



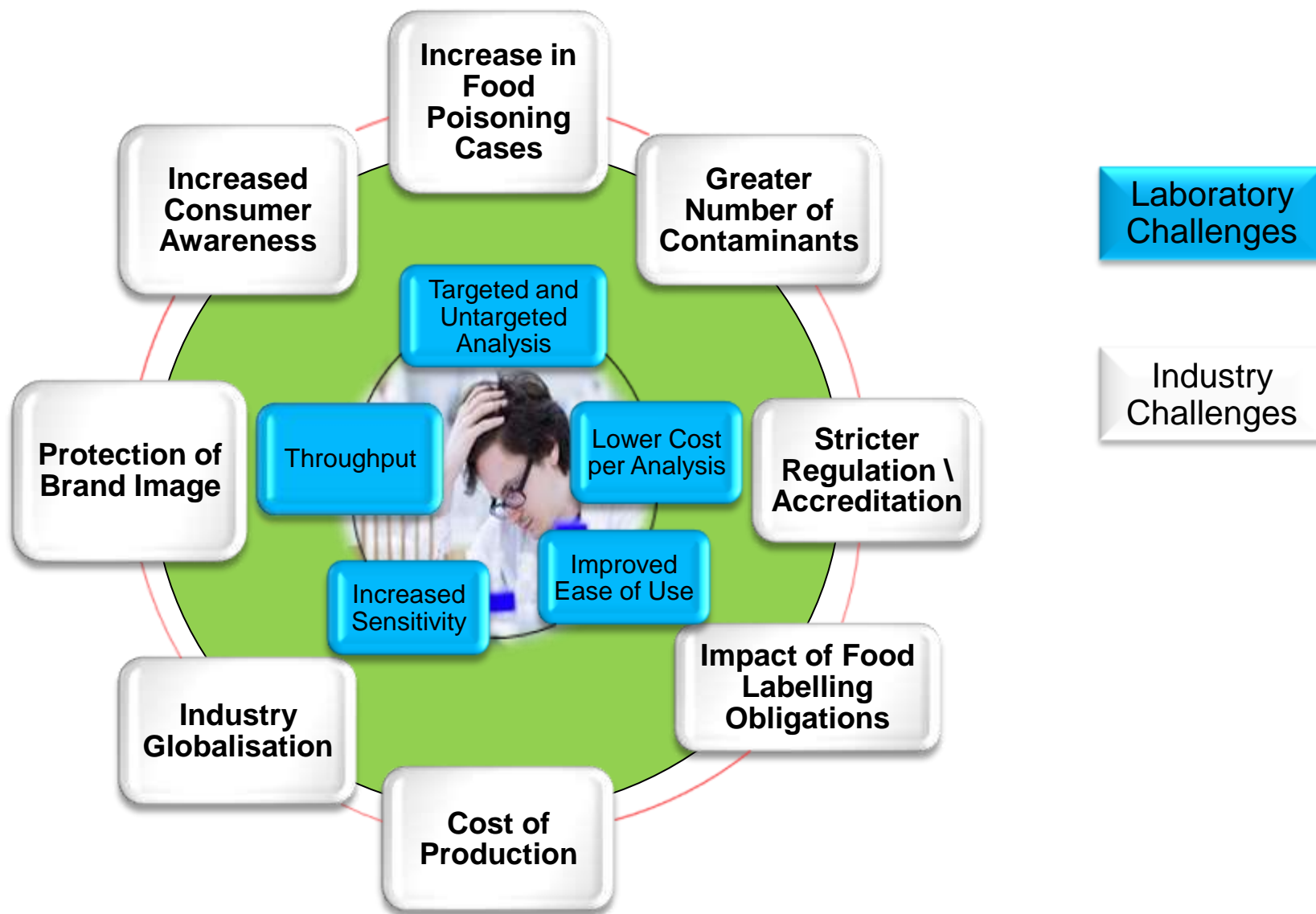
**ThermoFisher**  
SCIENTIFIC

# Food Hazards Screening and Food Fraud Analysis by Orbitrap HRAM MS

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LSMS Application Manager  
Sep. 13<sup>th</sup>, Shanghai

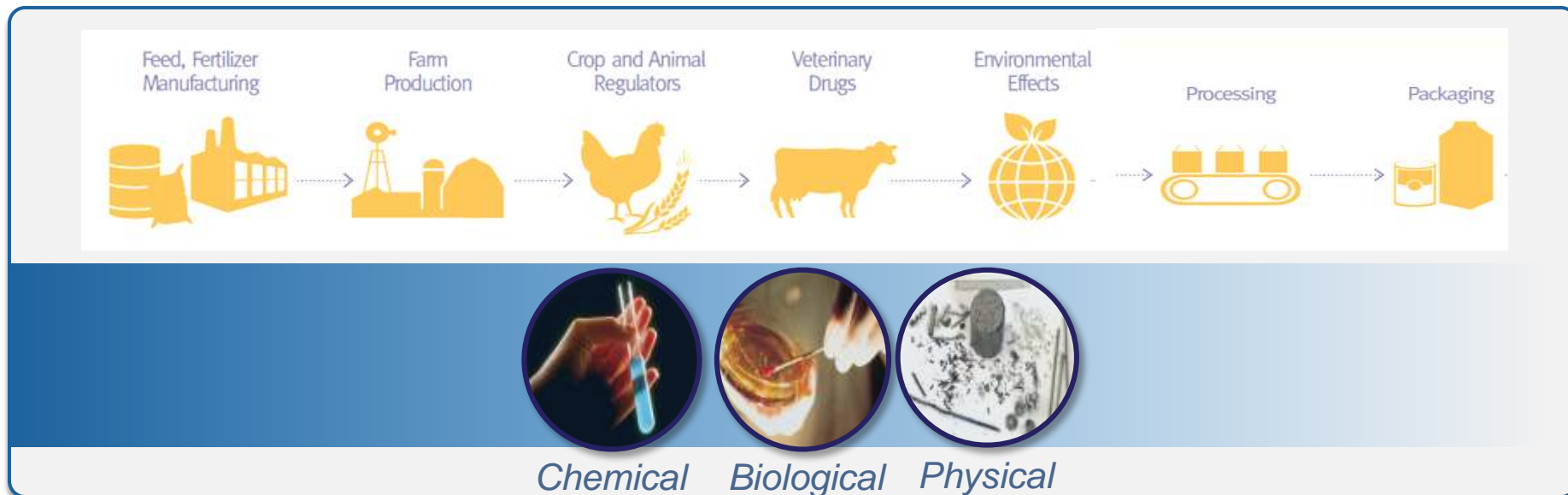
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# Safety and Authentication Challenges in Food Industry



# Key Requests from Food Manufacturers

## Food supply chain from source to consumer



- Produce safe food
- Prevent food fraud, adulteration
- Produce high quality products and monitor product variability
- Produce new products
- Meet legislative requirements for food safety, Labeling and GMP
- Produce and market healthier products

- Method Validation & Quality Control Procedures for Pesticide Residues Analysis in Food & Feed
- Implemented 1/1/2014

## HRAM criteria

≥2 ions (pref. including quasi molecular ion)

< 5 ppm mass accuracy

At least one fragment ion

Resolution typically >20000FWHM

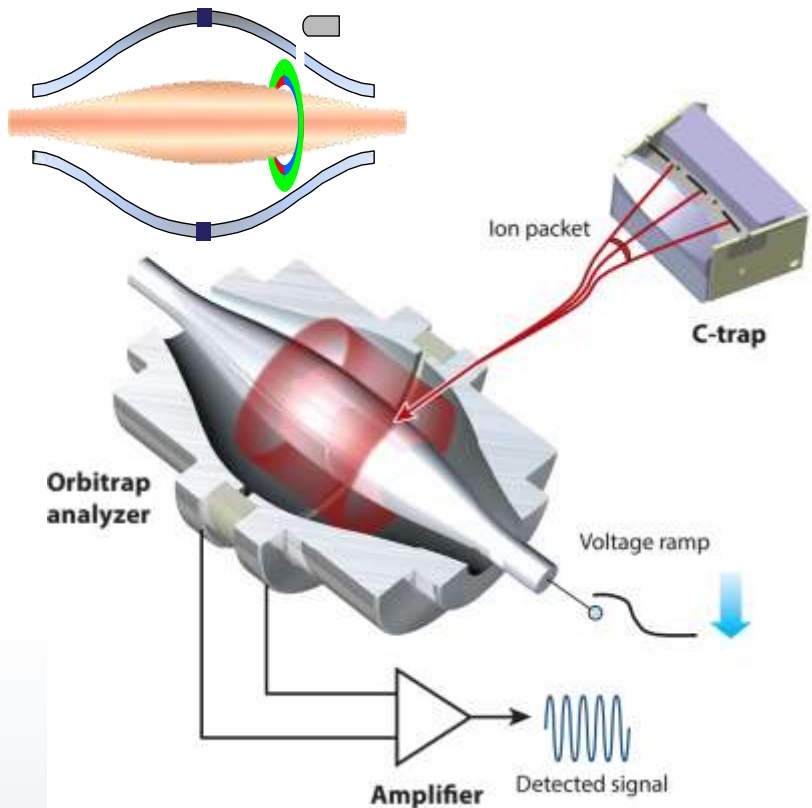
MS mode:	Single-stage MS (unit mass resolution)	Single-stage MS (high resolution/high mass accuracy)	MS/MS
Typical systems (examples):	Quadrupole, ion trap, time-of-flight (TOF)	TOF, Orbitrap, FTMS, magnetic sector	Triple quadrupole, ion trap, hybrid MS (e.g. Q-TOF, Q-trap)
Acquisition mode:	Full scan, Limited <i>m/z</i> range, Selected ion monitoring (SIM)	Full scan, Limited <i>m/z</i> range, Selected ion monitoring (SIM)	Selected/multiple reaction monitoring (SRM/MRM), full scan product-ion spectra
Requirements for identification:	≥ 3 diagnostic ions, preferably including the (quasi) molecular ion	≥ 2 diagnostic ions, preferably including the (quasi) molecular ion; mass accuracy < 5 ppm; at least one fragment ion	≥ 2 product ions
Ion ratio(s): according to Table 5			

# Identification Criteria - Other Examples

P. Lucci and C.P.B. Martins in *Fast Liquid Chromatography – Mass Spectrometry Methods in Food and Environmental Analysis* - World Scientific Publishing Company (March 2015)

	<b>EU 2002/657/EC</b>	<b>SANCO 12571/2013</b>	<b>EU-RL- MB SOP</b>	<b>Gerssen (2010)</b>	<b>Mol (2012)</b>	<b>Domènech (2014)</b>	<b>Kumar (2014)</b>	<b>Pitarch (2007)</b>
<b>Analytes</b>	-	Pesticides	Lipophilic Toxins	Lipophilic Toxins	Pesticides	Lipophilic Toxins	Ronidazole Nitroimidazoles	Priority organic micropollutants
<b>Matrix</b>	Food	Food and Feed	Molluscs	Shellfish	Vegetables &Fruits	Mussels	Muscle	Water
<b>Technique</b>	HRMS	HRMS	LC-MS/MS	LC-MS/MS	LC-HRMS/MS	LC-HRMS/MS	LC-HRMS/MS	GC-MS/MS
<b>Mass Accuracy</b>	-	< 5ppm	-	-	< 5ppm	< 5ppm	< 5ppm	-
<b>Mass Resolving Power (FWHM)</b>	-	≥20,000	-	-	≥20,000	≥20,000	≥70,000	-
<b>Retention Time (RT) Tolerance</b>	2.5%	2.5%	Not exceed 3%	5%	1%	Mean ±3SD (not relative to time)	±1%	Agreement - RT samples & standards
<b>Diagnostic Ions</b>	≥ 2	≥ 2	1 precursor	1 precursor	≥ 2	1	2	1 or 2 precursors
<b>Fragment Ions</b>	-	At least one	At least 2 precursor- product transitions	2 products	At least one	1	At least 1 >20,000FWHM	At least 2 precursor-product transitions
<b>Isotope Ions</b>	-	-	-	-	M+1 M+2	M+1	-	-
<b>Ion Ratio</b>	Relative intensity (% at base peak)	Relative intensity (% at base peak)	Must be recorded	As described 2002/657/EC	Fragment Ion Ratio: Diagnostic/Fragment Isotope Ion Ratio: Diagnostic/M+1 (M+2)	Fragment Ion Ratio: Diagnostic/Fragme nt Isotope Ion Ratio: Diagnostic/M+1	At least one	Ratio between quantitative and confirmation transition
<b>Fragment-Isotope Ion Ratio Tolerance</b>	2 IPs for precursor ion 2.5 IPs for a product	As described 2002/657/EC	-	As described 2002/657/EC	Independent of relative intensity between ions: ±50%	As described 2002/657/EC	-	As described 2002/657/EC

# Orbitrap HRAM MS with Unmatched Performance



Unmatched ultrahigh resolution,  
accurate mass performance

## ● What Orbitrap provides

- Ultra-high resolution
- Long-term mass accuracy
- Uncompromised sensitivity
- Scan speed and dynamic range
- High-quality MS/MS fragmentation
- - **at the same time**

## ● 10 Years Development in Product

- LTQ Orbitrap
- Q Exactive LC/GC
- Orbitrap Fusion

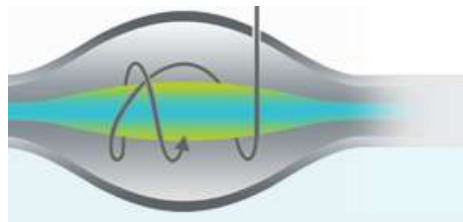
**Significantly superior  
analytical results for  
Multiple Omics**

**metabolomics/lipidomics/proteomics  
In foodomics testing and research**

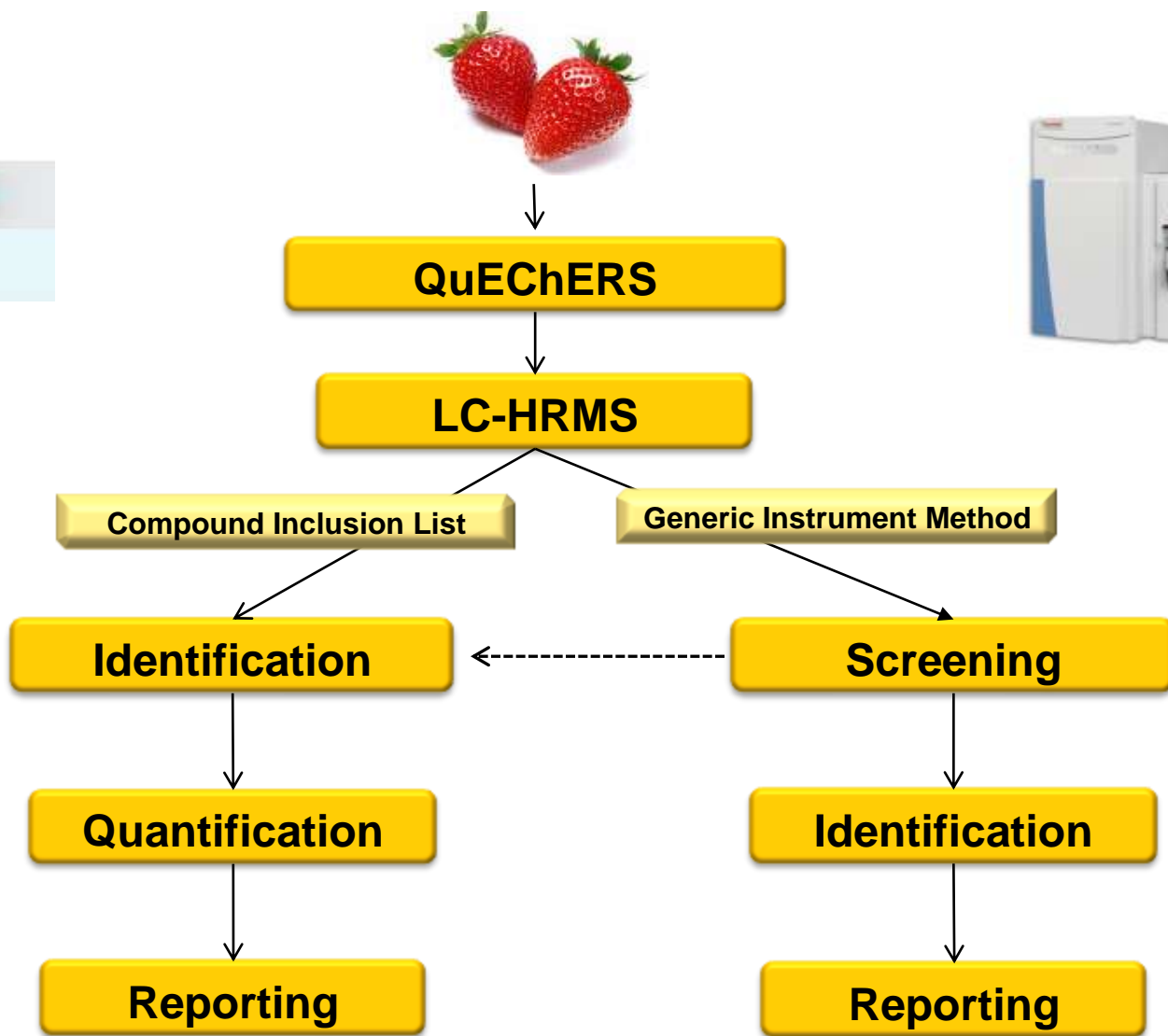




# Typical LC-HRMS Workflow for Pesticide Residue Analysis



Retention Time  
Exact Mass ( $\pm 5$  ppm)  
Product ions  
Ion ratios  
Isotopic Pattern  
Library Matching



# 3 ways of Screening/Quantitation for Routine Work

## Full MS or targeted SIM/ddMS2

- Post-acquisition - extracted ion chromatograms of parent ions of interest
- Relies on high resolution for selectivity
- Useful for less complex background
- No method development/preparation needed

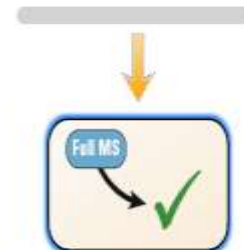
## Full MS/ All Ion Fragmentation – vDIA\*

- Post-acquisition - extracted ion chromatograms of parent ions of interest
- Scheduled target (inclusion) list (Rt,  $m/z$ )
- Minimum method development (e.g., predefine parent ions, tr)
- Also for screening purposes

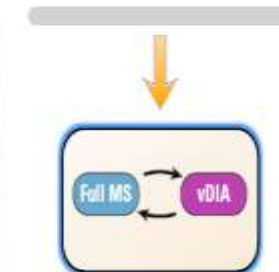
## PRM (Parallel Reaction Monitoring)

- Post-acquisition – extracted ion chromatograms of parent -> fragment transitions acquired
- Scheduled target list (Rt,  $m/z$ , collision energy)
- Most sensitive and selective even in highly complex matrices

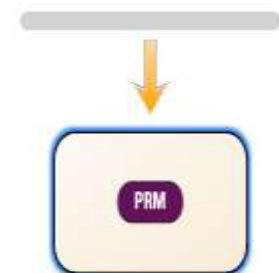
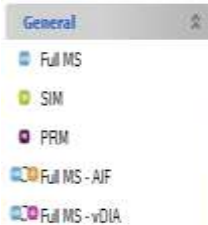
### Experiments



### Experiments



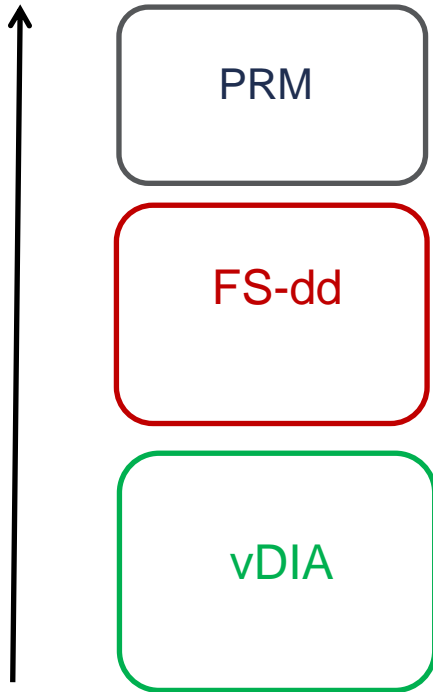
### Experiments



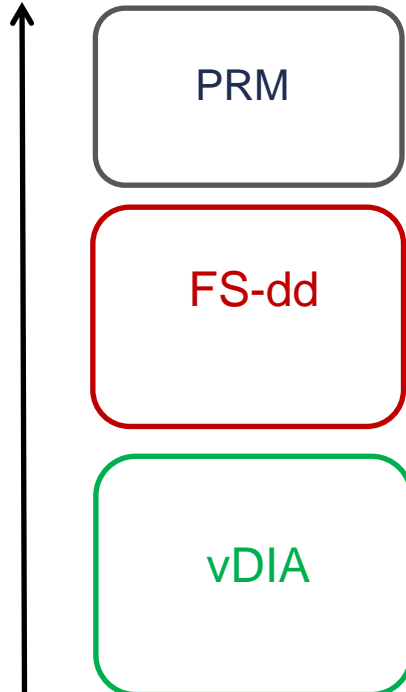


# Q Exactive Focus Scan Methods

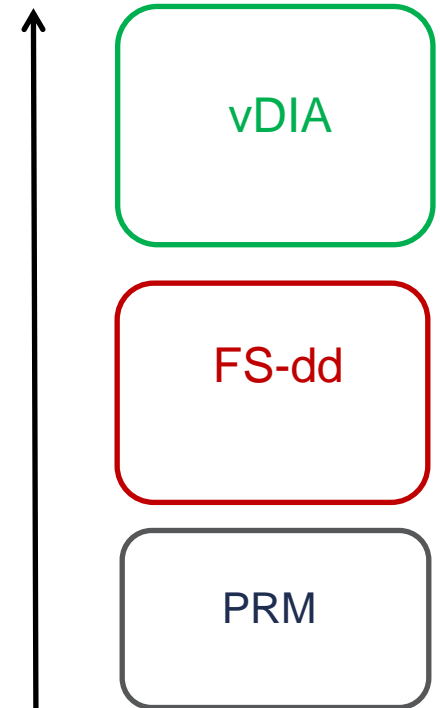
## Selectivity



## Sensitivity



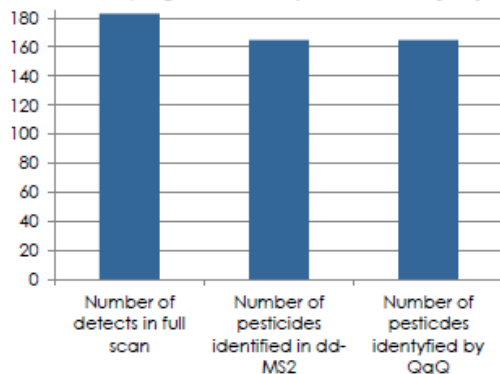
## Information



# What Scan Mode is Right for Your Workflow?

## Real samples

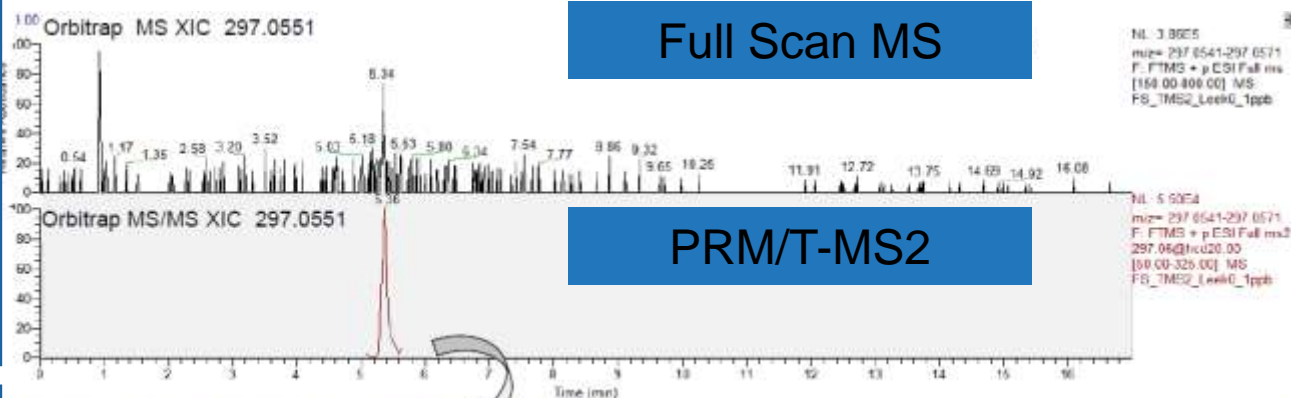
100 real samples analysed by QExactive™ FS/dd-MS<sup>2</sup> (single MS<sup>2</sup> scan\*) and verified by QqQ



Precursor ion @ R=70,000

Product ion @ R=70,000

Leek extract spiked with 0.1 ppb imazalil



Full Scan MS

PRM/T-MS2



## Thermo Scientific™ UltiMate™ XRS:

- Mobile phase:

A: Water:MeOH (98:2) + 5mM Ammonium formiate & 0.1% FA

B: MeOH:Water (98:2) + 5mM Ammonium formiate & 0.1% FA

- Injection volume: **1  $\mu$ l**

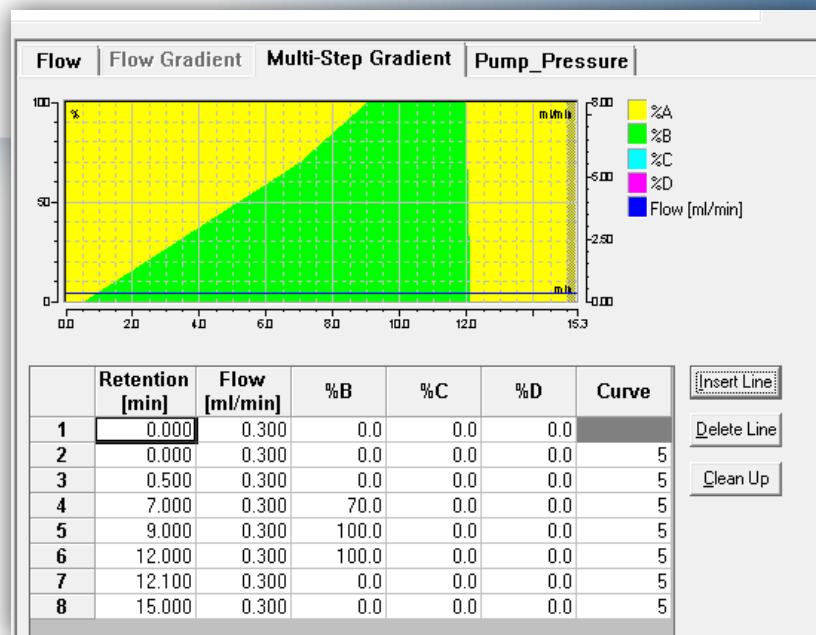
- Column: Accucore aQ column 100 mm x 2.1 mm x 2.6  $\mu$ m

- Column temperature: 25°C

- Flow rate: 300  $\mu$ l/min

- Run time: 15 min

- Gradient:



Thermo Scientific™ UltiMate™ XRS LC

# Recommended MS Tune Method Parameters

## Q Exactive Focus:

- Source: HESI
- Detection mode: variable Data Independent Analysis (vDIA)



Scan parameters	
History	→
Scan type	Full MS
Scan range	70.0 to 900.0 m/z
Fragmentation	None
Resolution	70,000
Polarity	Positive
Microscans	1
Lock masses	Off
AGC target	1e6
Maximum inject time	50

Apply Help  Hot link

HESI source	
	<b>actual</b>
Sheath gas flow rate	40
Aux gas flow rate	10
Sweep gas flow rate	2
Spray voltage ( kV )	3.50
Spray current (µA)	0.10
Capillary temp. (°C)	250
S-lens RF level	55.0
Aux gas heater temp (°C)	272

Source Auto-Defaults...

Apply Help  Hot link

# Recommended Method Parameters for Symmetric vDIA

## Properties

### Properties of the method

<input type="checkbox"/> <b>Global Settings</b>	
User Role	Advanced
Use lock masses	off
Lock mass injecti	—
Chrom. peak wid	12 s
<input type="checkbox"/> <b>Time</b>	
Method duration	15.00 min
<input type="checkbox"/> <b>Customized Tolerances (+/-)</b>	
Lock Masses	—
Inclusion	—
Exclusion	—
Dynamic Exclusion	—

### Properties of Full MS - vDIA

<input type="checkbox"/> <b>General</b>	
Polarity	positive
<input type="checkbox"/> <b>Full MS</b>	
Resolution	70,000
Scan range	120 to 1000 m/z
<input type="checkbox"/> <b>variable DIA</b>	
Resolution	17,500
# vDIA segment:	8
vDIA isolation range	50 to 150 m/z
vDIA isolation range	140 to 240 m/z
vDIA isolation range	230 to 330 m/z
vDIA isolation range	320 to 420 m/z
vDIA isolation range	410 to 510 m/z
vDIA isolation range	500 to 600 m/z
vDIA isolation range	590 to 690 m/z
vDIA isolation range	680 to 780 m/z
CE / stepped CE	30, 50, 70
Fixed first mass	—
AGC target	5e5
Spectrum data type	Centroid

# Alternative Recommended MS Method Settings

↓

↓

↓

**Properties of the method**

- Global Settings**
  - User Role: Advanced
  - Use lock masses: off
  - Lock mass inject: —
  - Chrom. peak wid: 12 s
- Time**
  - Method duration: 17.00 min
- Customized Tolerances (+/-)**

**Properties of Full MS - vDIA**

- General**
  - Polarity: positive
  - In-source CID: —
- Full MS**
  - Resolution: 70,000
  - Scan range: 120 to 1000 m/z
  - AGC target: 1e6
  - Maximum IT: auto
  - Microscans: 1
  - Spectrum data ty: Profile
- variable DIA**
  - Resolution: 17,500
  - # vDIA segment: 5
  - vDIA isolation rai: 120 to 205 m/z
  - vDIA isolation rai: 195 to 305 m/z
  - vDIA isolation rai: 295 to 405 m/z
  - vDIA isolation rai: 395 to 505 m/z
  - vDIA isolation rai: 495 to 1000 m/z
  - CE / stepped CE: 30, 50, 70
  - Fixed first mass: —
  - AGC target: 5e5
  - Maximum IT: auto
  - Microscans: 1
  - Spectrum data ty: Centroid

**Properties of the method**

- Global Settings**
  - User Role: Advanced
  - Use lock masses: off
  - Lock mass inject: —
  - Chrom. peak wid: 12 s
- Time**
  - Method duration: 17.00 min
- Customized Tolerances (+/-)**

**Properties of Full MS**

- General**
  - Polarity: positive
  - dd-MS<sup>2</sup>: Confirmation
  - In-source CID: —
- Full MS**
  - Resolution: 70,000
  - # Scan ranges: 1
  - Scan range: 80 to 750 m/z
  - AGC target: 1e6
  - Maximum IT: auto
  - Microscans: 1
  - Spectrum data ty: Profile
- dd-MS<sup>2</sup> Confirmation**
  - Resolution: 17,500
  - Isolation window: 3.0 m/z
  - Isolation offset: —
  - CE / stepped CE: 15, 30, 50
  - Fixed first mass: —
  - Default charge at 1
  - AGC target: 5e4
  - Maximum IT: auto
  - Loop count: 1
  - Underfill ratio: 1.0 %
  - Intensity thresh: auto
  - Apex trigger: —
  - Dynamic exclusi: auto
  - Spectrum data ty: Centroid

**Properties of the method**

- Global Settings**
  - User Role: Advanced
  - Use lock masses: off
  - Lock mass inject: —
  - Chrom. peak wid: 12 s
- Time**
  - Method duration: 17.00 min
- Customized Tolerances (+/-)**

**Properties of PRM**

- General**
  - In-source CID: —
- Targeted-MS<sup>2</sup>**
  - Resolution: 70,000
  - Isolation window: 3.0 m/z
  - Isolation offset: —
  - CE / stepped CE: 15, 30, 50
  - Fixed first mass: —
  - Default charge at 1
  - AGC target: 1e5
  - Maximum IT: auto
  - Microscans: 1
  - Spectrum data ty: Profile

\* Statistically identical results with symmetrical and asymmetrical vDIA settings



# Targeted 330 Compounds – vDIA Screening Method

Acephate	Bromacil	Cumyluron	Dinotefuran	Fenthion-sulfone	Heptenophos	Methabenzthiazuron	Penconazole	Pyridaben	Thiacloprid
Acetamiprid	Bromuconazole	Cyanazine	Dioxacarb H	Fenthion-sulfoxide	Hexaconazole	Methamidophos	Pencycuron	Pyridate e	Thiamethoxam
Acibenzolar-S-methyl	Bupirimate	Cyazofamid	Disulfoton	Fenuron	Hexaflumuron	Methidathion	Permethrin	Pyrimethanil	Thiazopyr H
Aclonifen	Buprofezin T	Cycloate	Dithiopyr	Fipronil	Hexazinone	Methiocarb	Phenmedipham	Pyroquilon	Thidiazuron T
Alachlor	Butachlor	Cyfluron	Diuron	Flazasulfuron	Hexythiazox T	Methiocarb sulfoxide	Phenthoate	Pyroxulam	Thiobencarb
Alanycarb	Butafenacil	Cyflufenamid	Dodemorph	Flonicamid T	Imazalil	Methiocarb-sulfone	Phoxim	Quinoxyfen	Thiodicarb
Aldicarb	Butocarboxim	Cymoxanil	Epoxiconazole	Florasulam	Imazaquin	Metholcarb	Picoxystrobin	Quizalofop T	Thiofanox
Aldicarb sulfone	Butoxyacboxim	Cypermethrin T	Esprocarb	Fluazifop	Imazethapyr	Methomyl	Piperonyl butoxide	Quizalofop-p-ethyl	Thionazin
Aldicarb sulfoxide	Carbaryl	Cyproconazole	Etaconazole	Flufenacet	Imidacloprid	Methoprotryne	Piperophos	Resmethrin	Tolfenpyrad
Allethrin	Carbendazim	Cyprodinil	Ethiofencarb	Flufenoxuron	Indoxacarb	Methoxyfenozide	Pirimicarb	Rimsulfuron	Tralkoxydim
Ametryn	Carbetamide	Cyromazine	Ethiofencarb_sulfoxide	Flumetsulam	lprovalicarb	Metobromuron	Pirimiphos-ethyl	Rotenone	Triadimefon
Aminocarb	Carbofuran	Deltamethrin	Ethiofencarb-sulfone	Flumioxazin	Isocarbophos	Metolachlor	Pirimiphos-methyl	Schradan	Triadimenol
Ancymidol	Carbofuran-3-hydroxy	Demeton-S-methylsulfone	Ethiprole	Fluometuron	Isofenphos	Metosulam	Pretilachlor	Sethoxydim	Triazophos
Anilofos	Carbosulfan	Desmedipham	Ethirimol	Fluopicolide	Isoprocarb	Metoxuron	Primsulfuron-methyl	Simeconazole T	Trichlorfon
Aramite H	Carboxin	Desmethyl-pirimicarb	Ethofumesate	Fluopyram	Isoprothiolane	Metrafenone	Prochloraz	Simecyn	Tricyclazole
Atrazine	Carfentrazone-ethyl	Desmetryn	Ethoxyquin	Fluoxastrobin	Isoproturon	Metribuzin	Profenofos	Spinosyn A	Tridemorph T
Azaconazole	Carpropamide	Dichlofenthion	Etofenprox	Fluquinconazole T	Isoxaben	Metsulfuron-methyl	Promecarb	Spiromesifen	Trietazine
Azamethiphos	Chlorantraniliprole	Dichlorvos	Etoxazole	Flurochloridone	Isoxadifen-ethyl	Mevinphos	Prometon	Spiroxamine	Trifloxystrobin
Azinphos-ethyl	Chlorbromuron	Diclobutrazol	Etrimfos	Fluroxypyr	Isoxaflutole	Mexacarbate	Prometryn	Sulfotep	Triflumizole
Azinphos-methyl	Chlorfenvinphos	Diclotophos	Famoxadone	Flusilazole	Isoxathion	Monocrotophos	Propamocarb	Sulprofos HT	Triflumuron
Azoxystrobin	Chlorfluazuron	Diethofencarb	Fenamidone	Flutriafol	Kresoxim-methyl	Monolinuron	Propanil	Tebuconazole	Triforine
Barban	Chloridazon	Difenacoum	Fenamiphos	Fonofos	Lenacil	Napropamide	Propargite	Tebufenozide	Triticonazole
Bendiocarb	Chlorotoluron	Difenoconazole	Fenarimol	Forchlorfenuron	Malaoxon	Naptalam	Propazine	Tebufenpyrad	Vamidothion
Benfuracarb	Chloroxuron	Diflubenzuron	Fenazaquin	Formetanate	Malathion	Neburon	Propetamphos H	Tebuthiuron	Zoxamide
Benodanil	Chlorpyrifos	Dimefuron	Fenbuconazole	Formetanate hydrochlorid	Mandipropamide	Nicosulfuron	Propiconazole	Teflubenzuron	24D (neg)
Benoxacor	Cinosulfuron	Dimethachlor	Fenhexamid	Formothion	Mefenacet	Nitenpyram	Propoxur	Terbufos	Bentazone (neg)
Bensulfuron-methyl	Clethodim	Dimethametryn	Fenobucarb	Fosthiazate	Mepanipyrim	Nuarimol	Propyzamide	Terbumeton	Bromoxynil (neg)
Benzoximate	Clofentezine	Dimethenamide	Fenoxanil	Fuberidazole	Mepronil	Ofurace	Prosulfocarb	Terbuthylazine	DNOC (neg)
Benzoylprop-ethyl	Clomazone	Dimethoate	Fenoxycarb	Furathiocarb	Mesotrione	Omethoate	Pymetrozine	Terbutryn	Fluazinam (neg) H
Bifenazate	Clopyralid	Dimethomorph	Fenpiclonil	Griseofulvin	Metalaxyl	Oxadixyl	Pyraclostrobin	Tetrachlorvinphos H	Flubendiamide (neg)
Bitertanol	Clothianidin	Dimetilan	Fenpyroximate	Halofenozide	Metamitron	Oxamyl	Pyrazophos	Tetraconazole	MCPA (neg)
Boscalid	Coumaphos	Dimoxystrobin	Fensulfothion	Haloxypop	Metazachlor	Oxyfluorfen	Pyrethrin I	Tetramethrin	Tepraloxymid (neg)
Brodifacoum	Crotoxyphos	Diniconazole	Fenthion	Haloxypop-methyl	Metconazole	Paclbutrazol	Pyrethrin II	Thiabendazole	Terbacil (neg)

Constant peak area    
  No peak in tea at 10ppb    
  Peak neither in honey nor in tea at 10 ppb    
  Not found at all  
 Bad peak shape    
 No peak in honey at 10 ppb    
Compound H or T    
Missing fragment ion confirmation in one of the matrices at 10 ppb

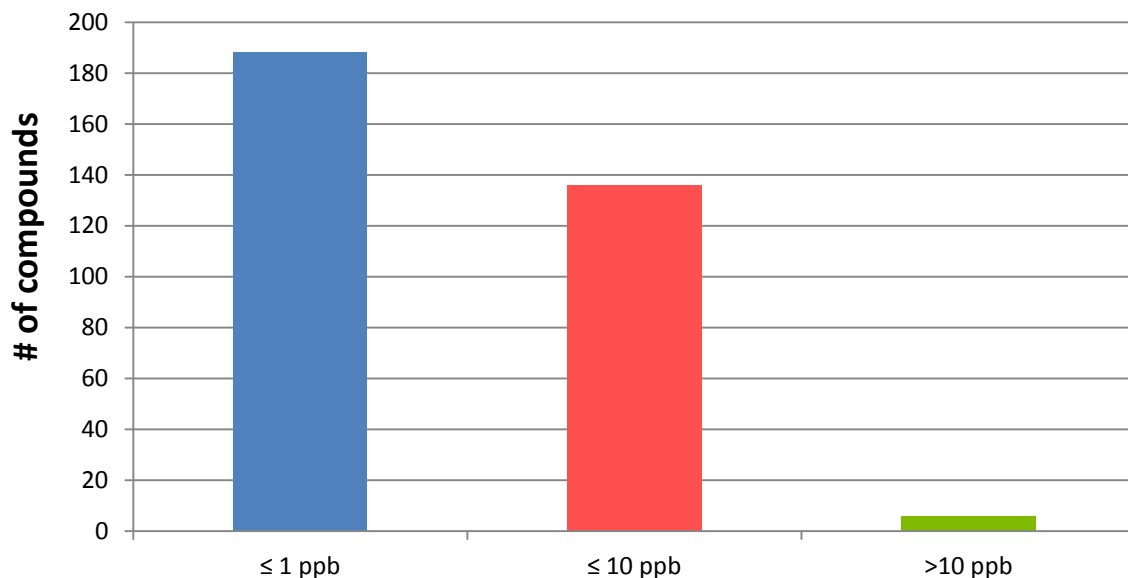
# Validation Parameters

- Selectivity criteria: RT, min. detection of 1 HRAM quan and 1 HRAM qual ion with 5 ppm mass accuracy, ion ratio
- False negative and positive evaluation
- Recovery & repeatability at two concentration levels: 10 & 100 µg/kg
- Injection precision – 10 repeated injections of standard 100 µg/kg
- Screening detection limit (SDL), LOD/LOQ definition
- Linearity: 5 levels matrix matched calibration, duplicate measurement

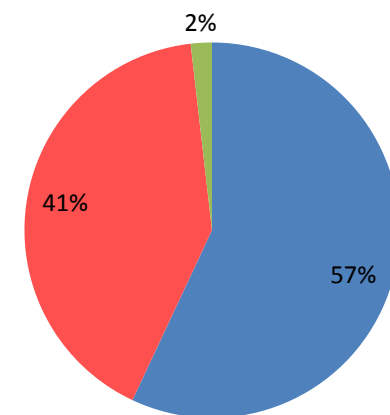
# Results vDIA – Sensitivity Overview Honey

- Screening Detection Limits (SDL) in honey matrix

SDL values in Honey



■ ≤ 1 ppb ■ ≤ 10 ppb ■ >10 ppb

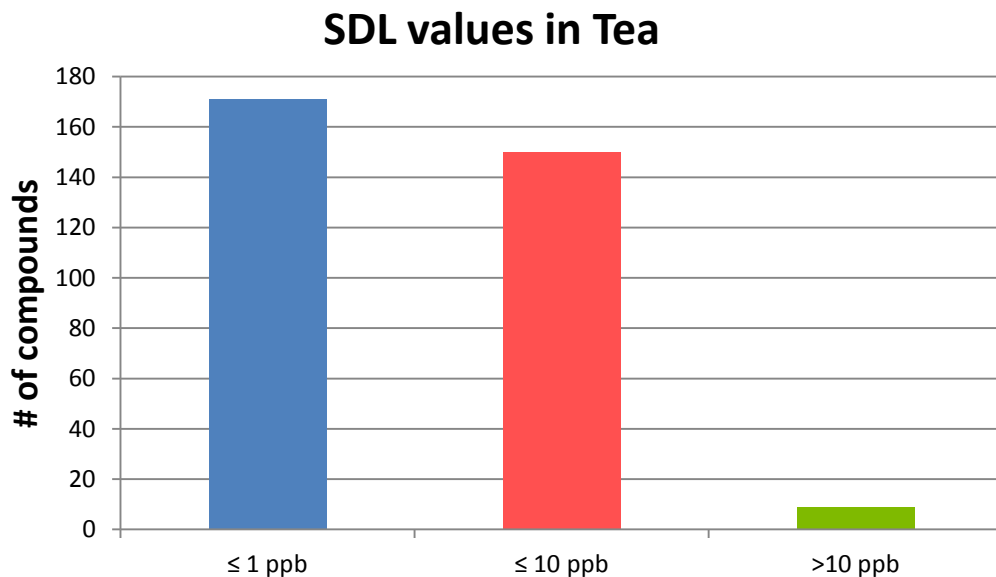


**SUM = 330 compounds; 324 compounds SDL ≤ 10 µg/kg**

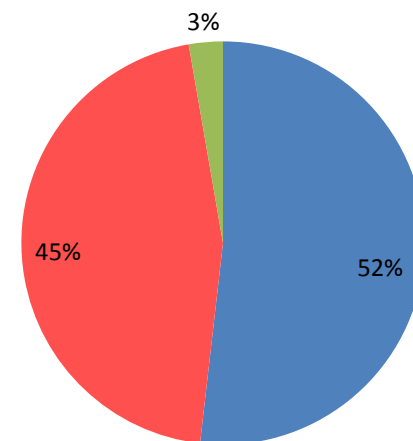
SDL determination according to SANCO12571/2013

# Results vDIA – Sensitivity Overview Tea

- Screening Detection Limits – Tea matrix



■ ≤ 1 ppb ■ ≤ 10 ppb ■ >10 ppb



**SUM = 330 compounds; 321 compounds SDL ≤ 10 µg/kg**

SDL determination according to SANCO12571/2013

# Missing Compounds at 10 ppb

SDL and MRL values for compounds not seen at 10 ppb

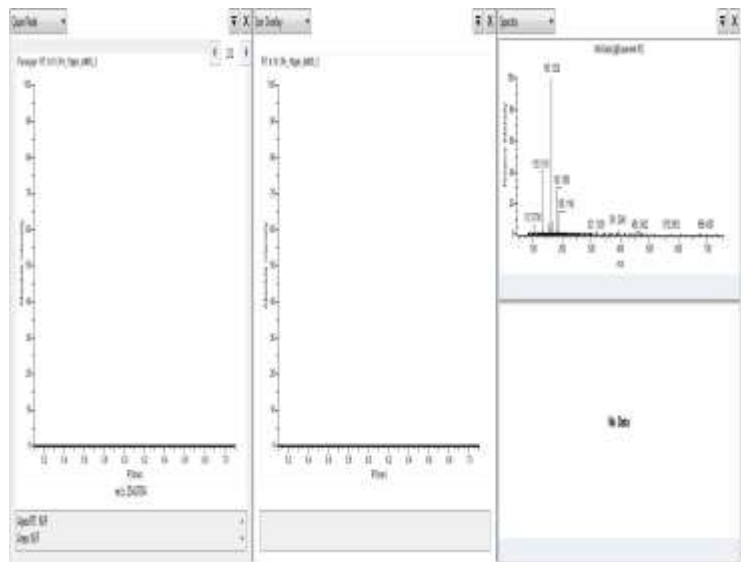
Compound name	SDL [ug/kg] in Honey	MRL for Honey [ug/kg]	SDL [ug/kg] in Tea	MRL for Tea [ug/kg]
Aramite	10	100	>100	n.d.
Bentazone	>100	50	0.5	100
Butafenacil	5	n.d.	60	n.d.
Dimethachlor	50	n.d.	40	20
Fenthion-sulfone	5	10	70	50
Hexaflumuron	3	n.d.	30	n.d.
Isoxathion	50	n.d.	50	n.d.
Mesotrione	>100	n.d.	>100	100
Pyridate	15	50	20	50
Sethoxym	40	50	50	100
Thiazopyr	10	n.d.	50	n.d.

n.d. - not defined

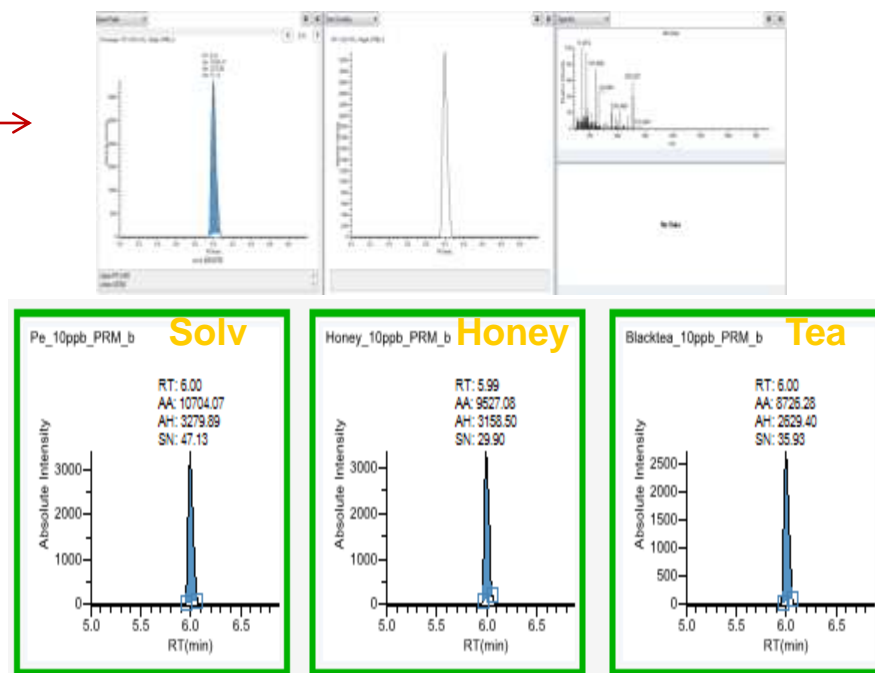
# Sensitivity Improvement at MRL – ddMS vs PRM

Most of compounds **fulfill EU MRL criteria**, but in 16 cases not → PRM recommended

## Fluroxypyr 10 ppb ddMS



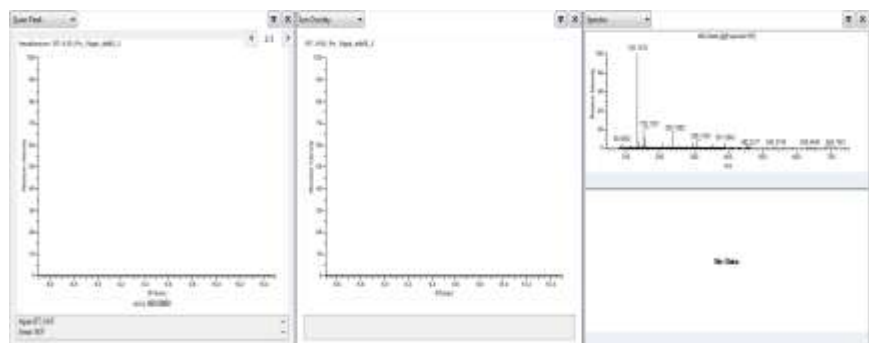
## Fluroxypyr 10 ppb PRM



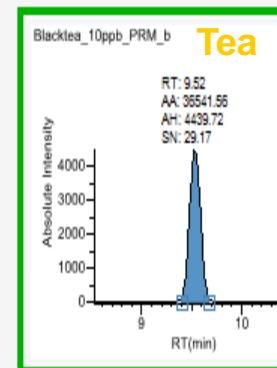
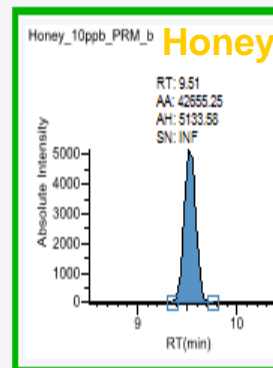
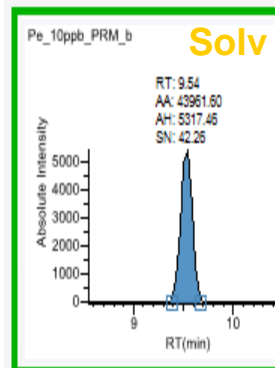
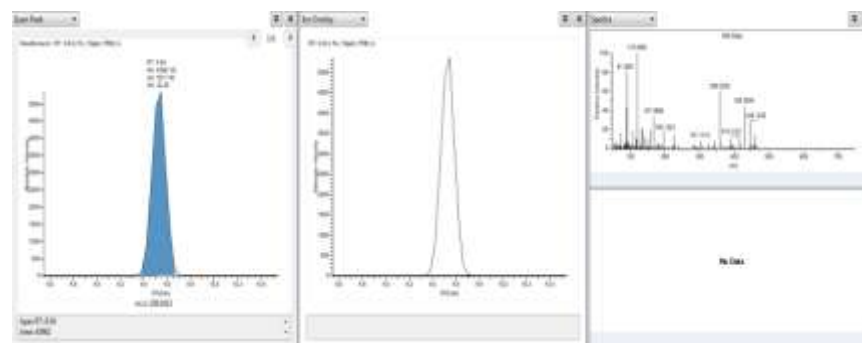


# Sensitivity Improvement at MRL – ddMS vs PRM

Hexaflumuron 10 ppb ddMS

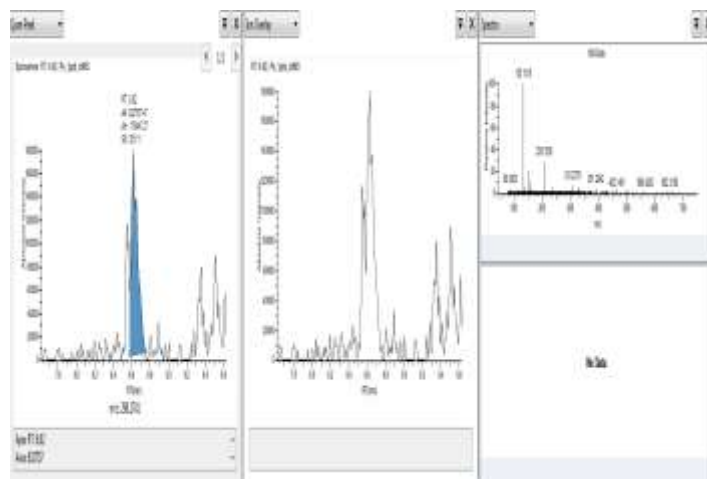


Hexaflumuron 10 ppb PRM

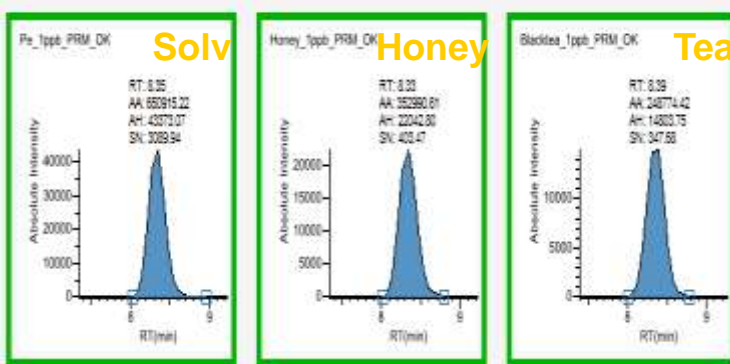
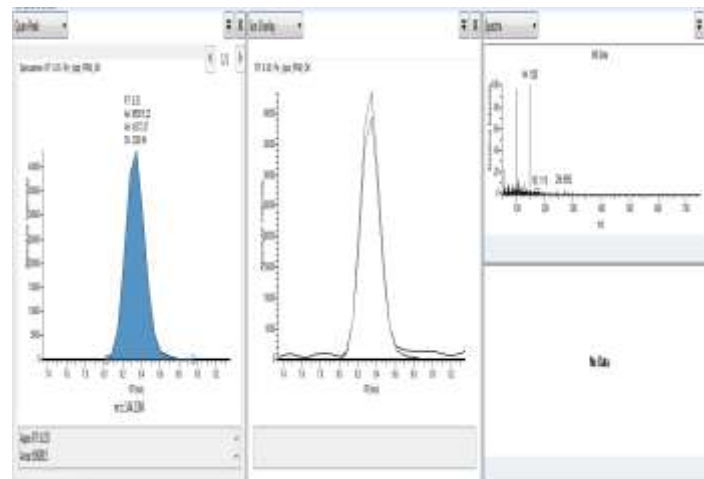


# Sensitivity Improvement – With Parallel Reaction Monitoring

## Spiroxamine 1ppb in ddMS



## Spiroxamine 1ppb in PRM

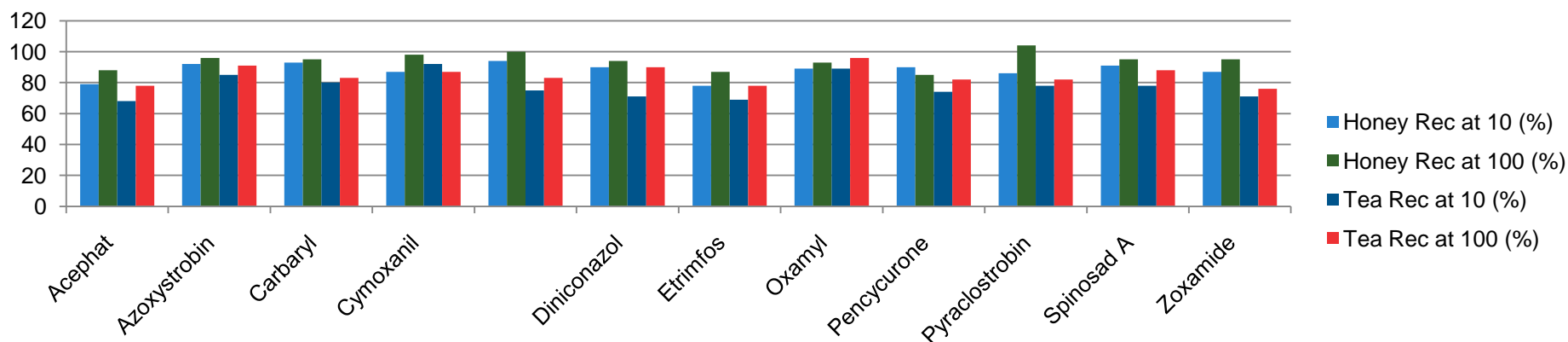


# Method Sensitivity For 12 Representative Compounds - vDIA

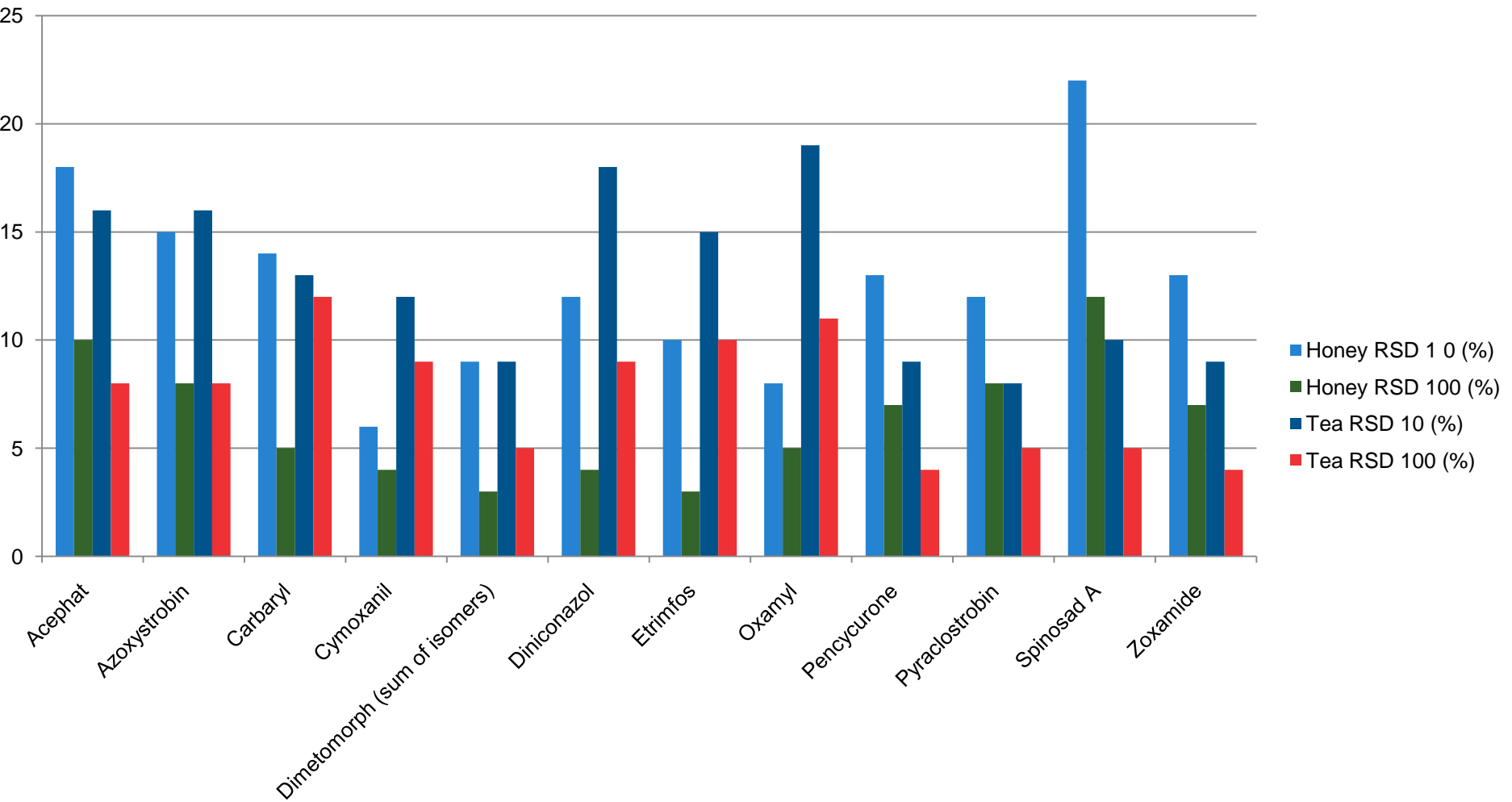
Analyte	Solvent (µg/kg)	Honey (µg/kg)	Tea (µg/kg)	MRL's (µg/kg)	
	LOQ	LOQ	LOQ	Honey	Tea
Acephate	2.5	10	10	20	50
Azoxystrobin	0.25	0.75	1.25	50	100
Carbaryl	0.25	2.5	3	50	50
Cymoxanil	0.25	1.25	5	50	50
Dimetomorph (sum of isomers)	12.5	12.5	25	50	50
Diniconazol	2	2.5	10	50	50
Etrimfos	1.25	1.25	2.5		
Oxamyl	1.25	2.5	5	50	50
Pencycurone	0.25	0.25	1.75	50	10
Pyraclostrobin	0.25	0.25	1.25	50	100
Spinosad A	12.5	12.5	25	50	100
Zoxamide	1.25	1.25	1.75	50	50

# Method Recovery, Repeatability, Linearity - vDIA

Analyte	Sp. Level 1 (µg/kg)	Sp. Level 2 (µg/kg)	Honey				Tea				Linearity in solvent
			RSD 1 (%)	RSD 2 (%)	Rec 1 (%)	Rec 2 (%)	RSD 1 (%)	RSD 2 (%)	Rec 1 (%)	Rec 2 (%)	
Acephat	10	100	18	10	79	88	16	8	68	78	0.9902
Azoxystrobin	10	100	15	8	92	96	16	8	85	91	0.9879
Carbaryl	10	100	14	5	93	95	13	12	80	83	0.9906
Cymoxanil	10	100	6	4	87	98	12	9	92	87	0.9894
Dimetomorph (sum of isomers)	10	100	9	3	94	100	9	5	75	83	0.9855
Diniconazol	10	100	12	4	90	94	18	9	71	90	0.9872
Etrifos	10	100	10	3	78	87	15	10	69	78	0.9992
Oxamyl	10	100	8	5	89	93	19	11	89	96	0.9875
Pencycurone	10	100	13	7	90	85	9	4	74	82	0.991
Pyraclostrobin	10	100	12	8	86	104	8	5	78	82	0.9896
Spinosad A	10	100	22	12	91	95	10	5	78	88	0.9899
Zoxamide	10	100	13	7	87	95	9	4	71	76	0.9913



# vDIA - Repeatability – For 12 Representatives

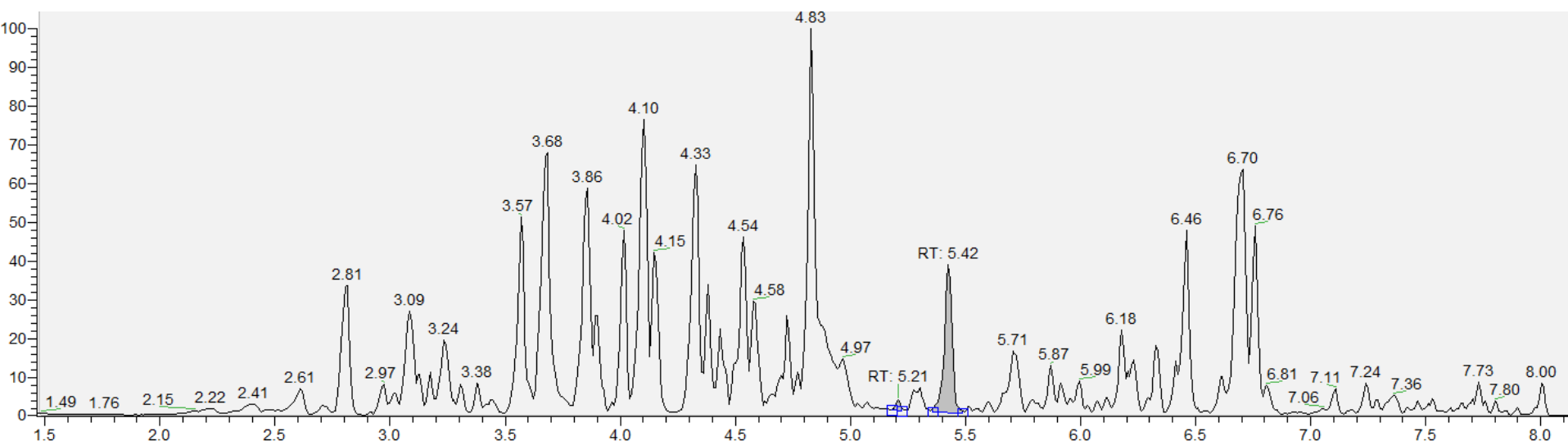


# Non-target Screening of Food Contaminants

Targeted analysis has its limits... its targeted

How do we detect all the other compounds in a sample?

Do we want/need to detect all compounds in a sample?



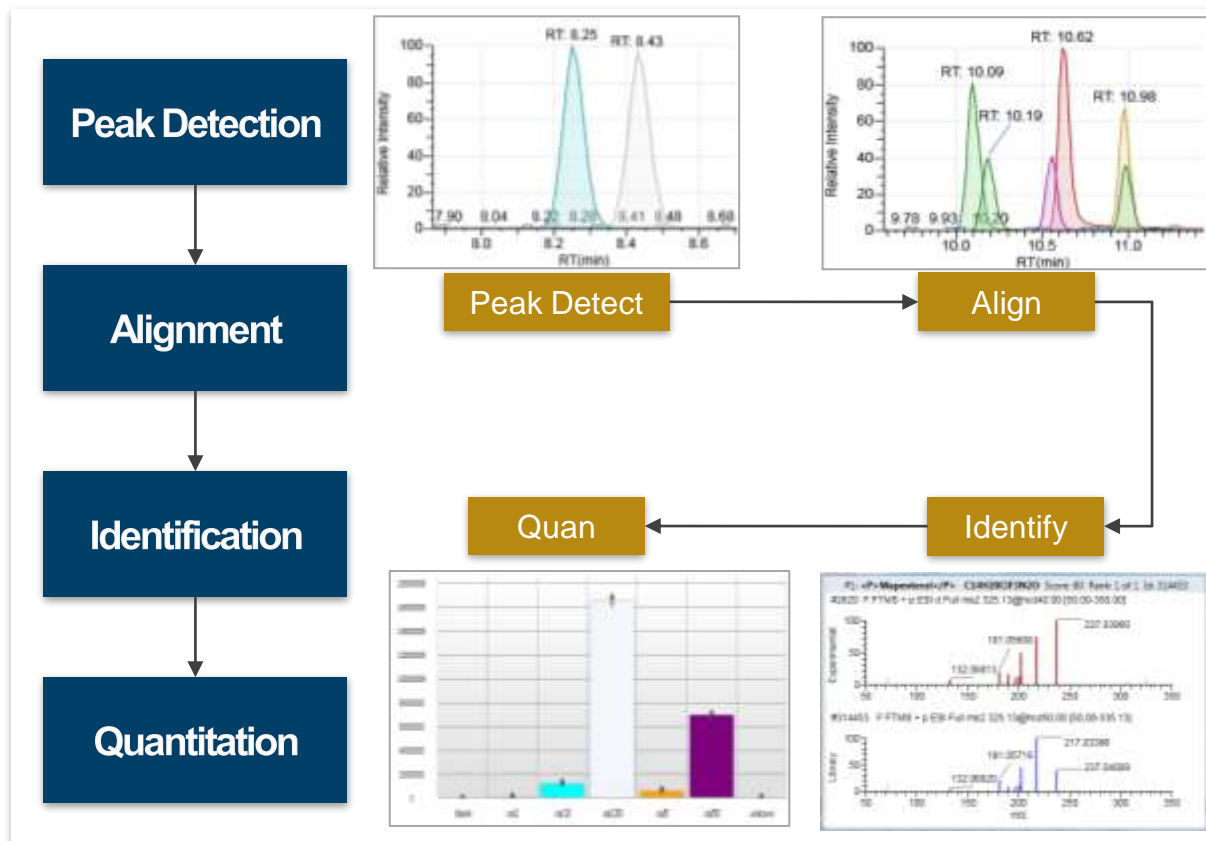


# Unknown screening workflow

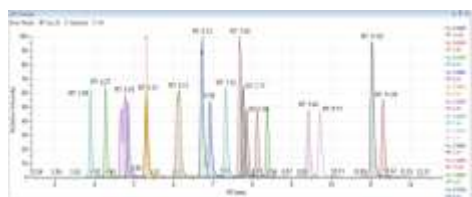
All-in-One solution

## Trace Finder

- Targeted Quan
- Targeted Screening
- Unknown Screening
- Targeted Analysis with Unknown Screening

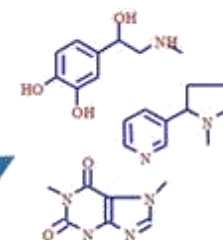


# Orbitrap-based Unknown Screening and ID Workflow

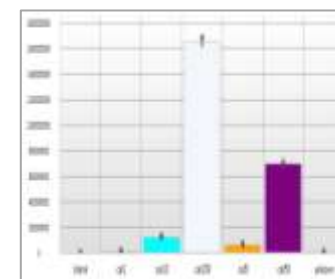
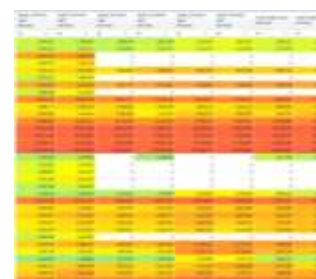


Aliphatic Chemicals Database (36)	Biological Properties (14)
Chemical Reactions (7)	Chemical Safety Data (7)
Class of Compounds in Development (16)	Cloning Agents (3)
Information Aggregators (33)	Control Policies via Mobile App
Lipidology/Protein Structure Database (7)	Metabolic Pathways (7)
Molecular Libraries Screening Center Network (2)	Natural Products (12)
MS Reference Spectra (5)	Physical Properties (including SARFAAS Database) (7)
Protein 1D Database (3)	Spectroscopy Database (13)
Substance Vendors (22)	Theoretical Properties (7)
Toxicology/Environmental Database (7)	Virtual Library (3)
Patents (3)	Physical Database (34)
Publications of Negative Advice (4)	Web-based Advice (only in commenting) (2)

Chemspider: 高针对性的食品安全、环境污染、药物毒物数据库



mzVault: 契合离线mzcloud数据库的高准确性风险物质鉴定谱库



差异未知风险物质定量、半定量统计分析手段

## Introducing The EFS HRAM MS/MS Spectral Libraries:

- OPTON-30386 EFS HRAM MS/MS Spectral Library 1.0

EFS	No. of Spectra
Pesticides	698
Veterinary Drugs	108
Emerging Contaminants	756
PFCs	21
Mycotoxins	44

# Food authentication challenges

- Chemically identical foods or identical chemical entities
- Unique marker compounds rarely found - more often small analytical differences (isotopic patterns)
- Large natural variability based on climatic conditions, fertilizers used, variety, processing.....
- Techniques must be able to distinguish small differences

- **Protected Designation of Origin (PDO)**

*covers agricultural products and foodstuffs which are produced, processed and prepared in a given geographical area using recognized know-how.*

- **Protected Geographical Indication (PGI)**

*covers agricultural products and foodstuffs closely linked to the geographical area. At least one of the stages of production, processing or preparation takes place in the area.*

- **Traditional Speciality Guaranteed (TSG)**

*highlights traditional character*

<http://ec.europa.eu/agriculture/quality/>



**Council Regulation (EC)  
No 510/2006 of 20 March 2006**



**Council Regulation (EC)  
No 509/2006 of 20 March 2006**

## Why Adulterate

- Mainly for financial gain
- Improve perceived quality attributes
- Brand / product substitution
- Reduce manufacturing cost

## May also be malicious

- Reputation damage
- Terrorism

### Top Adulterated Matrix:

Milk, Olive Oil, Honey, Orange Juice, Coffee, Seafood, Flour, Meat

## High Profile Food Fraud



### Olive Oil Authenticity

[Determination of Olive Oil Adulteration by Principal Component Analysis with HPLC-Charged Aerosol Detector](#)



### Orange Juice Adulteration

[Fruit Juice Adulteration Notebook](#)



### Melamine in Milk

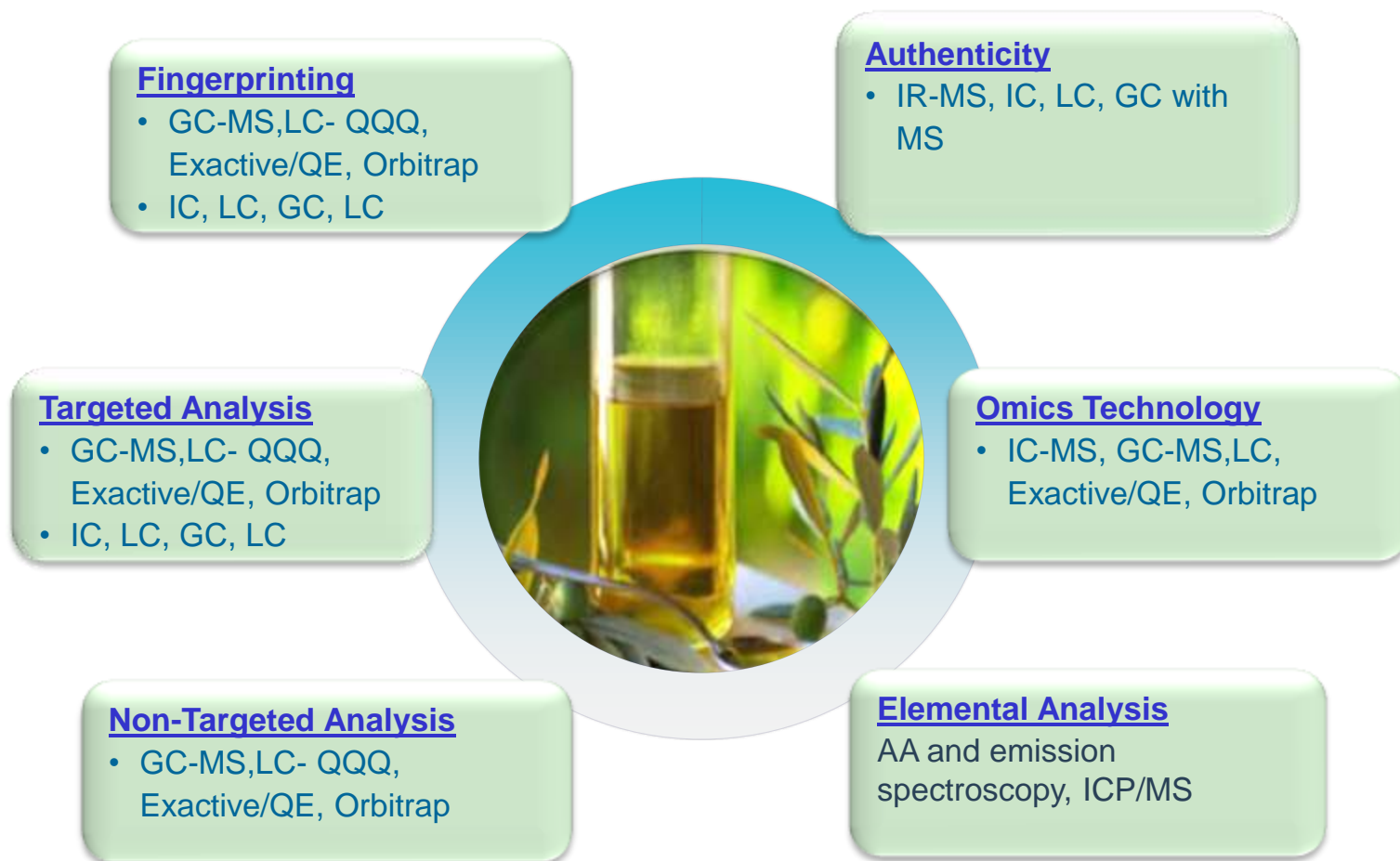
- [AN424: Simple and Rapid Screening of Melamine in Milk Products with LC-MS](#)
- [AN502: Determination of Melamine in Powdered Milk by LC-MS/MS Using a Core Enhanced Technology Solid Core HPLC Column -](#)



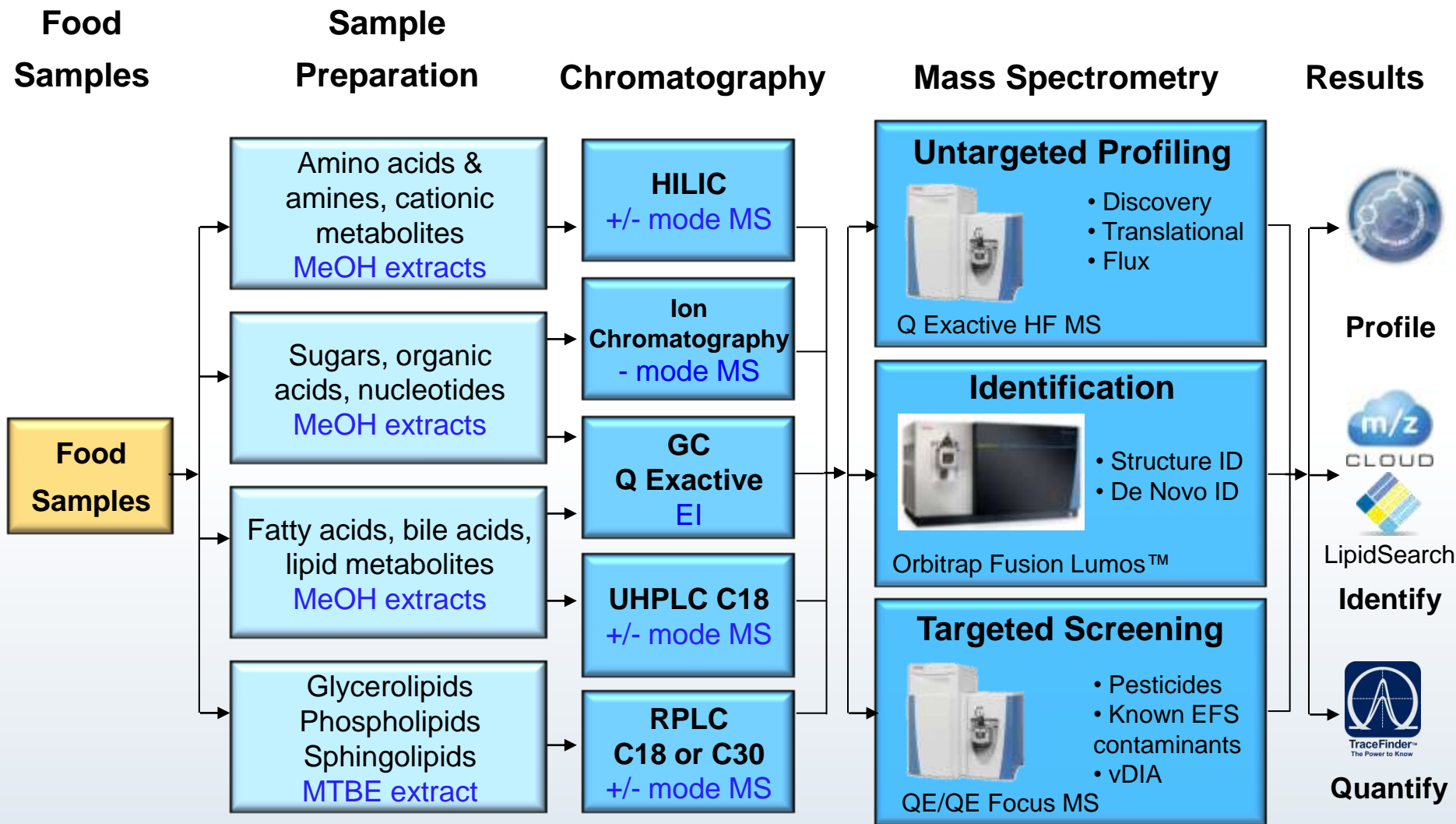
### Honey

[Isotope Ratio Mass Spectrometry: Authenticity Control, Fraud And Forensics In Food -](#)

# Food Fraud Analysis Solutions



# Comprehensive Workflow for Foodomics Analysis



# Hot Topics Food Fraud

- [Olive oil](#)
- Wine adulteration
- [Halal food testing](#)
- General untargeted screening for adulterants
- Country of origin (COO)
  - [Isotope Ratio Mass Spectrometry: Authenticity Control, Fraud And Forensics In Food](#)
- Honey
  - [AN30177:Detection of Honey Adulteration with FlashEA Elemental Analyzer and DELTA V Isotope Ratio Mass Spectrometer](#)





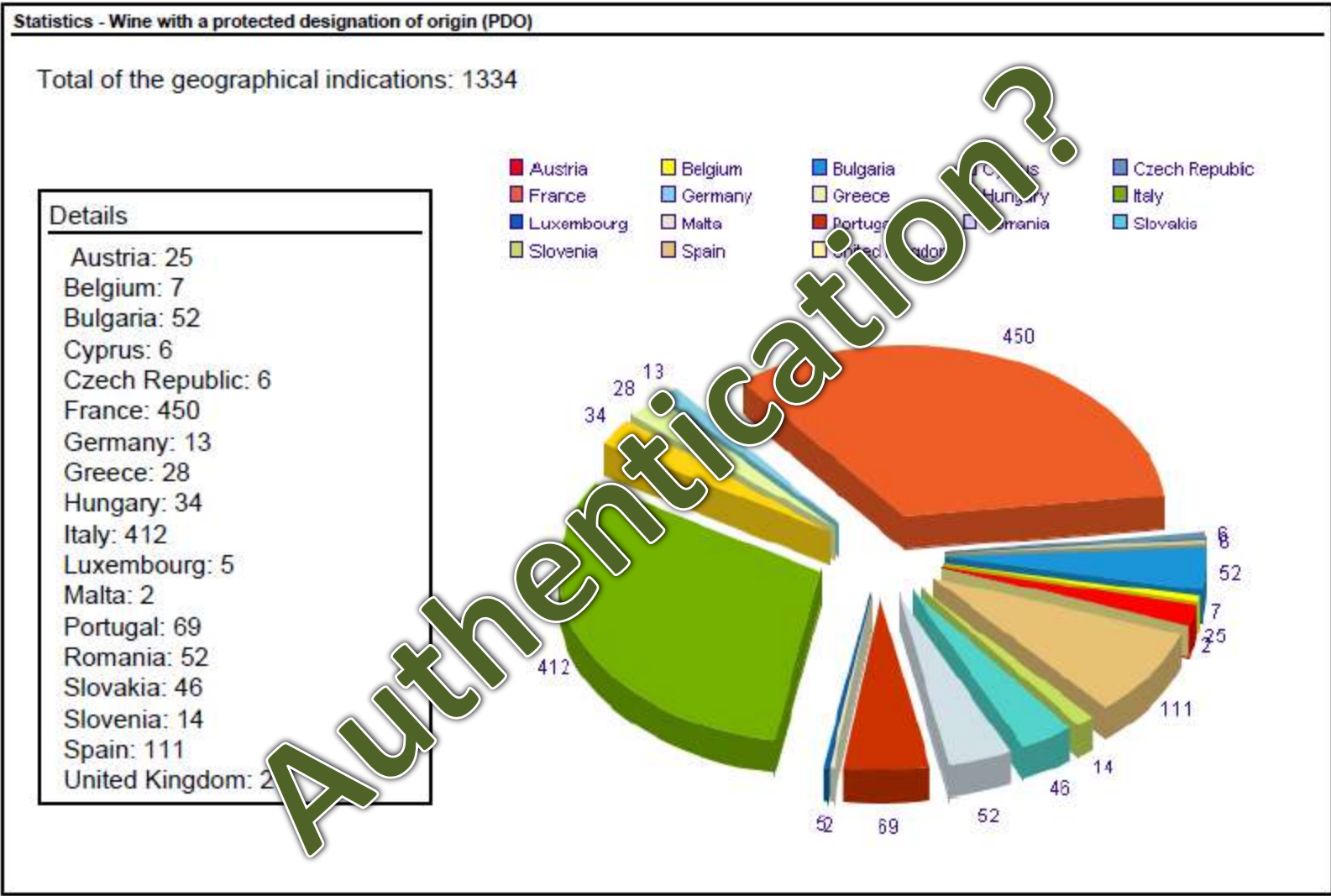
# UHPLC-HRMS UNTARGETED METABOLOMICS APPROACH APPLIED TO THE AUTHENTICATION OF ALCOHOLIC BEVERAGES

C. Martins; A. Checa; H. Gallart-Ayala; O.  
Núñez; X. Saurina; S. Hernández-Cassou

# INTRODUCTION

E-Bacchus Database: <http://ec.europa.eu/agriculture/markets/wine/e-bacchus/>  
As of May 2012

## PROTECTED DESIGNATION OF ORIGIN – WINE



# INTRODUCTION

## Analytical methodologies in food authentication

- LC (UV, FLD)

- LC-MS

- NMR

- CE-MS

METABOLOMIC

- ELECTRONIC NOSE

- AAS/ICP-MS (Mineral profile)

- IR

- UV-spectra

- Stable isotope analysis

- IMS....

# INTRODUCTION

## METABOLOMIC

Comprehensive analysis of low molecular weight compounds (< 1000 Da) in a biological sample

## TARGETED METABOLOMIC

Measurement of defined groups of chemically characterized metabolites

**Advantages: Sensitivity**

**Disadvantages: Discriminant compounds can remain undetected**

## UNTARGETED METABOLOMIC

Comprehensive analysis of all the measurable analytes in a sample including chemical unknowns

**Advantages: Wider spectra of compounds detectable**

**Disadvantages: Sensitivity**

# EXPERIMENTAL CONDITIONS

## Sample Treatment

Beer and wine samples were diluted with water 1:1

## Liquid Chromatography

Column: Hypersil Gold aQ (100 mm x 2.1 mm ID, 1.9  $\mu\text{m}$ )

Mobile Phase: ACN:0.1% Formic Acid

Flow: 600  $\mu\text{L min}^{-1}$

## Mass Spectrometry

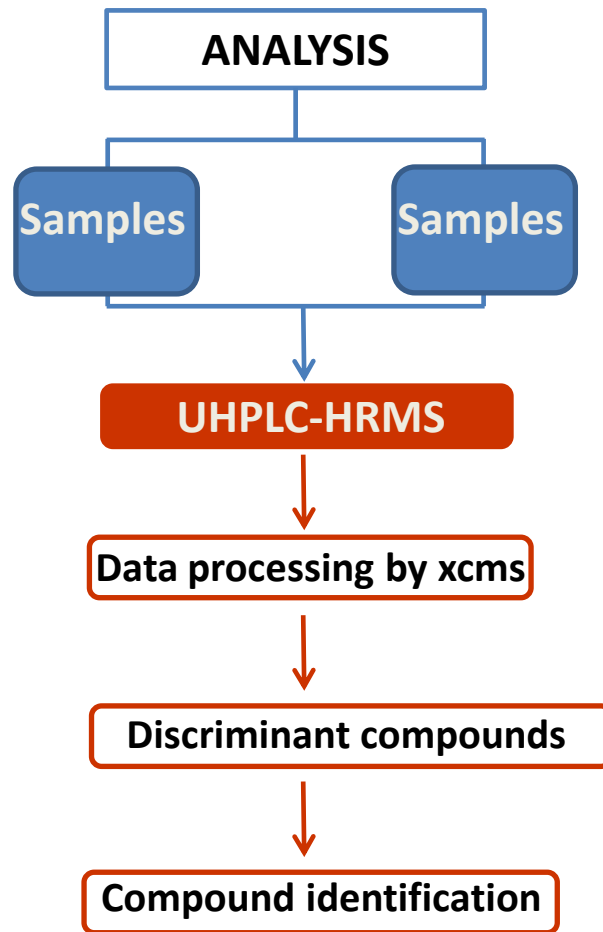
LITQ-Orbitrap Velos

ESI (+) and (-)

Mass Resolving Power: 60,000 FWHM

Acquisition mode: Scan event 1: Full scan

Scan event 2: MS/MS using CID





## **CASE STUDY**

# **Spanish wines classification**



# Spanish wine classification

## Wine composition

→ Water, Alcohols, Sugars, Polysaccharides, Aminoacids, Biogenic amines, Organic acids, Phenolic compounds, Metals

## Factors that affect wine composition

→ Climate

→ Grape growing area

→ Grape variety

→ Winemaking process



Can we  
differentiate  
between PDO?



**2.a**

**Two class comparison**

**Somontano vs Rioja wines**





# POTENTIAL DESCRIPTORS

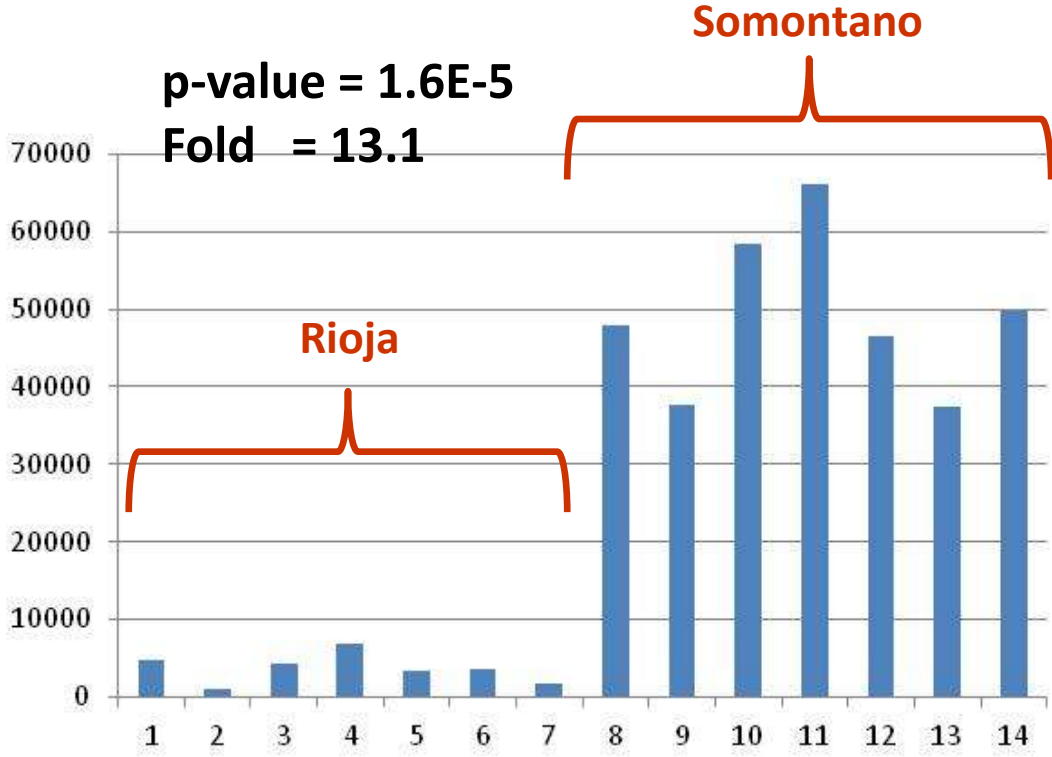
ESI (-)

14 PEAKS  
DETECTED  
WITH XCMS



6  
POTENTIAL DISCRIMINANT  
COMPOUNDS

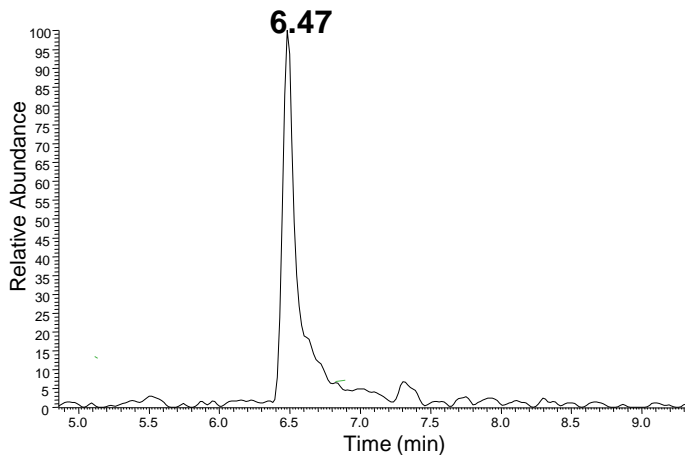
Name	m/z value	Retention time (sec)	Elemental composition	RDB value	Mass error (ppm)
M301T334	301.0343	334	C15H9O7	11.5	-3.71
M151T334	151.0030	334	C7H3O4	6.5	-4.12



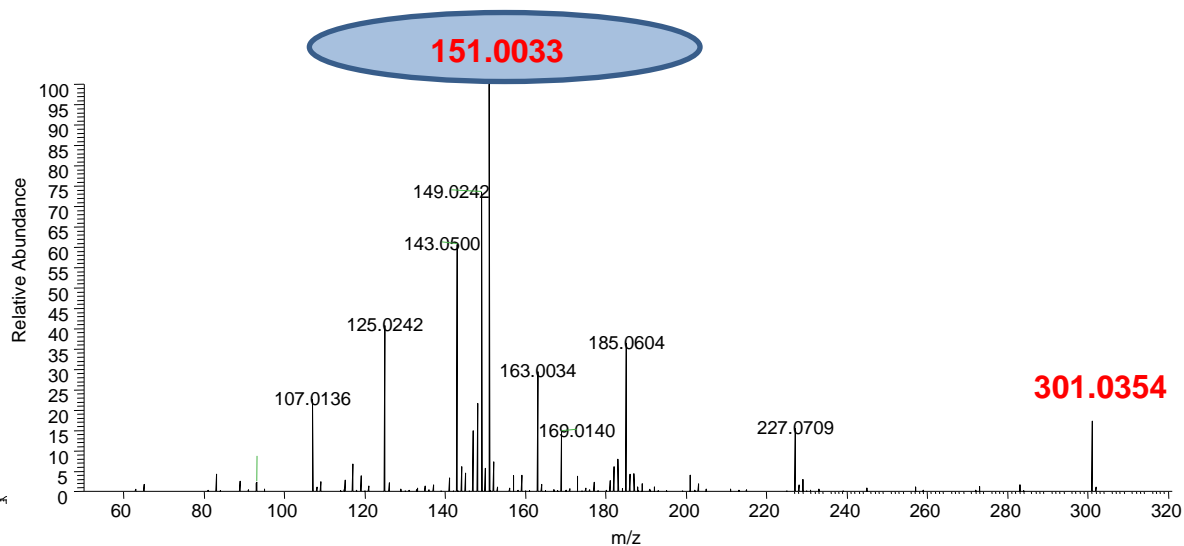
# POTENTIAL DESCRIPTORS

ESI (-)

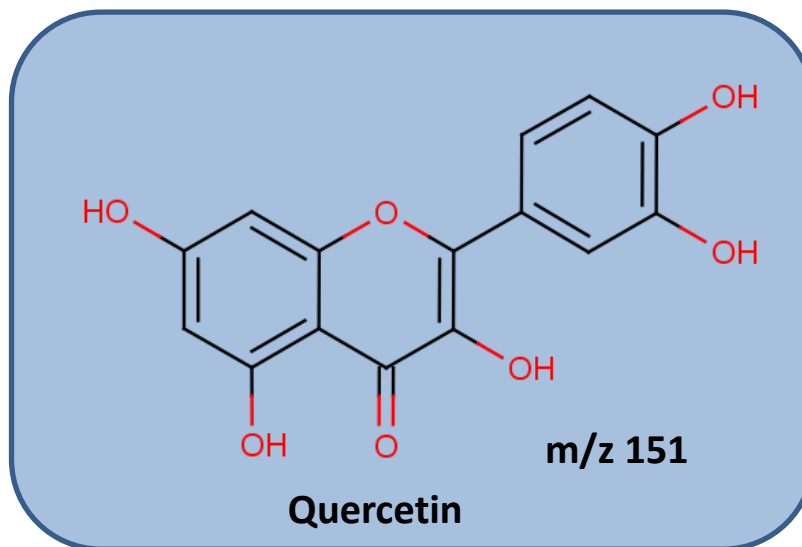
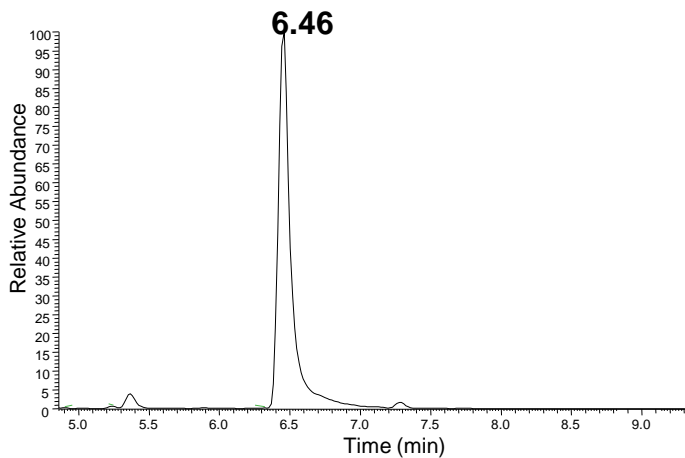
## Somontano wine



## All Ion Fragmentation MS/MS



## Quercetin standard





**2.b**

**Multi-class comparison**

**Penedes vs Ribera vs Rioja wines**

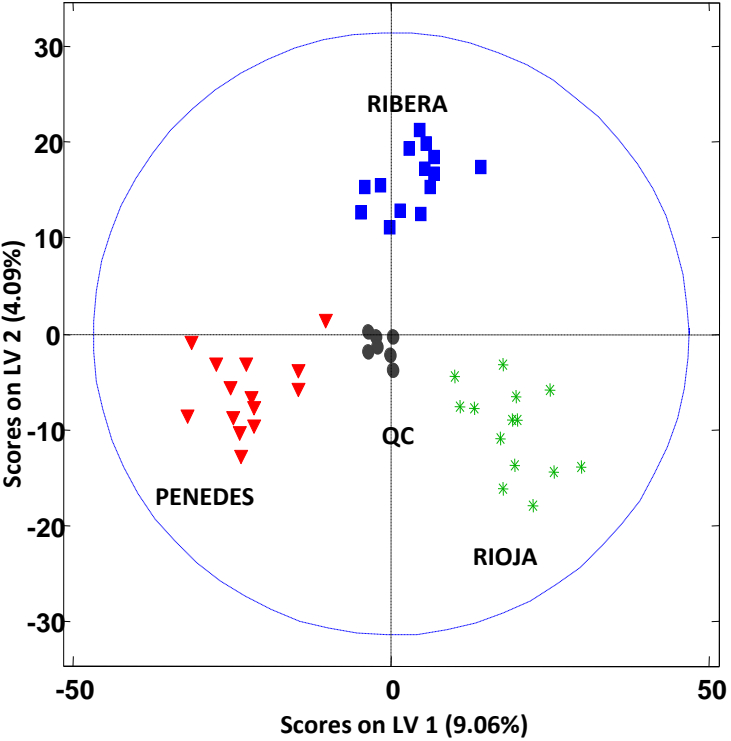


# Spanish wines classification



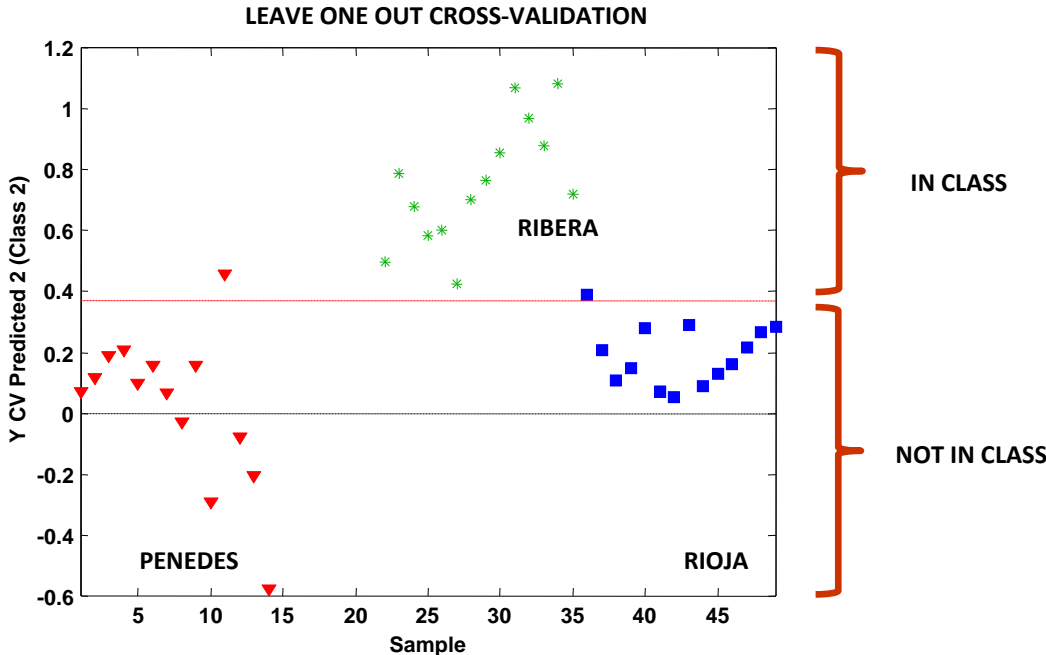
**Positive mode**

**PLS-DA**



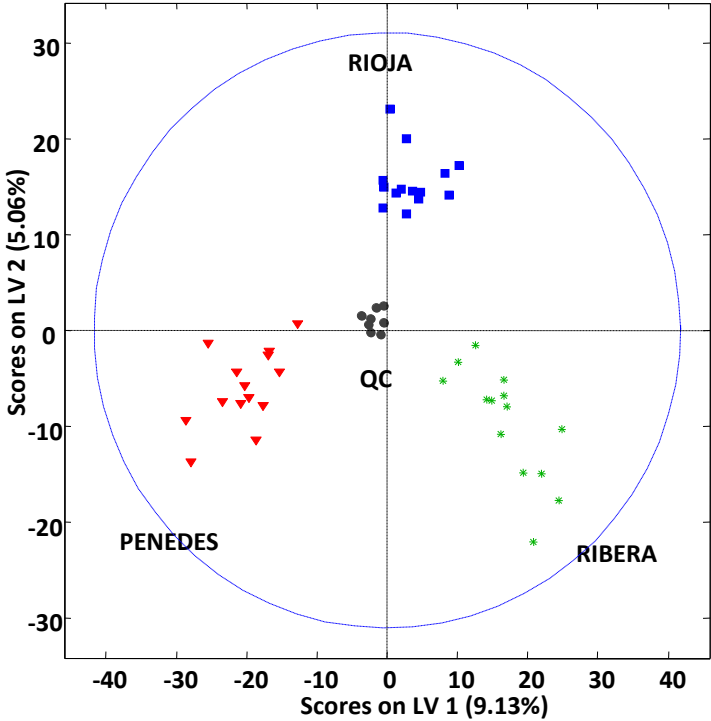
**3 Latent Variables**

	PENEDES	RIBERA	RIOJA
$R^2$	0.95	0.91	0.94
$Q^2$	0.62	0.63	0.54
FP (CV)	0	2	0
FN (CV)	1	0	1

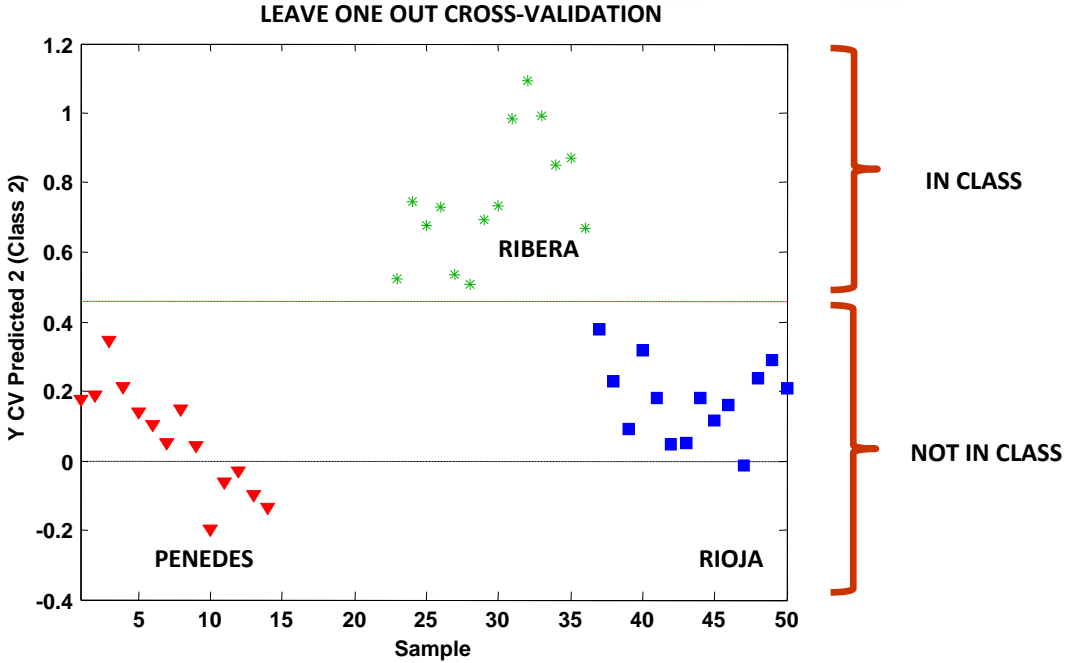


# Spanish wines classification

Negative mode    3 Latent Variables

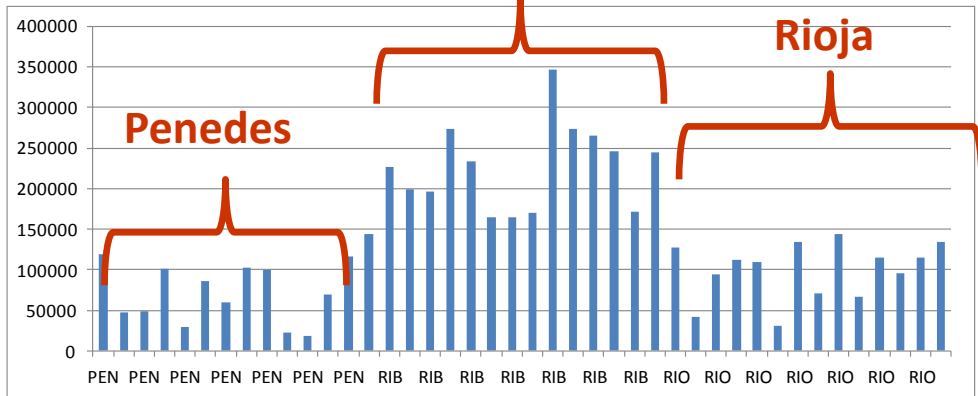


	PENEDES	RIBERA	RIOJA
$R^2$	0.93	0.92	0.93
$Q^2$	0.86	0.78	0.74
Specificity(CV)	1.00	1.00	1.00
Sensitivity (CV)	1.00	1.00	1.00

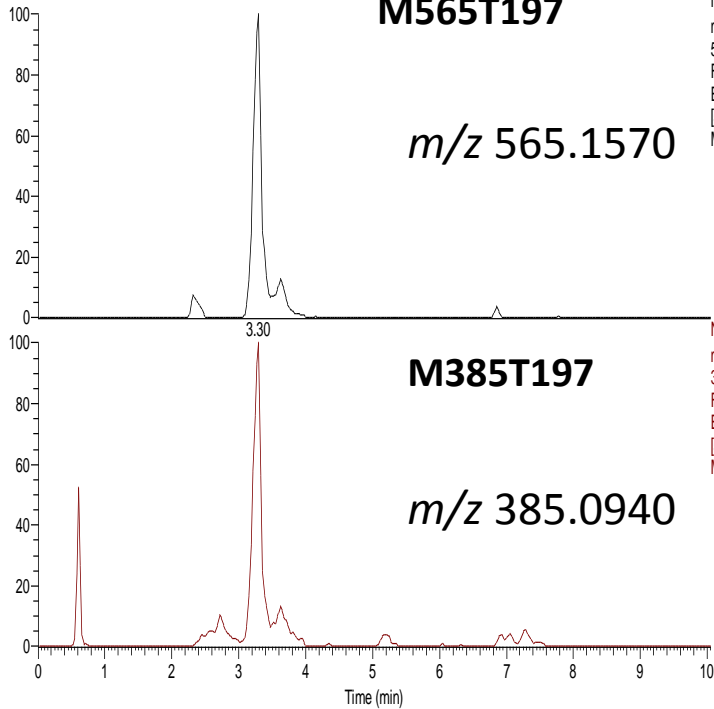


# Spanish wines classification

M565T197



RT: 0.00-10.05 SM: 5G



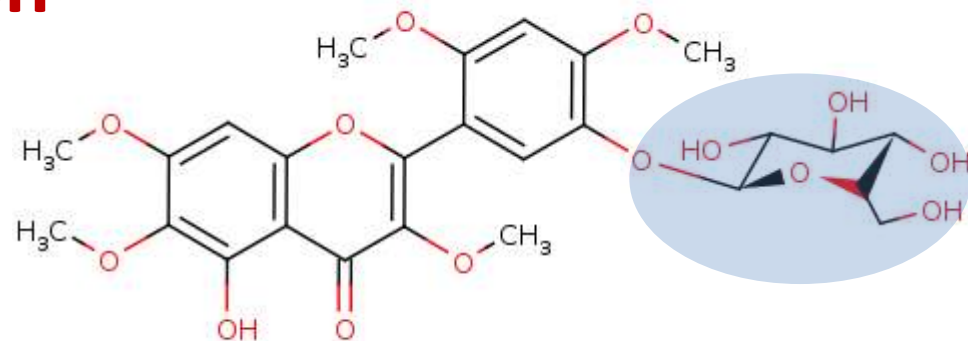
NL: 9.95E4  
m/z=  
565.0644-565.2319  
F: FTMS (1,2) -p  
ESI Full ms  
[50.00-1000.00]  
MS Vi\_24

NL: 2.26E4  
m/z=  
385.0823-385.1055  
F: FTMS (1,2) -p  
ESI Full ms  
[50.00-1000.00]  
MS Vi\_24

# Spanish wines classification

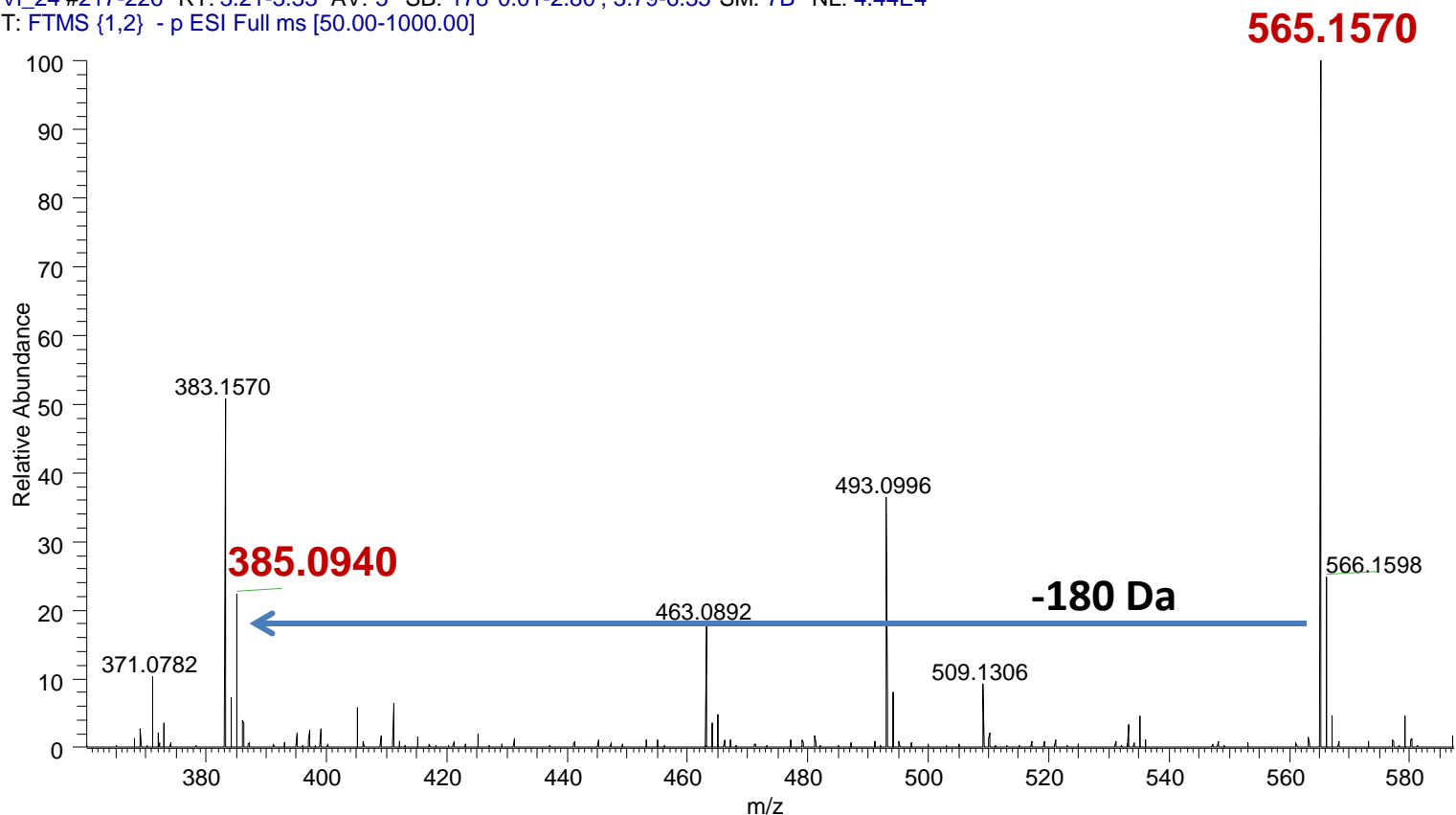
## Compound identification

### Database Search

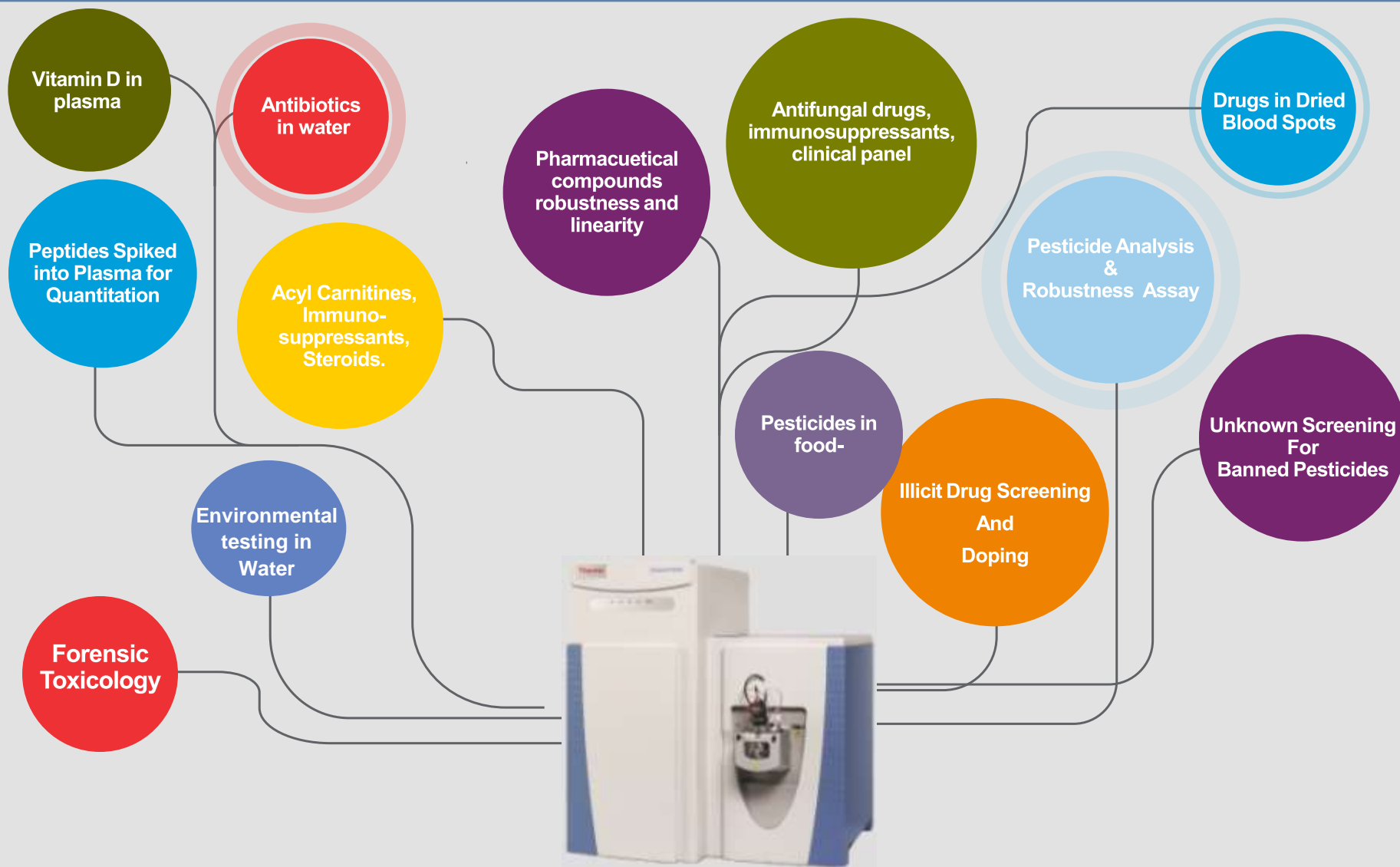


**5,5'-Dihydroxy-3,6,7,2',4'-  
pentamethoxyflavone 5'-glucoside**

Vi\_24 #217-226 RT: 3.21-3.33 AV: 5 SB: 178 0.01-2.80 , 3.79-6.35 SM: 7B NL: 4.44E4  
T: FTMS {1,2} - p ESI Full ms [50.00-1000.00]



# Q Exactive Focus Applications Universe





The background is a dark blue gradient with vertical light blue streaks. A bright white horizontal lens flare is on the left, and a white wave graphic with small square markers is in the center. The text 'Thermo' is in a large, bold, white sans-serif font, and 'SCIENTIFIC' is in a smaller, white, spaced-out sans-serif font below it.

**Thermo**  
SCIENTIFIC

**Transform Your Science**