



**UNIVERSITY OF
CHEMISTRY AND TECHNOLOGY
PRAGUE**



Challenges in Mass Spectrometry Based Non-Targeted Analysis

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FOOD: A COMPLEX COCKTAIL OF CHEMICALS

Food fiber

Natural toxins /
antinutrients

Primary flavour
compounds

Biologically
active
compounds,
beneficial
(antioxidants...)

Nutrients

Minerals
Vitamins

**NATURAL COMPONENTS –
primary and secondary
metabolites**

**Products originated during
processing**

Food additives

**Fraudulent components -
'UNKNOWN'**

CONTAMINANS

Environmental
contaminants

Pesticide / vet
drugs residues

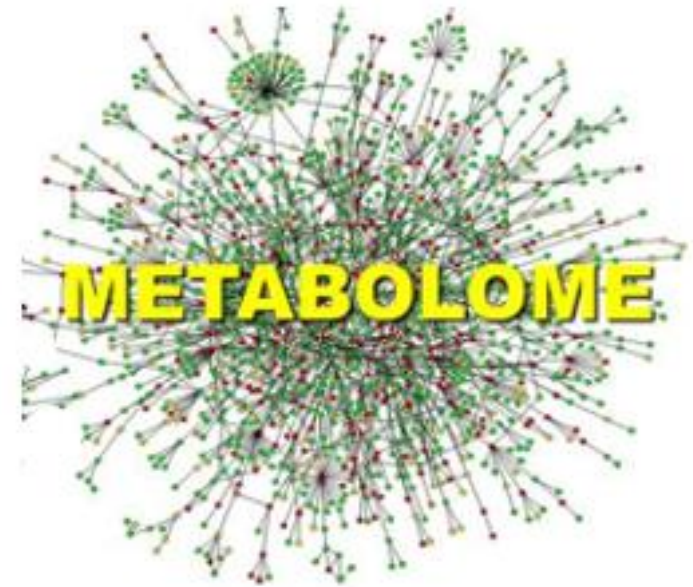
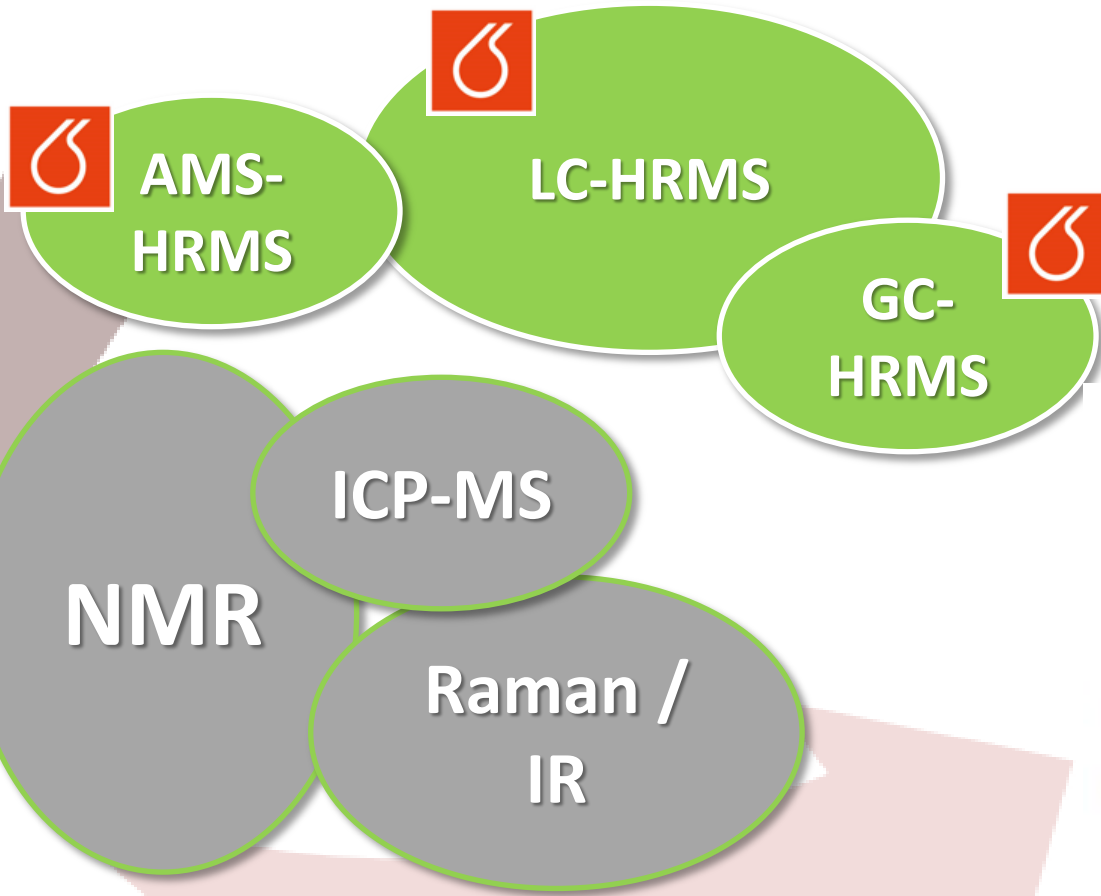
Migrants from food
contact materials

Toxic metals



Need for new analytical approaches to detect food fraud

FINGERPRINTING TECHNIQUES FOR FOOD FRAUD DETECTING





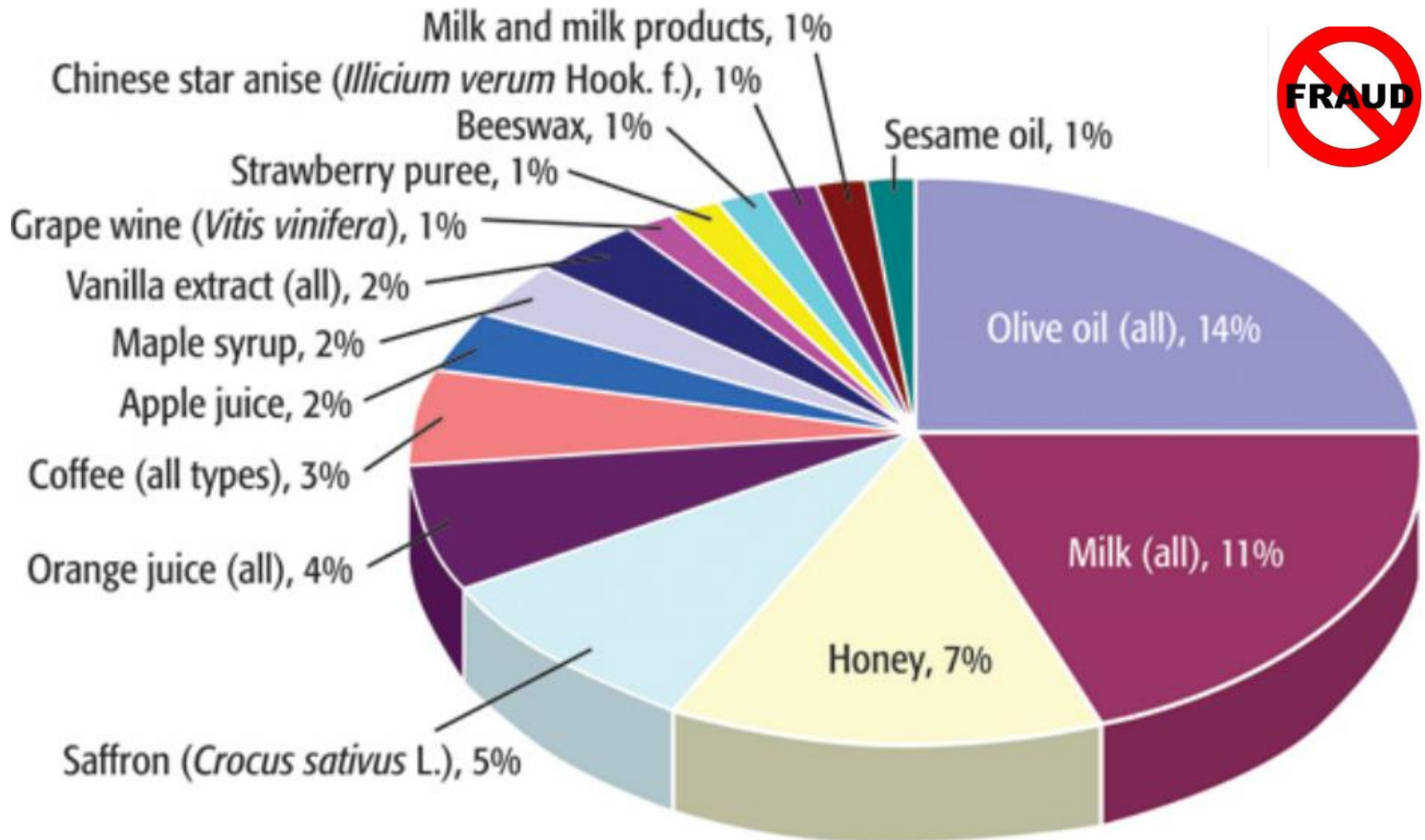
UCT Prague – wide platform of cutting edge HRMS technologies for authenticity / safety testing

TOP INSTRUMENTATION FOR EXCELLENT RESEARCH

Continuous integration of emerging technologies into food fraud fight



Fifteen most problematic food commodities for economical adulteration in the previous decade





APPLICATION POTENTIAL OF HIGH RESOLUTION MASS SPECTROMETRY

Analytica Chimica Acta 645 (2009) 56–63

Metabolomics (2012) 8:793–803

DOI 10.1007/s11306-011-0371-7

Anal Bioanal Chem (2014) 406:6791–6803

DOI 10.1007/s00216-014-7864-y

PAPER IN FOREFRONT

Metabolic fingerprinting based on high-resolution tandem mass spectrometry: a reliable tool for wine authentication?

Josep Rubert · Ondrej Lacina · Carsten Faulstich ·
Jana Hajslova

origin recognition

Tomas Cajka · Katerina Ridelova ·
Monika Tomaniova · Jana Hajslova

Metabolomics (2011) 7:500–508

DOI 10.1007/s11306-010-0266-z

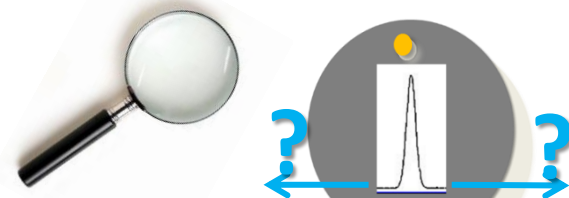


Department of Food Analysis and Nutrition, Faculty of Food and Biochemical Technology, University of Chemistry and Technology, Prague, Technická 3, 166 28 Prague 6, Czech Republic

Moving ahead with HRMS fingerprints...

'CLASSIC' APPROACH

A set of physico-chemical and/or biochemical measurements → **TARGET ANALYSIS** of one or few markers

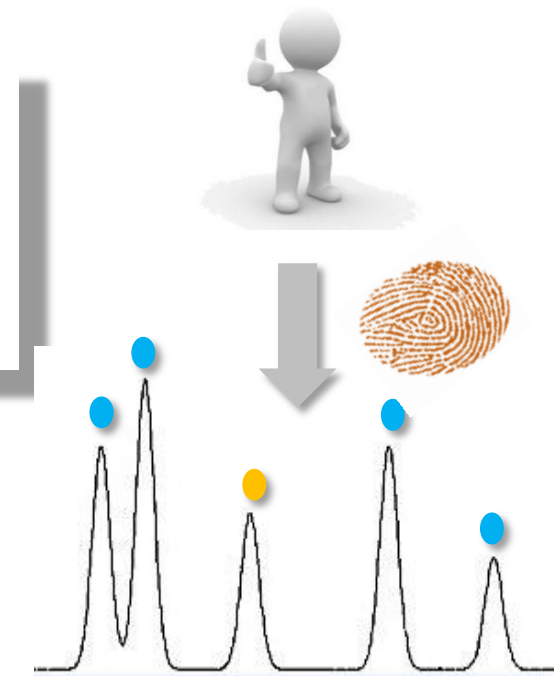


NOVEL STRATEGY

Metabolomic fingerprinting

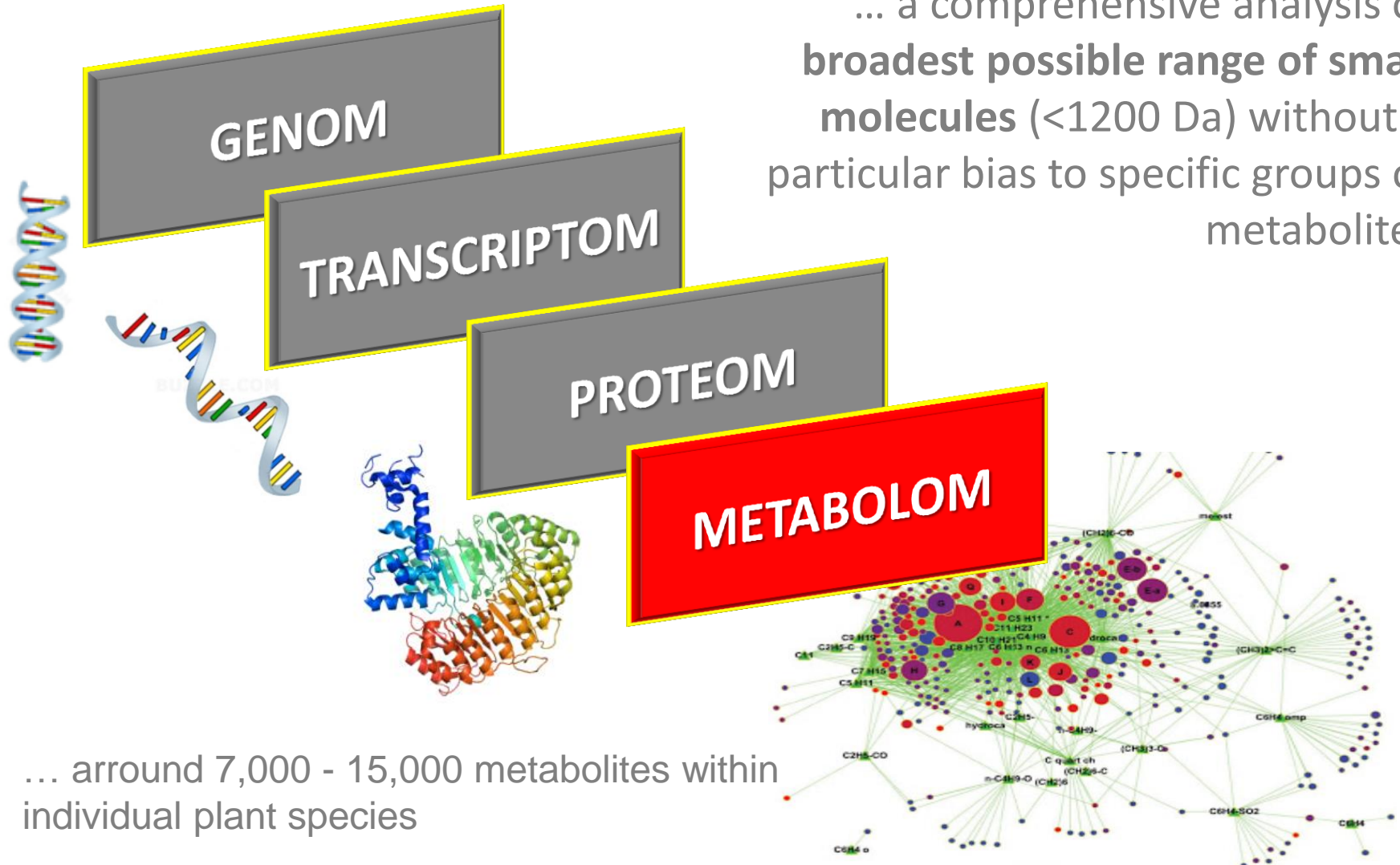
→ **NON TARGET SCREENING**

- ➔ complex characterization of matrix pattern
- ➔ detection / identification of 'unknown' components (even retrospective)
- ➔ identification of a set of composition markers



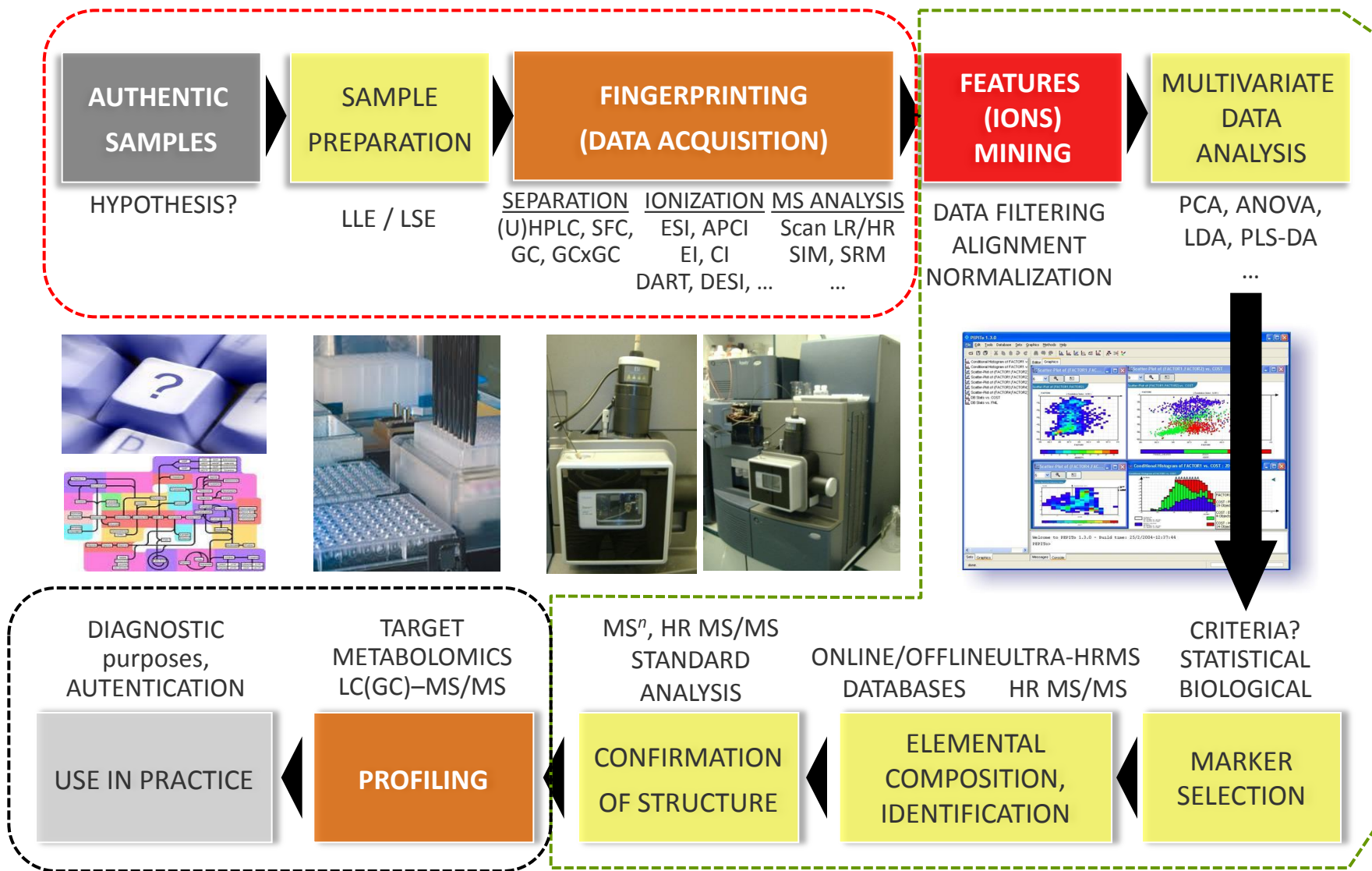
METABOLOMIC FINGERPRINTING

... a comprehensive analysis of **broadest possible range of small molecules (<1200 Da)** without a particular bias to specific groups of metabolites



... around 7,000 - 15,000 metabolites within individual plant species

TYPICAL METABOLOMIC WORKFLOW



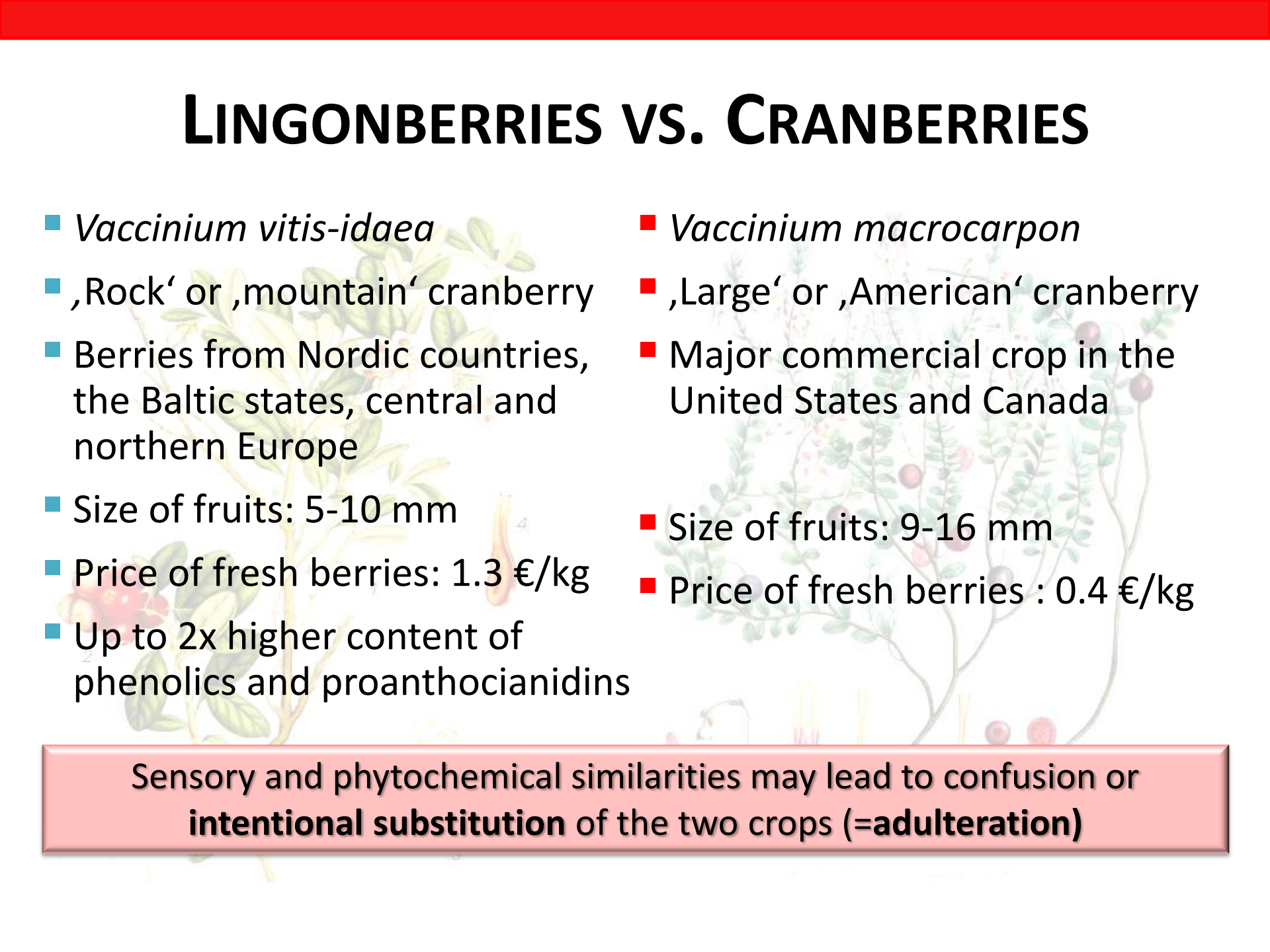
Case study # 1



DISTINGUISHING OF BOTANICAL ORIGIN OF BERRIES (LINGONBERRIES VS. CRANBERRIES)



LINGONBERRIES VS. CRANBERRIES

- 
- *Vaccinium vitis-idaea*
 - ‚Rock‘ or ‚mountain‘ cranberry
 - Berries from Nordic countries, the Baltic states, central and northern Europe
 - Size of fruits: 5-10 mm
 - Price of fresh berries: 1.3 €/kg
 - Up to 2x higher content of phenolics and proanthocyanidins
 - *Vaccinium macrocarpon*
 - ‚Large‘ or ‚American‘ cranberry
 - Major commercial crop in the United States and Canada
 - Size of fruits: 9-16 mm
 - Price of fresh berries : 0.4 €/kg

Sensory and phytochemical similarities may lead to confusion or **intentional substitution** of the two crops (=adulteration)

AIM OF THE STUDY



FINGERPRINTING

- Development of extraction procedure

- LC-HRMS analysis, data processing, markers characterization

- Assessment of stability of markers during food processing

- Analysis of markers in processed foods

- Assessment of authenticity of the lingonberry-based products

PROFILING

SAMPLES ANALYZED WITHIN THE INITIAL METABOLOMIC FINGERPRINTING (1st step)

- **Fresh samples** (stored in freezer)
 - 15 authentic samples of **lingonberries** (*Vaccinium vitis-idaea*)
 - 10 authentic samples of **cranberries** (*Vaccinium macrocarpon*)

SAMPLES ANALYZED WITHIN THE ASSESSMENT OF SUITABILITY OF MARKERS FOR PROCESSED FOODS (2nd step)

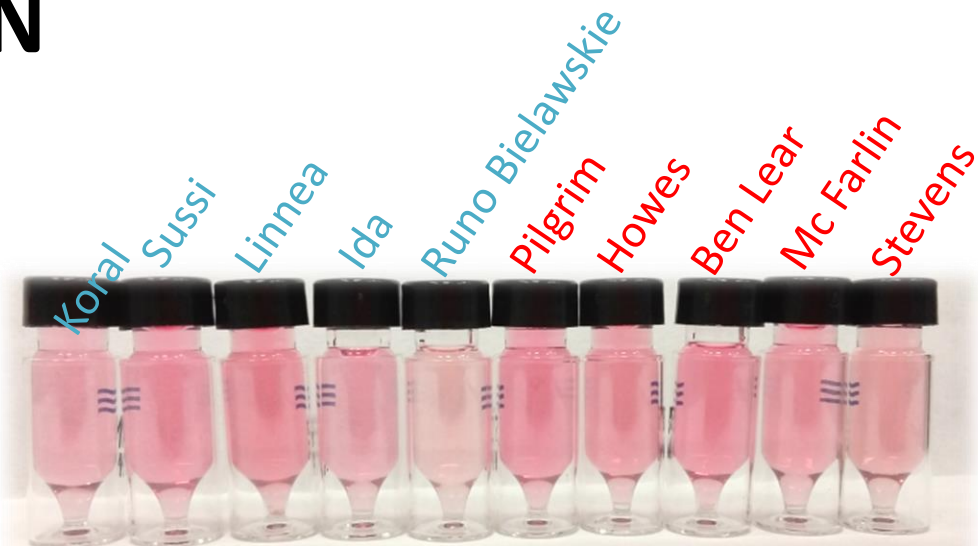
- **Dried samples**
 - 3 samples of **lingonberries** (*Vaccinium vitis-idaea*)
 - 16 samples of **cranberries** (*Vaccinium macrocarpon*)



SAMPLE PREPARATION

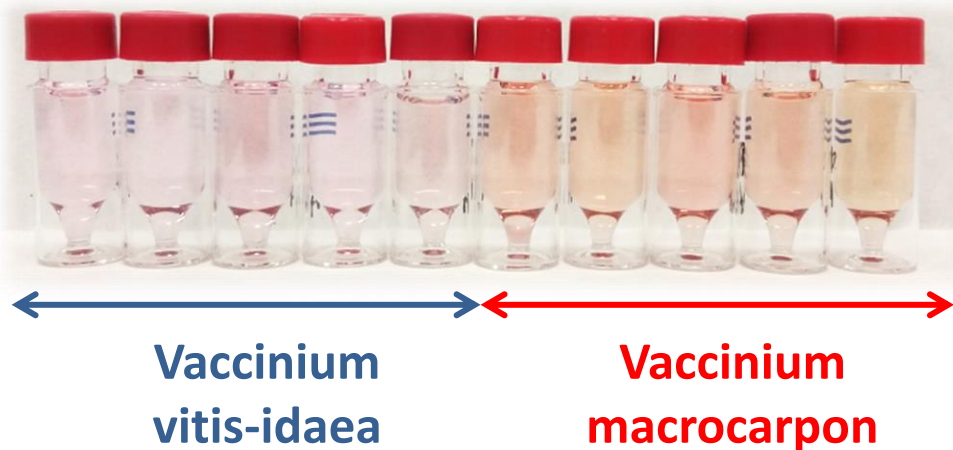
1st STEP Polar extract

1 g of sample with 5 mL of methanol
Ultraturrax
Centrifugation



2nd STEP Non-polar extract

Repeated extraction of solid residue
with 5 mL of hexane/2-propanol
Vortex
Centrifugation



UHPLC-HRMS analysis

UHPLC-Thermo Dionex UltiMate 3000

Non-polar extract

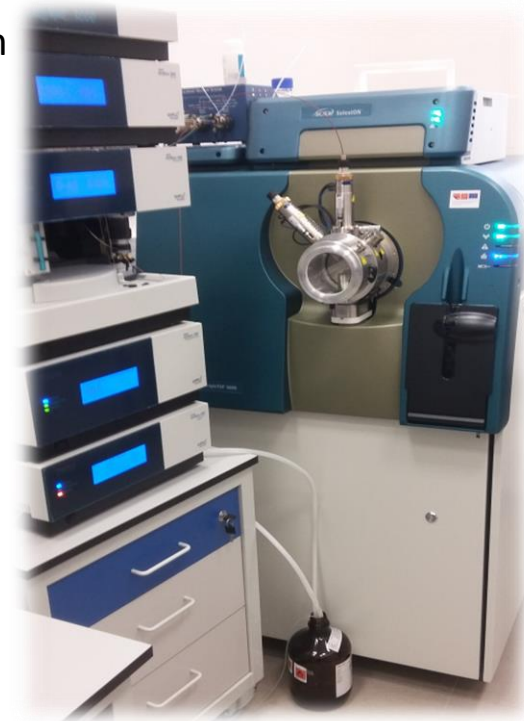
- **Column:** BEH C18
(2.1x100 mm, 1.7 μ m)
Column temperature: 60 °C
- **Mobile phase:**
A: 5 mM ammonium formate in
H₂O:MeOH (95:5) + 0,1% formic acid
B: 5 mM ammonium formate in
iPrOH:MeOH:H₂O (65:30:5) + 0,1% formic acid

Polar extract

- **Column:** HSS T3
(2.1x100 mm, 1.8 μ m)
Column temperature: 40 °C
- **Mobile phase:**
A: 5 mM ammonium
formate in H₂O
B: MeOH

MS(MS/MS)-TripleTOF™ 6600 (Sciex)

- m/z range: 100-1200
- Ionisation technique: ESI +/-
- This technology allowed: Sensitivity, Speed, Resolution and Mass Accuracy



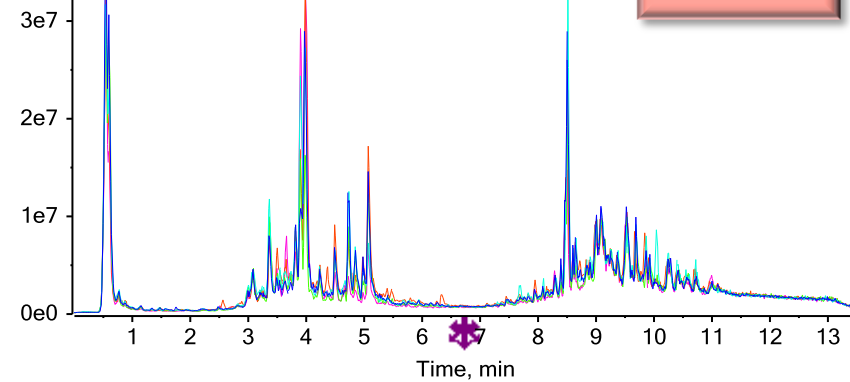
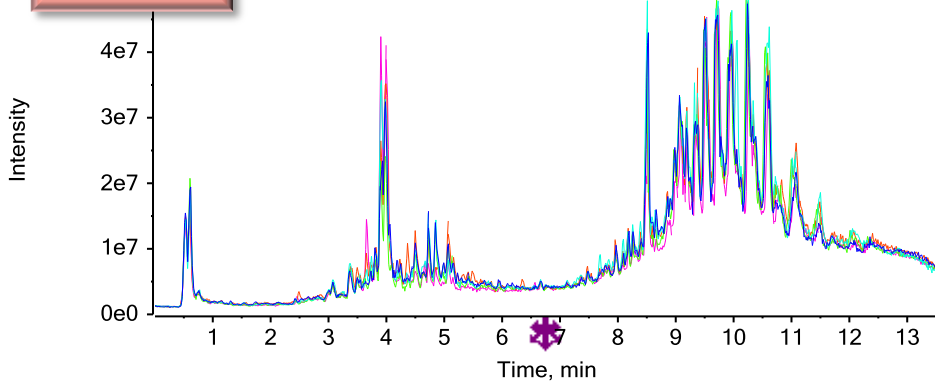
U-HPLC-HRMS/MS metabolomic fingerprint of polar extracts of fruits

ESI+

Lingonberries

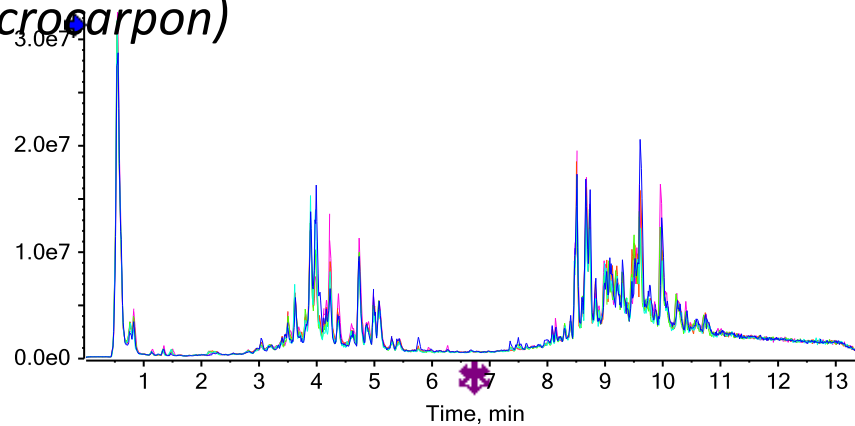
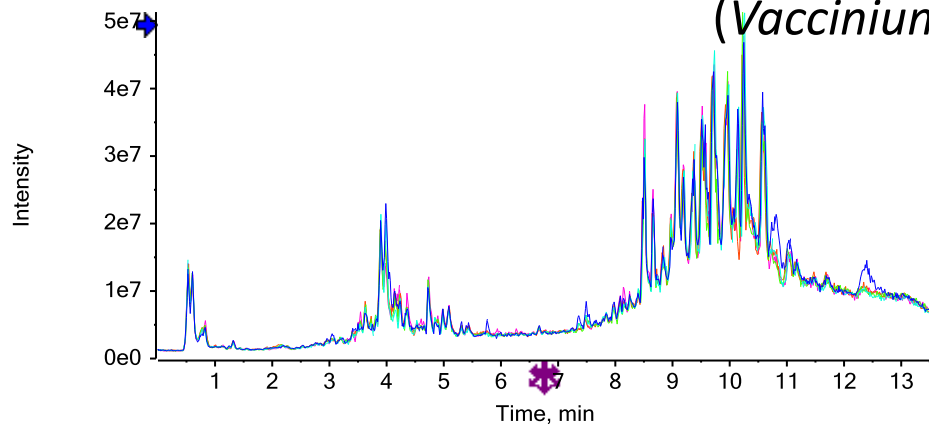
(*Vaccinium vitis-idaea*)

ESI-



Cranberries

(*Vaccinium macrocarpon*)

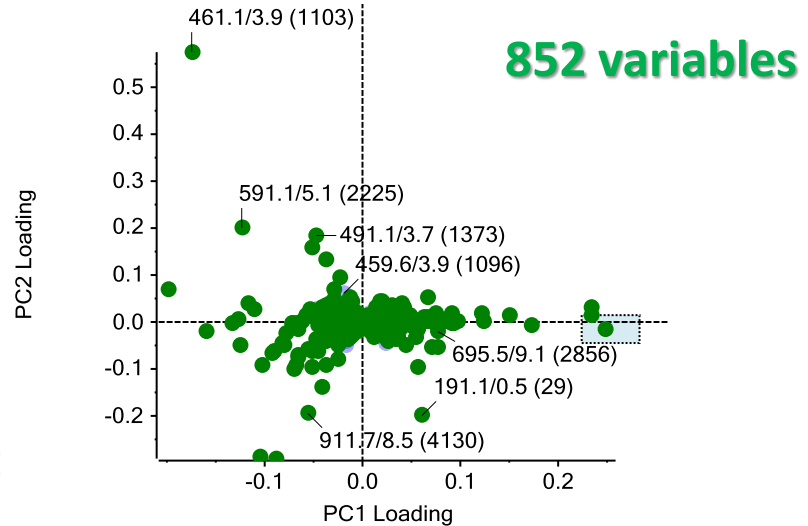
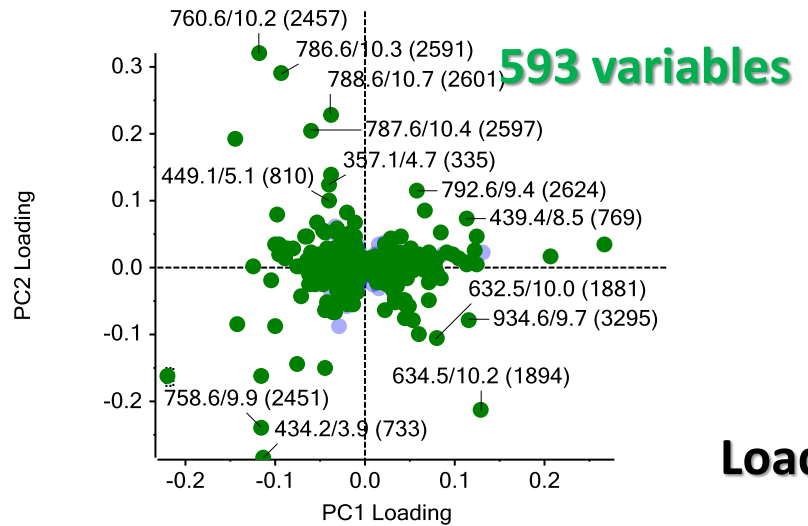
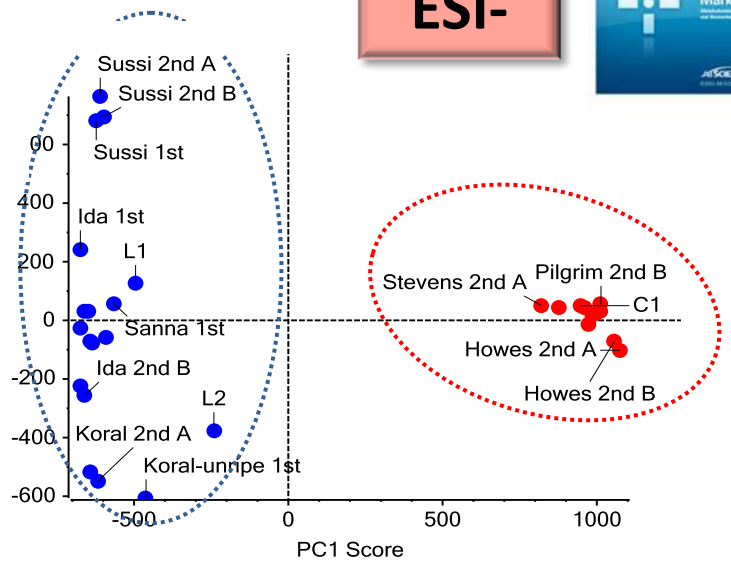
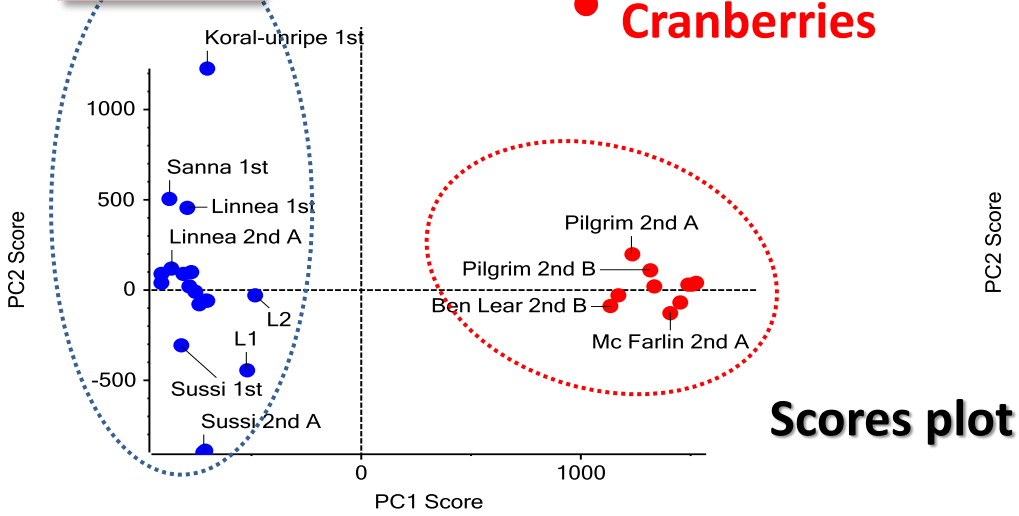


Multivariate PCA analysis of polar extract

ESI+

● **Lingonberries**
● **Cranberries**

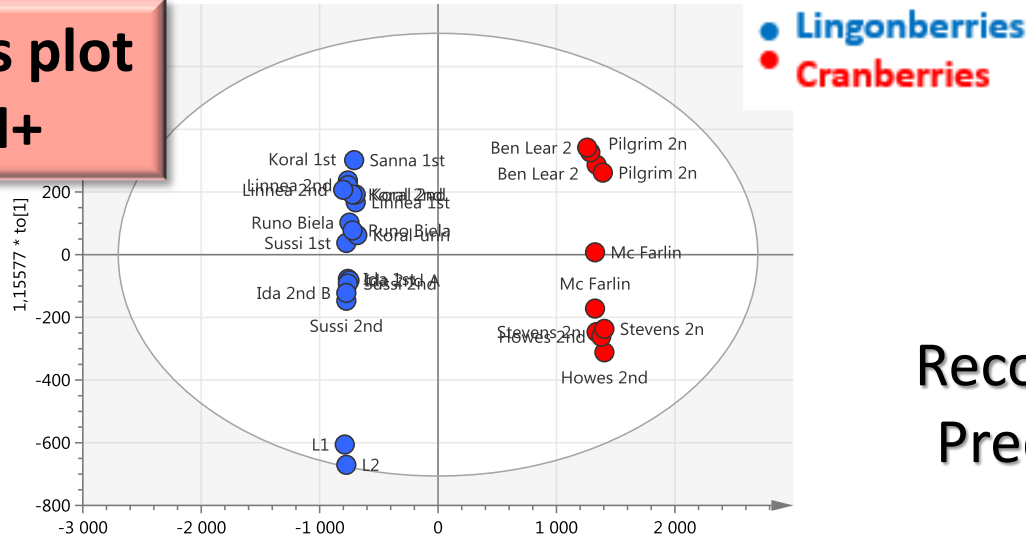
ESI-



Multivariate OPLS-DA analysis of polar extract

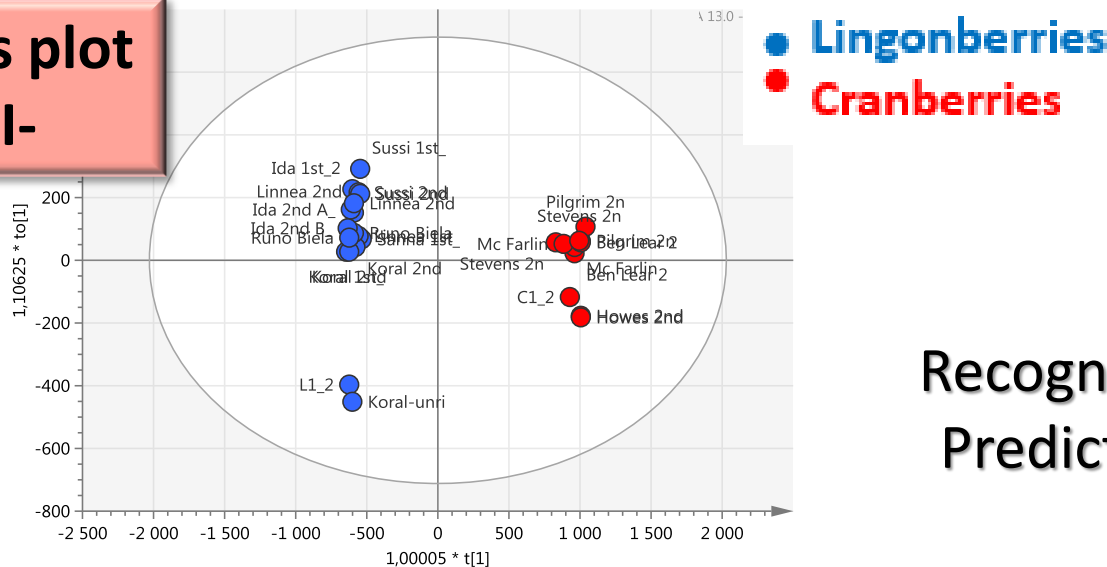


Scores plot
ESI+



Recognition ability: 100 %
Prediction ability: 100 %

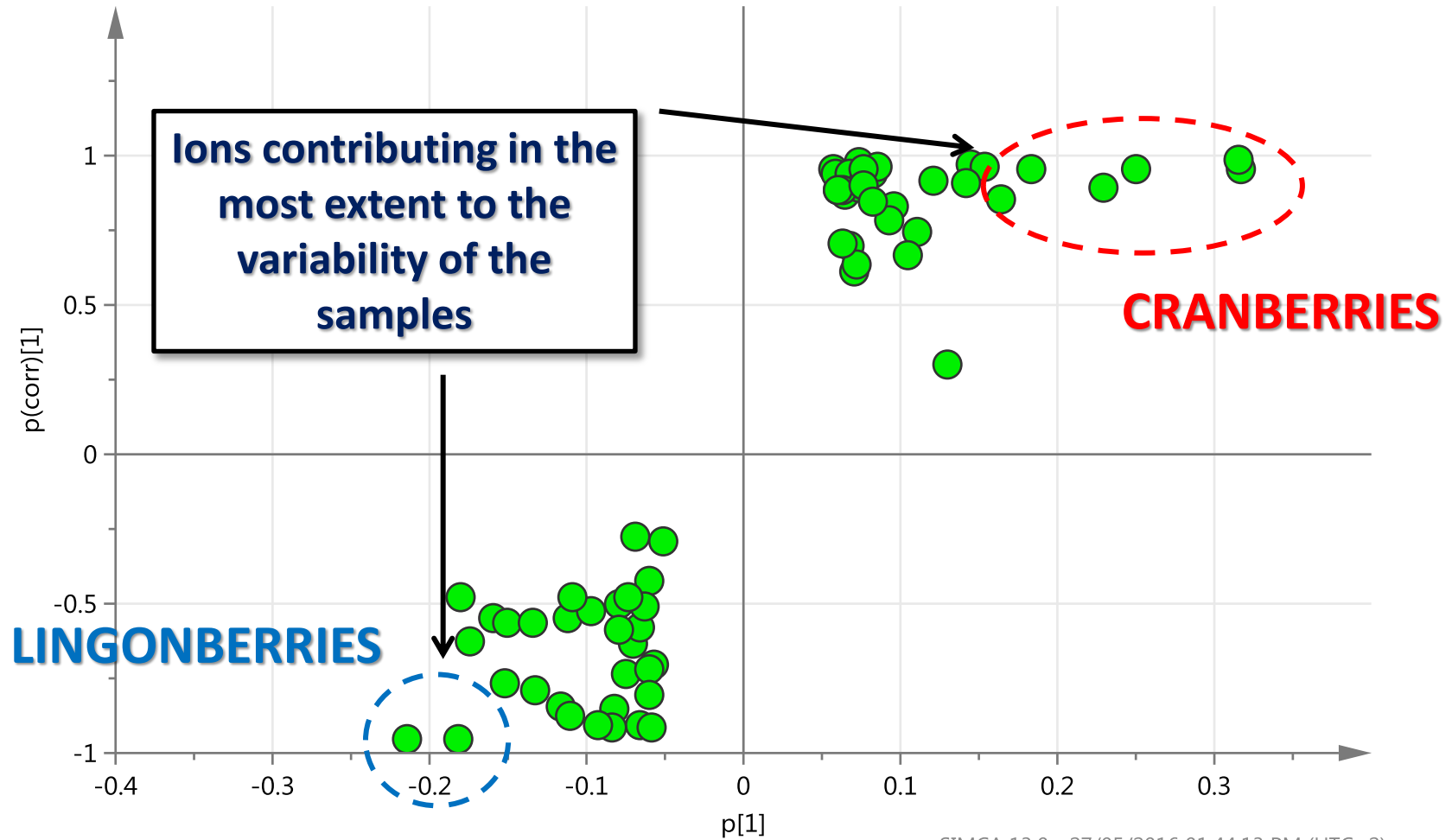
Scores plot
ESI-



Recognition ability: 100 %
Prediction ability: 100 %

Multivariate analysis of polar extract

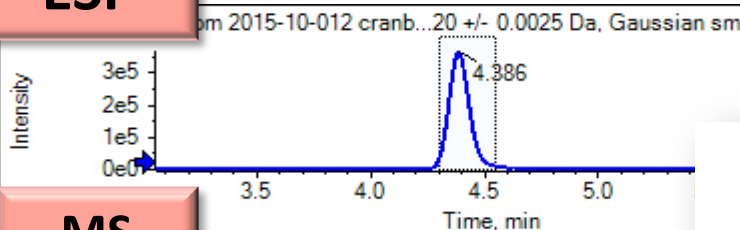
S-plot (ESI-)



Marker identification: cranberries (*Vaccinium macrocarpon*)

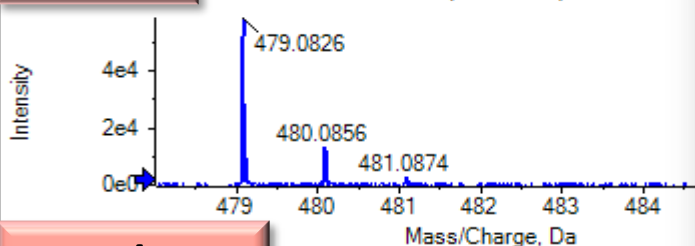


ESI-

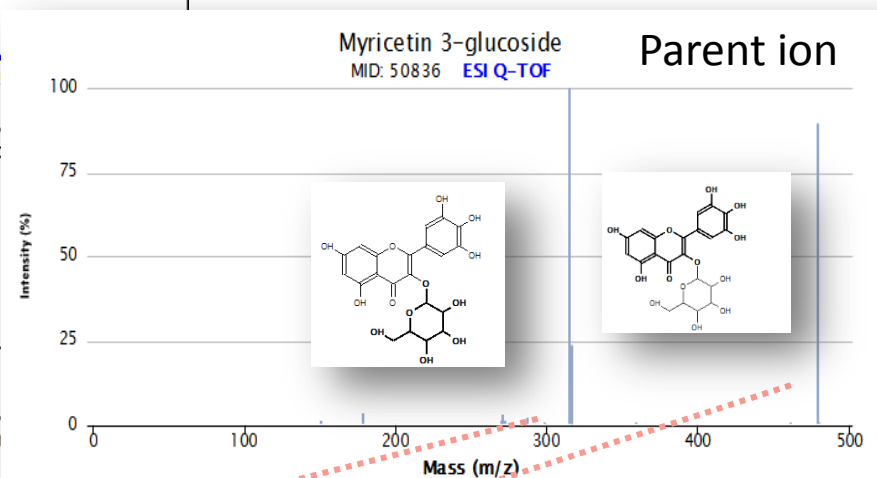
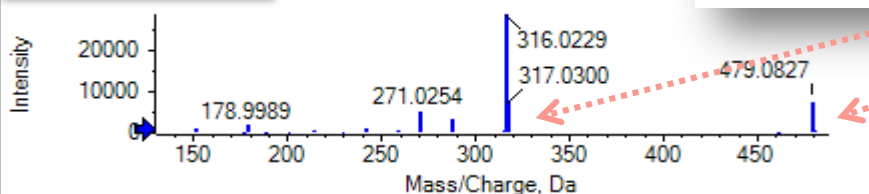


XIC
m/z 479.0857

MS



MS/MS



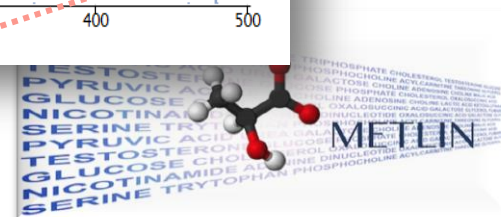
Found elemental compositions

Find Any Find

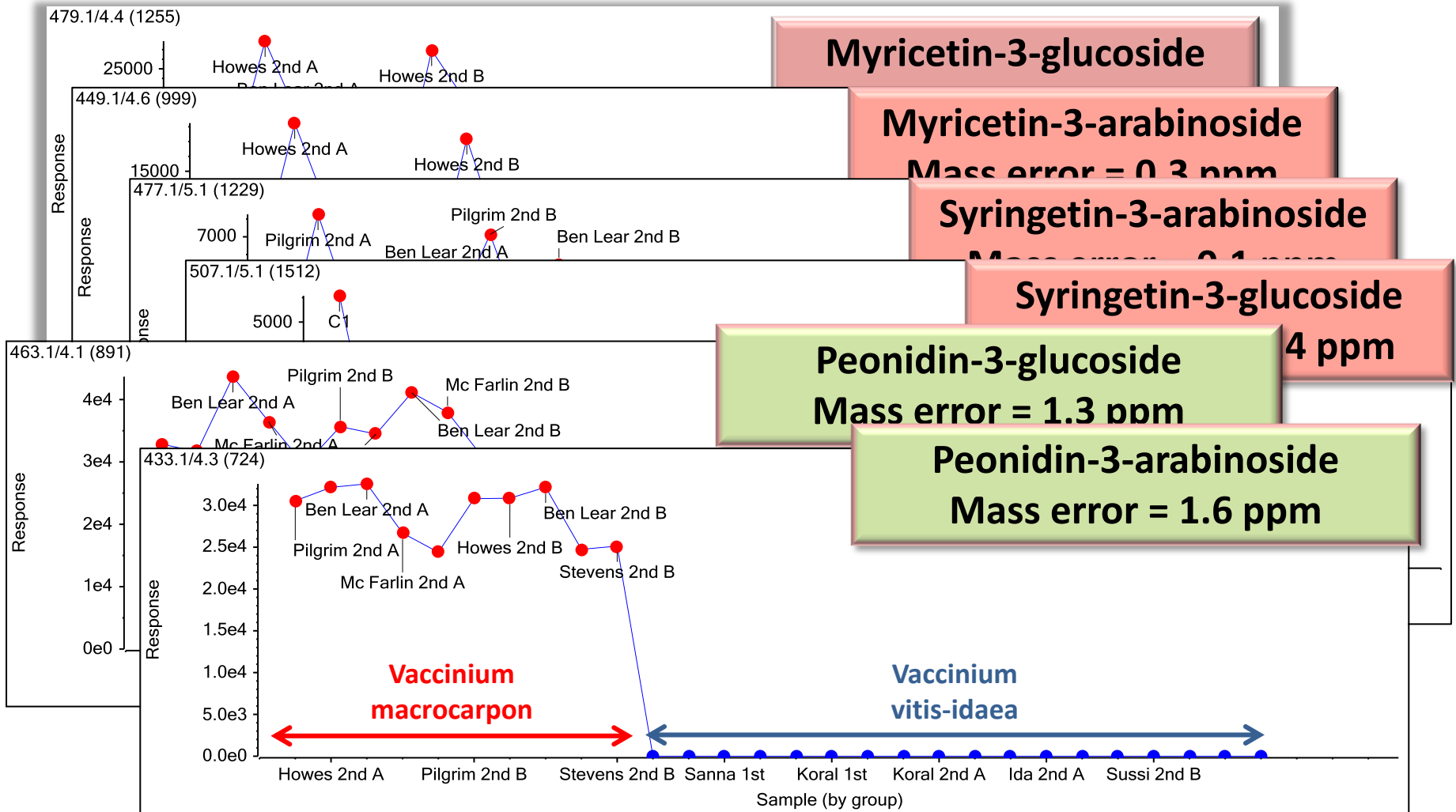
Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
4	C ₂₁ H ₂₀ O ₁₃	479.0831	12.0	-1.1	5 (2)	1.3 (8)	9	NA/N
5	C ₁₇ H ₁₆ N ₆ O ₁₁	479.0804	13.0	4.5	7	2.0 (9)	10	NA/N

Formula finder

C₂₁H₂₀O₁₃
myricetin-3-glucoside
Mass error <3 ppm



Plot profiles of markers of cranberries (*Vaccinium macrocarpon*), ESI- / ESI+



CHARACTERISTIC MARKERS FOR DIFFERENTIATION OF CRANBERRIES AND LINGONBERRIES

Markers for <i>Vaccinium macrocarpon</i> (cranberry)	Summary formula	Ion	m/z
Myricetin-3-arabinoside	$C_{20}H_{18}O_{12}$	[M-H] ⁻	449,0726
Myricetin 3-glucoside	$C_{21}H_{20}O_{13}$	[M-H] ⁻	479,0831
Syringetin-3-glucoside	$C_{23}H_{24}O_{13}$	[M-H] ⁻	507,1144
Peonidin 3-arabinoside	$C_{21}H_{20}O_{10}$	[M+H] ⁺	428,0726
Peonidin 3-glucoside	$C_{22}H_{22}O_{11}$	[M+H] ⁺	456,0831

Markers for <i>Vaccinium vitis-idaea</i> (lingonberry)	Summary formula	Ion	m/z
Catechin	$C_{15}H_{14}O_6$	[M-H] ⁻	290,0790
Procyanidin B	$C_{30}H_{26}O_{12}$	[M-H] ⁻	577,1351
1-O-feruloylglucose	$C_{16}H_{20}O_9$	[M-H] ⁻	355,1034

MARKERS SHOULD BE PRESENT IN THE TYPICAL RATIOS

CHARACTERISTIC MARKERS FOR DIFFERENTIATION OF CRANBERRIES AND LINGONBERRIES

Markers for <i>Vaccinium macrocarpon</i> (cranberry)	Summary formula	Ion	m/z
Myricetin-3-arabinoside	$C_{21}H_{20}O_{10}$	$[M+H]^+$	433,1129
Myricetin 3-glucoside	$C_{22}H_{22}O_{11}$	$[M+H]^+$	463,1235
Syringetin-3-glucoside	$C_{23}H_{24}O_{11}$	$[M+H]^+$	477,1253
Peonidin 3-arabinoside	$C_{21}H_{20}O_{10}$	$[M+H]^+$	433,1129
Peonidin 3-glucoside	$C_{22}H_{22}O_{11}$	$[M+H]^+$	463,1235

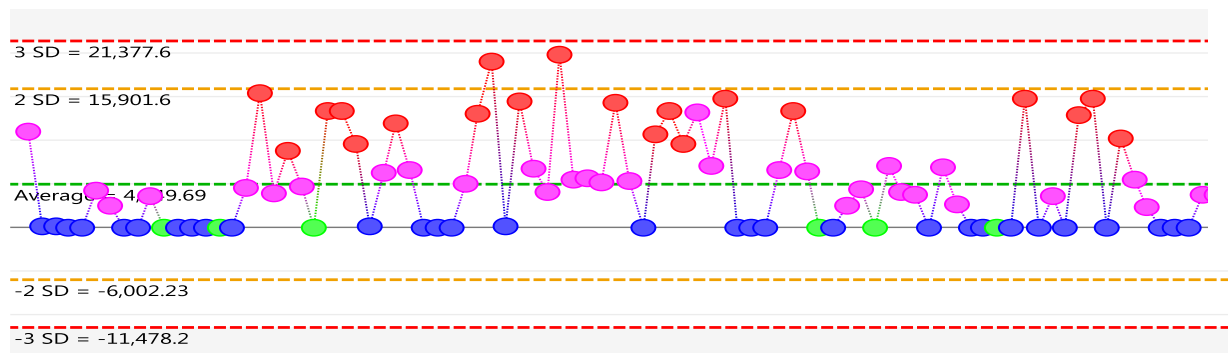
SOME MARKERS CAN BE PRESENT IN OTHER BERRIES

Markers for <i>Vaccinium vitis-idaea</i> (lingonberry)	Summary formula	Ion	m/z
Catechin	$C_{15}H_{14}O_6$	$[M-H]^-$	290,0790
Procyanidin B	$C_{30}H_{26}O_{12}$	$[M-H]^-$	577,1351
1-O-feruloylglucose	$C_{16}H_{20}O_9$	$[M-H]^-$	355,1034

FRESH VS. DRIED FRUITS, markers stability during the food processing



FRESH VS. DRIED FRUITS, markers stability



Lingonberries fresh

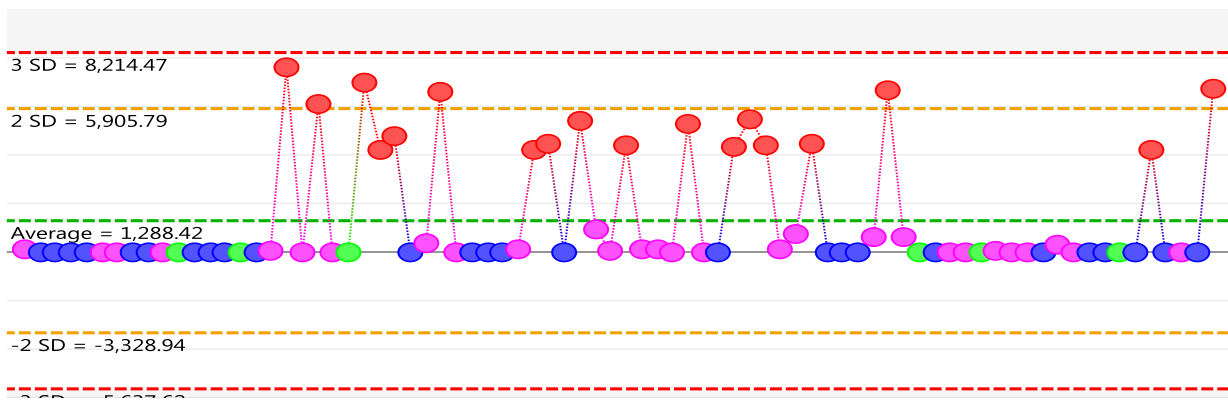
Lingonberries dried

Cranberries fresh

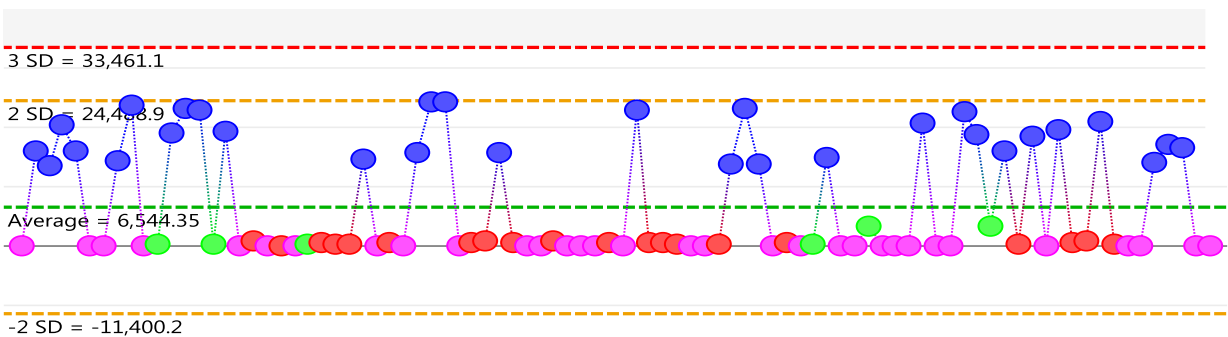
Cranberries dried



m/z 479.0836
Myricetin-3-glucoside



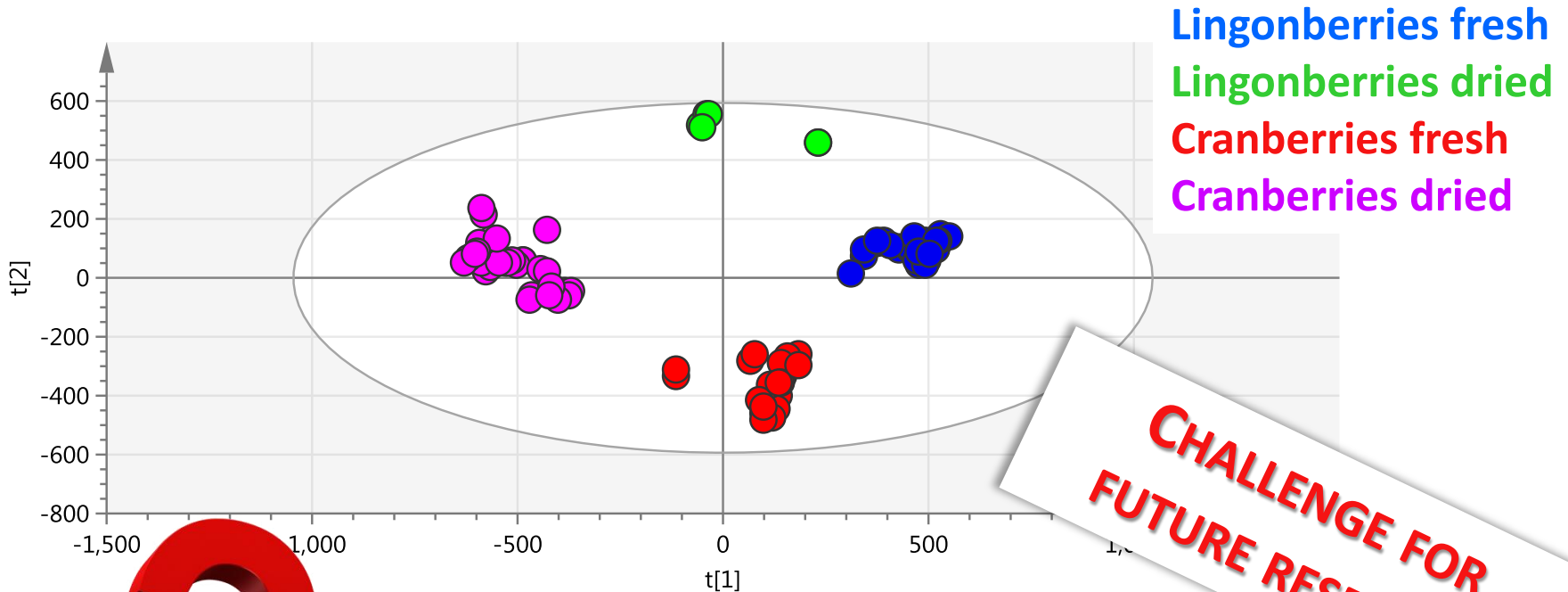
m/z 463.1240
Peonidin-3-glucoside



m/z 289.0730
Katechin

FRESH VS. DRIED FRUITS

Multivariate PCA analysis of polar extract



**CHALLENGE FOR
FUTURE RESEARCH**

- Different markers for dried fruits?
- Reaction (thermal degradation) products of original markers?



METABOLOMIC PROFILING (MARKERS ANALYSIS) IN PROCESSED FOODS



Metabolomic fingerprinting rather unsuitable for multicomponent foods

- Cranberries / lingonberries are often the minor component of foods
- High variability in food composition - difficult to create multivariate models from fingerprints obtained

RESULTS: CONFORMITY WITH DECLARATION, REVEALING OF FRAUD (fruit juices)

Lingonberries 30% ✓

Lingonberries 10% ✗

Cranberries 4% ✓

Lingonberries 12% ?

Cranberries 4% ?

Lingonberries 100% ?

Lingonberries ✓

Lingonberries ✓

Lingonberries 8% ✗

Cranberries 17% ✗

Cranberries 10% ?

Cranberries 20% ?

lingonberries ?

Markers in atypical range as a consequence of processing techniques

CHALLENGE FOR FUTURE RESEARCH

Case study # 2



AUTHENTICATION OF COLD PRESSED OILS OF VARIOUS BOTANICAL ORIGIN



Cold pressed (virgin) oils


- Gentle pressing at low temperatures
- Advantages compared to refined oils
 - Delicious taste and smell
 - Natural color
 - Presence of valuable biologically active components
- Composition:
 - Lipids - mainly TAGs, free FA, minority of DAGs, MAGs
 - Lipid accompanying compounds: alcohols, terpenoids, vitamins



Cold pressed (virgin) oils

- Protective effects against lifestyle diseases (e.g. Cardiovascular)
- Results of epidemiological studies - healthy Mediterranean diet with a high use of olive oil

Products and services



Natural Remedies and Conventional Medicine

MAYO CLINIC
Book of Alternative Medicine

Learn More ►►

Mediterranean diet: A heart-healthy eating plan

The heart-healthy Mediterranean diet is a healthy eating plan based on typical foods and recipes of Mediterranean-style cooking. Here's how to adopt the Mediterranean diet.

By Mayo Clinic Staff

If you're looking for a heart-healthy eating plan, the Mediterranean diet might be right for you.



- **Expensive commodity – subject for frauds!**

AIM OF THE STUDY




- Development of extraction procedure



- LC-HRMS analysis, multivariate analysis



- Markers characterization



- Assessment of stability of markers during oils oxidation

FINGERPRINTING



SAMPLES ANALYZED



12 types of cold pressed plant oils:

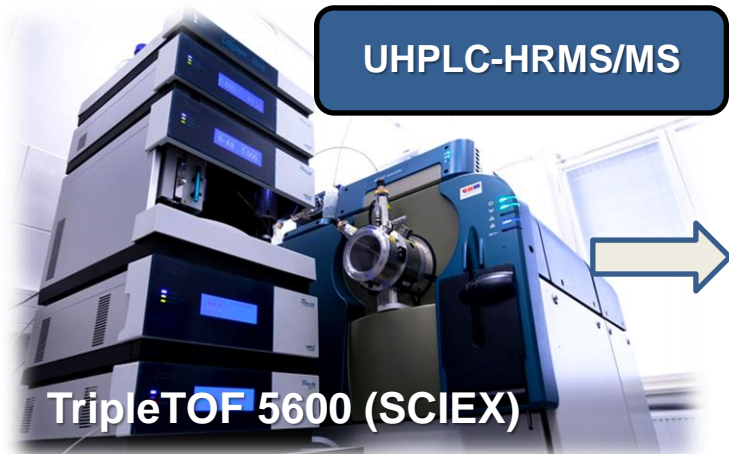
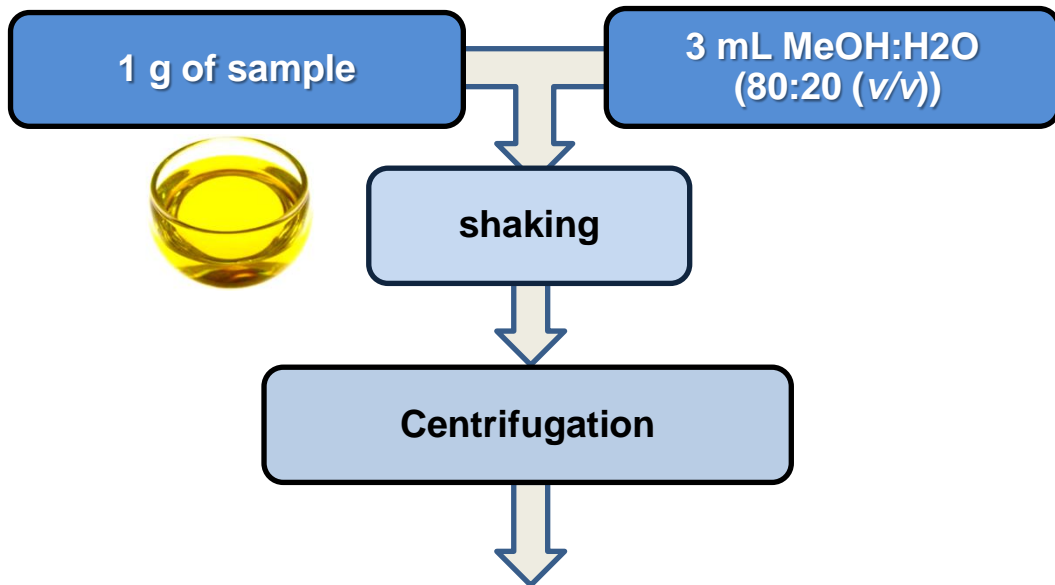
1. Argan (roasted seeds) (ROA)
2. Argan (raw seeds) (RA)
3. Gold flax (GF)
4. Brown flax (BF)
5. White poppy (WP)
6. Blue poppy (BP)
7. Nigella sativa (NS)
8. Red pumpkin (RP)
9. White sesame (WS)
10. Black sesame (BS)
11. Milk thistle (MT)
12. Hemp (HE)



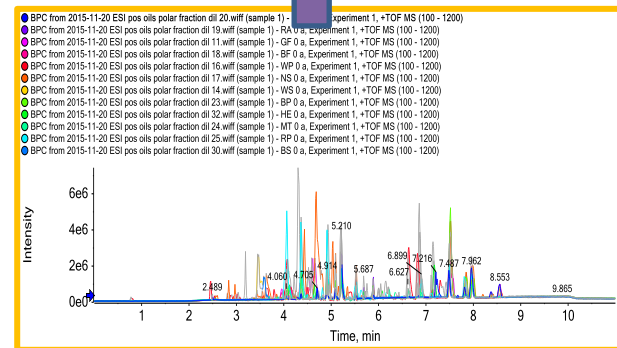
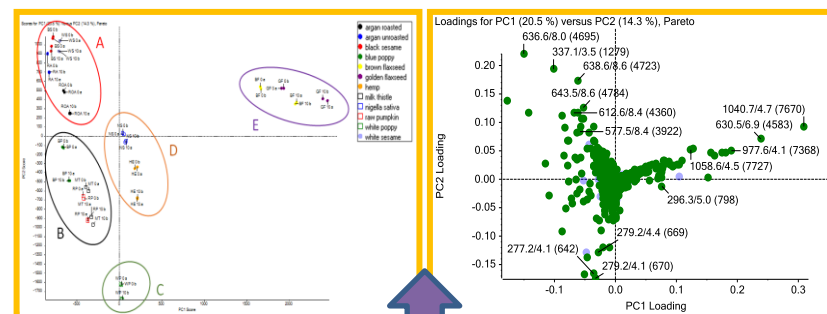
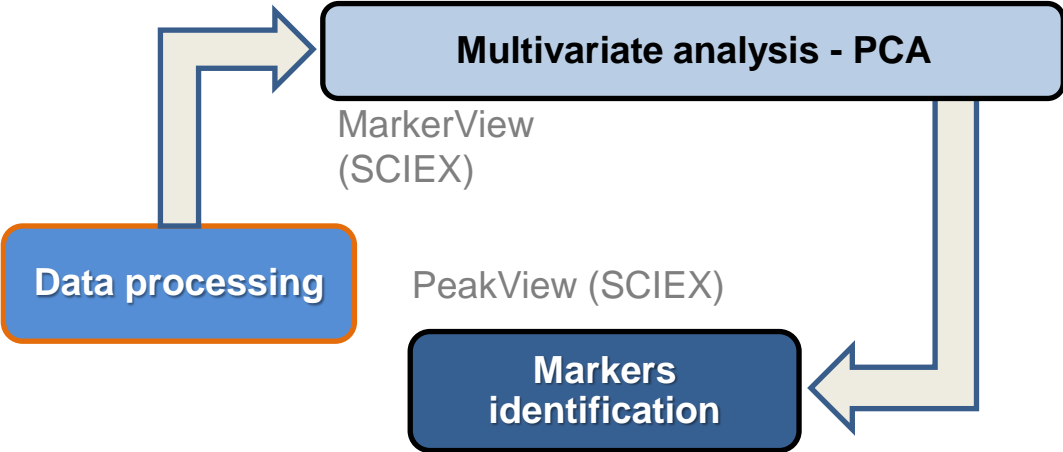
Fresh vs. oxidized (10 days at 60°C)



WORKFLOW OF SAMPLE PREPARATION AND DATA PROCESSING

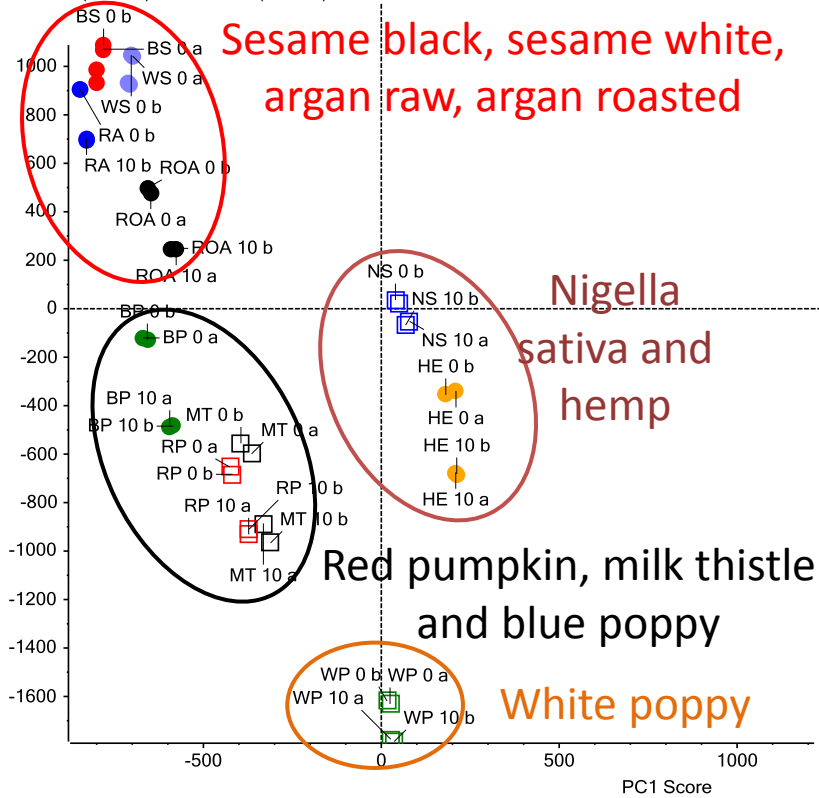


UHPLC-HRMS/MS



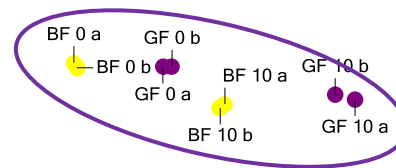
Multivariate PCA analysis (ESI+)

Scores for PC1 (20.5 %) versus PC2 (14.3 %), Pareto

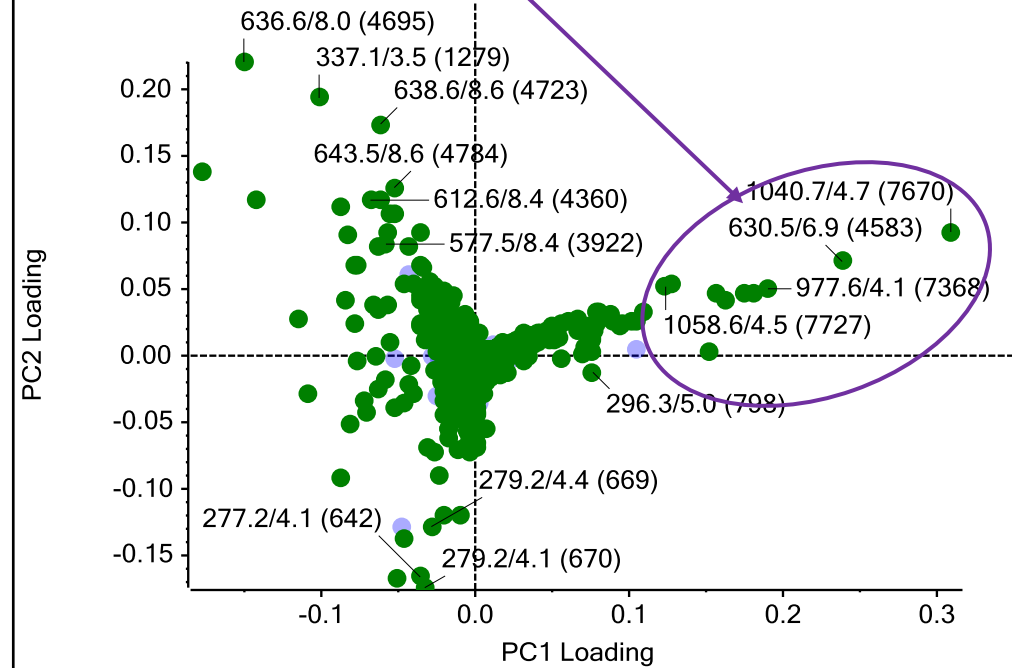


↑ Score plot

Gold flax, brown flax



Loadings for PC1 (20.5 %) versus PC2 (14.3 %), Pareto



Loadings plot →

Markers identification

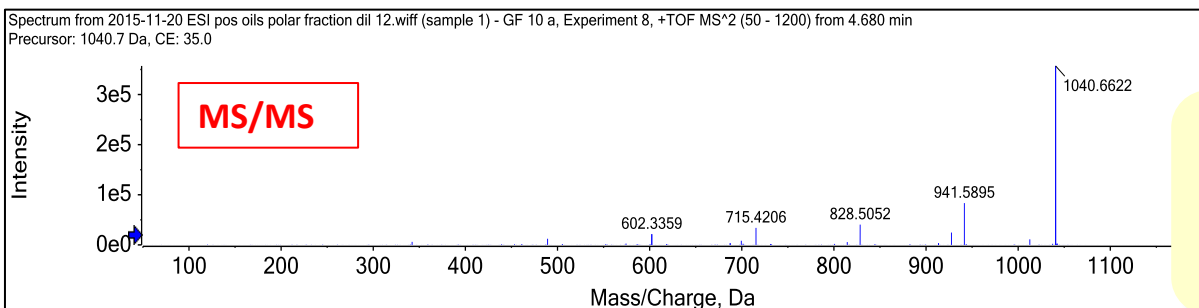
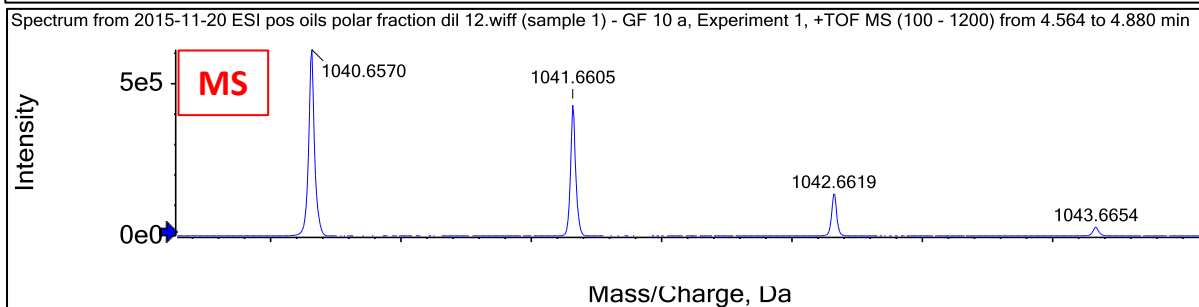
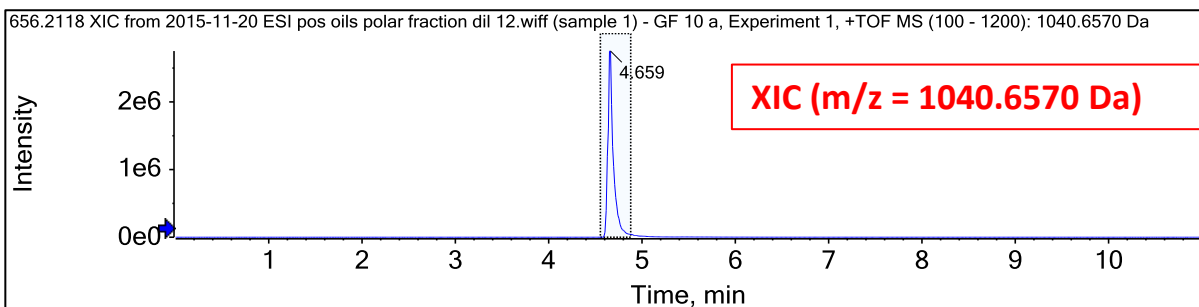
Example for gold flax seed oil



PeakView

Navržené sumární vzorce

Hit	Formula	m/z	RDB	ppm	MS Rank
73	C54H93N3O16	1040.6...	10.0	-5.6	73
74	C49H101O20P	1040.6...	0.5	-4.6	74
75	C57H85N9O9	1040.6...	20.0	2.6	75
76	C47H99N3O17P2	1040.6...	1.0	4.6	76 ...
77	C50H99N5O11...	1040.6...	5.0	-3.8	76 ...
78	C49H100N6O9P4	1040.6...	5.5	7.0	78
79	C49H88N10O14	1040.6...	11.5	9.0	79



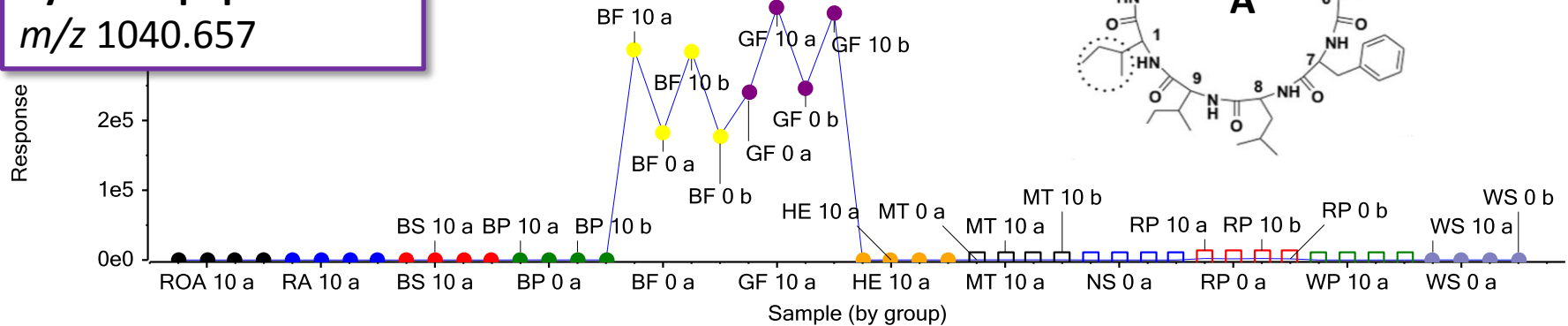
ChemSpider
Metlin

Formula finder
 $C_{47}H_{85}N_9O_9$
Cyclolinopeptide A
Mass error = 2.6 ppm

Plot profiles of markers of gold flax seed oil

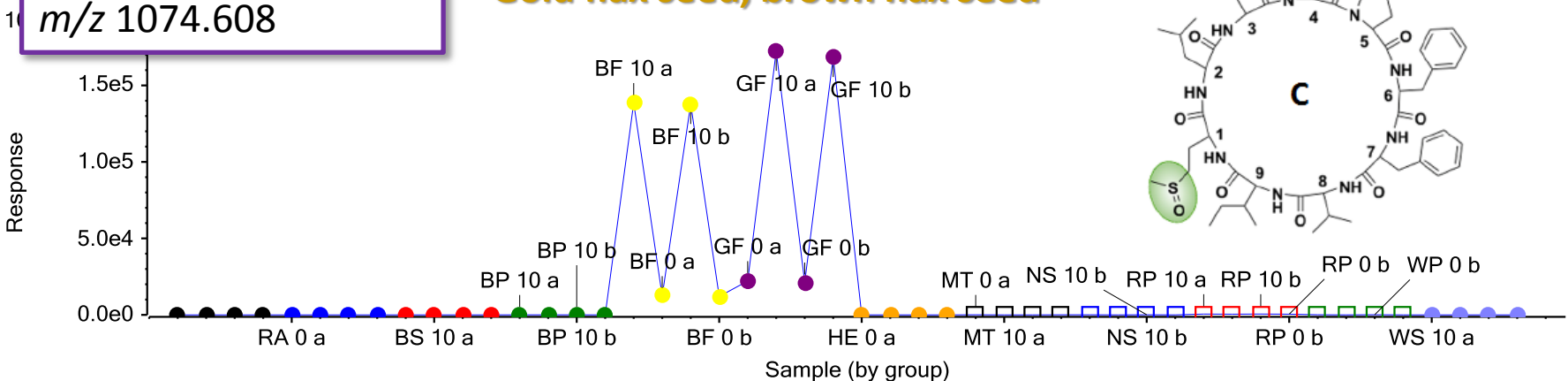
Cyclinopeptide A
 m/z 1040.657

Gold flax seed, brown flax seed

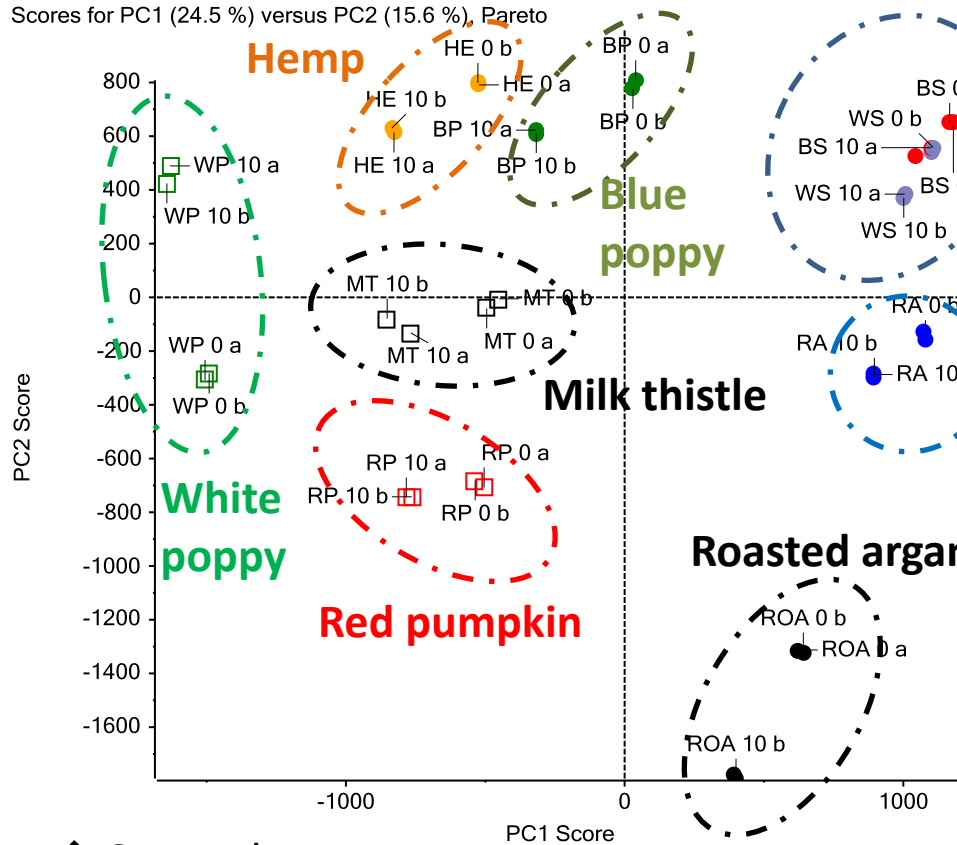


Cyclinopeptide C
 m/z 1074.608

Gold flax seed, brown flax seed



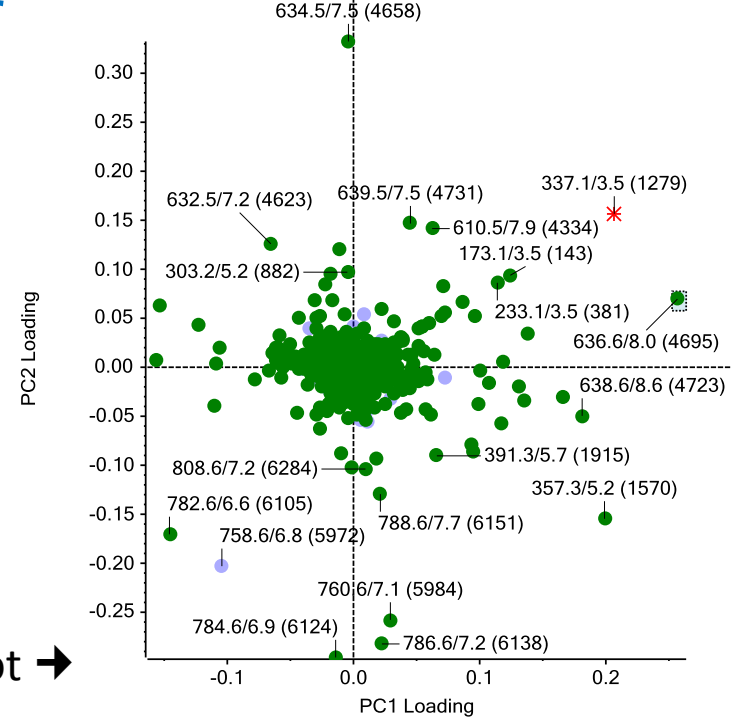
Multivariate PCA analysis, without flax seed oils (ESI+)



Black sesame, white sesame

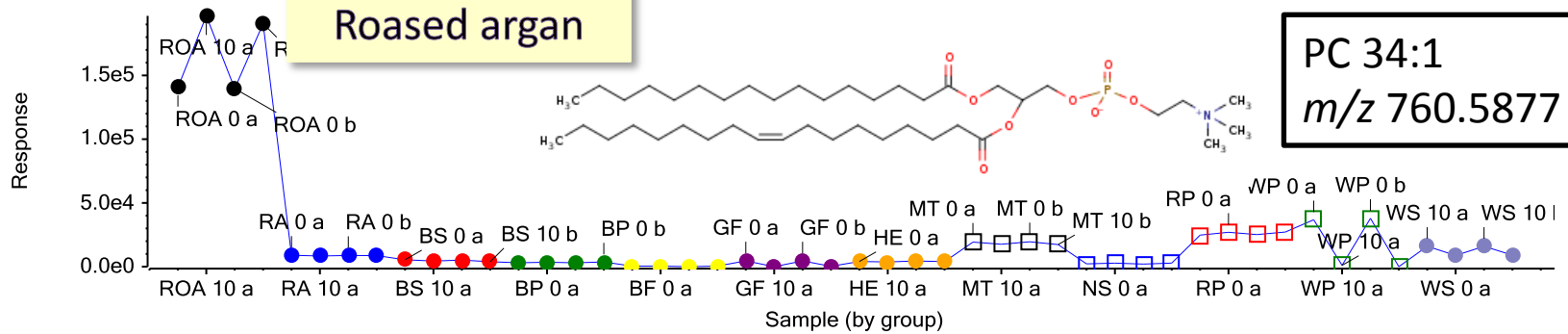
Row argan

Loadings for PC1 (24.5 %) versus PC2 (15.6 %), Pareto

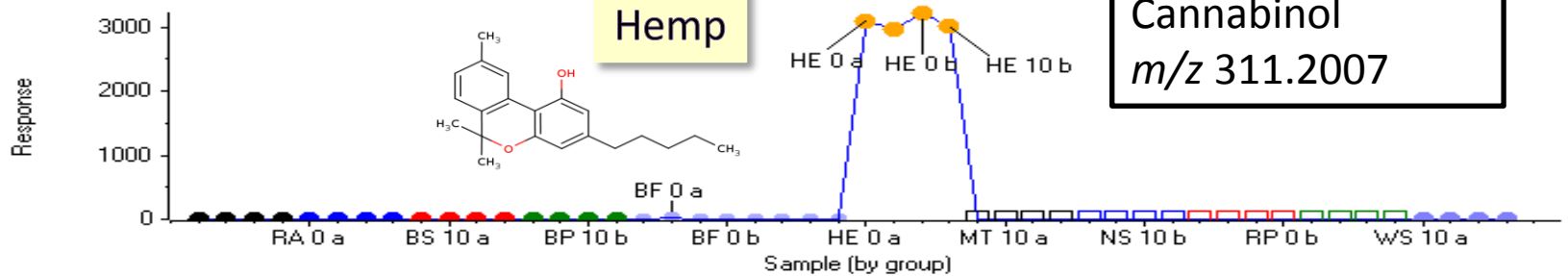


Plot profiles of markers of other cold pressed oils

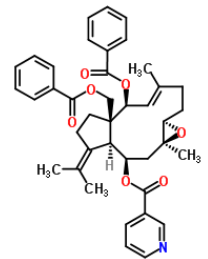
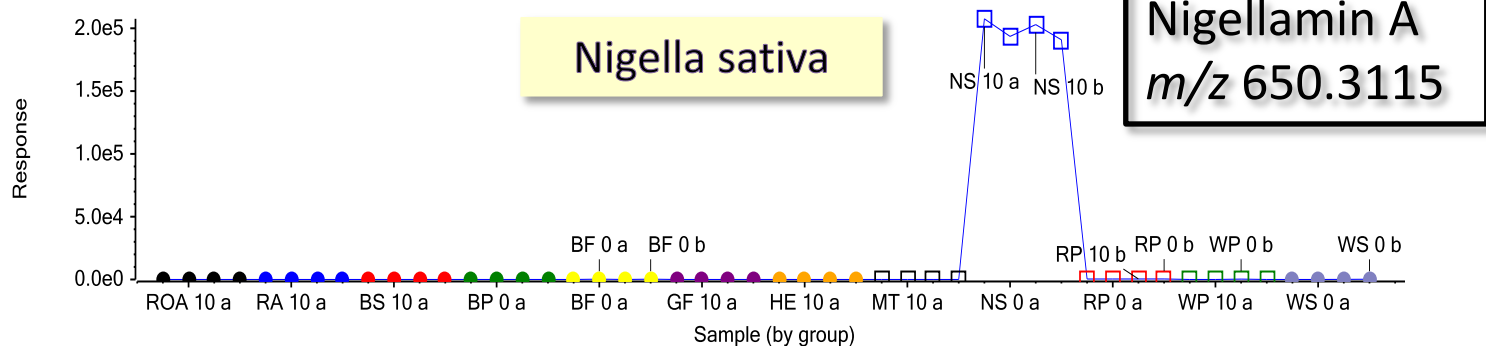
760.6/7.1 (5984)



311.2/4.6 (936)



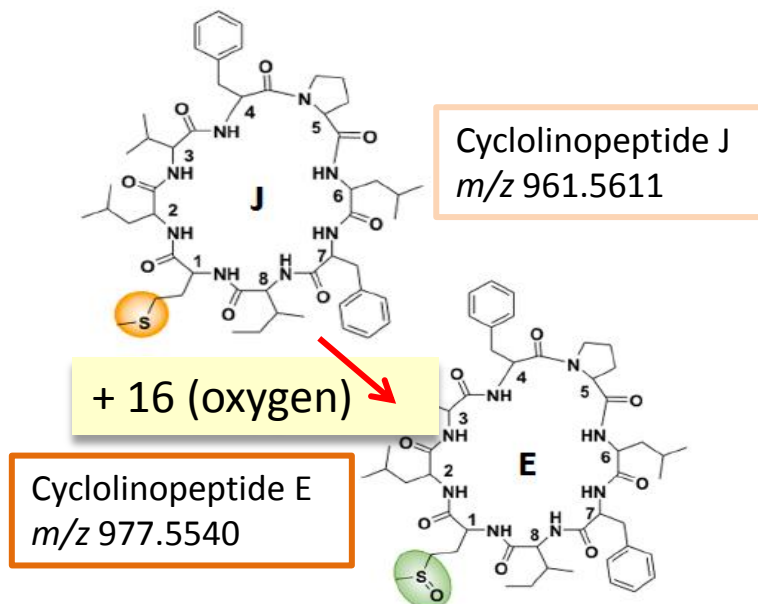
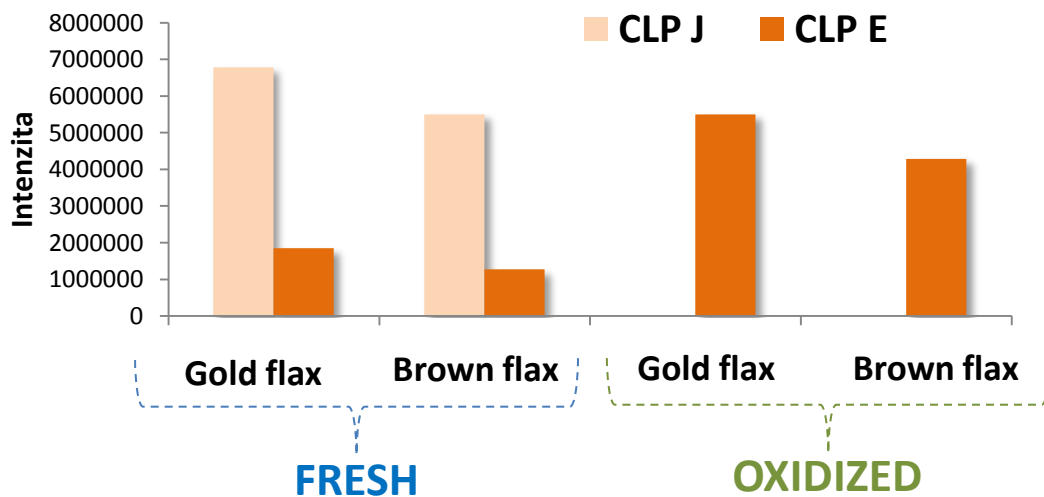
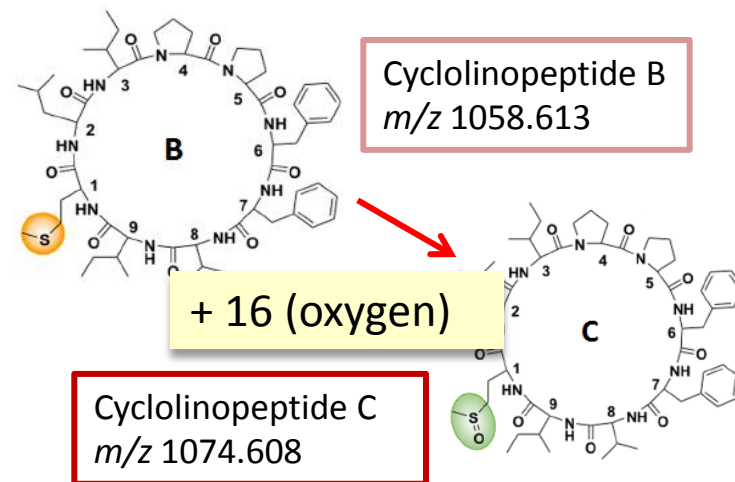
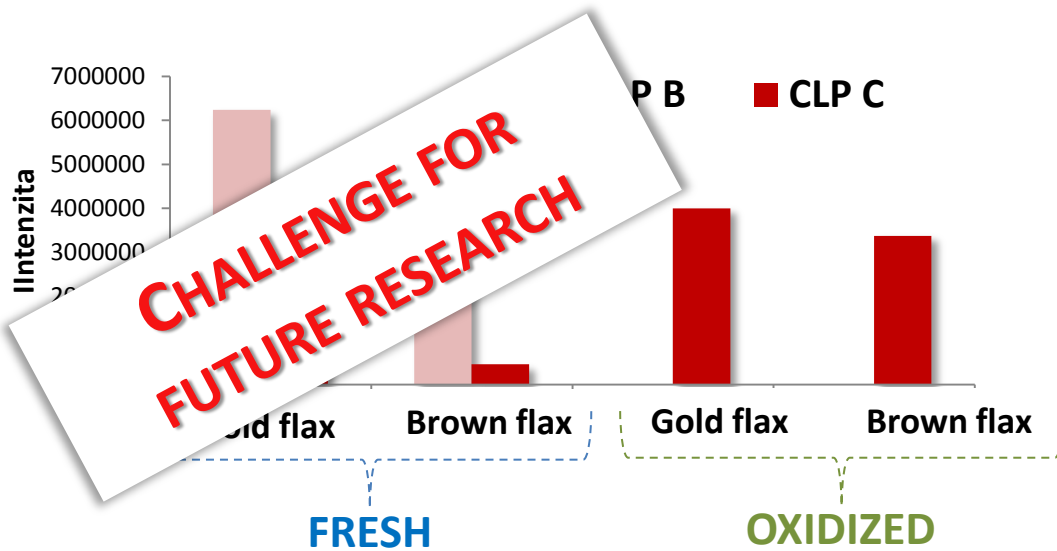
650.3/4.9 (4872)



FRESH VS. OXIDIZED LIPIDS, stability of the markers during the flax seed oil oxidation



FRESH VS. OXIDIZED LIPIDS, stability of the markers

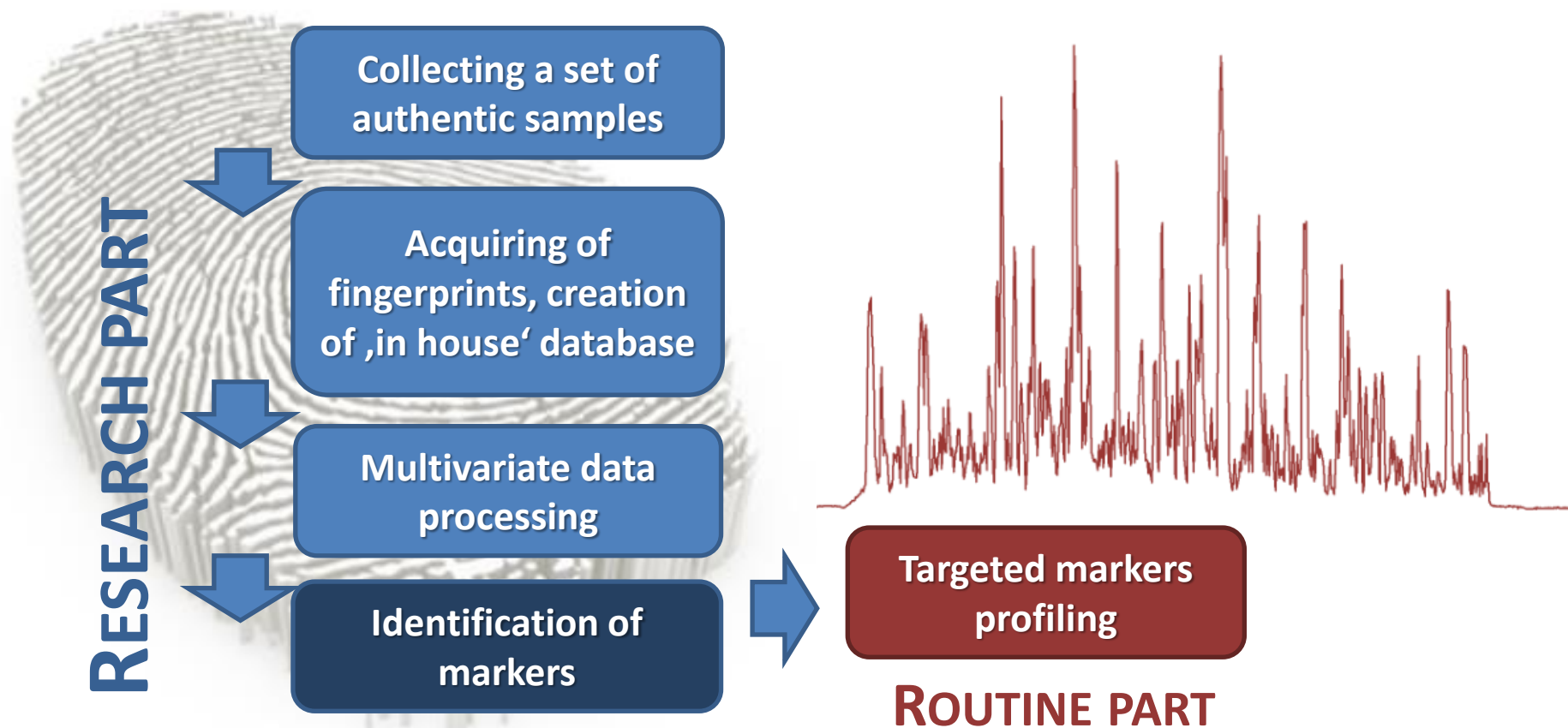


APPROACHING TO THE END...



CONCLUDING REMARKS

- Broad application potential of LC-HRMS non-target metabolomic fingerprinting of small molecules



CONCLUDING REMARKS

■ MAIN ADVANTAGES OF LC-HRMS:

- **Low detection limits achievable** compared to other instrumental methods (IR or NMR), hence possibility to identify unique markers occurring at trace level
- **Structural identification of markers is feasible** in most cases



CONCLUDING REMARKS

■ MAIN BOTTLENECKS AND CHALLENGES – BUILDING OF HARMONIZED LC-HRMS DATABASES:

■ DATA HANDLING

- ‚In house‘ databases - inter-temporal data transfer
- ‚Joint‘ databases - interlaboratory data harmonization

■ METHODS TRANSFER

- Harmonized sample preparation protocols assuring uniformity and representativeness of metabolomic fingerprints are necessary





**UNIVERSITY OF
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PRAGUE**



Challenges in Mass Spectrometry Based Non-Targeted Analysis

Thank you for your attention...

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CONCLUDING REMARKS

■ MAIN CHALLENGES FOR OVERCOMING THE HURDLES:



To assure harmonization of input data as much as possible

- Define requirements for MS instrumentation used (e.g. minimal mass resolving power)
- Assure high quality data alignment
- Define minimal threshold for intensity of input features
- Develop the algorithms for compensating the differences in response of different instruments (by using of suitable matrix reference material)

