

High resolution mass spectrometry based metabolomics - an effective tool for fruit products authentication

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PRAGUE**



Red and blue berries

■ Source of bioactive substances

- Flavonoids (anthocyanins, flavonols, flavanols)
- Tannins (proanthocyanidins)
- Phenolic acids

Health promoting properties

- ▶ Antioxidant activity
- ▶ Anti-inflammatory activity
- ▶ Prevention of weight gain
- ▶ Preventive and therapeutic effects against UTI



**Popular high value
fruit products**

Lingonberries vs. Cranberries

Vaccinium vitis-idaea

- “ Rock or mountain cranberry “
- Berries from North and Central Europe, Baltic countries
- Size of berries: 5-10 mm
- **Price of fresh berries: 1.3 €/kg**
- Up to twice the total phenolics and proanthocyanidins content (prevention of urinary tract infection)

Vaccinium macrocarpon

- “ Large or American cranberry “
- Major commercial crop in the United States and Canada
- Size of berries: 9-16 mm
- **Price of fresh berries : 0.4 €/kg**
- More prevalent A-type proanthocyanidins

Sensory and phytochemical similarities may lead to confusion or intentional substitution of the two crops

Adulteration practices: partial or total replacement of lingonberries by cranberries

Adulteration of red and blue berries

- **Substitution** of high value berries (lingonberry, bilberry or black currant) by less expensive ingredients (swamp cranberry, chokeberry)
- **Admixtures** were reported mainly in case of
 - Fruit juices
 - Jams and purees
 - Food supplements
- ➔ **Determination of fruit authenticity**
 - **Quality control**
 - **Consumer protection**
 - **Unfair competition avoidance**

Aim of the study

To investigate the applicability of HRMS based

METABOLOMIC FINGERPRINTING



for berries authentication (according to species)

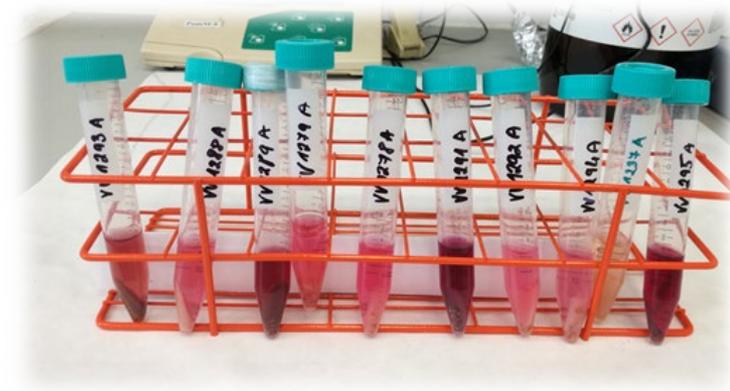


Analyzed samples

11 fruit species

altogether **90** authentic samples
(frozen and dried samples)

- Cranberry
- Lingonberry
- Raspberry
- Red currant
- Strawberry
- Swamp cranberry
- Bilberry
- Black currant
- Blackberry
- Chokeberry
- Elderberry



Optimization of sample preparation

1st STEP Polar extract

1 g of sample extracted with 5 mL of **methanol**

Ultraturrax (1 min)

Centrifugation (5 min, 5 °C, 10,000 rpm)

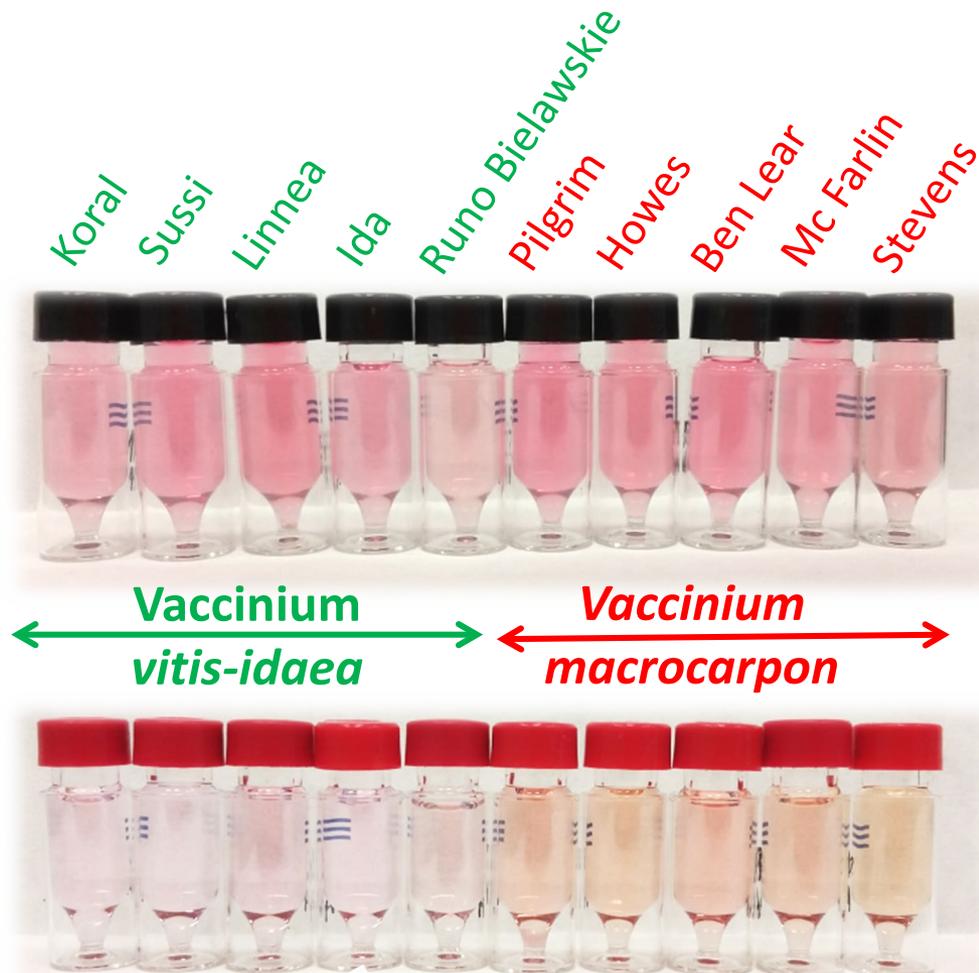
2nd STEP Non-polar extract

repeated extraction of solid residue with 5 mL of **hexane/2-propanol**

(50:50, v/v)

Vortex (1 min)

Centrifugation (5 min, 5 °C, 10,000 rpm)



U-HPLC-HRMS/MS conditions

U-HPLC-Thermo Dionex UltiMate 3000

- **Column:** HSS T3

(2.1x100 mm, 1.8 μm)

Column temperature: 40 °C

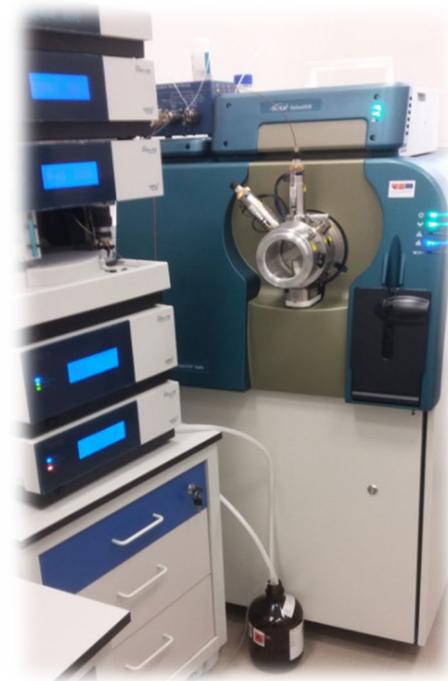
- **Mobile phase:**

A: 5 mM ammonium formate in H₂O

B: 5 mM ammonium formate in MeOH

- **Injection volume:** 4 μL

HSS
HIGH STRENGTH SILICA
HPLC COLUMNS



HRMS(MS/MS)-TripleTOF™ 6600 (Sciex)

- *m/z* range: 100-1200
- Ionization technique: ESI +/-
- Ion source temperature: 480 °C
- Capillary voltage: +5000 V/-4500 V
- Collision energy: 35 eV (+/- 15 eV)

SCIEX

Multivariate analysis of polar extract:

PCA

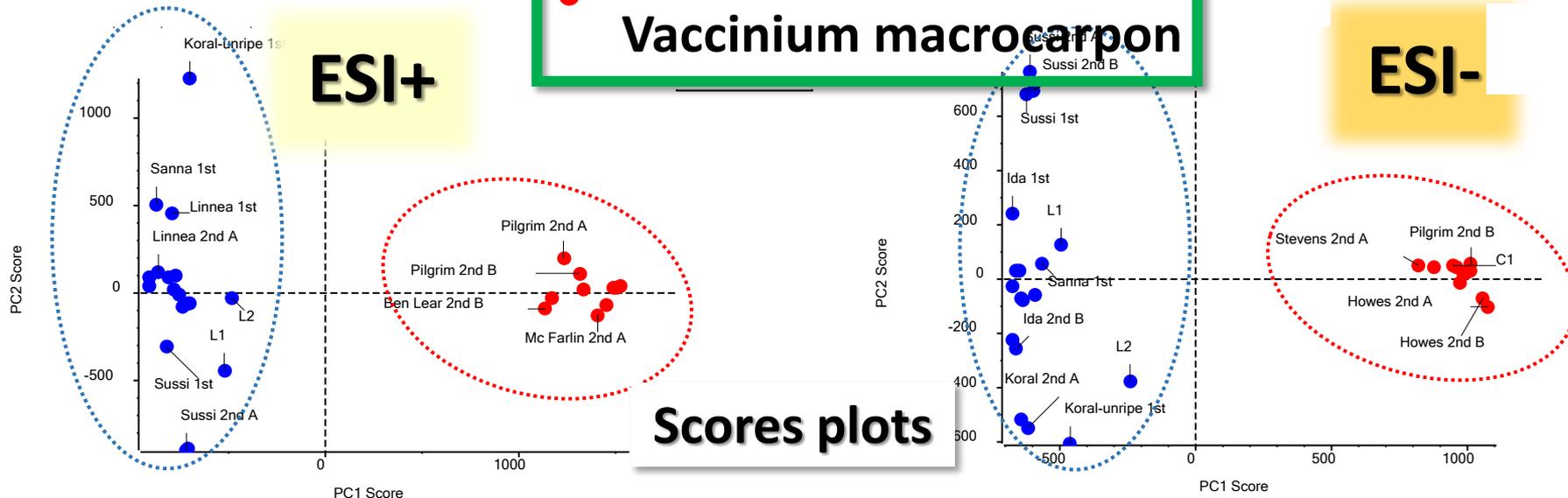
● **Vaccinium vitis-idaea**
● **Vaccinium macrocarpon**

ESI+

ESI-

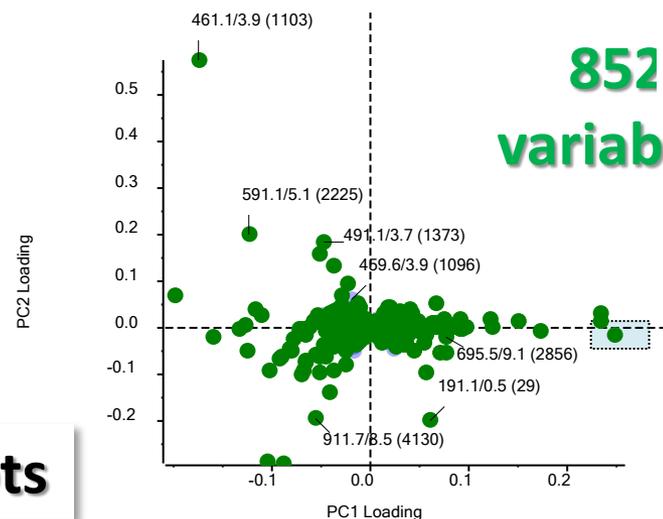
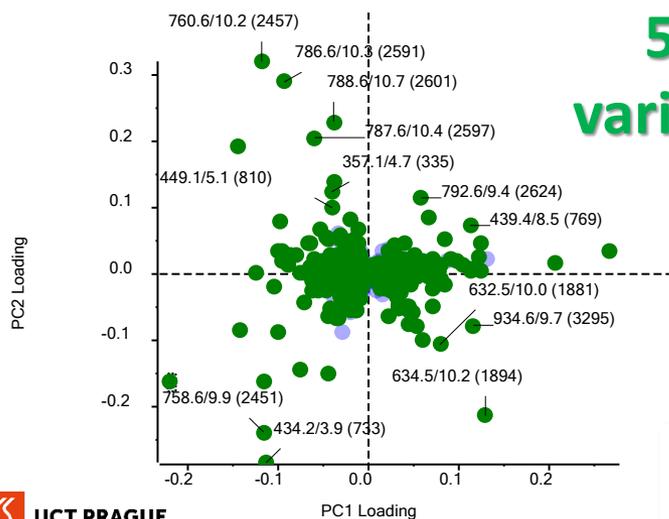
Scores plots

Loadings plots

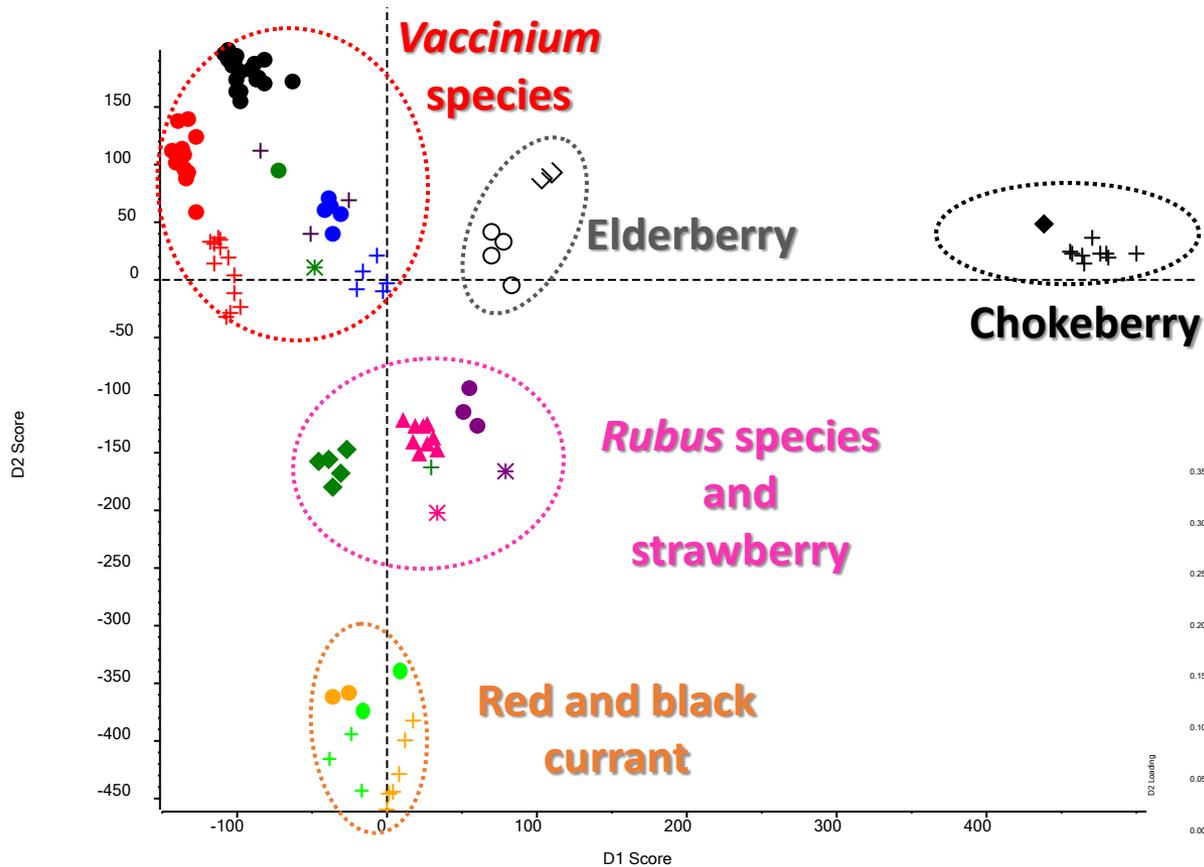


593 variables

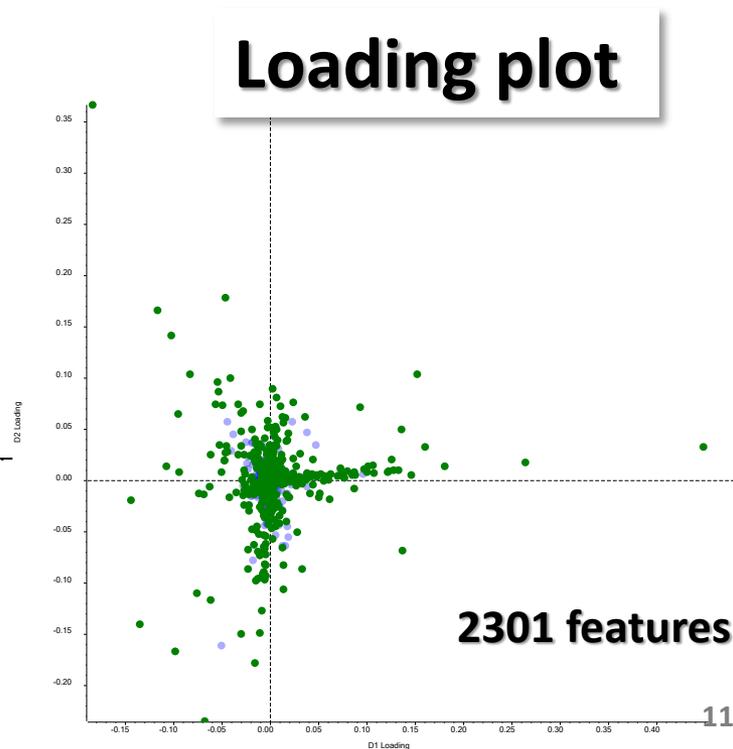
852 variables



Multivariate analysis of polar extract (ESI-) PCA-DA

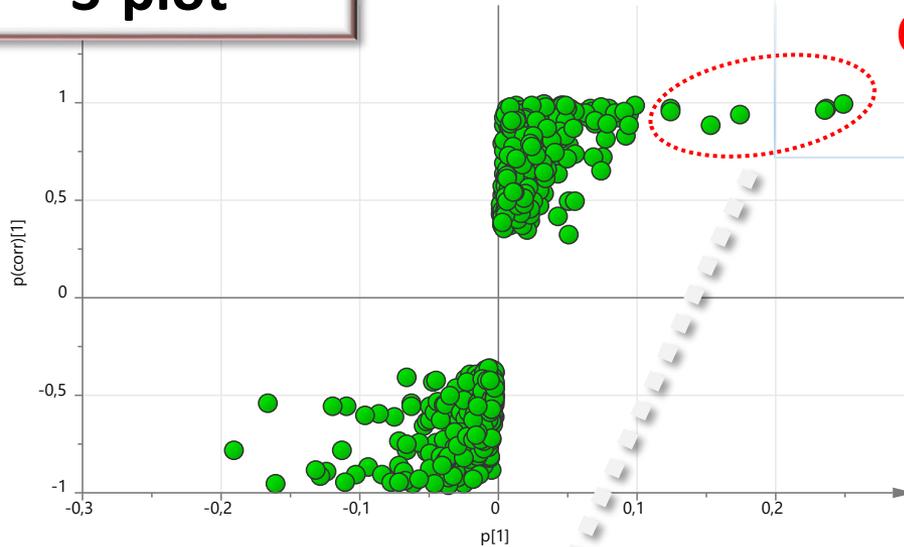


Score plot



Marker identification: *Vaccinium macrocarpon*

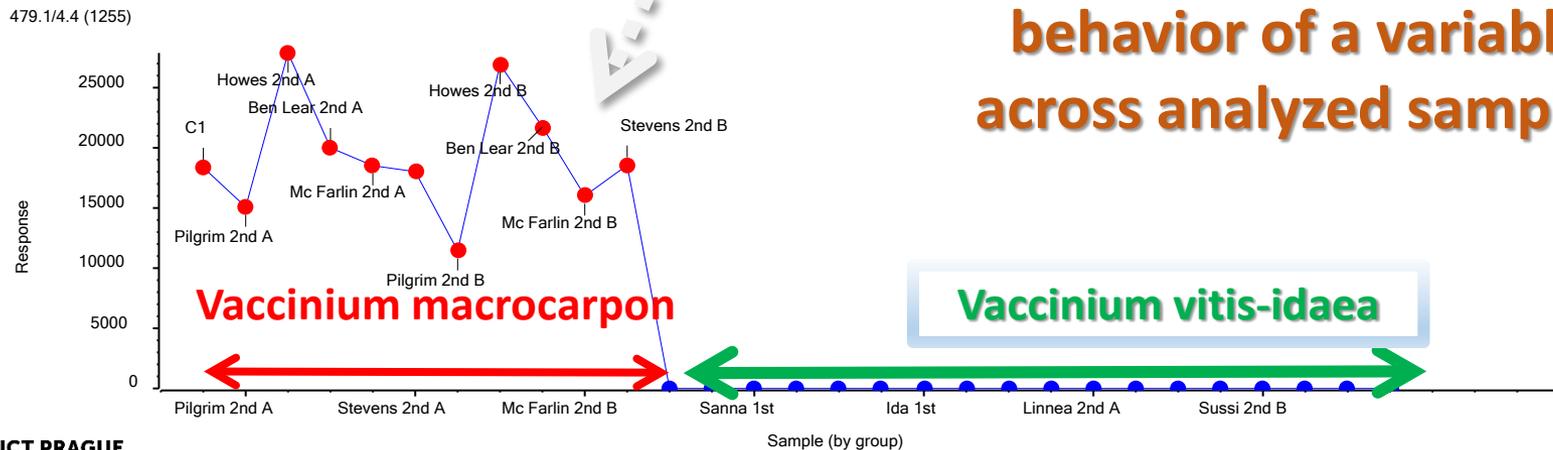
S-plot



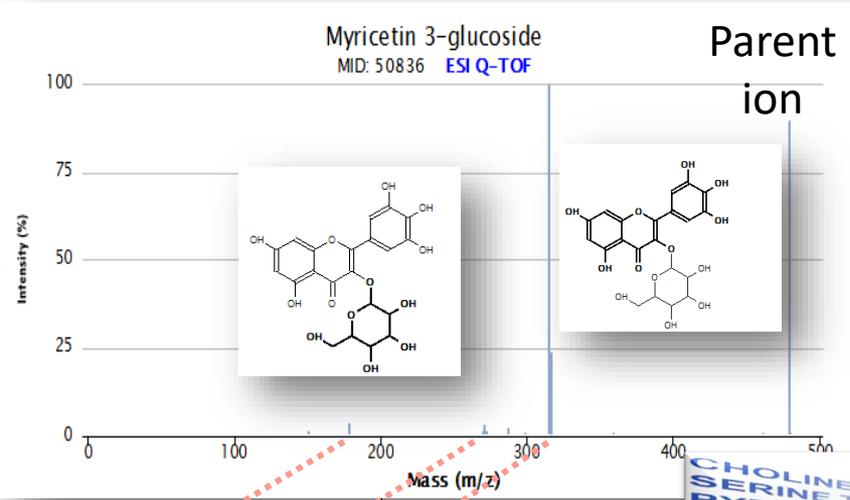
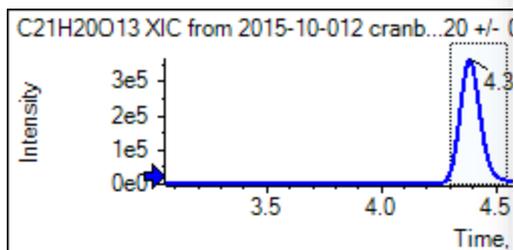
Characteristic markers for
Vaccinium macrocarpon

Plot profile
m/z 479.0857
RT: 4.4 min

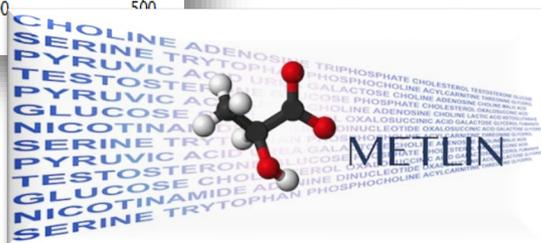
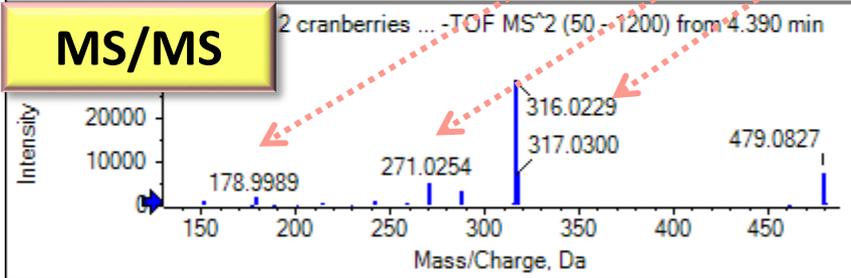
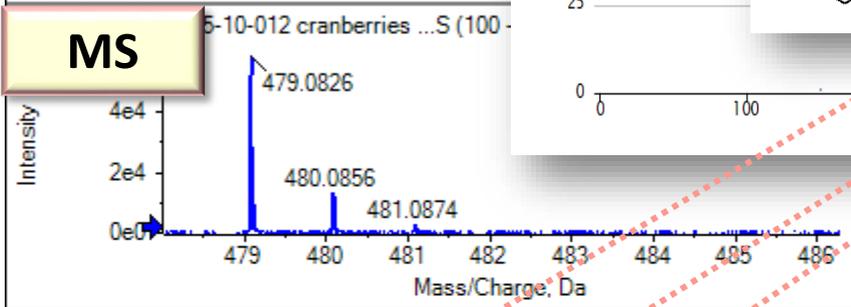
Plot profile expresses
behavior of a variable
across analyzed samples



Marker identification: *Vaccinium macrocarpon*



ESI-



Found elemental compositions

Find Any Find

| Hit | Formula | m/z | RDB | ppm | MS Rank | MSMS ppm | MSMS Rank | Found |
|-----|-------------|----------|------|------|---------|----------|-----------|-------|
| 4 | C21H20O13 | 479.0831 | 12.0 | -1.1 | 5 (2) | 1.3 (8) | 9 | NA/N |
| 5 | C17H16N6O11 | 479.0804 | 13.0 | 4.5 | 7 | 2.0 (9) | 10 | NA/N |

Formula finder
 $C_{21}H_{20}O_{13}$
 myricetin-3-glucoside
 Mass error <3 ppm

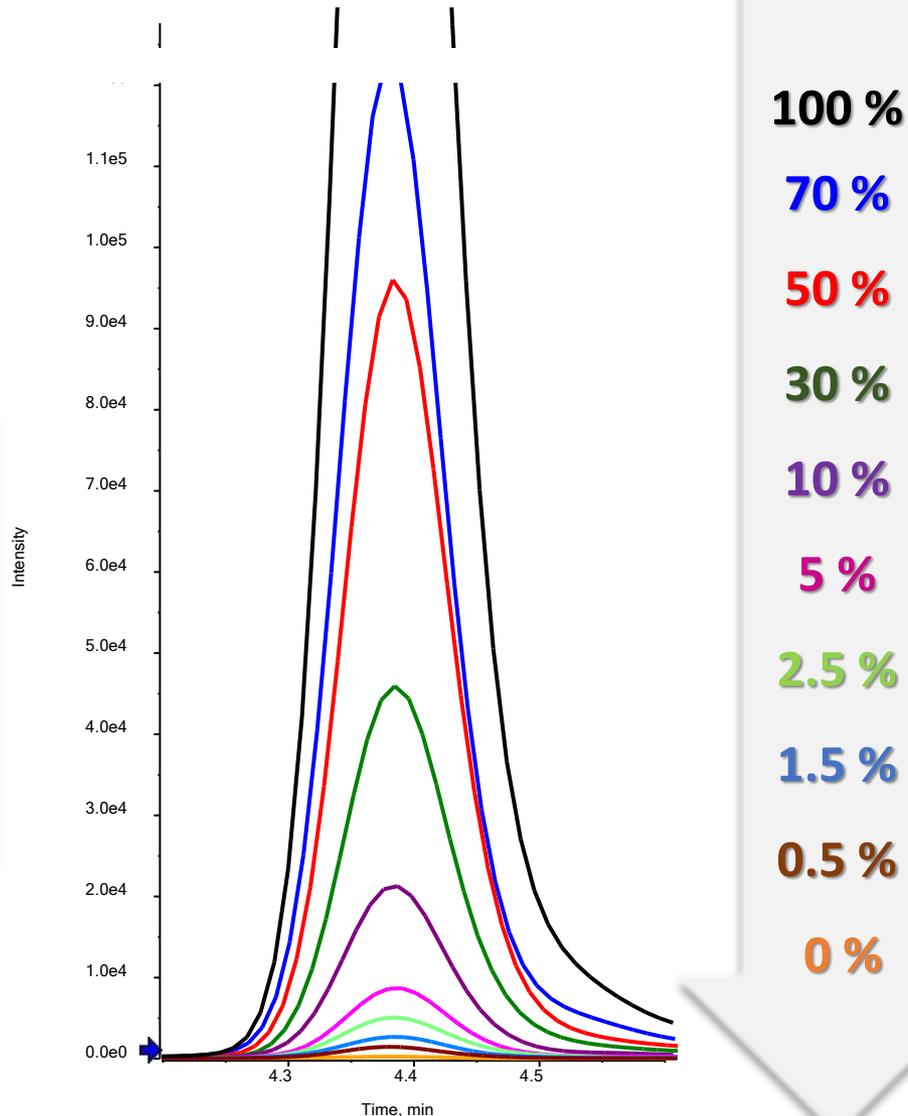
Admixtures 100- 0 % of *Vaccinium macrocarpon*

XIC of myricetin-3-glucoside/galactoside, C₂₁H₂₀O₁₃

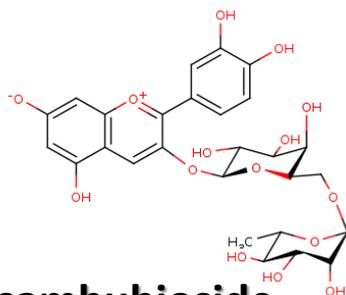
ESI-

marker detected
with addition of
only 0.5 %
of *Vaccinium
macrocarpon*

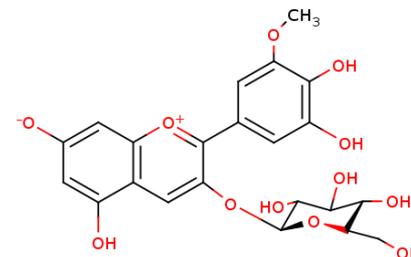
C21H20O13 XIC from 2015-10...0025 Da, Gaussian smoothed



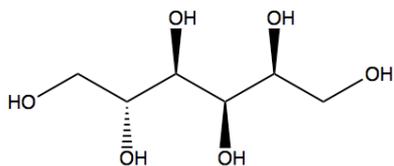
Identified characteristic markers



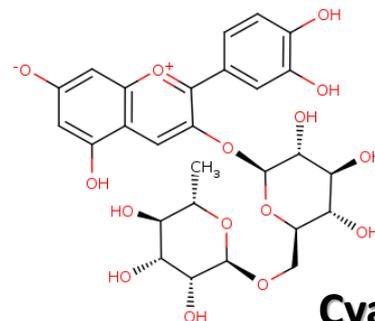
Cyanidin-3-O-sambubioside
m/z 579.1368; [M-H]⁻



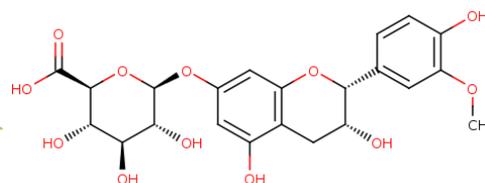
Petunidin-3-O-glucoside
m/z 479.1185; [M+H]⁺



Sorbitol
m/z 181.0732; [M-H]⁻



Cyanidin-3-O-rutinoside
m/z 593.1525; [M-H]⁻

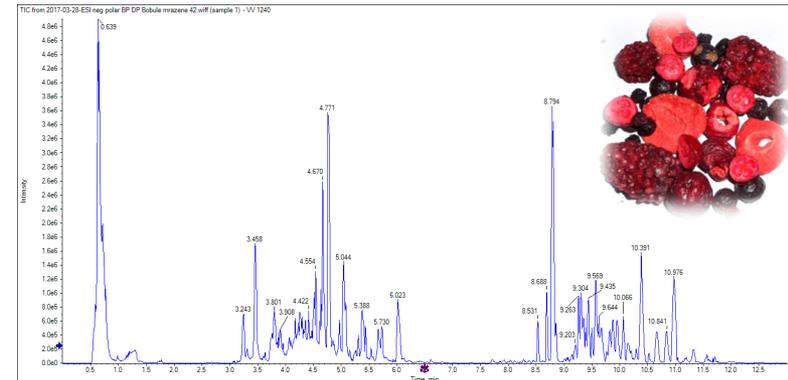
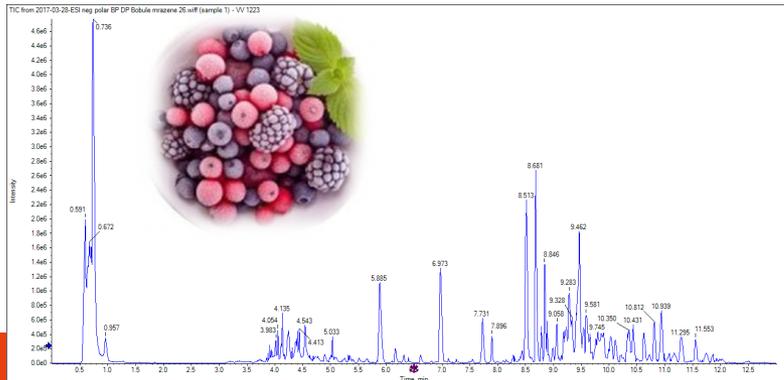


3-O-methyl-epicatechin-7-O-glucuronide
m/z 525.1256; [M-H]⁻

How does the drying process influence the metabolome?

Thermal treatment induces a lot of changes (interactions with amino acids, Maillard reaction...)

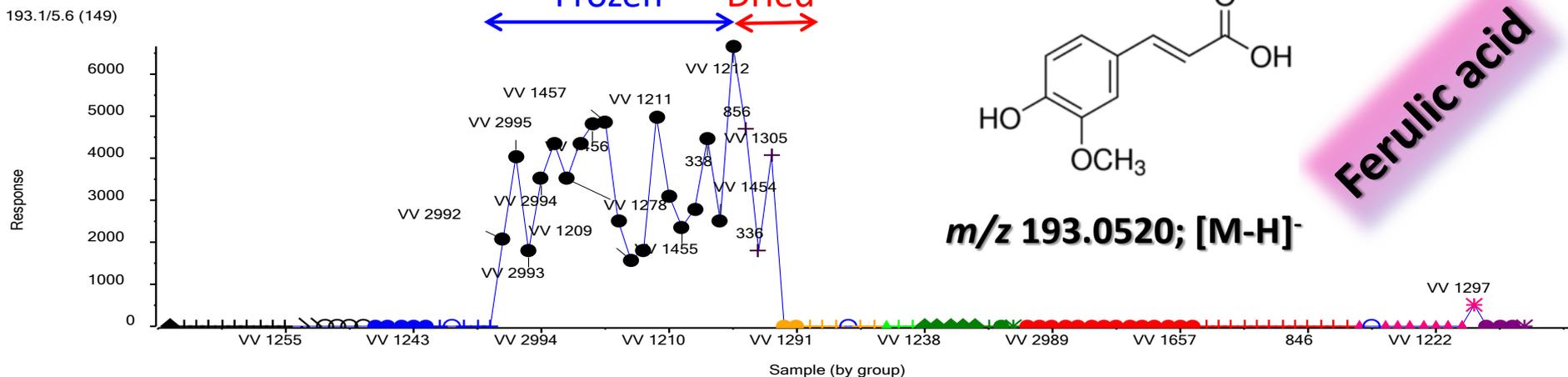
Does it affect stability of identified characteristic markers?



Markers stability: trend diagrams

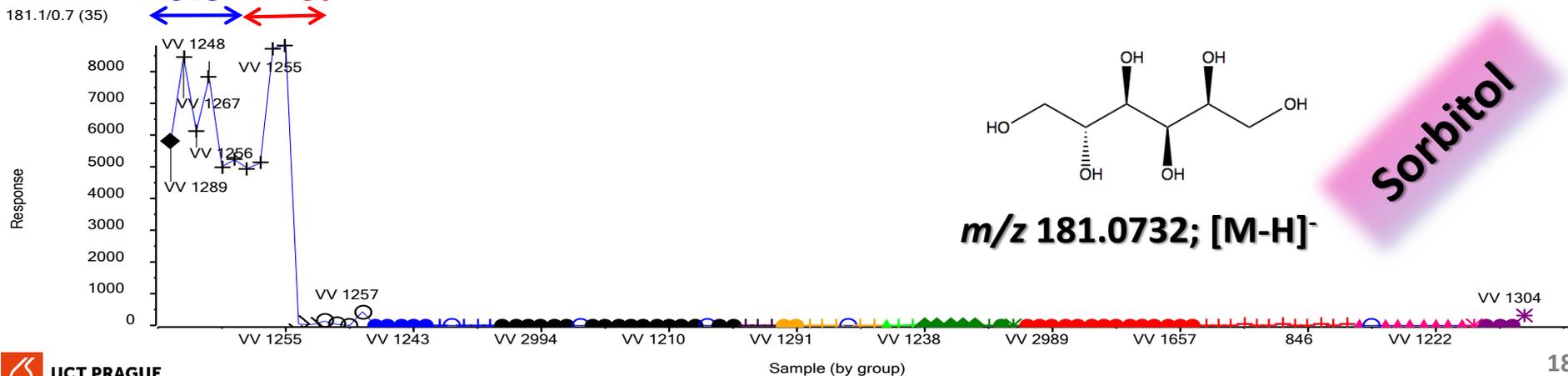
Lingonberry

Frozen Dried

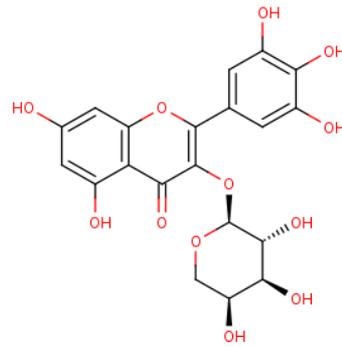


Chokeberry

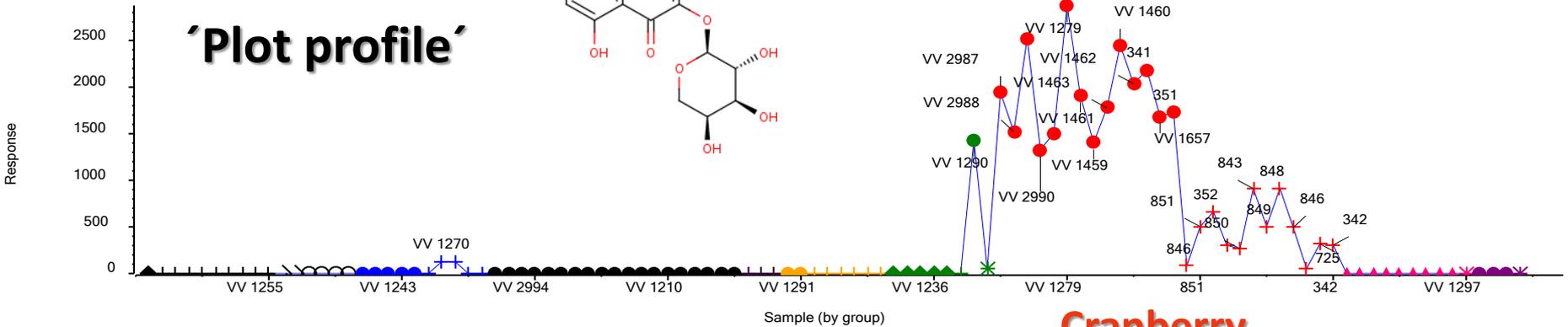
Frozen Dried



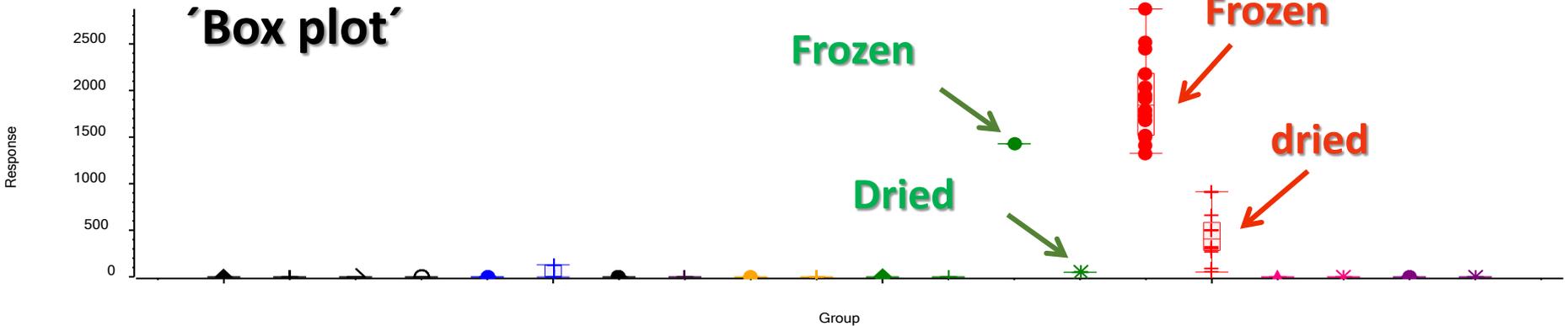
Trend diagrams: Myricetin-arabinoside



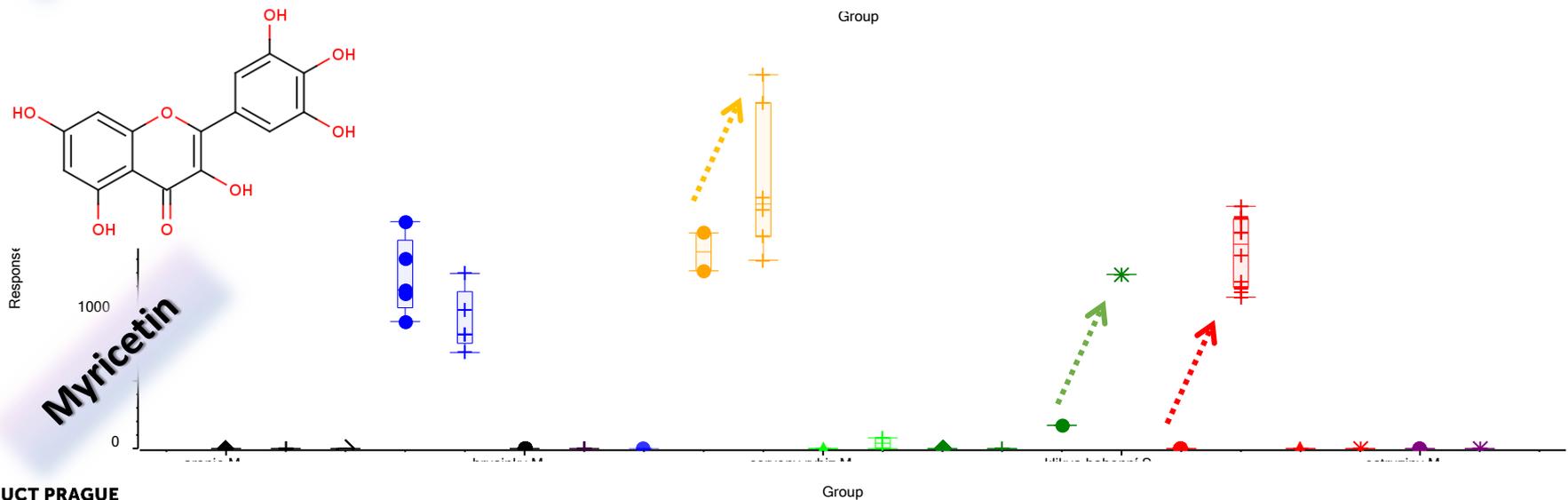
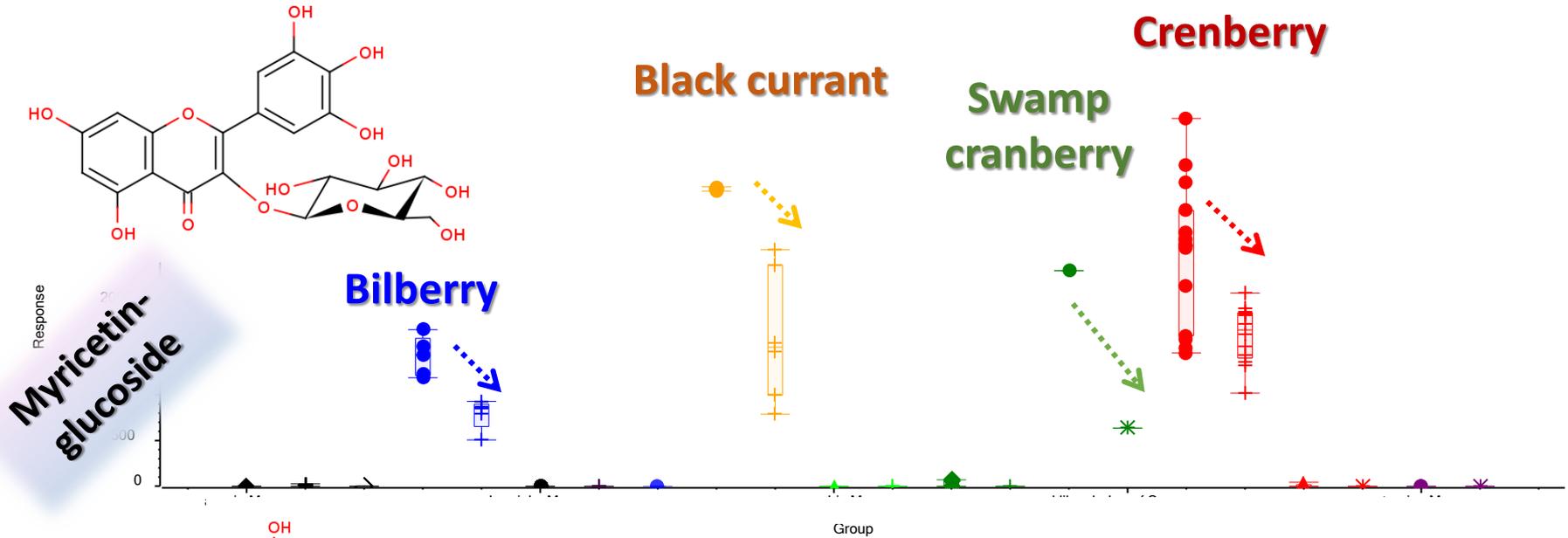
449.1/5.3 (1049)



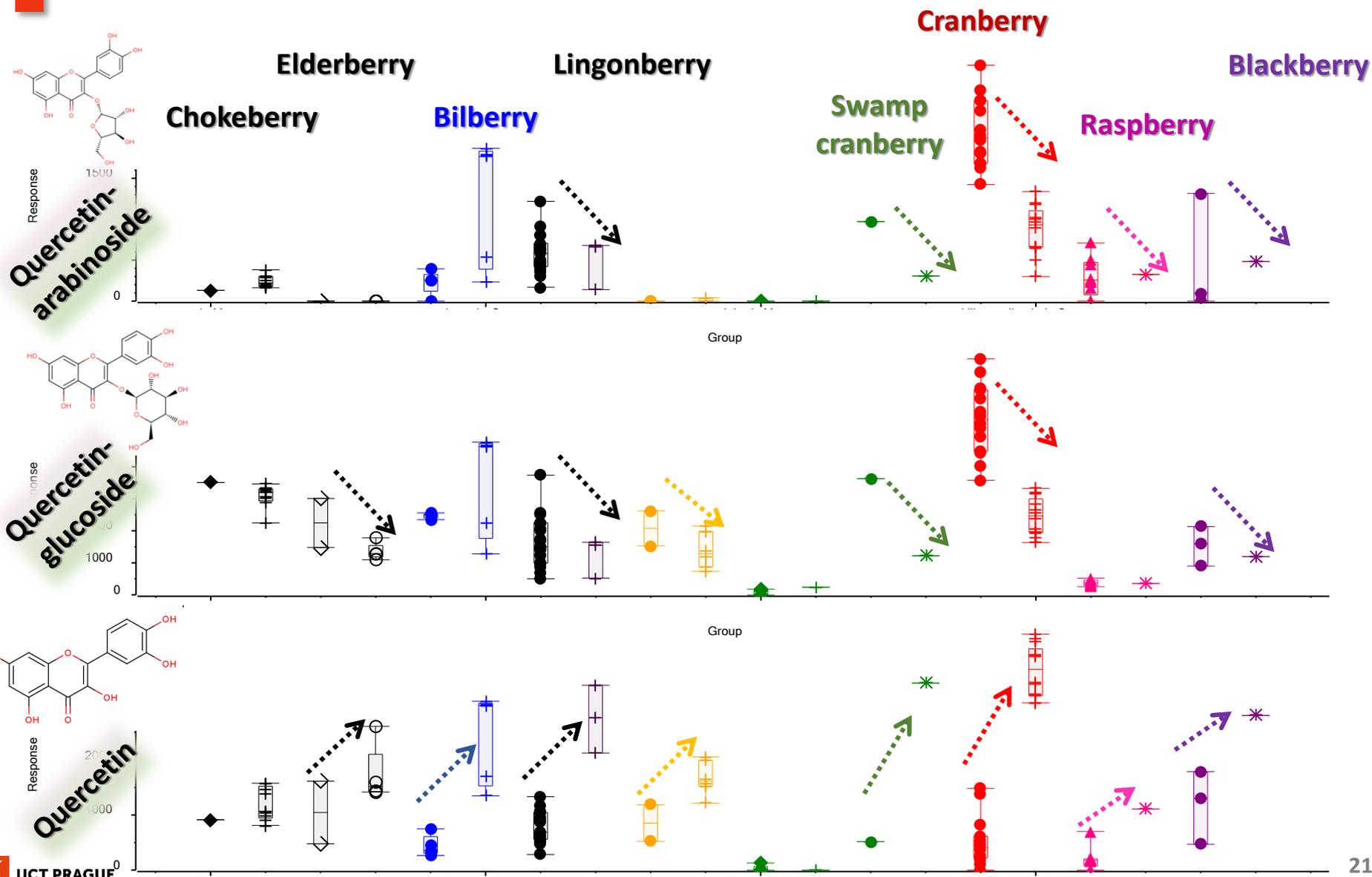
449.1/5.3 (1049)



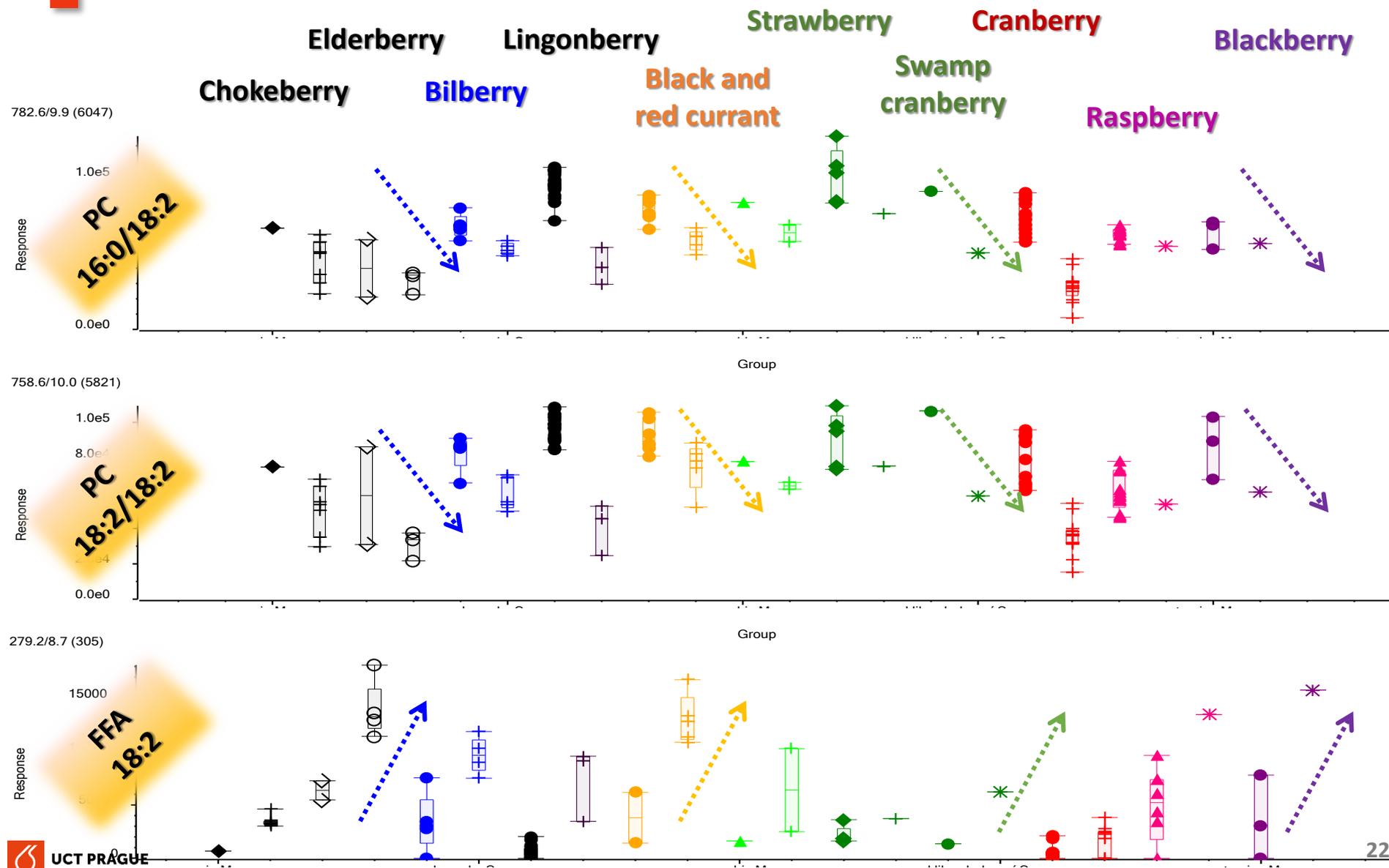
Influence of drying on flavonoids: Myricetin



Influence of drying on flavonoids : Quercetin



Influence of drying on phospholipids



Conclusions



- **U-HPLC-HRMS/MS** (6600 Sciex) technique was successfully used for metabolomic fingerprinting, **11 species** of red and blue berries involved
- **MarkerView (Sciex)** allowed data processing, pre-treatment and analysis to be carried out in order to find characteristic markers.
- **Phenolic compounds and lipids** proved to be the most significant markers enabling discrimination of berries (methanolic extracts)
- Drying process was shown to **influence the stability** of certain markers
- As several markers disappear there is a great chance of finding **new markers**, which are originated during the heat treatment