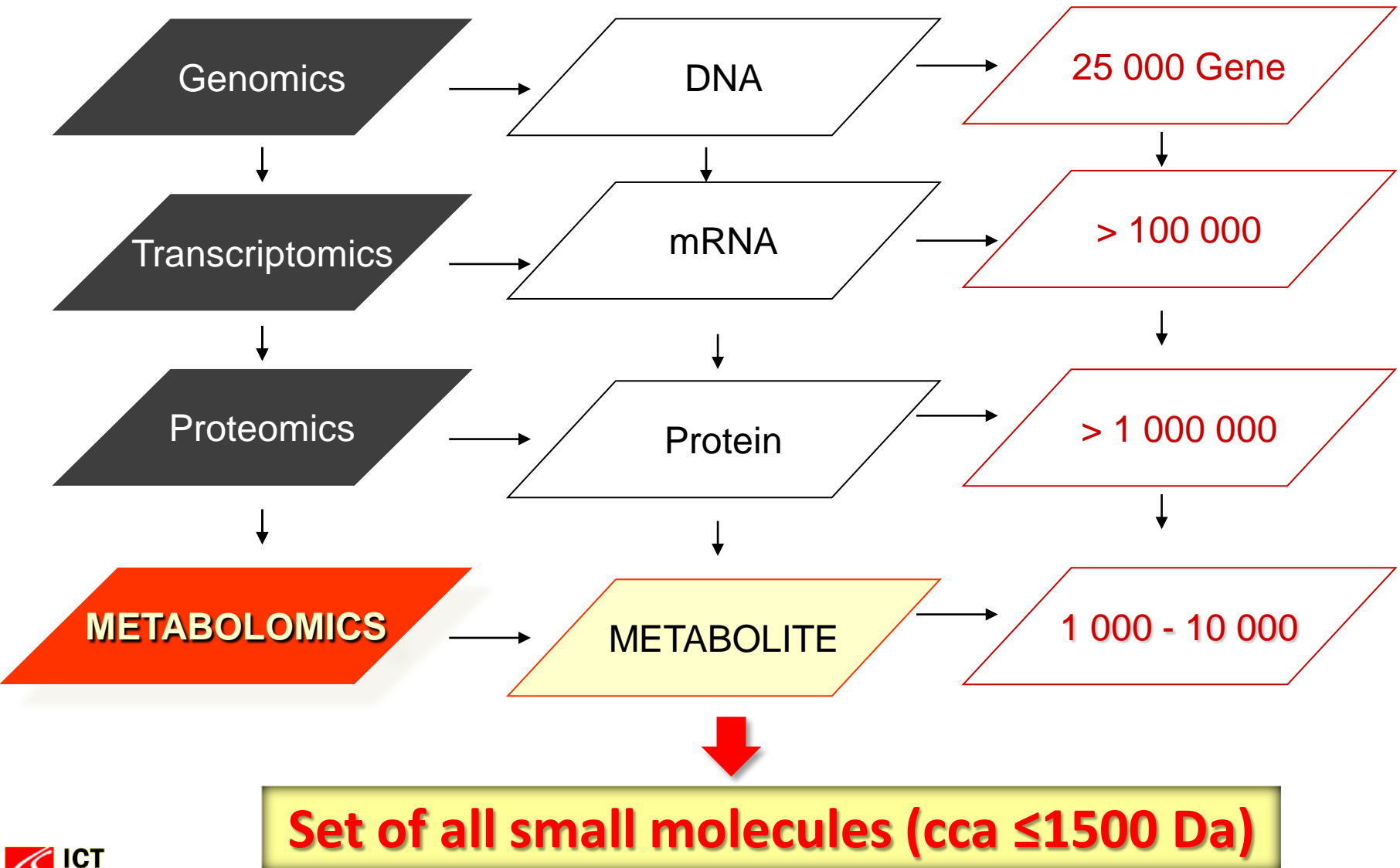
**COMPREHENSIVE SAMPLE ANALYSIS**

- detection / identification of 'unknown' components
- Identification of set of composition markers



**METABOLOMICS →**

# „OMICS“ hierarchy





# Application scope of metabolomics

2008



TRENDS IN  
FOOD SCIENCE  
& TECHNOLOGY

Trends in Food Science & Technology 19 (2008) 482–493

Review

## Metabolomics: applications to food science and nutrition research

David S. Wishart<sup>a,b,c,\*</sup>

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<sup>b</sup>Department of Biological Sciences, University of Alberta, Edmonton, AB, Canada T6C 2E8

<sup>c</sup>National Institute for Nanotechnology, 11421 Saskatchewan Drive, Edmonton, AB, Canada T6G 2M9  
(Department of Computing Science, University of Alberta, 2-21 Athabasca Hall, Edmonton, AB T6C 2E8, Canada. Tel.: +1-780-492-0383; fax: +1-780-492-1071; e-mail: david.wishart@ualberta.ca)

molecules can include a range of endogenous and exogenous chemical entities such as peptides, amino acids, nucleic acids, carbohydrates, organic acids, vitamins, polyphenols, alkaloids, minerals and just about any other chemical that can be used, ingested or synthesized by a given cell or organism.

Metabolomics, which (Nicholson, Lindon & Holmes, 2006), on the technological breakthrough and identification. The mass spectrometry (MS) termination, high-resolution (nuclear magnetic resonance) trophoresis (CE) and ultragraphy (UPLC and HPLC) separation, as well as process spectral or chromatography (Johnson & Johnson, 2005; Wishart et al., 2005) to the development of a nation of electronic databases spectral information on different metabolomes (Subramaniam, 2006; Kishore et al., 2006; Smith et al., 2005) ware and software innovations detect and characterize at a time but dozens of minutes (Dunn et al., 2005; Dunn et al., 2001).

As with many new technologies, there is a tendency towards rapid technological proliferation followed

*This review focuses on the recent trends and potential applications of metabolomics in four areas of food science and technology:*

- (1) food component analysis;
- (2) food quality/authenticity assessment;
- (3) food consumption monitoring
- (4) physiological monitoring in food intervention or diet challenge studies

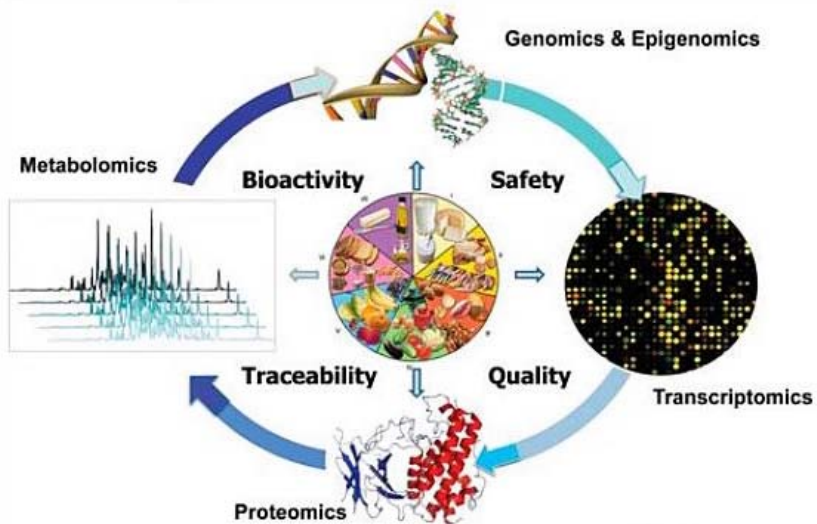
Metabolomics is an emerging field of “omics” research that focuses on high-throughput characterization of small molecule metabolites in biological matrices. As such, metabolomics is ideally positioned to be used in many areas of food science and nutrition research. This review focuses on the recent trends and potential applications of metabolomics in four areas of food science and technology: (1) food component analysis; (2) food quality/authenticity assessment; (3) food consumption monitoring; and (4) physiological monitoring in food intervention or diet challenge studies.

# Foodomics: a new keyword in food analysis

# Foodomics

Advanced Mass Spectrometry in  
Modern Food Science and Nutrition

EDITED BY **Alejandro Cifuentes**



WILEY

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2013



International Conference  
on FoodOmicS  
3<sup>rd</sup> Edition



CESENA, ITALY

MAY 22-24, 2013

Photos by: Gianfranco Pione

2013 "FoodOmicS, the science for discovering"

Countdown

76 10 45 04

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**Foodomics: a new comprehensive approach to food and nutrition**

Francesco Capozzi and Alessandra Bordonì

In the past 20 years, the scientific community has faced a great development in different fields due to the development of high-throughput, omics technologies. Starting from the four major types of omics measurements (genomics, transcriptomics, proteomics, and metabolomics), a variety of omics subdisciplines (epigenomics, lipidomics, interactomics, metallomics, diseasomics, etc.) has emerged.

Thanks to the omics approach, researchers are now facing the possibility of connecting food components, foods, the diet, the individual, the health, and the diseases, but this broad vision needs not only the application of advanced technologies, but mainly the ability of looking at the problem with a different approach, a "foodomics approach".

Foodomics is the comprehensive, high-throughput



Click on the picture

PREVIOUS EDITIONS

[2nd International Conference on FOOD-OMICs](#)

A science for nutrition, health and wellness in the post-genomic era  
Cesena (Italy), June 22-24, 2011

[1st International Conference on FOOD-OMICs](#)

A science for nutrition, health and wellness in the post-genomic era  
Cesena (Italy), May 28-29, 2009

OTHER MEETINGS

[2nd International Conference on Food Digestion](#)

Madrid (Spain) March 6-8, 2013

# METABOLOMICS (‘FOODOMICS’?)



# WORKFLOW: profiling / fingerprinting

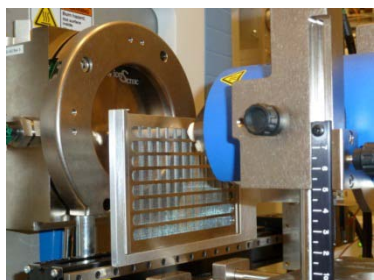
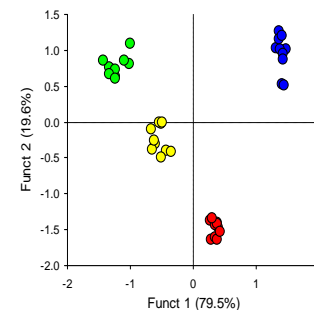
Samples



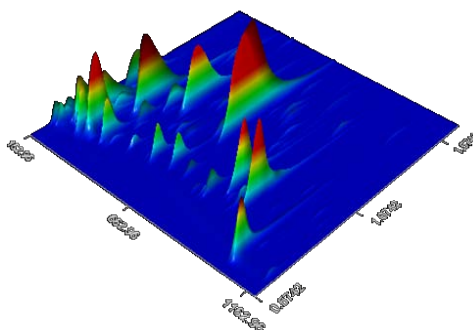
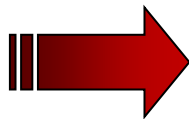
Output (confirmation of rejection of claim/identity)

Act. Group	Pred. Group (Holdout)				Correctly Classified
	Group 1	Group 2	Group 3	Group 4	
Group 1	10	0	0	0	1.000
Group 2	0	10	0	0	1.000
Group 3	0	0	10	0	1.000
Group 4	0	0	0	10	1.000
Overall Correct Classification Rate					1.000

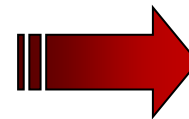
Data analysis



Fingerprinting technique

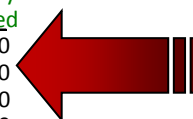


Recorded output

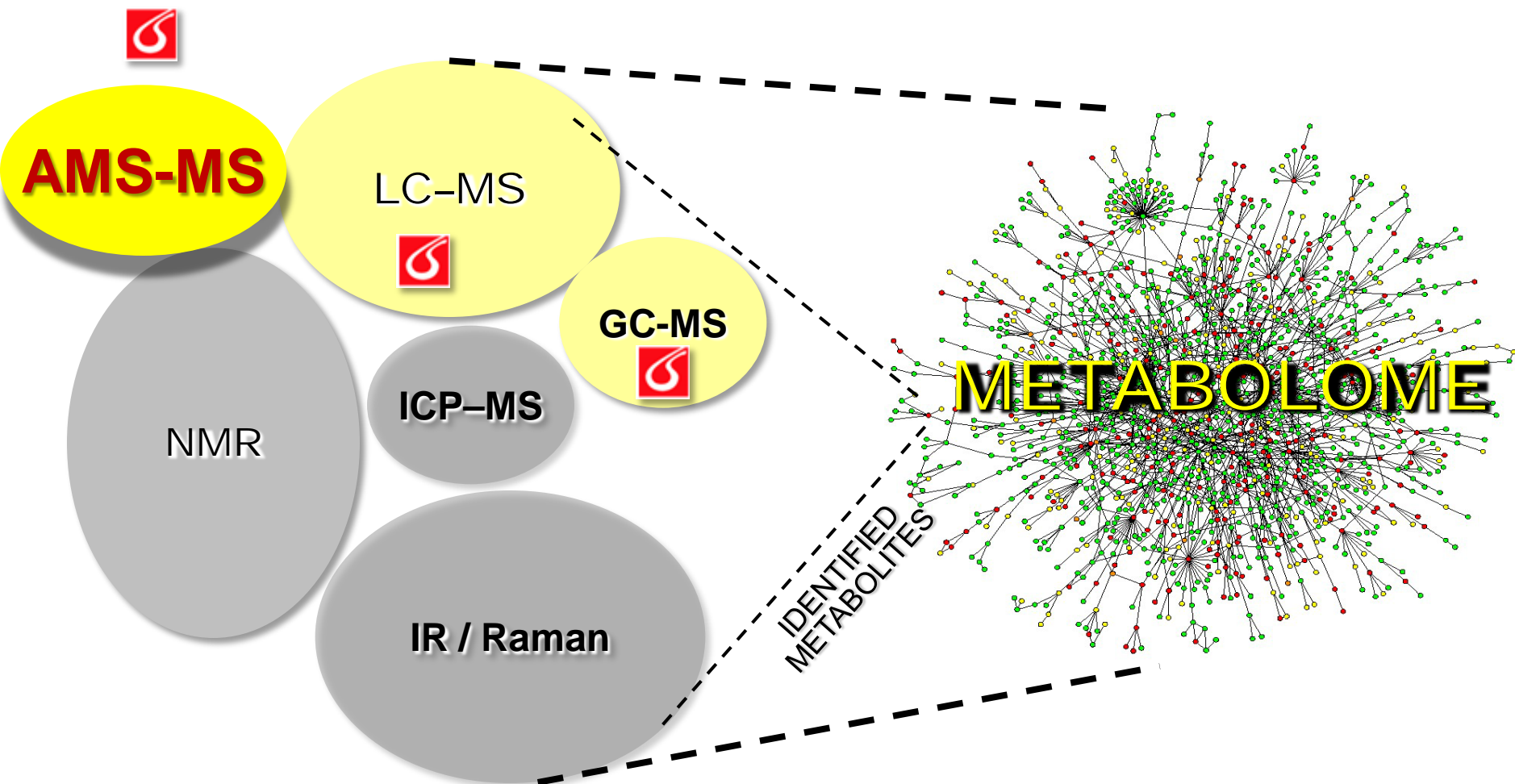


No.	Group	Var1	Var2	Var3	Var4	Var5	Var6	Var7	Var8	Var9	Var33
1	Group 1	0.4880	1.580	2.170	5.640	0.690	2.420	1.150	0.640	0.019...	0.005
2	Group 1	0.4960	1.660	2.220	5.800	0.700	2.490	0.570	0.330	0.020...	0.013
3	Group 1	0.0830	0.043	0.0320	0.0010	0.0220	0.140	0.0010	0.0010	0.008...	0.002
...	...	...	...	...	...	...	...	...	...	...	...
11	Group 2	0.8660	0.450	0.7360	0.290	0.260	0.250	0.0000	0.0040	0.010...	0.011
12	Group 2	0.8530	0.0430	0.7030	0.0280	0.0260	0.0240	0.0000	0.0040	0.009...	0.000
13	Group 2	0.7240	0.271	0.0000	0.160	0.0430	0.0490	0.0000	0.0030	0.007...	0.014
...	...	...	...	...	...	...	...	...	...	...	...
21	Group 3	0.2640	0.980	0.0600	0.150	0.280	1.530	0.0000	0.0010	0.015...	0.006
22	Group 3	0.2730	0.970	0.0610	0.150	0.290	1.540	0.0000	0.0010	0.015...	0.005
23	Group 3	0.4350	0.160	0.1000	0.0000	0.120	1.650	0.0000	0.0010	0.002...	0.000
...	...	...	...	...	...	...	...	...	...	...	...
38	Group 4	1.0000	0.0080	0.0020	0.0000	0.0200	0.920	0.0050	0.0040	0.001...	0.014
39	Group 4	0.7830	0.070	0.0000	0.0050	0.0450	0.360	0.0000	0.0020	0.000...	0.022
40	Group 4	0.8490	0.0000	0.0000	0.0060	0.0510	0.380	0.0000	0.0040	0.000...	0.026

Data matrix



# Instrumental platforms for metabolomics

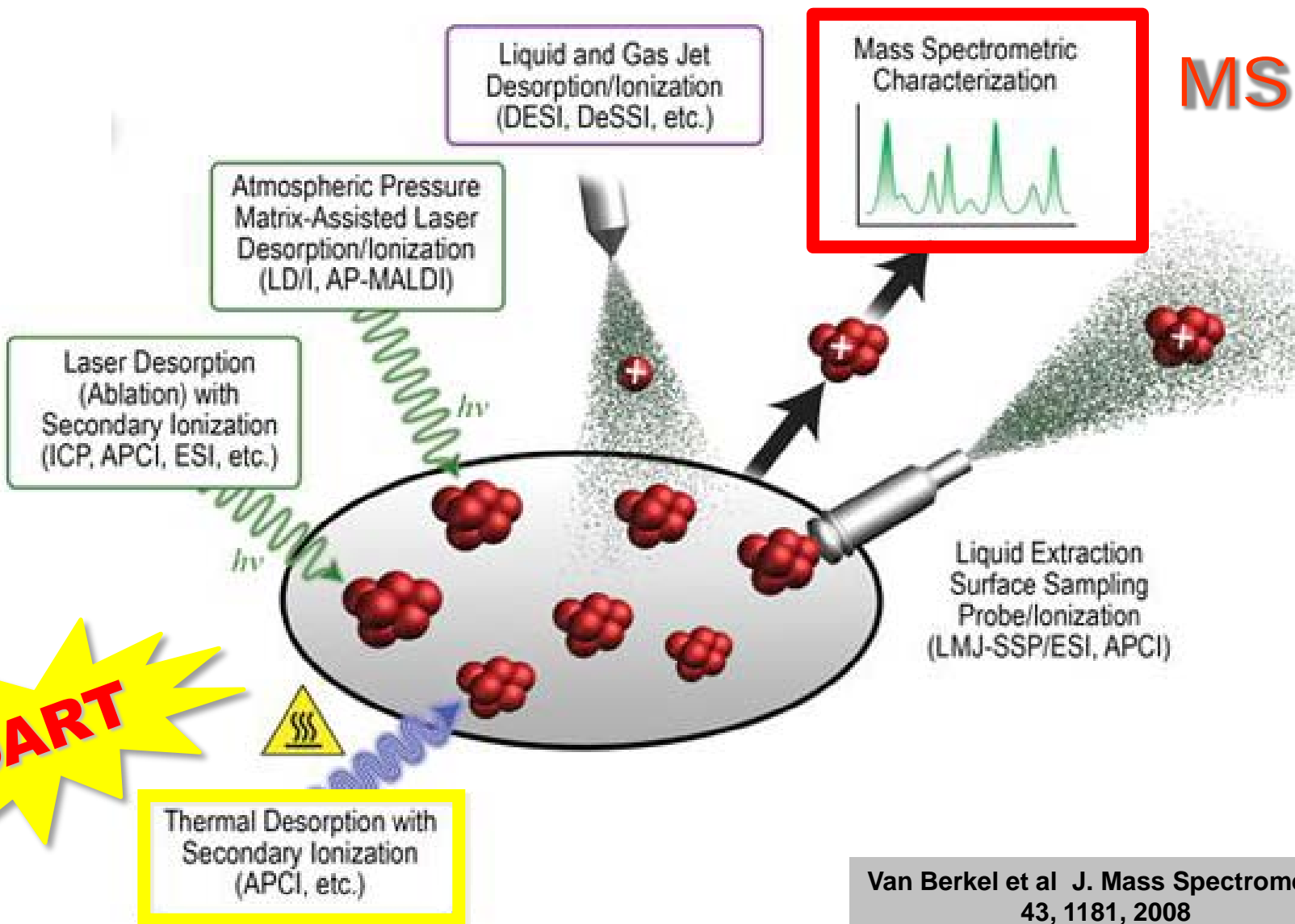


# AMBIENT MASS SPECTROMETRY (AMS)

A new challenge for our labs?



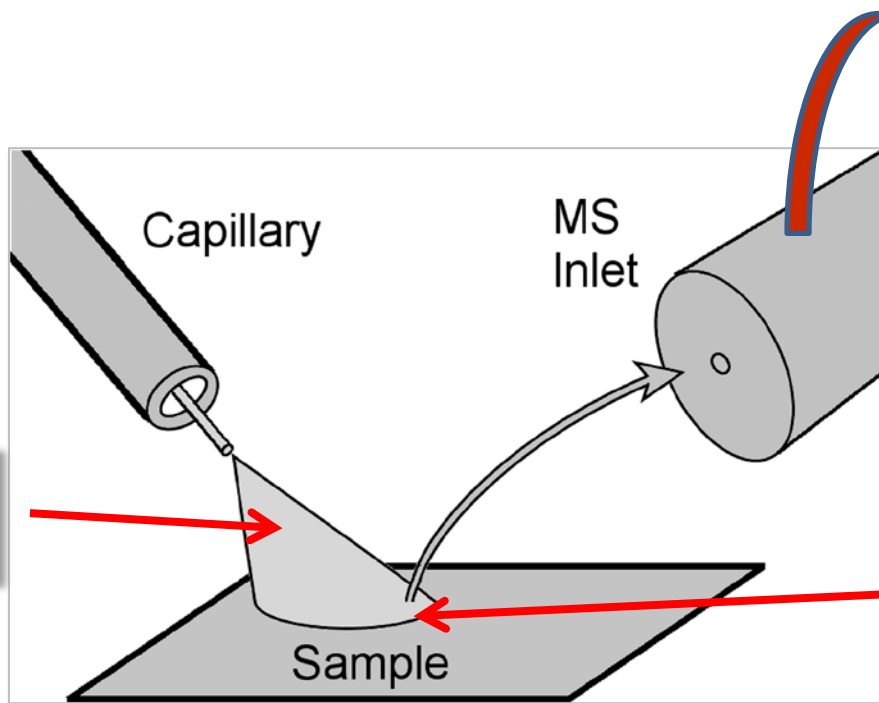
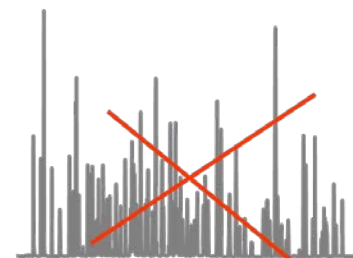
# AMBIENT MASS SPECTROMETRY OPTIONS



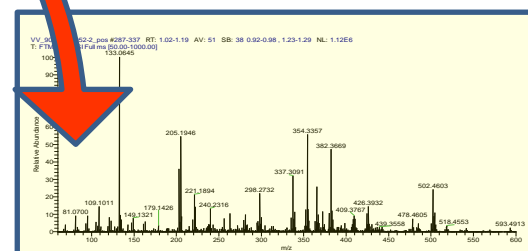
Van Berkel et al J. Mass Spectrometry, 43, 1181, 2008

# Novel fingerprinting technique:

## AMBIENT MASS SPECTROMETRY



Mass spectrum



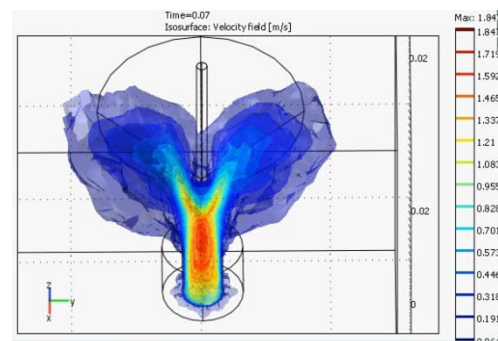
excited He  
200- 400°C

THERMO DESORPTION,  
APCI ionization



Steps occurring in DART together, or separately, under the influence of a particular agent:

- **DESORPTION** – a change in phase (e.g., solid to vapor)
- **IONIZATION** - an acquisition of charge by neutral analyte molecules



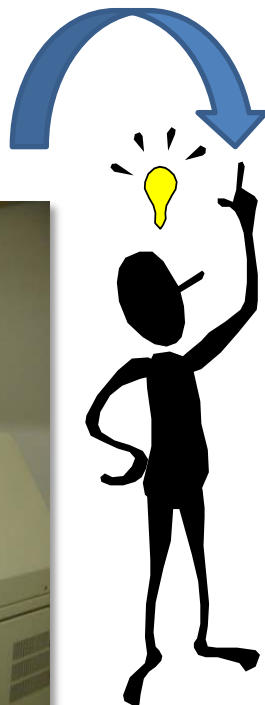
# DART – HRMS has become an important tool in routine food analysis / food research



## DART-TOFMS



AccuTOF LP (Jeol)  
Time-of-flight mass spectrometer  
~ 5000 – 7000 fwhm



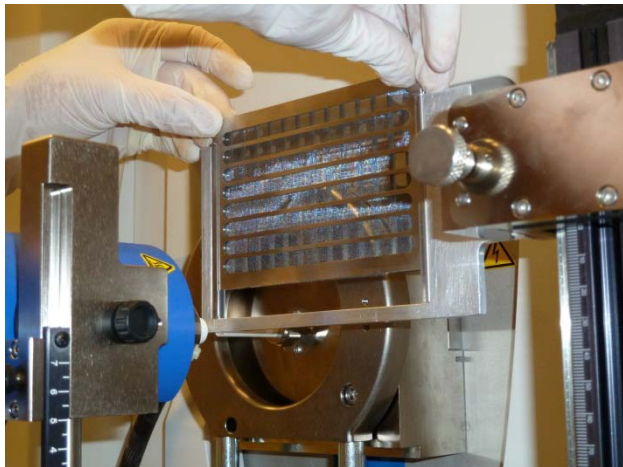
## DART-orbitrapMS



Exactive (Thermo Scientific)  
Orbitrap mass spectrometer  
~ 10,000 – 100,000 fwhm

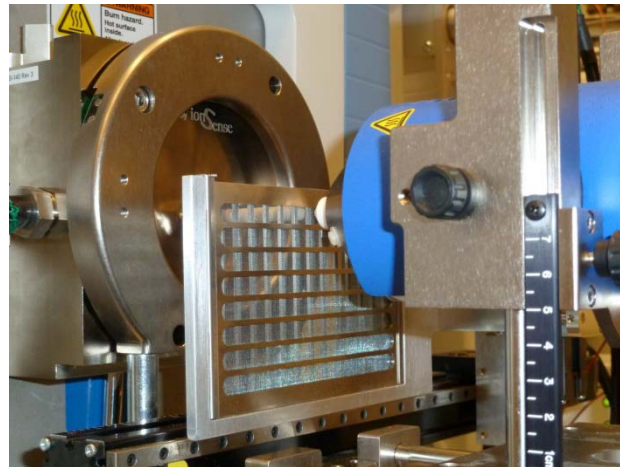
# X-Z TRANSMISSION MODULE AUTOSAMPLER

(2)



Metal mesh is placed into a holder

(3)

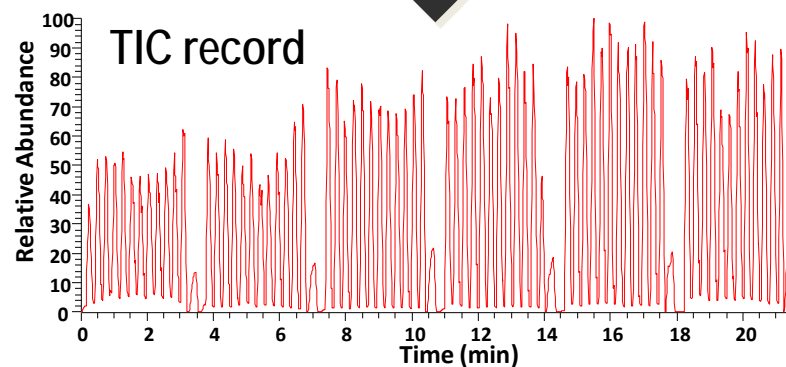


Gas flows through the metal mesh, desorbs and ionizes sample components.

(1)



Sample (5  $\mu$ l) is placed on metal mesh.



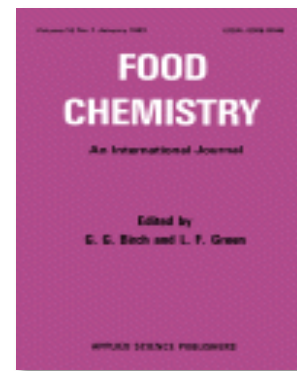
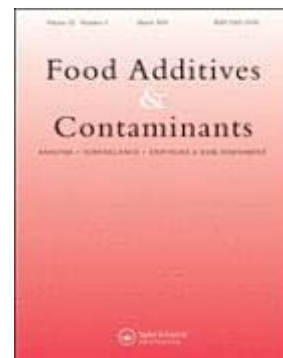
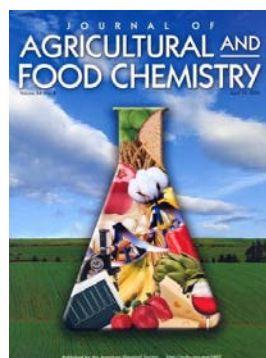
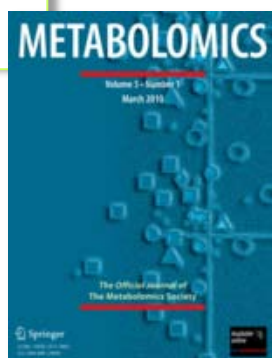
# METABOLOMICS-BASED CASE STUDIES



# Metabolomics based studies we have performed



- ➔ Adulteration of olive oil by hazelnut oil
- ➔ Fish feeding history authentication
- ➔ Classification of bear origin, traceability
- ➔ Identification of geografic origin of honey
- ➔ Authentication of fruit juice freshness
- ➔ Wines authentication
- ➔ Authentication of apple variety and the way of farming





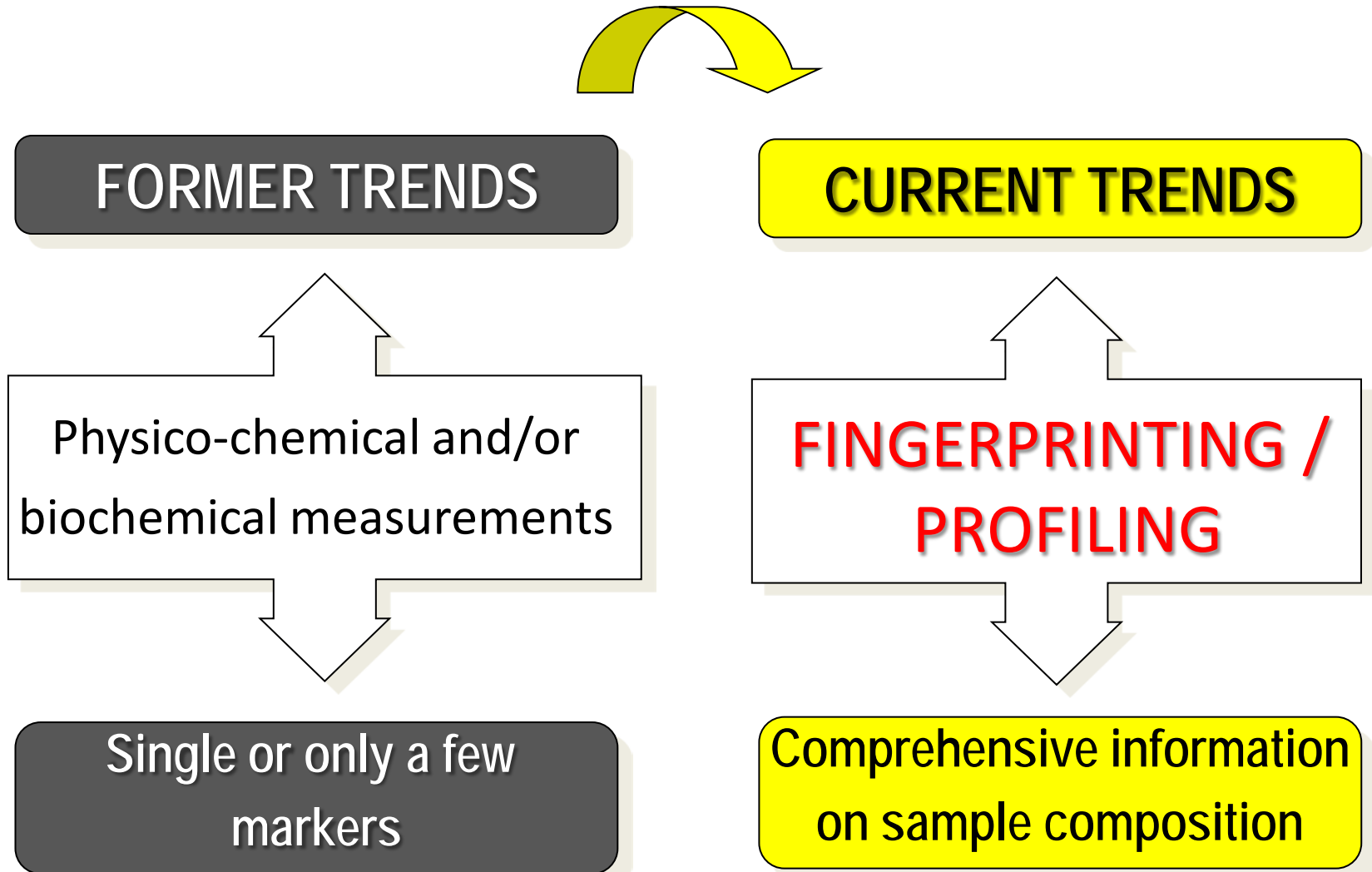
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## **NON-TARGET ANALYSIS:**

### **Wine authentication**



# WINE AUTHENTICATION TOOLS



# METHODS FOR WINE AUTHENTICATION

HPLC–UV-VIS /–MS  
(phenolic compounds)

Infrared  
spectroscopy

DNA (biomethods)

$^{13}\text{C}$  NMR

AAS nebo ICP-MS  
(minerals)



SPME–GC–FID  
SPME–GC–MS  
**SPME–GC×GC–MS**  
(volatile compounds)

**ASAP–MS**  
**DART–MS**  
(ionizable compounds)

GC–FID  
(alcohols)

**LC–MS**  
(polar compounds)



# AN OVERVIEW OF ANALYZED SAMPLES (WHITE WINES)

- Chardonnay ( $n = 20$ )
- Müller Thurgau ( $n = 20$ )
- Rhine Riesling ( $n = 15$ )



- Origin of the samples: **Australia, Austria, Bulgaria, Chile, Czech Republic, Hungary, Italy, Moldavia, South Africa, USA**
- Years of the production: **2005–2008**

# AN OVERVIEW OF ANALYZED SAMPLES (RED WINES)

- **Cabernet Sauvignon** ( $n = 15$ )

- **Merlot** ( $n = 16$ )

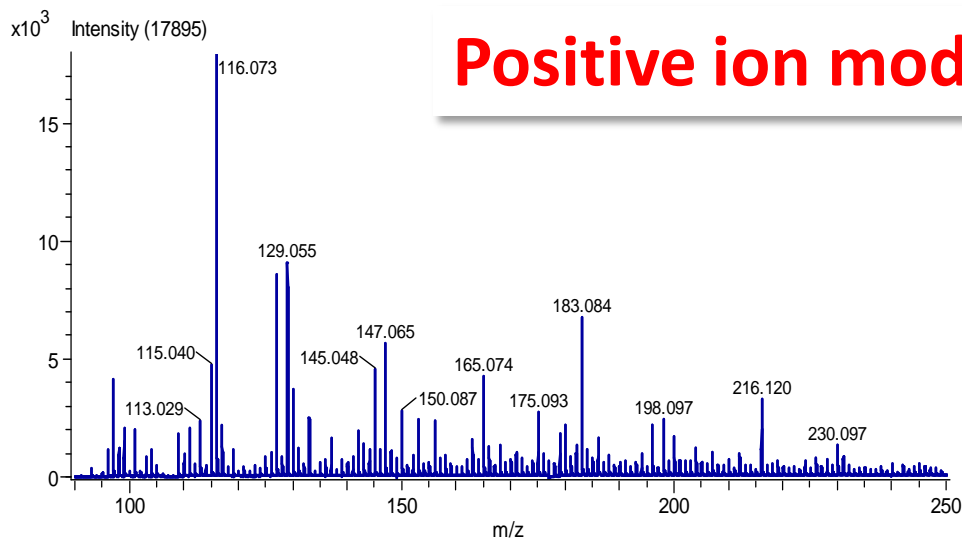
- **Pinot Noir** ( $n = 14$ )



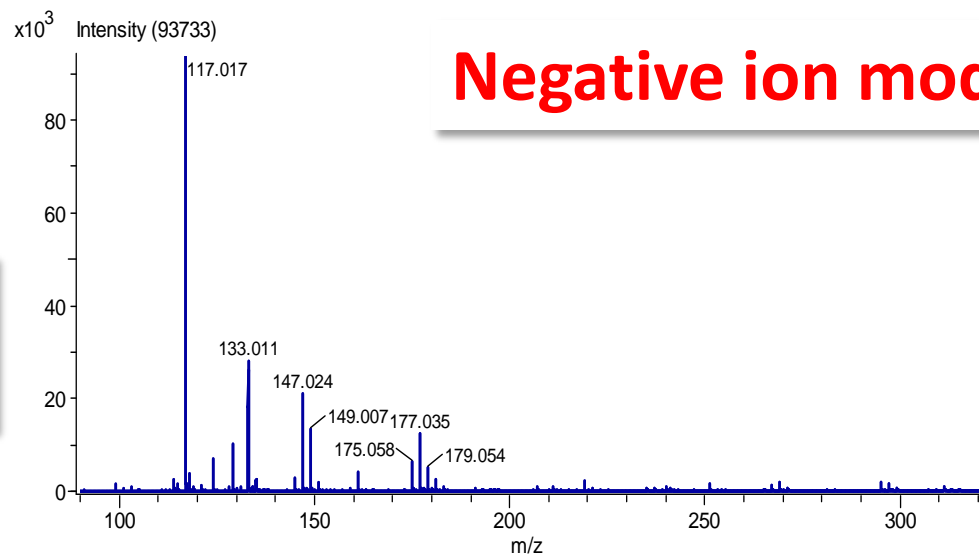
- **Origin of the samples: Australia, Bulgaria, Czech Republic, France, Germany, Hungary, Chile, Makedonia, Slovakia, Spain, USA**

- **Years of the production: 2004–2008**

# DART-MS: Direct measurement of wine



*Example: Cabernet  
Sauvignon*



# CHEMOMETRIC ANALYSIS (wine directly measured)

Model: Cabernet Sauvignon vs. Merlot

**25 markers**

(DART(+)-MS, dir. meas.)

( $m/z$ )



PLS-LDA + model validation  
(LOOCV)



6 latent variables (LVs)



Recognition ability: 100%  
Prediction ability: 96.0%

**14 markers (**

(DART(-)-MS, dir. meas.)

( $m/z$ )



PLS-LDA + model validation  
(LOOCV)

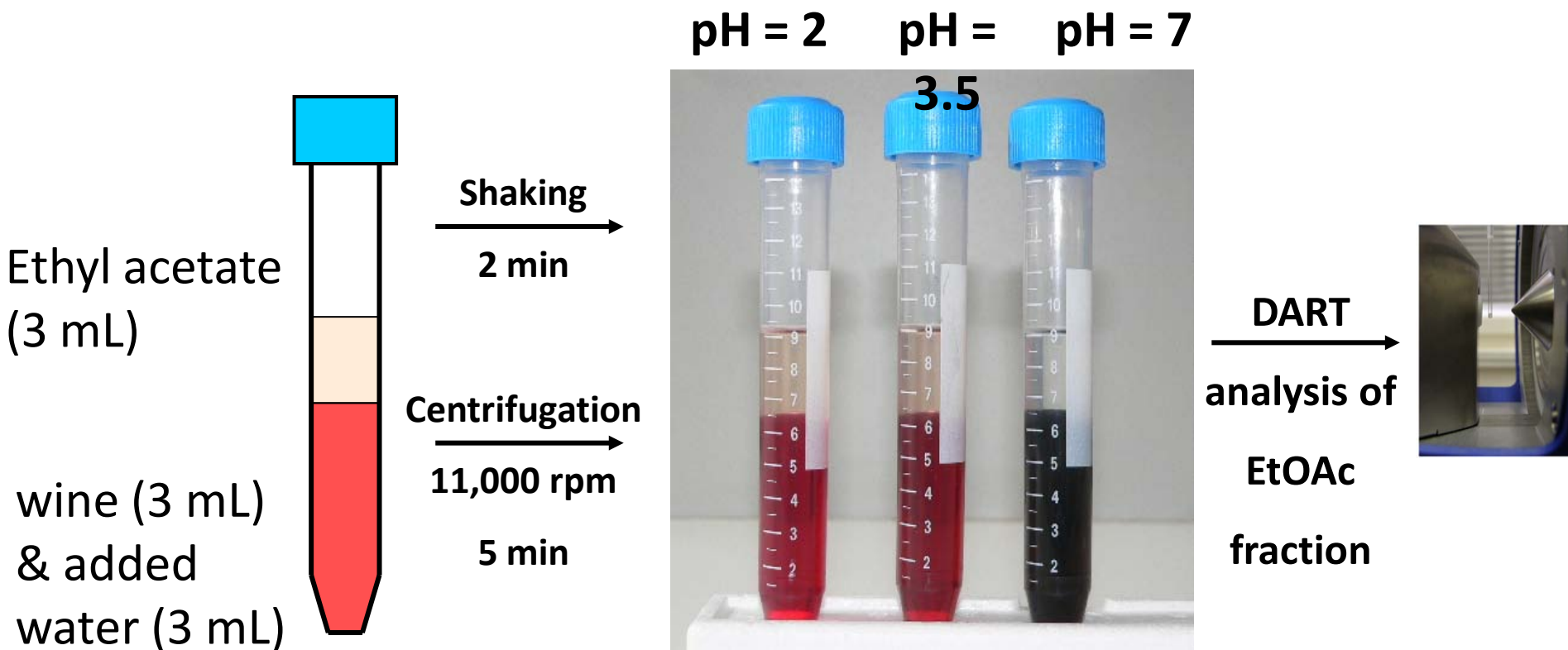


5 latent variables (LVs)



Recognition ability: 88.0%  
Prediction ability: 88.0%

# DART: Direct measurement of phenolics fraction isolated by EtOAc extraction



# PHENOLIC COMPOUNDS

## Phenolic acids

*p*-Hydroxybenzoic; *o*-hydroxybenzoic; salicylic; gallic; cinnamic; *p*-coumaroyltartaric (= coutaric); caffeoyltartaric (= caftaric); feruloyltartaric (= fertaric); *p*-coumaroyl glucose; feruloylglucose; glucose ester of coutaric acid

## Anthocyanins

Cyanidin 3-glucoside; cy 3-acetylglucoside; cy 3-*p*-coumaryl-glucoside; peonidin 3-glucoside; pn 3-acetylglucoside; pn 3-*p*-coumarylglucoside; pn 3-caffeoylglucoside (?); delphinidin 3-glucoside; dp 3-acetylglucoside; dp 3-*p*-coumarylglucoside; petunidin 3-glucoside; pt 3-*p*-coumarylglucoside; malvidin 3-glucoside; mv 3-acetylglucoside; mv 3-*p*-coumarylglucoside; mv 3-caffeoylglucoside

## Flavonols

Kaempferol 3-glucoside; k 3-glucuronide; k 3-glucosylarabinoside (?); k 3-galactoside; quercetin 3-glucoside; q 3-glucuronide; q 3-rutinoside; q 3-glucosylgalactoside (?); q 3-glucosylxyloside (?); iso-rhamnetic 3-glucoside

## Flavan-3-ols and tannins

(+)Catechin; (-)epicatechin; (+)gallocatechin; (-)epigallocatechin; epicatechin-3-O-gallate; procyanidins B1, B2, B3, B4, C1, C2, polymeric forms of condensed tannins

## Flavanonols

Dihydroquercetin 3-rhamnoside (= astilbin); dihydrokaempferol 3-rhamnoside (= engeletin)

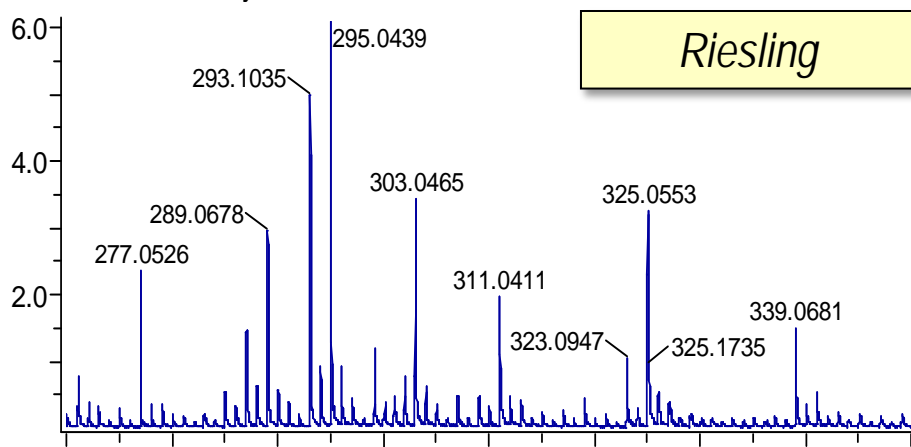
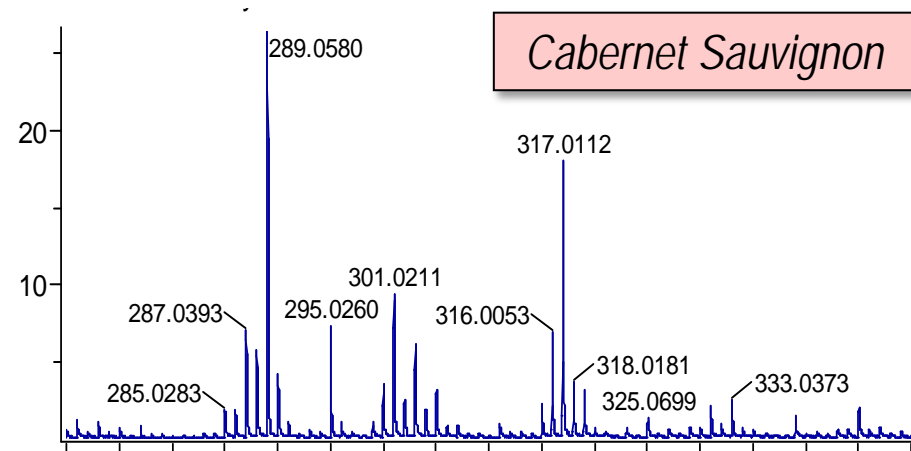
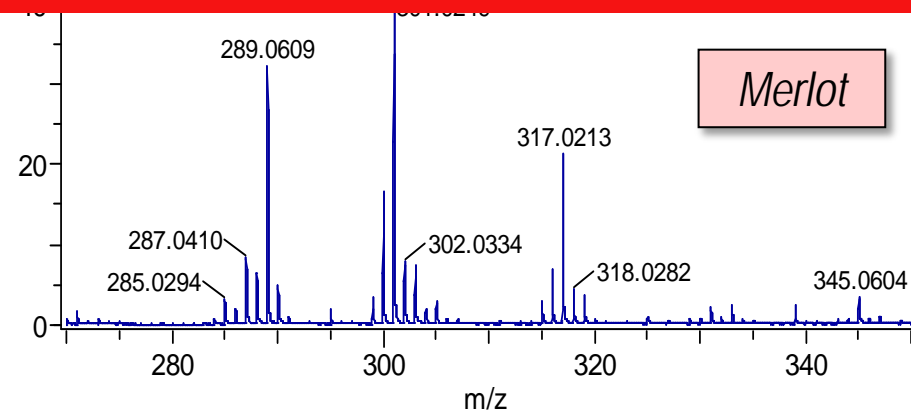
# DART [-]: PROFILING of phenolics fraction isolated by EtOAc extraction



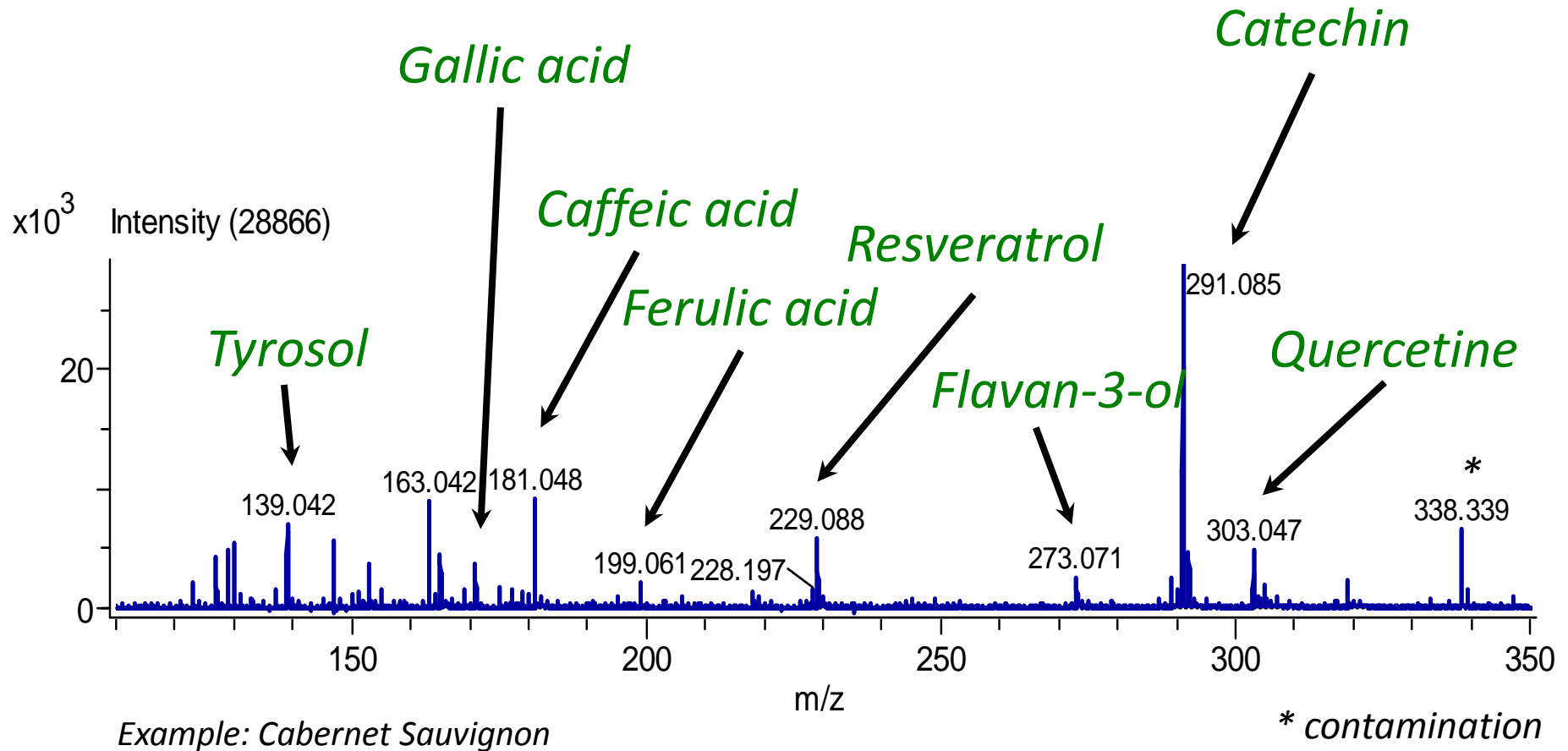
$m/z$  260–350



High-MW phenolics fraction

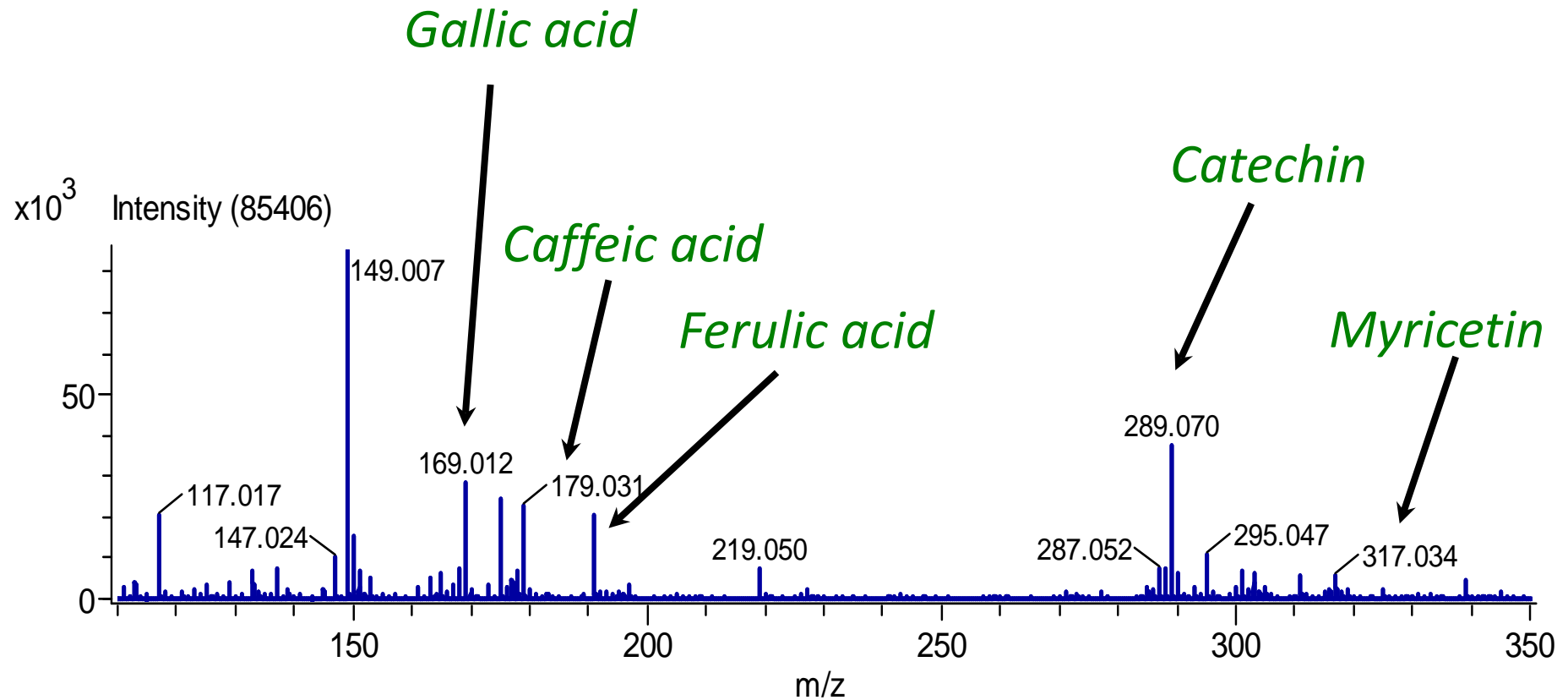


# DART(+) measurement of phenolics fraction - identification based on exact mass





# DART(-) measurement of phenolics fraction - identification based on exact mass



Example: Cabernet Sauvignon

# CHEMOMETRIC ANALYSIS (based on phenolics)

Model: Cabernet Sauvignon vs. Merlot

**25 markers** (DART(+)-MS, EtOAc  
fr.)

( $m/z$ )



**PLS-LDA + model validation  
(LOOCV)**



**5 latent variables (LVs)**



**Recognition ability: 100%**  
**Prediction ability: 100%**

**27 markers** (DART(-)-MS, EtOAc  
fr.)

( $m/z$ )



**PLS-LDA + model validation  
(LOOCV)**



**5 latent variables (LVs)**



**Recognition ability: 100%**  
**Prediction ability: 100%**

# CHEMOMETRIC ANALYSIS

Model: Chardonnay vs. Müller Thurgau vs. Rhine Riesling

**28 markers** (DART(-)-MS, EtOAc fr.)

**( $m/z$ )**



**PLS-LDA + model validation**  
**(LOOCV)**

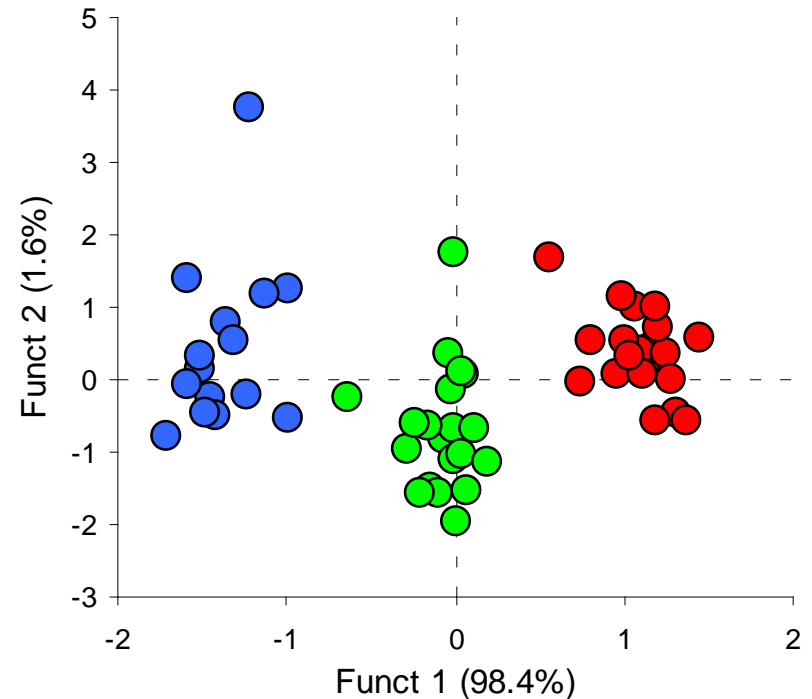


**9 latent variables (LVs)**



**Recognition ability: 98.9%**

**Prediction ability: 93.6%**



● Chardonnay ● Müller Thurgau  
● Rhine Riesling

# RECENT REVIEW SUMMARIZING ICT EXPERIENCE



8 December 2010; Disk Used

Trends in Analytical Chemistry, Vol. xxx, No. x, 2010

Trends

## 2 Challenging applications offered by 3 direct analysis in real time (DART) 4 in food quality and safety analysis

5 Jana Hajslova, Tomas Cajka, Lukas Vaclavik

7 Direct analysis in real time (DART) is an ambient ionization technique undergoing a rapid development. With a minimal sample  
8 pre-treatment, ionization of analyte molecules outside the mass spectrometer (MS) in the ordinary atmosphere is feasible. This  
9 ionization approach relies upon fundamental principles of atmospheric pressure chemical ionization (APCI). The current review

# ANOTHER CHALLENGE : LC-MS, direct injection

- To avoid any possible discrimination of metabolites a **direct injection** of wine samples was carried out.



# Liquid Chromatography (UHPLC Ultimate 3000 Dionex-Thermo)

- Different mobile phases and modifiers were tested:
  - Formic acid
  - Ammonium acetate
  - Ammonium formate

Two aqueous mobile phase components were used with methanol for elution during acquisition in positive and negative mode.

- **ESI +** The lower pH (3.8) eluent containing **ammonium formate** was employed in positive ion mode to support protonation of eluted molecules and ammonium adducts  $[M+NH_4]$ .
- **ESI -** Higher pH (4.5) value was used in negative ion mode (**ammonium acetate**) and  $[M+CH_3 COO]^-$  could be detected.

# Liquid Chromatography (UHPLC Ultimate 3000 Dionex-Thermo)

- Different columns were tested:
  - C18 (RPLC)
  - HILIC
- 1. The reversed phase chromatographic system enabled rapid separation of a wide range of compounds present in wine samples.
- 2. The use of stationary phase with sub-2  $\mu\text{m}$  particles significantly increased column separation efficiency and chromatographic resolution.

# UHPLC-Q Exactive

The Thermo Scientific Q Exactive benchtop UHPLC-MS/MS combines high-performance quadrupole precursor selection with high-resolution, accurate-mass (HRAM) Orbitrap detection to deliver high performance and tremendous versatility.

- Resolving power up to 140,000
- Maximum scan speed 12Hz
- Intra-scan dynamic range > 5000:1
- Quadrupole mass filter
- Spectral multiplexing for enhanced duty cycle
- S-Lens ion source for increased sensitivity





# UHPLC-Q Exactive: Method development

2013\_05\_14\_METABOLOMICS\_ESI+ - WINE.meth - Thermo Xcalibur Instrument Setup

File Help

Global Lists  
Tune Files  
External Hardware  
Chromatogram  
Scan Groups

Top10  
0 1 2 3 4 5 6 7 8 9 10 11 12 13  
time (min)

Experiments

Templates

System templates

- Full MS - SIM
- AIF
- Full MS / AIF
- Full MS / dd-MS<sup>2</sup> (TopN)
- Targeted-SIM
- Targeted-MS<sup>2</sup>
- Targeted-SIM / dd-MS<sup>2</sup>
- Full MS / AIF / NL dd-MS<sup>2</sup>

Properties

Properties of the method

Global Settings	best
use lock masses	best
show all properties	False

Time

Method duration	13.50 min
-----------------	-----------

Properties of Full MS / dd-MS<sup>2</sup> (TopN)

General	
Runtime	0 to 13.5 min
Polarity	positive
Default charge state	1
Inclusion	-
Exclusion	-
Tags	-

Full MS	
Resolution	70,000
AGC target	1e6
Maximum IT	250 ms
Scan range	80 to 1200 m/z

dd-MS <sup>2</sup> / dd-SIM	
Resolution	17,500
AGC target	5e4
Maximum IT	50 ms
Loop count	10
TopN	10
Isolation window	6.0 m/z
Fixed first mass	-
NCE	30.0

dd Settings	
Underfill ratio	5.0 %
Intensity threshold	5.0e4
Apex trigger	-
Charge exclusion	-
Peptide match	-
Exclude isotopes	-
Dynamic exclusion	10.0 s

LC settings

MS settings

Acquisition modes

Full MS

ddMS<sup>2</sup>

## FULL MS-ddMS<sup>2</sup>

This acquisition method consists in a high resolution MS survey scan followed by MS/MS using a data depending settings.

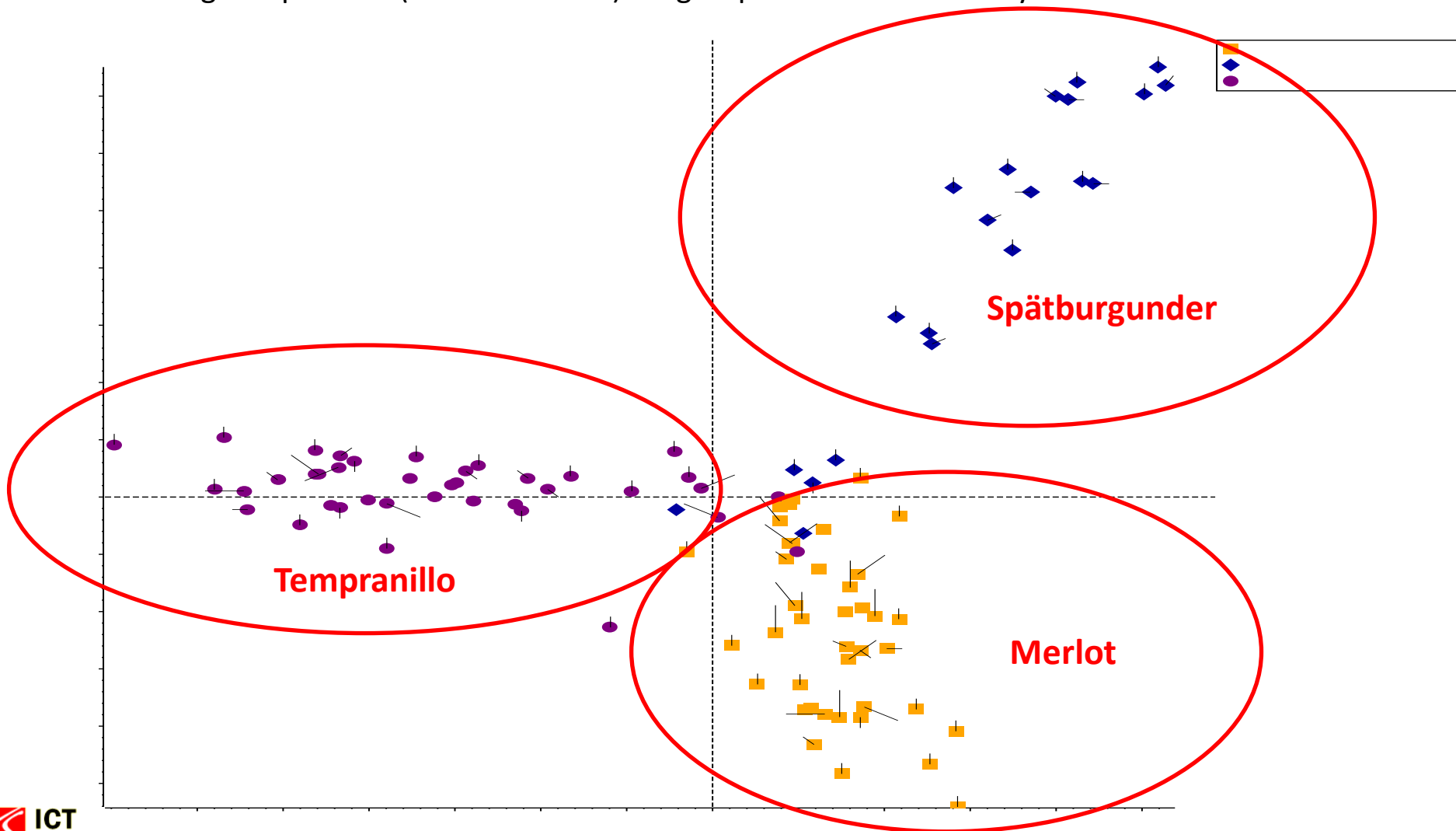
## Benefits of Q-Exactive: Metabolomics

- The superior quality of Q Exactive MS/MS data enables identification and quantitation of more compounds with greater confidence.
- Find the real differences in components across multiple sample groups with SIEVE 2.0

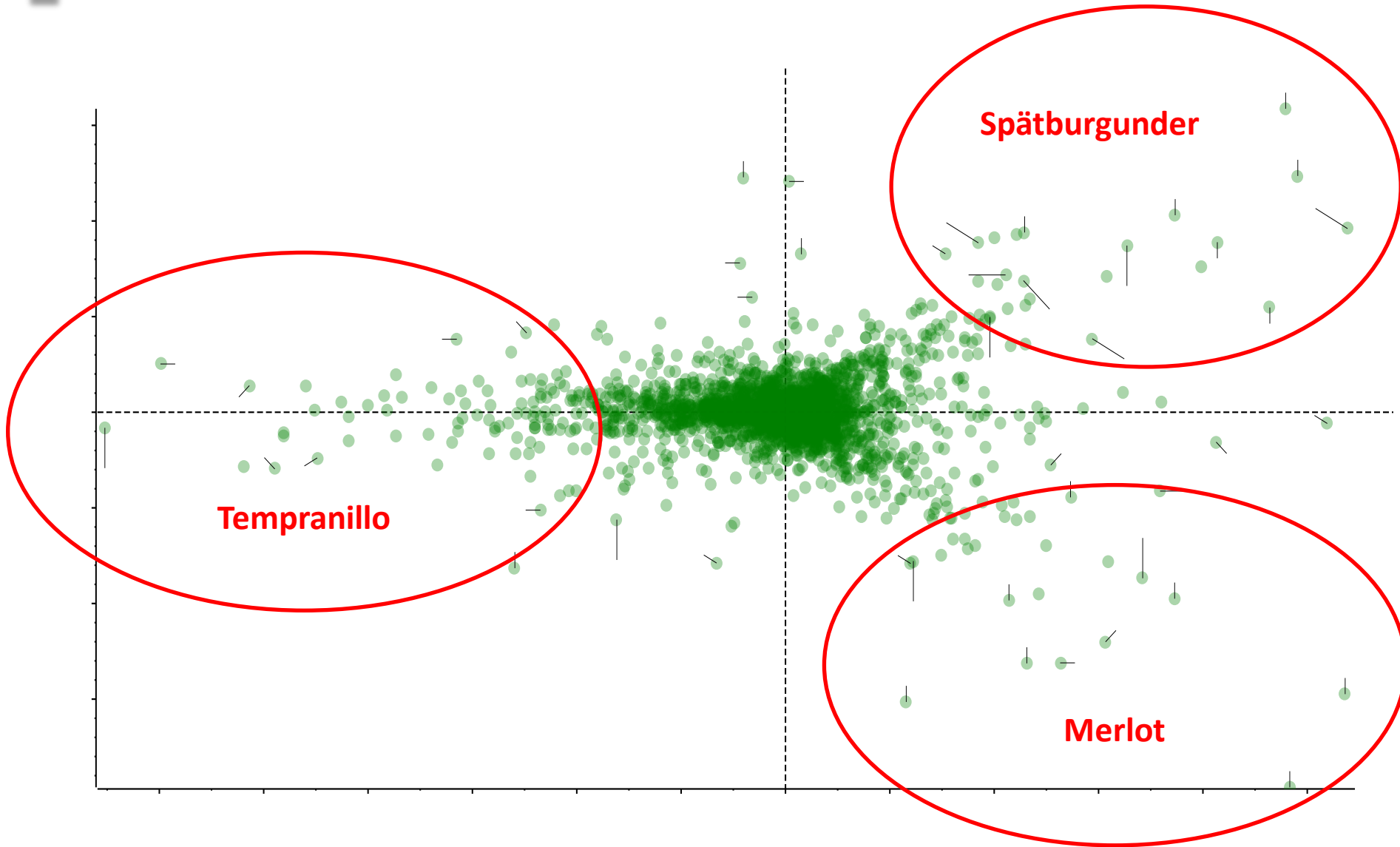


# Chemometrics analysis: Red wine samples (PCA-DA ESI+)

**Principal Component Analysis-Discriminant Analysis (PCA-DA).** PCA calculates the best discriminating components without foreknowledge about groups, whereas discriminant analysis calculates the best discriminating components (= discriminants) for groups that are defined by the user.



# Chemometrics analysis: red wine samples (PCA-DA ESI+)



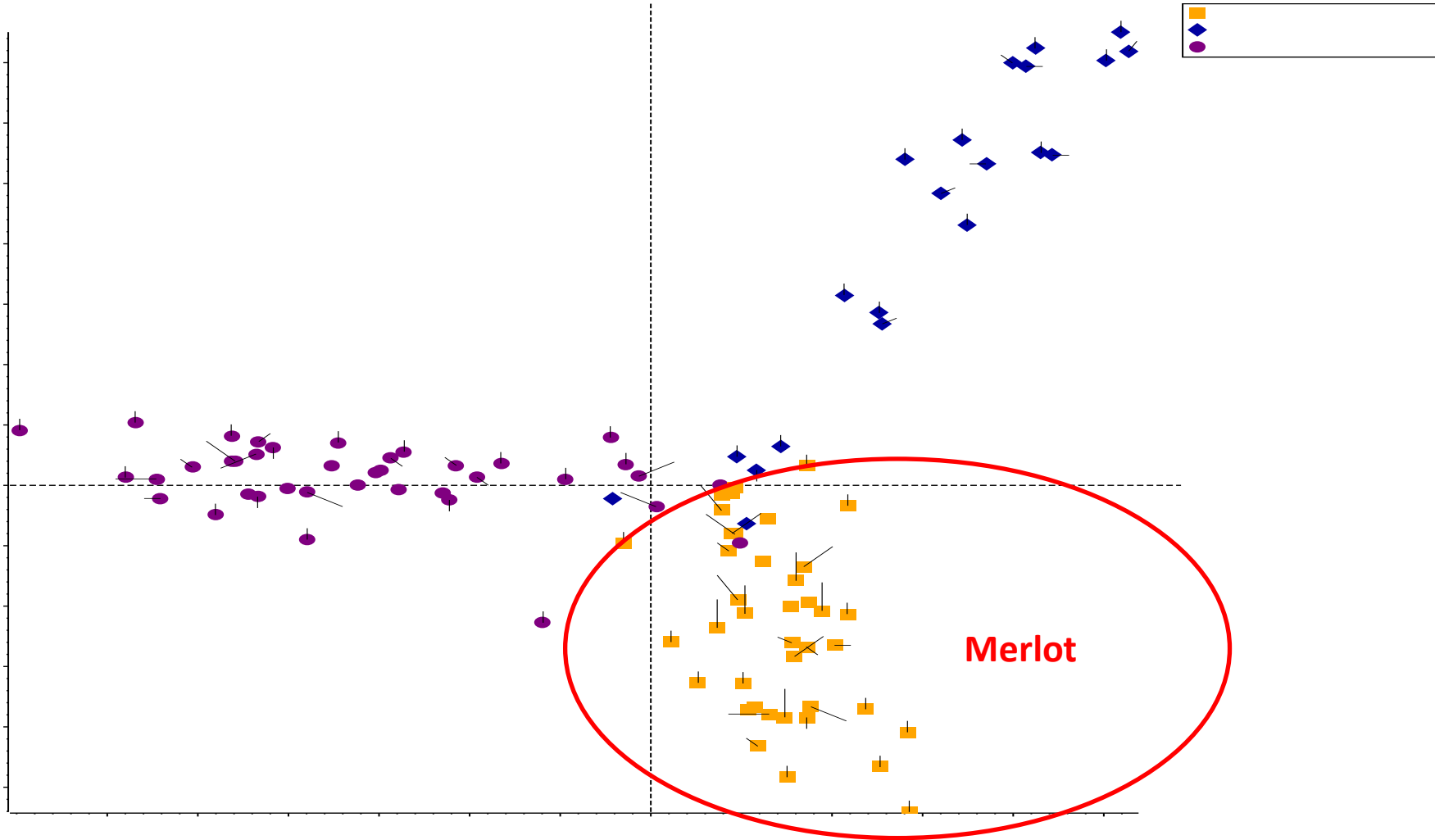
# Identification of markers: Workflow

- Identification of markers using PCA discriminant analysis.
- Xcalibur<sup>®</sup>:
  - **Full MS** extraction ion chromatogram and estimation of molecular formula.
  - **dd-MS<sup>2</sup>** data depending acquisition (Pathway)

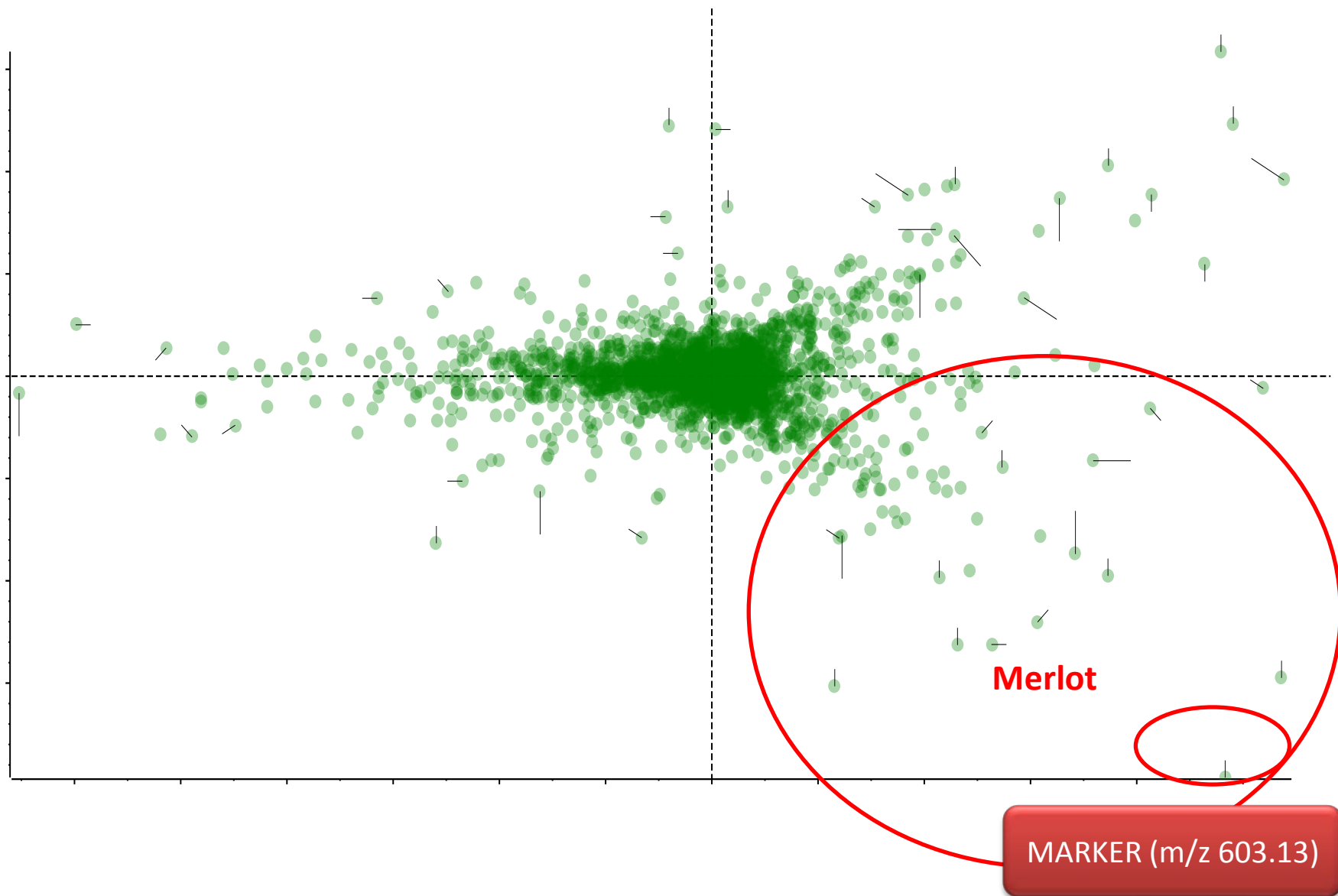
- Libraries:



# Chemometrics analysis: Red wine samples (PCA-DA ESI+)



# Chemometrics analysis: Merlot markers



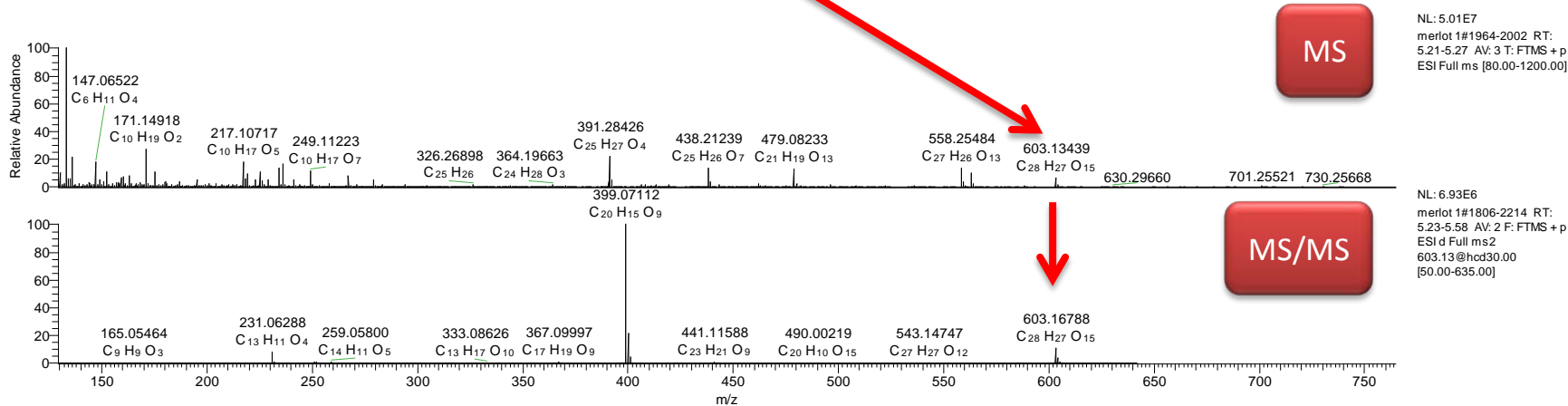
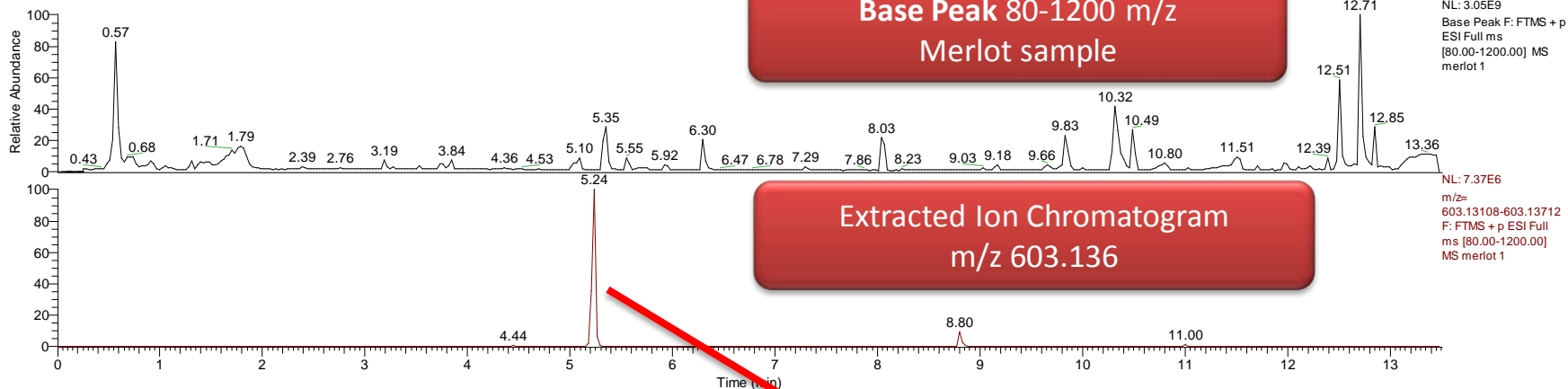
# Chemometrics analysis: Merlot markers (Workflow)

C:\Thermo\...\dd settings\Data\merlot 1

5/15/2013 11:41:40 AM

m1

RT: 0.00 - 13.50





# Chemometrics analysis: Merlot markers (Workflow)

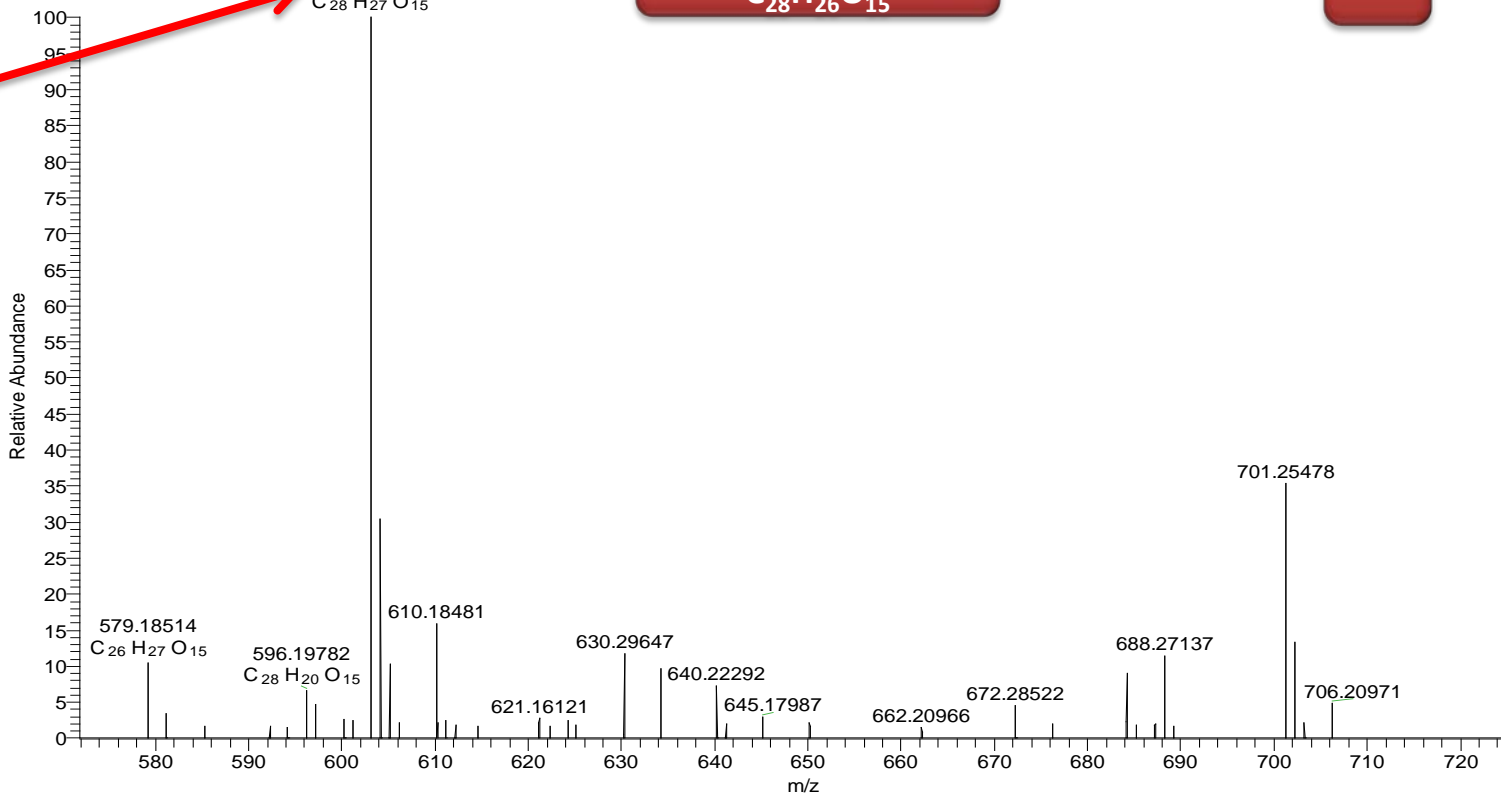
Molecular Formula



MS

merlot 1 #1960-1979 RT: 5.18-5.21 AV: 2 NL: 1.26E6  
T: FTMS + p ESI Full ms [80.00-1200.00]

603.13464  
 $\text{C}_{28}\text{H}_{27}\text{O}_{15}$



Elemental composition

Single mass

Mass: 603.1343

Max. results 10

Calculate

Idx	Formula	RDB	Delta ppm
1	C <sub>27</sub> H <sub>21</sub> O <sub>10</sub> N <sub>7</sub>	21.0	-0.231
2	C <sub>28</sub> H <sub>27</sub> O <sub>15</sub>	15.0	-0.243
3	C <sub>28</sub> H <sub>25</sub> O <sub>14</sub> N <sub>3</sub>	16.0	1.983
4	C <sub>25</sub> H <sub>19</sub> O <sub>9</sub> N <sub>10</sub>	21.5	1.992
5	C <sub>29</sub> H <sub>23</sub> O <sub>11</sub> N <sub>4</sub>	20.5	-2.460
6	C <sub>24</sub> H <sub>23</sub> O <sub>13</sub> N <sub>8</sub>	16.5	4.210
7	C <sub>30</sub> H <sub>19</sub> O <sub>7</sub> N <sub>8</sub>	25.5	-4.678
8	C <sub>22</sub> H <sub>21</sub> O <sub>12</sub> N <sub>9</sub>	17.0	6.436

File... List Simulate

Limits

Charge: 1

Nitrogen-Rule: Do not use

Mass tolerance: 10.00 ppm

RDB equiv: -1.0-100.0

Elements in use

Isotope	Min	Max	DB eq.	Mass
14 N	0	10	0.5	14.003
16 O	0	15	0.0	15.995
12 C	0	30	1.0	12.000
1 H	0	60	-0.5	1.008

Load... Save as... Apply Help

# Chemometrics analysis: Merlot markers

**Eriodictyol 7-(6-galloylglucoside)** is found in fruits. Eriodictyol 7-(6-galloylglucoside) could be a constituent of the leaves and branches of *Vitis Vinifera*.

Scripps Center For Metabolomics  
METLIN: Metabolite and Tandem MS Database

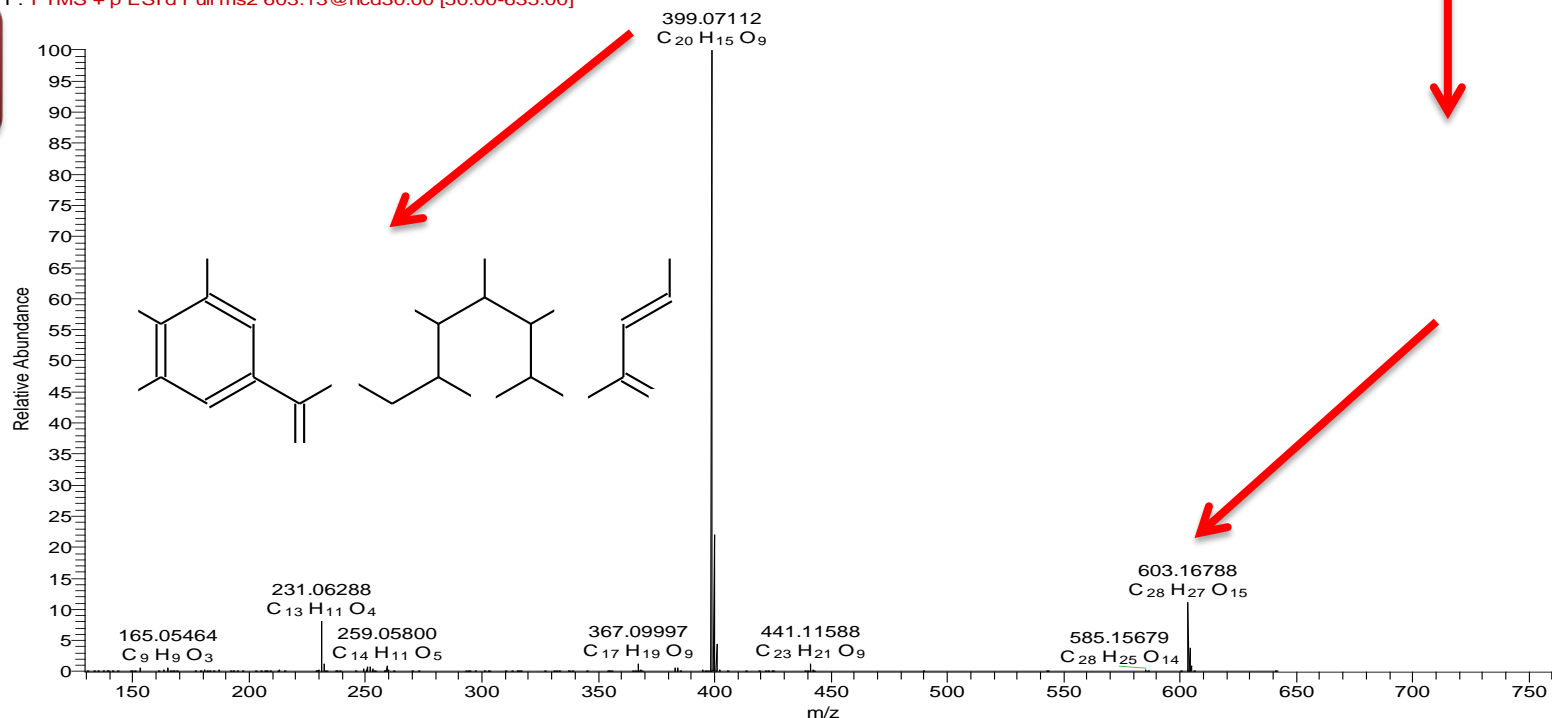


**Eriodictyol 7-(6-galloylglucoside)**

$C_{28}H_{26}O_{15}$   
Flavonoid Glycosides

dd-MS<sup>2</sup>

merlot 1 #1806-2214 RT: 5.23-5.58 AV: 2 NL: 6.93E6  
F: FTMS +p ESI d Full ms2 603.13@hcd30.00 [50.00-635.00]



# Conclusions

- Q Exactive is well suited for routine in-depth metabolome analysis with fast high-resolution MS and MS/MS scan rate, and high quality HR/AM HCD spectrum.
- The applicability of UHPLC-Q Exactive for WINE (-omics) has been demonstrated, providing authentication of grape varieties.

# Future trends

- Chemiometrics analysis:
  - ESI- data PCA and PCA-DA
  - ESI+ and ESI- data using SIMCA (PLS-DA, OPLS-DA and O2PLS-DA)
- Authentication:
  - Grape varieties
  - Origin and/or area
  - Quality

# ASSESSMENT OF OILS AND FAT OXIDATION EXTENT



# CHANGES OF FATS /OILS DURING PROCESSING

ALTERATION	CAUSATIVE AGENT	COMPOUNDS ORIGINATED
HYDROLYSIS	moisture	<ul style="list-style-type: none"><li>◆ Fatty acids</li><li>◆ Diacylglycerols</li></ul>
OXIDATION	air	<ul style="list-style-type: none"><li>◆ Oxidized monomeric triacylglycerol</li><li>◆ Oxidized dimeric and oligomeric triacylglycerols</li><li>◆ Volatile compounds (aldehydes, ketones, alcohols, hydrocarbons, etc.)</li></ul>
THERMAL CHANGES	temperature	<ul style="list-style-type: none"><li>◆ Cyclic monomeric triacylglycerols</li><li>◆ Isomeric monomeric triacylglycerols</li><li>◆ Nonpolar dimeric and oligomeric triacylglycerols</li></ul>

# AUTOOXIDATION OF FATS / OILS

- ➔ Accumulation of oxidation products in oils / fats and fried foods
- ➔ **Adverse health effects of oxidation products upon ingestion**

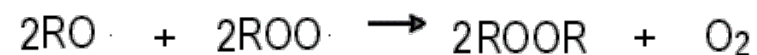
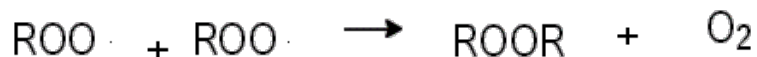
## Initiation



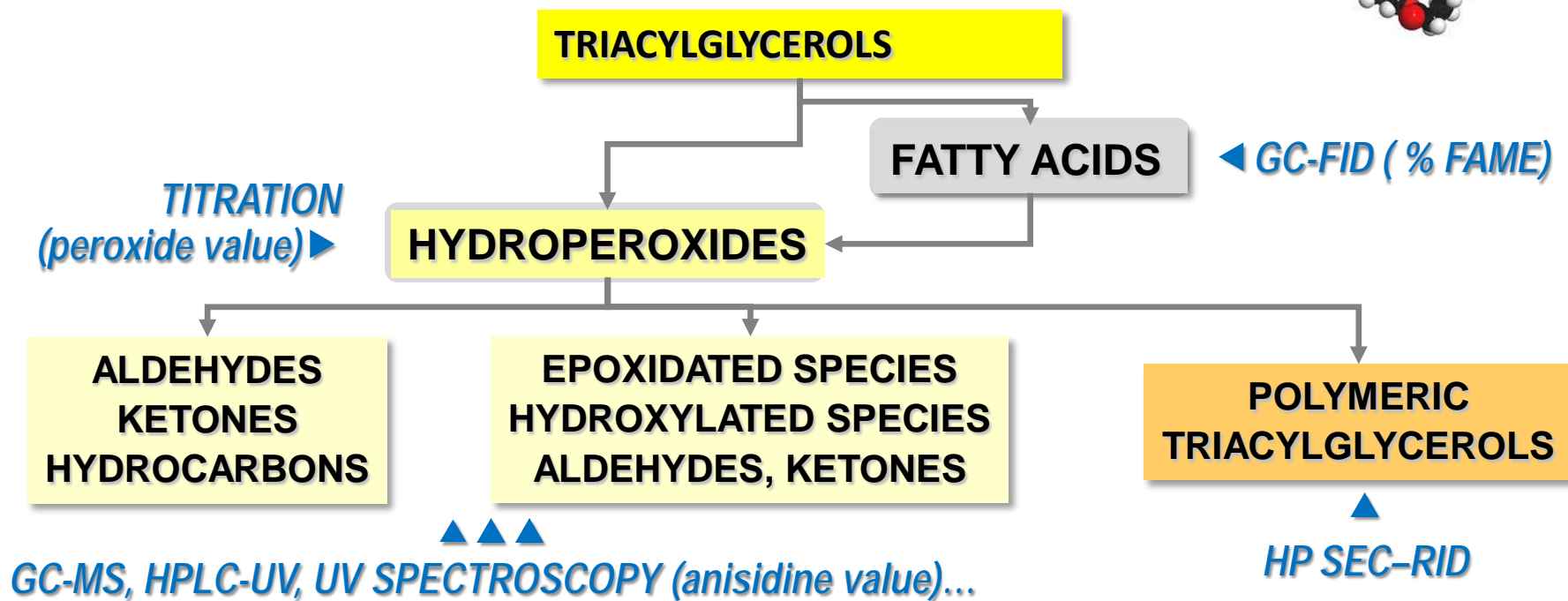
## Propagation



## Termination



# Conventional strategies in oils / fats quality assessment



High-throughput analytical methods for monitoring of oxidative degradation some innovation needed....



# Samples

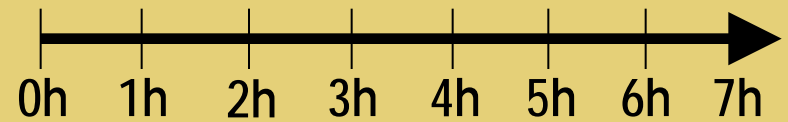
various fresh refined vegetable oils were sourced from retail market

- OLIVE OIL (PV 4.6)\*
- RAPESEED OIL (PV 2.0)\*
- SOYBEAN OIL (PV 2.2)\*
- SUNFLOWER OIL (PV 2.3)\*
- PALM OIL (PV 0.6)\*



## HEAT-TREATMENT

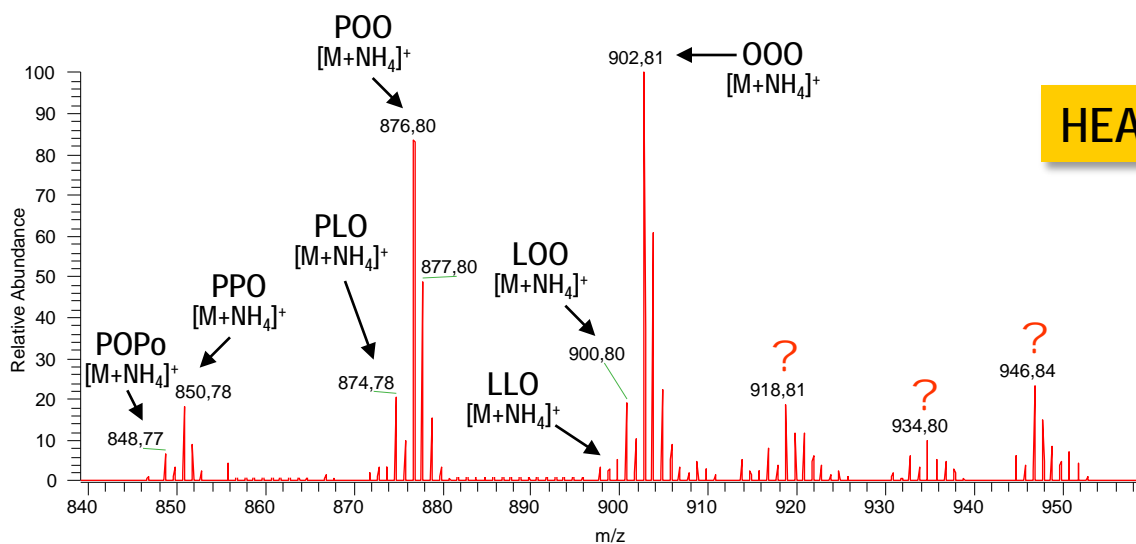
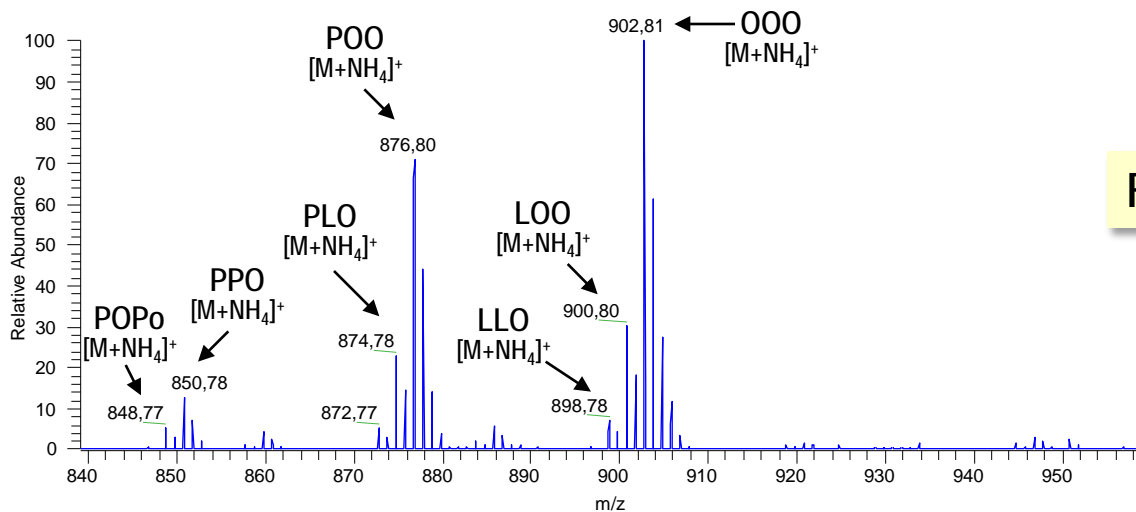
- Heating of 25g of oil in a beaker ( $n=3$ )
- Heating time 0 – 7 hours
- Controlled temperature  $180 \pm 2^\circ\text{C}$



\* PEROXIDE VALUE (mekv. of active oxygen per kg)

# OXIDATION OF TRIACYLGLYCEROLS

DART-MS SPECTRA (POSITIVE ION MODE): DILUTED **OLIVE OIL**

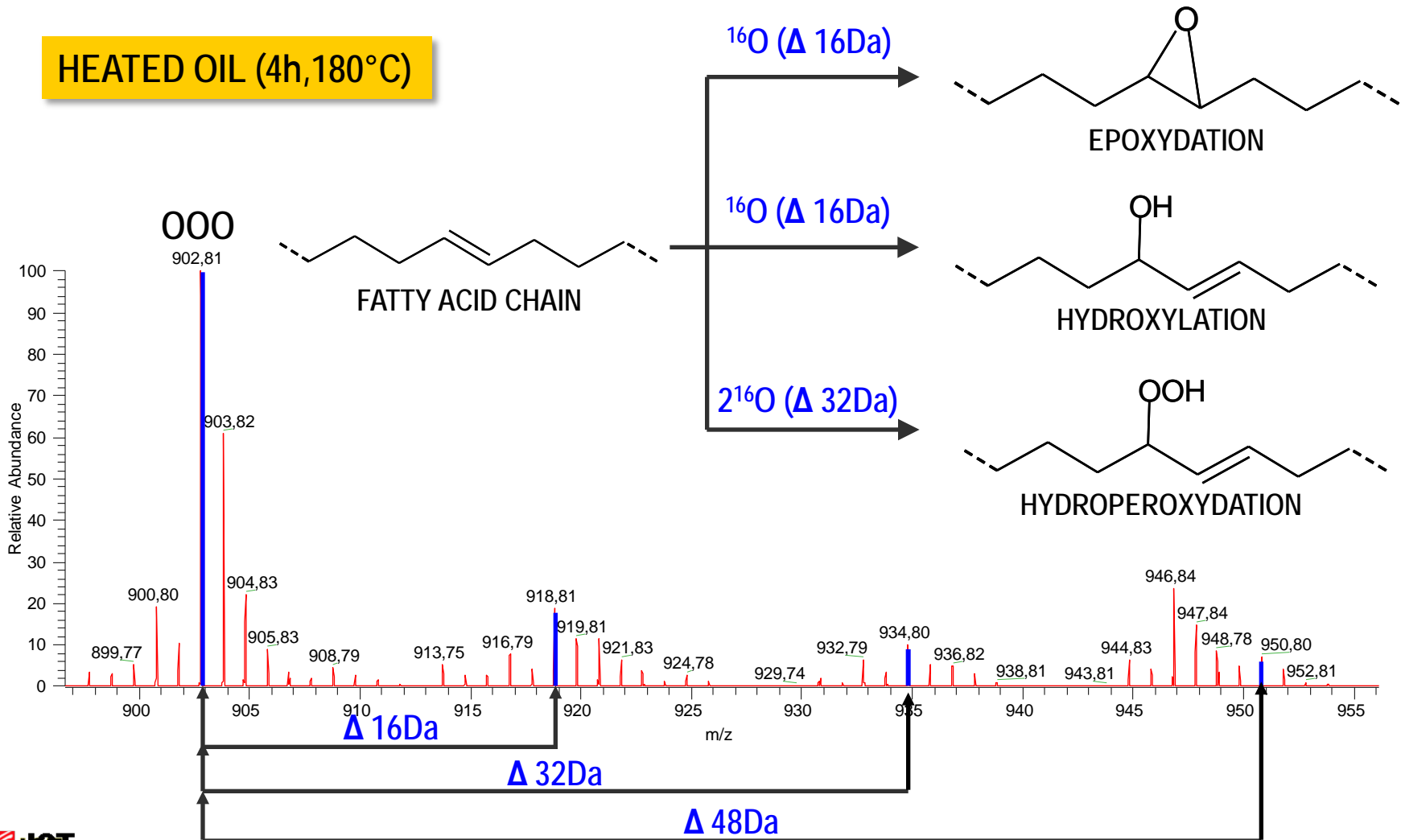


# OXIDATION OF TRIACYLGLYCEROLS



DART-MS SPECTRA (POSITIVE ION MODE): DILUTED **OLIVE OIL**

HEATED OIL (4h, 180°C)

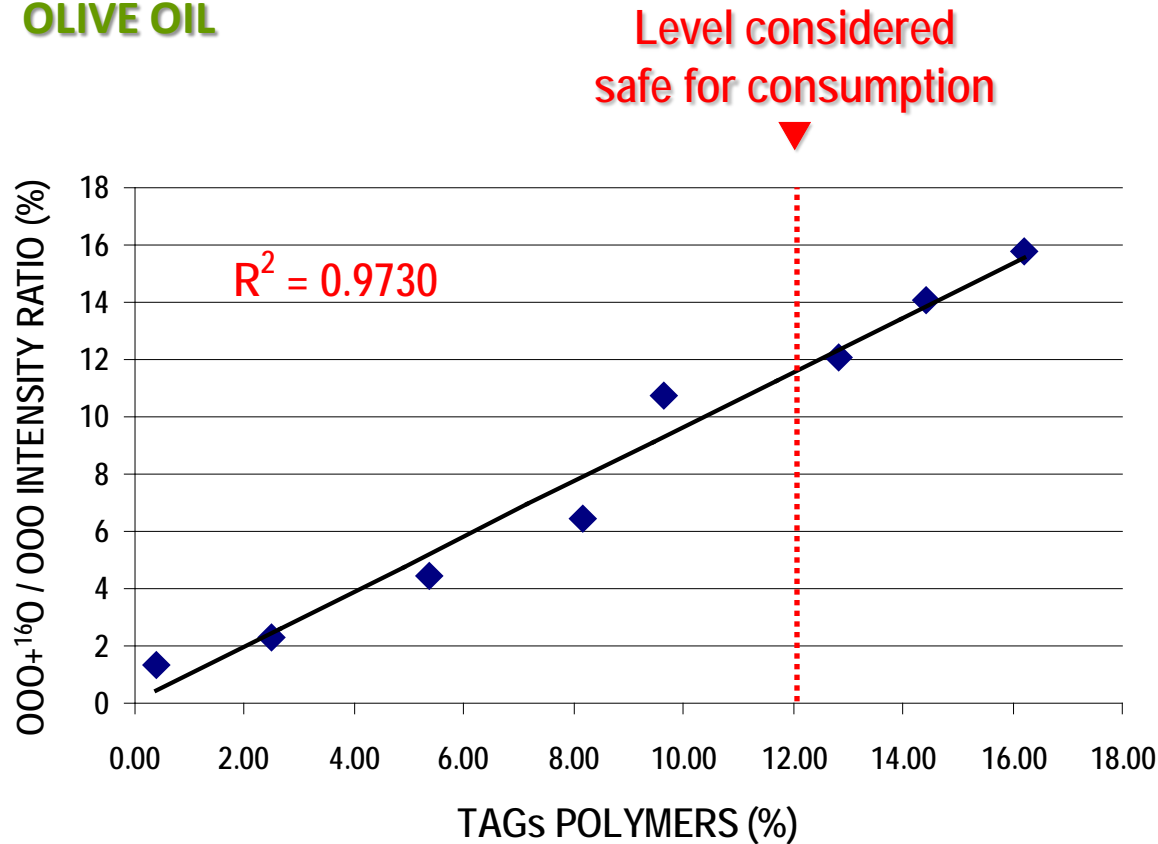


# Oxidation of triacylglycerols



## ➔ CORRELATION WITH TAG POLYMERS CONTENT

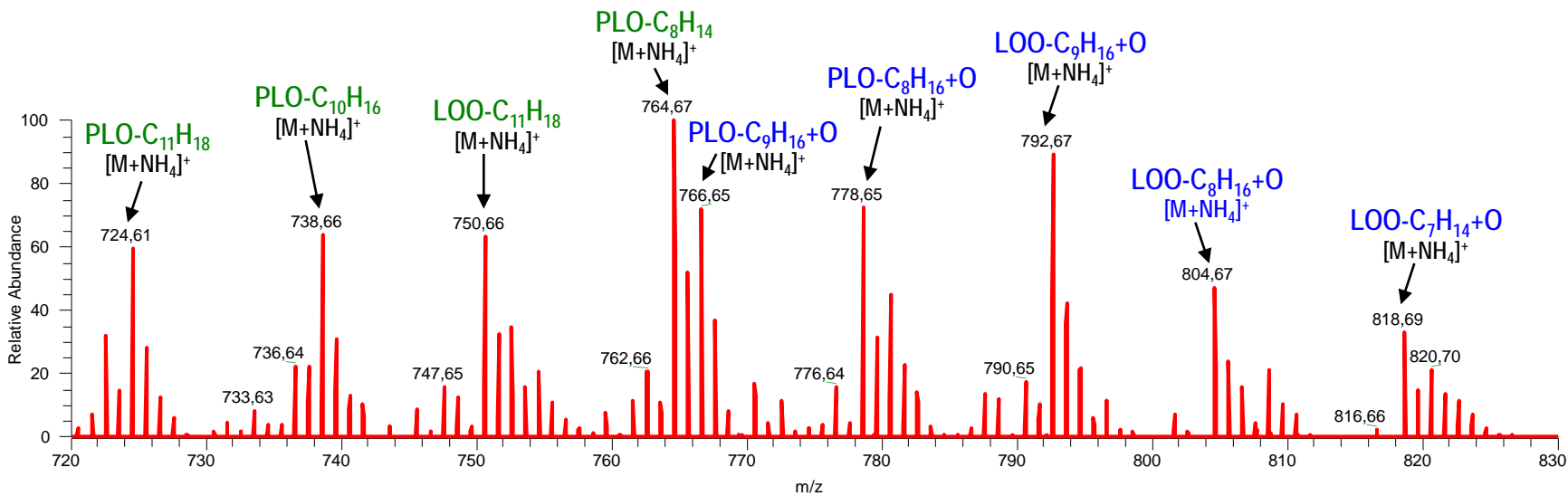
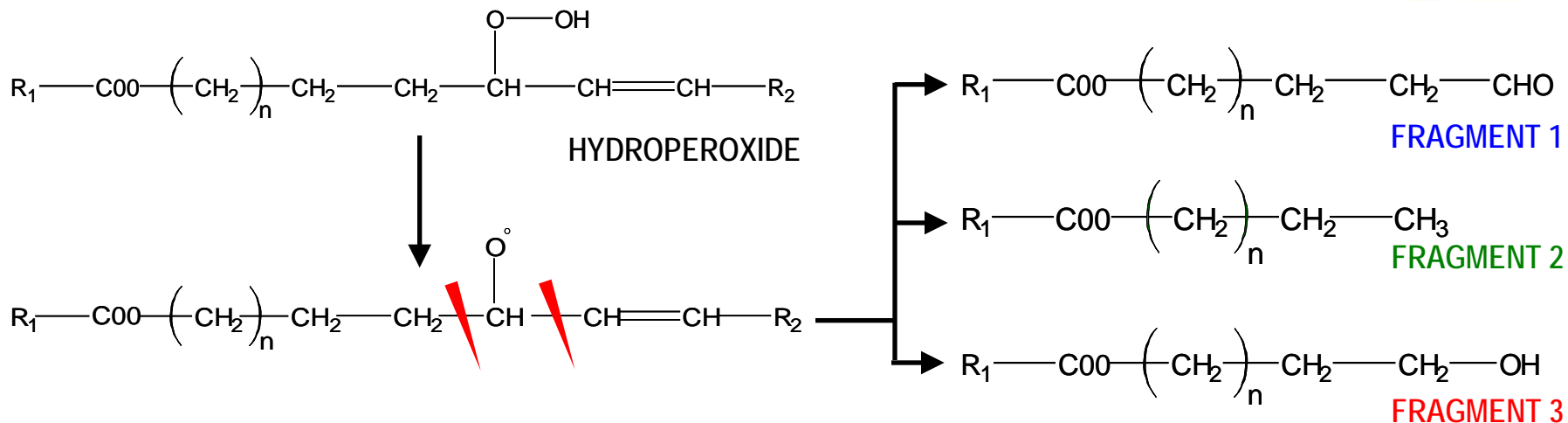
### OLIVE OIL



# β-SCISSION FRAGMENTS OF TAGs



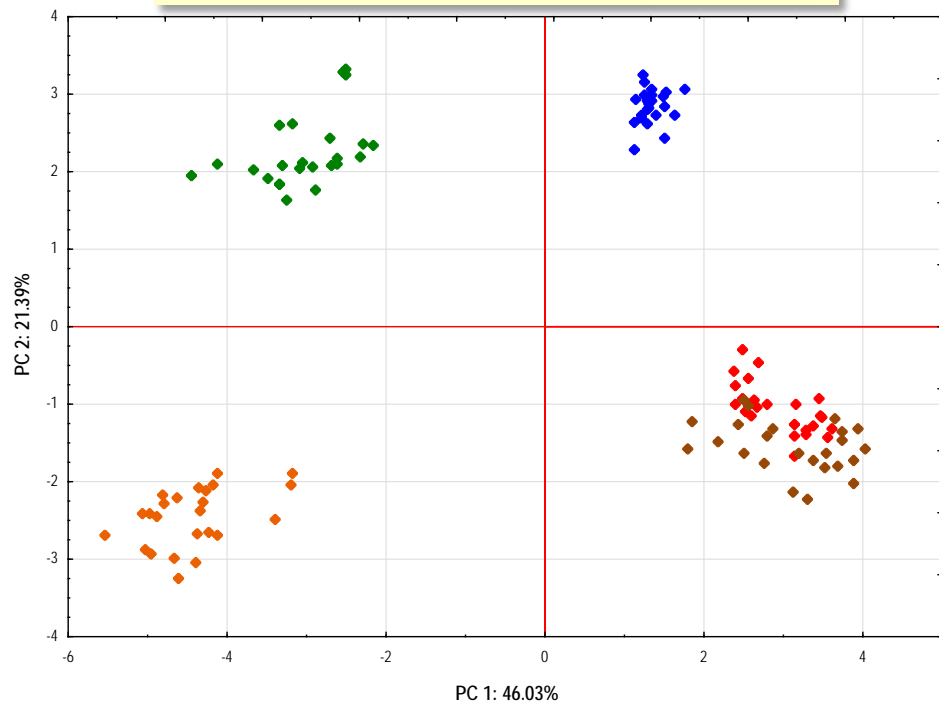
DART-MS SPECTRA (POSITIVE ION MODE): DILUTED **OLIVE OIL**



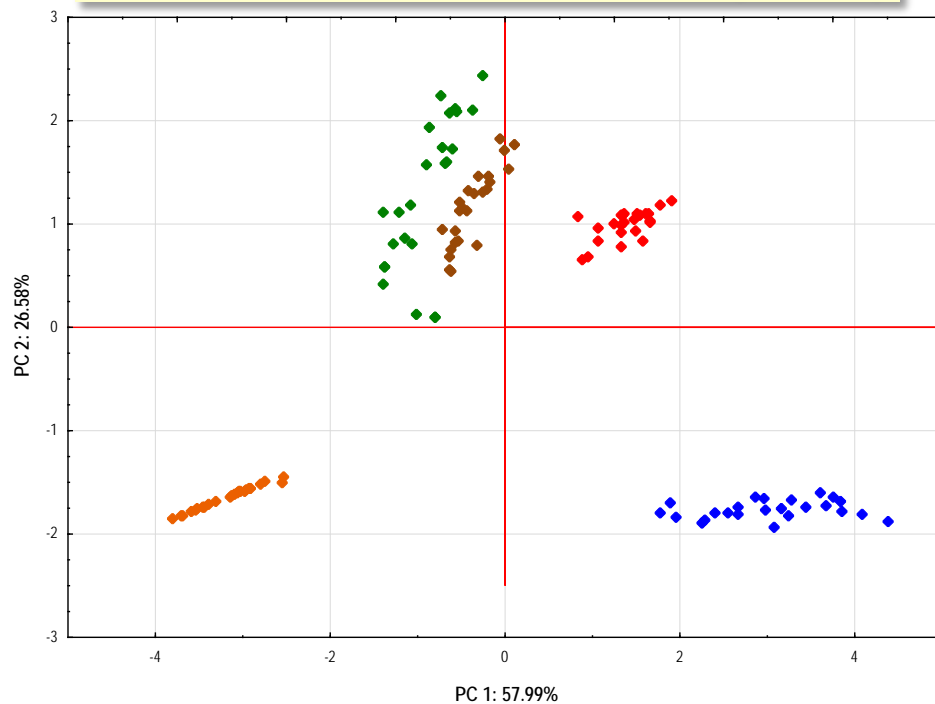
# COMPREHENSIVE VIEW ON THE DATA

## PRINCIPAL COMPONENT ANALYSIS

TAGs & OXIDIZED TAGs (22 IONS)



STEROLS & OXIDIZED STEROLS (8 IONS)



- SOYBEAN OIL
- OLIVE OIL
- RAPESEED OIL
- SUNFLOWER OIL
- PALM OIL

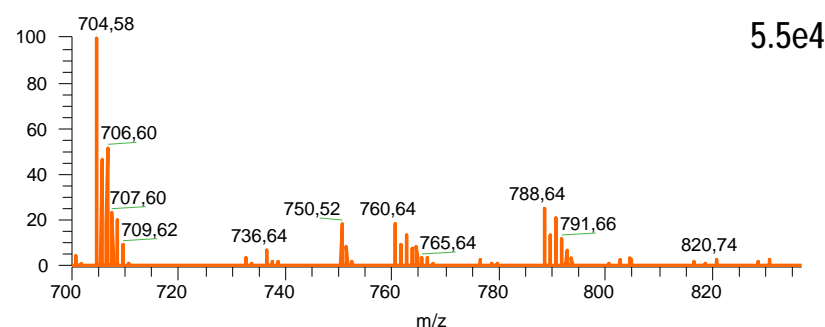
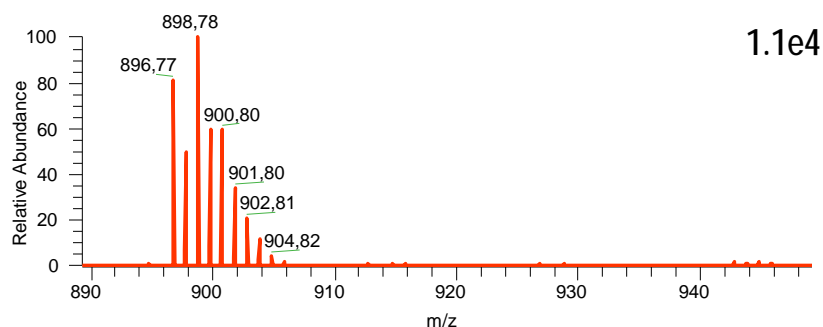
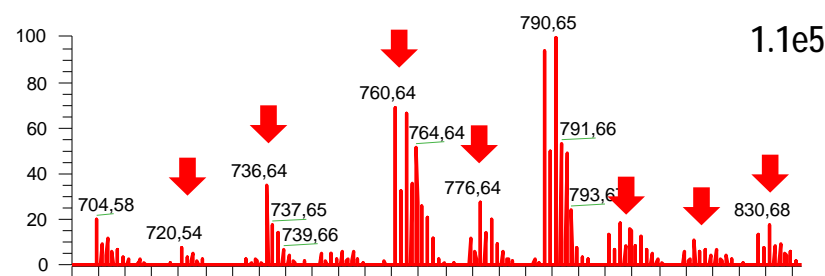
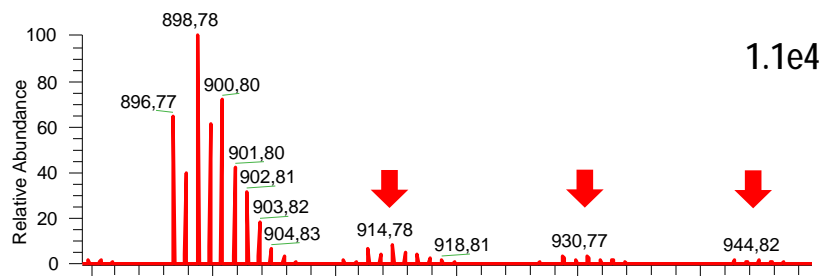
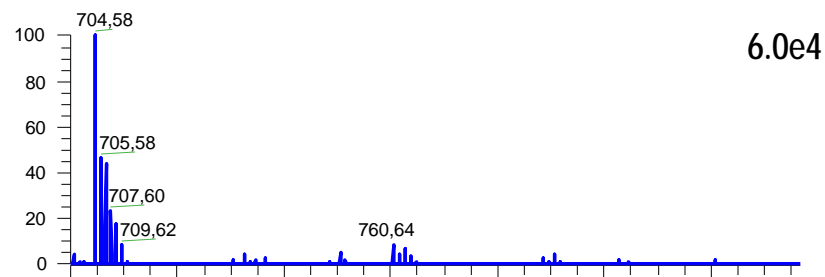
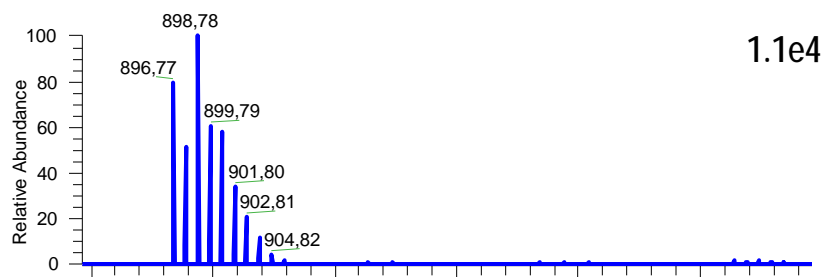
# MONITORING OF ANTIOXIDANT ACTIVITY

- ▶ OXIDATION OF SUNFLOWER OIL (120°C, 2h) WITH AND WITHOUT ADDITION OF ANTIOXIDANT (GALLIC ACID, 100ppm)...

DART-MS SPECTRA (POSITIVE ION MODE)

NATIVE / OXIDATED TRIACYLGLYCEROLS

β-SCISSION FRAGMENTS

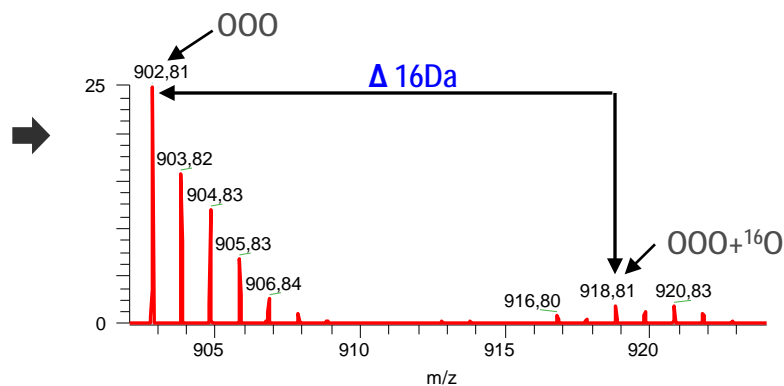
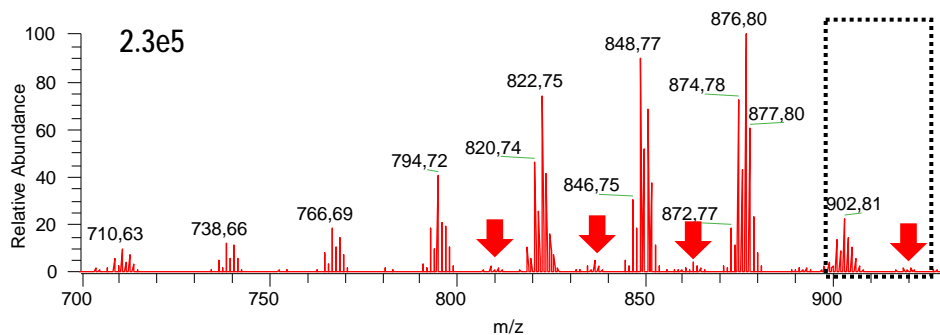


# OXIDIZED TAGs IN HUMAN ADIPOSE TISSUE

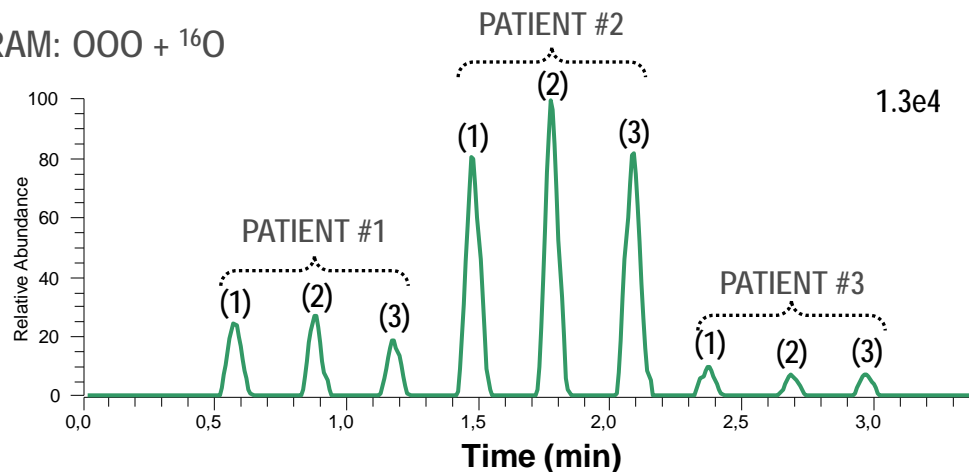
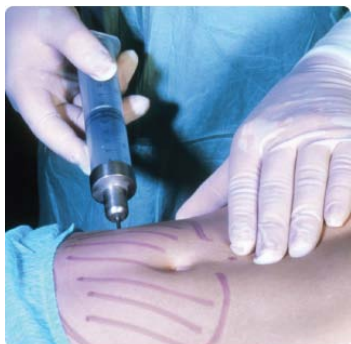
➔ HUMAN ADIPOSE TISSUE SOURCED FROM LIPOSUCTION SURGERY  
(3 SAMPLES COLLECTED FROM 3 DIFFERENT PATIENTS)

DART-MS SPECTRA (POSITIVE ION MODE)

NATIVE / OXIDATED TRIACYLGLYCEROLS



EXTRACTED ION CHRONOGRAM: 000 + 160





# CONCLUSIONS: FUTURE CHALLENGES

- **Databases vs. standardization**
- **Diversity/variation of metabolomic data**
- **More efficient ways of identifying markers**
- **More efficient ways of coupling:**
  - ↳ Instruments → Data processing → Chemometric tools
- **New instrumentation platforms**

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- Food-Omics
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- Organic crops & foodstuffs
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- Nanoparticles
- Packaging contaminants
- Pesticide residues
- Processing contaminants
- Veterinary drugs



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<b>Prof. Franz Ulberth</b>	JRC, Institute for Reference Materials and Measurements, Geel, B
<b>Dr. Frans Verstraete</b>	European Commission, DG Health and Consumers (DG SANCO), B
<b>Dr. Zhihua Ye</b>	Chinese Academy of Agricultural Sciences, Beijing, CN

## KEYNOTE SPEAKERS INTRODUCING SYMPOSIUM TOPICS:

- Elke ANKLAM (European Commission, DG Joint Research Centre Director, Institute for Reference Materials and Measurements, Geel, B)

### **NEW RESEARCH CHALLENGES IN FOOD QUALITY AND SAFETY CONTROL**

- Paul BRERETON (The Food and Environment Research Agency, York, UK)

### **ANALYTICAL CHALLENGES AND OPPORTUNITIES FOR ASSURING INTEGRITY OF THE FOOD SUPPLY**

- Hermann BROLL (Federal Institute for Risk Assessment, Berlin, D)

### **RECENT TRENDS IN APPLICATION OF MOLECULAR BIOLOGY BASED METHODS IN FOOD ANALYSIS**

- Alejandro CIFUENTES (National Research Council of Spain, Institute of Food Science Research, Laboratory of Foodomics, Madrid, ES)

### **FOOD-OMICS: PRESENT AND FUTURE CHALLENGES IN FOOD ANALYSIS**

- Christopher ELLIOTT (Queen's University Belfast, Belfast, UK)

### **BIOASSAYS FOR BIOMARKERS: TOOLS FOR DETERMINING HUMAN EXPOSURE TO FOOD TOXINS**

- Jacob DE JONG (RIKILT Wageningen UR, Wageningen, NL)

### **LESSONS LEARNT FROM THE CONFIDENCE PROJECT: CONTAMINANTS IN FOOD AND FEED – INEXPENSIVE DETECTION FOR CONTROL OF EXPOSURE**

- Rudolf KRŠKA (University of Natural Resources and Life Sciences, Vienna, IFA-Tulln, A)

### **MYCOTOXINS AND THEIR METABOLITES: FROM TARGETED TO UNTARGETED ANALYSIS**

- Erich LEITNER (Graz University of Technology, Graz, A)

### **STRATEGIES FOR FLAVOUR AND OFF-FLAVOUR DETECTION**

- Ryszard LOBINSKI (CNRS-University of Pau, Pau, FR)

### **METALS AND METALLOIDS IN FOOD: SPECIATION AND –OMICS**

# RAFA 2013 keynote speakers, cont.

- Bert POPPING (Eurofins Scientific Group, Hamburg, D)  
**THE EVOLUTION OF ALLERGEN METHODS: WILL TOMORROWS METHOD JUST BE BETTER OR ALSO FASTER?**
- Peter SCHIEBERLE (Food Chemistry, Technical University of Munich, Garching, D)  
**THE CHEMISTRY OF FOOD FLAVOURS: SIMPLY PLEASURE OR BEYOND?**
- Yang SHUMING (Institute of Quality Standards & Testing Technology for Agro-Products, Chinese Academy of Agricultural Sciences, Beijing, CN)  
**Title to be confirmed**
- Michele SUMAN (Barilla Food Research Labs, Parma, I)  
**SUMMARY & DISCUSSION PLATFORM: INDUSTRY PERSPECTIVES**
- Bert VAN BAVEL (Örebro University, Örebro, S)  
**THE IMPORTANCE OF QA/QC AND THE LATEST DEVELOPMENTS IN POP ANALYSIS REGULATED BY THE STOCKHOLM CONVENTION**
- Frans VERSTRAETE (European Commission, DG Health and Consumers (DG SANCO), B)  
**RECENT AND FUTURE DEVELOPMENTS IN THE REGULATION OF FOOD CONTAMINANTS**
- Stefan WEIGEL (RIKILT Wageningen UR, Wageningen, NL)  
**ANALYSIS OF NANOPARTICLES IN FOOD: FROM CHALLENGE TO ROUTINE?**

# Acknowledgement:

all the nice colleagues from Sicily

**ThermoFisher**  
S C I E N T I F I C