



Routine Pesticide Residue Analysis Using Thermo Scientific High Resolution MS Technology

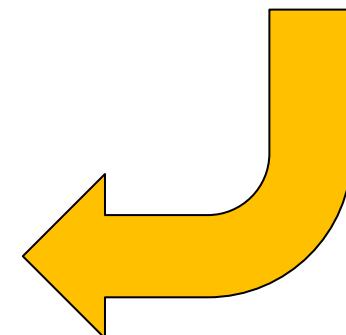
Laszlo Hollosi, Michal Godula, Maciej Bromirski
January, 2016

Part#: PP64679 – EN0216S

The world leader in serving science

Challenges of Pesticide-Residues Analysis

- Sample variability (matrix)
- Different compound characteristics
- Large number of samples
- Hundreds of analytes monitored
- Low levels controlled
 - Baby food
(MRL for all pesticides = 0.01 mg/kg)
- Fast response required



Former Pesticide Multi-Residue Method Setup



■ Extraction

Acetonitrile, Ethyl acetate, Methanol...



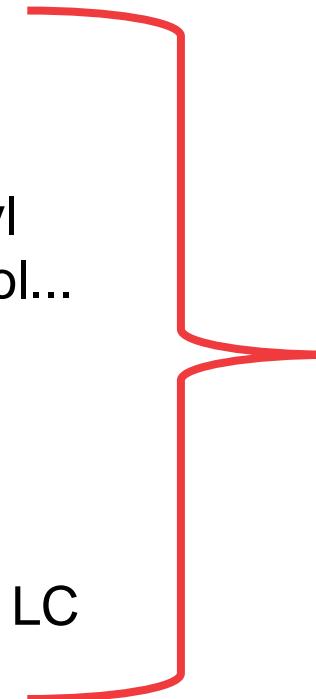
■ Clean-up

GPC, SPE, LLE, LC



■ Determination

GC, LC, GC-MS, LC-MS, GC-MS/MS, LC-MS/MS...



Mostly replaced by **QuEChERS** today



Thermo Scientific™ QuEChERS™
method

Simplified Extraction Procedure Applied



10 g of sample is weighed into Quechers extraction tube

+ 20 mL of water

+ 10 mL of ACN

shaking 10 min

Centrifugation 5 min @ 5000 rpm

Injection to LC-HRAM

Consumables Used

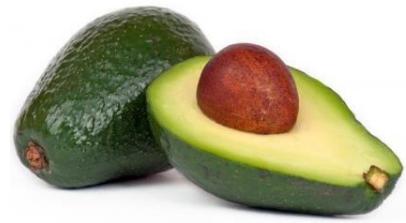
Consumables/Chemicals	Part Number
Acetonitrile	A/0638/17
QuEChERS extraction tube, 50 mL, 250 pack	60105-216
QuEChERS pouches, 50 pack	60105-344
Apparatus/Columns	Part Number
Horizontal shaker	1069-3391
Horizontal shaker plate	1053-0102
Thermo Scientific™ Barnstead™ EASYpure™ II water	3125753
Thermo Scientific™ Heraeus™ Fresco™ 17 micro centrifuge	3208590
Thermo Scientific™ Accucore™ aQ column 100x2.1, 2.6 µm	17326-102130



www.thermoscientific.com/chromatography

Improving QuEChERS extraction tips & tricks:

- **Dry food (cereals/dried food, < 25 % water content):**
 - Addition of water to enable adequate partitioning and reducing interaction of pesticides with matrix
- **Food containing fat/wax (avocado/oil):**
 - After extraction step add a freezing out step and transfer supernatant to clean-up tube
 - More clean-up might be needed of raw extract (PSA+C18)
- **Food containing complex matrix (tea/spices)**
 - Additional clean-up with GCB might be necessary (potential loss of planar structure pesticides like thiabendazole)
- **Acidic food (citrus):**
 - Adjust pH (5-5.5) to increase recovery (e.g. citrate buffering salts in QuEChERS extraction tube) and reduce coextraction of matrix interferences (Note: acid labile compounds require higher pH 8)



Improving QuEChERS Clean-up Tips & Tricks:

- **QuEChERS clean-up tube additives:**

Product Selection

Matrix Type	Examples	Sorbent Requirements
General Matrices	Apples Cucumber Melon	Magnesium Sulfate, PSA
Fatty Matrices	Milk Cereals Fish	Magnesium Sulfate, PSA, C18
Pigmented Matrices	Lettuce Carrot Wine	Magnesium Sulfate, PSA, C18, GCB
High Pigmented Matrices	Spinach Red Peppers	Magnesium Sulfate, PSA, C18, GCB

[Click here for more information](#)

How To Use QuEChERS ?

The screenshot shows a Thermo Fisher website section titled "Productivity simplified". On the left, there's a call-to-action for speaking with an expert, a sidebar with navigation links, and a large blue button labeled "Link to video". On the right, there's a search and filter interface, a video player showing a pesticide analysis tip, and a navigation bar.

Productivity simplified

Video Library
Pesticide Analysis

Search: Search

Filter by: -- All Techniques -- -- All Products --

previous 1 next

Sample Preparation Tips to Improve Your Pesticide Anal... 1 next

Sample Preparation Tips To Improve Your Pesticide Analysis
Presented by Ebru Sarikaya

00:00 06:26

• [Link to video](#)

- ▶ View All
- ▶ Compound Detection
- ▶ Compound Separation
- ▶ Data Processing
- ▶ Sample Preparation

Pesticide Analysis by HRMS



Start-to-finish
workflows for pesticide analysis

Thermo
SCIENTIFIC

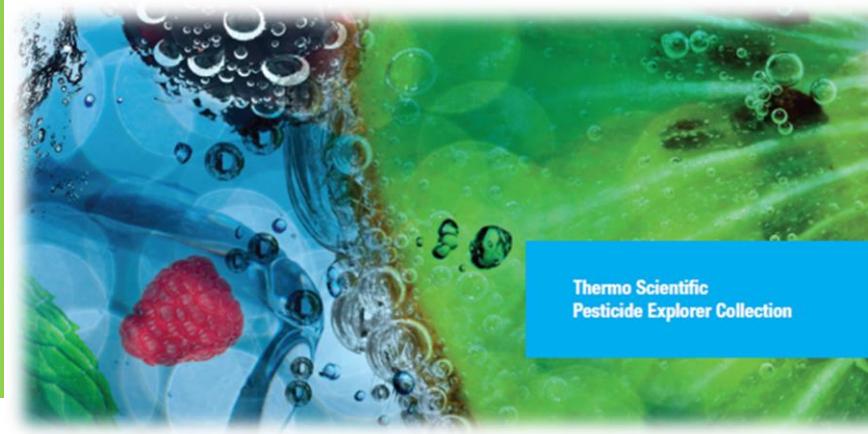
Configuration and complete start-up kit provides everything needed to perform robust, high resolution routine workflows for rapid screening and quantitation of pesticides, from the QuEChERS sample extraction kit to proven multi-class pesticide residue analysis methods.



Thermo Scientific™ Q Exactive™ Focus MS

Quantitation Package

- Thermo Scientific™ TraceFinder™ 3.2 SP2 software
- EFS HRAM MS/MS Spectral Library 2.0
- USB (Methods and User guide)



Thermo Scientific
Pesticide Explorer Collection

- Pesticide Explorer Collection comes with the software installation instructions as you open the box. It is an easy to follow guide that walks you through the process from installation to activation of the software.
- Pesticide Explorer Collection also comes with an usb drive which includes a Method Installation guide as well as methods and libraries. This document outlines the necessary steps that will be needed to install the Pesticide methods and it's library within TraceFinder software as well as using TraceFinder software.

Total Pesticide Solution Workflow

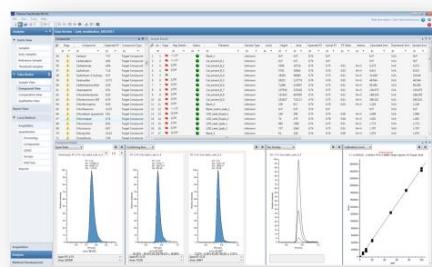


QuEChERS sample preparation kits



Thermo Scientific™ Q Exactive™
Hybrid Quadrupole-Orbitrap Mass
Spectrometer

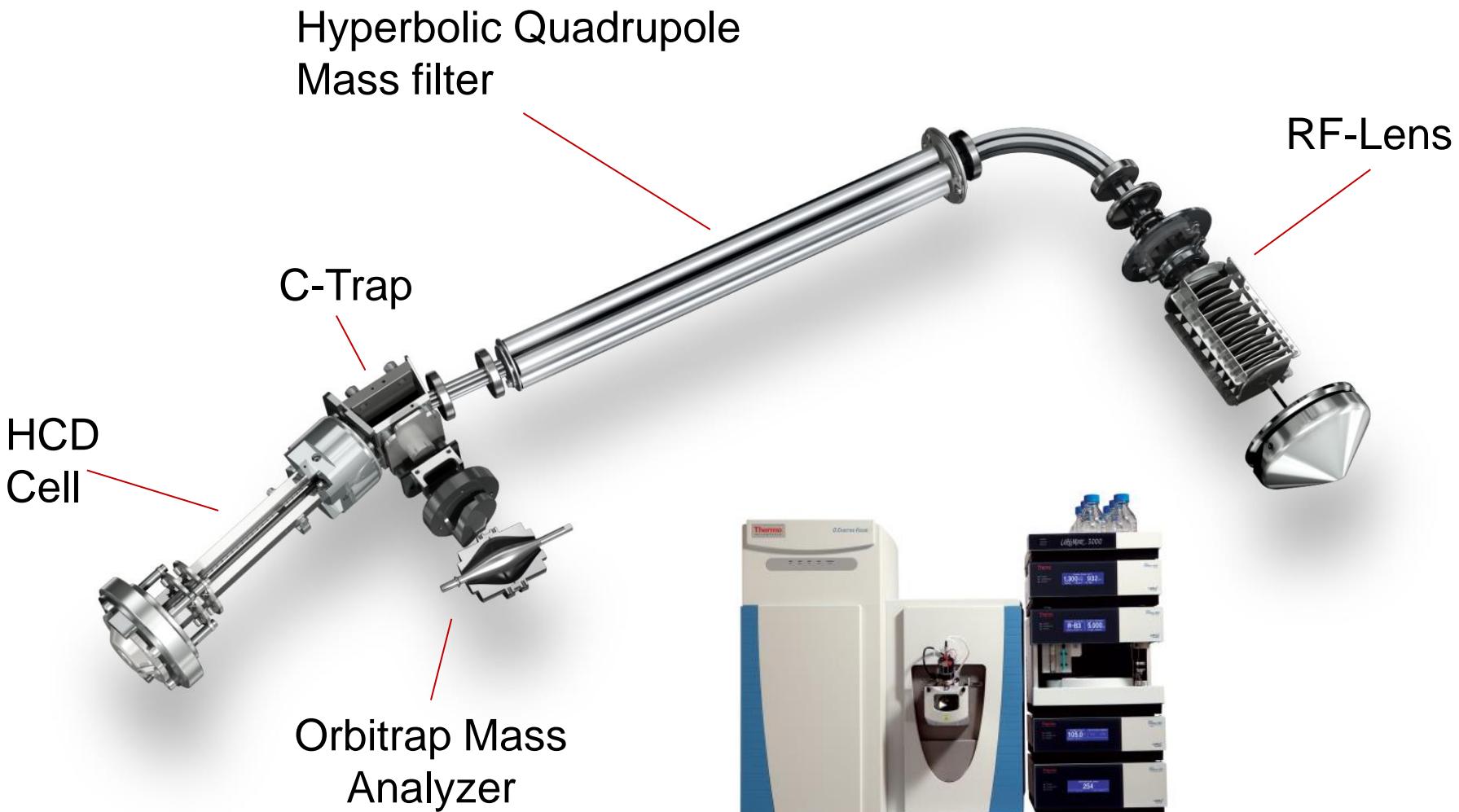
On-site method
by experienced
Special Solutions
Center chemists



TraceFinder Data Processing



Q Exactive™ Focus - What is New?



3 ways of Quantitation/Screening 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

Experiments

General
Full MS
SIM
PRM
Full MS - AIF
Full MS - vDIA



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

Experiments

General
Full MS
SIM
PRM
Full MS - AIF
Full MS - vDIA



PRM (Parallel Reaction Monitoring)

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

Experiments

General
Full MS
SIM
PRM
Full MS - AIF
Full MS - vDIA



Q Exactive Focus Scan Methods

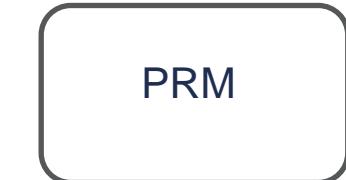
Selectivity



Sensitivity



Information



LC Instrumental Method

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 µl**

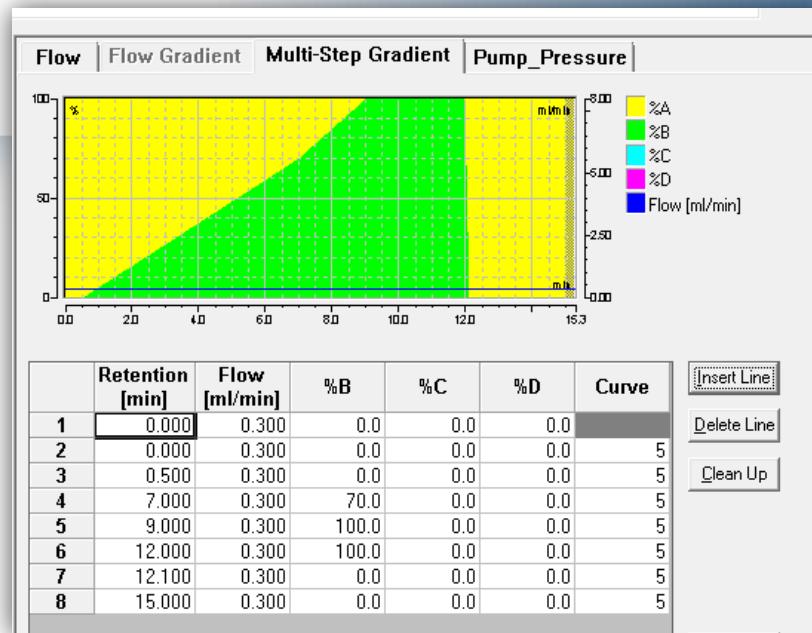
- Column: Accucore aQ column 100 mm x 2.1 mm x 2.6 µm

- Column temperature: 25°C

- Flow rate: 300 µl/min

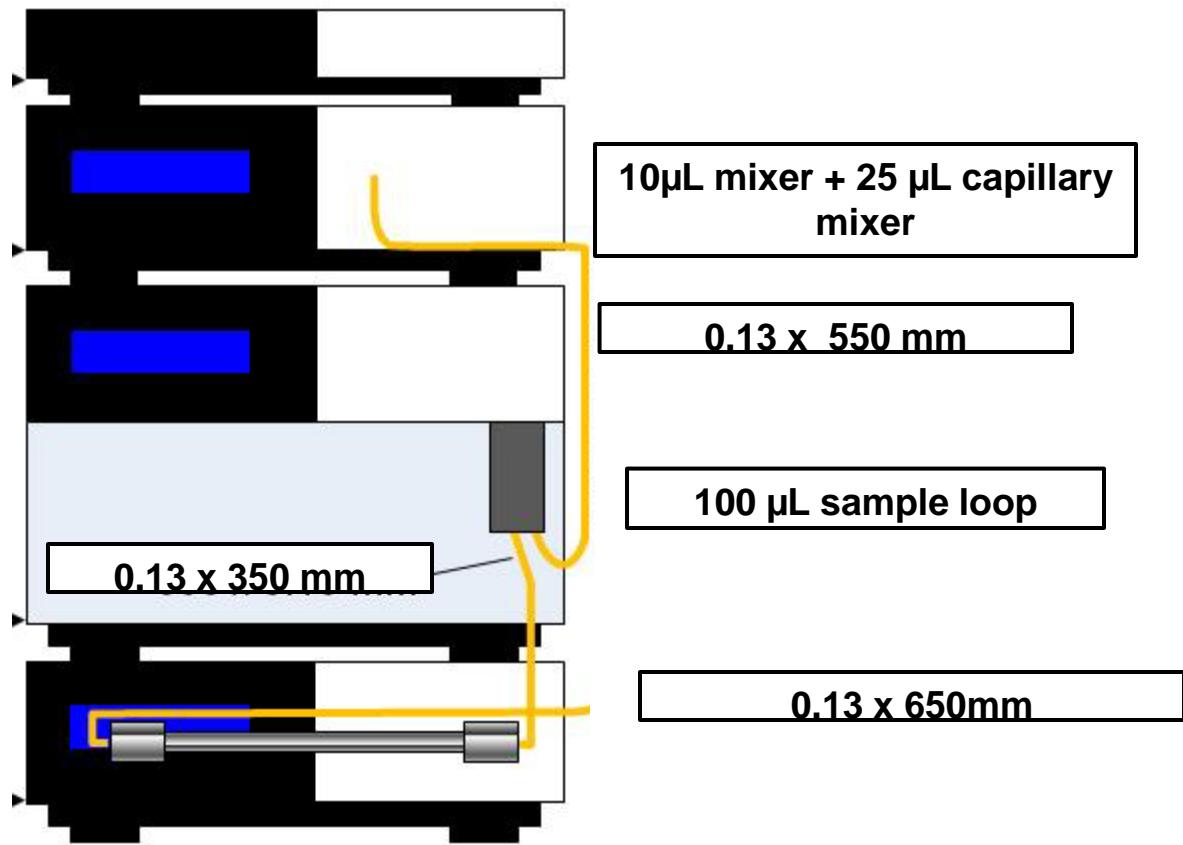
- Run time: 15 min

- Gradient:



Thermo Scientific™ UltiMate™
XRS LC

Recommended Configuration of the LC System



Thermo Scientific™ UltiMate™
XRS Liquid Chromatography
System

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		actual
Sheath gas flow rate	40	40
Aux gas flow rate	10	10
Sweep gas flow rate	2	2
Spray voltage (kV)	3.50	3.47
Spray current (μA)		0.10
Capillary temp. (°C)	250	250
S-lens RF level	55.0	
Aux gas heater temp (°C)	270	272

Source Auto-Defaults...

Apply

Help

Hot link

Recommended Method Parameters for Symmetric vDIA

Properties

Properties of the method

Global Settings

User Role Advanced

Use lock masses off

Lock mass injecti —

Chrom. peak wid 12 s

Time

Method duration 15.00 min

Customized Tolerances (+/-)

Lock Masses —

Inclusion —

Exclusion —

Dynamic Exclusio —

Properties of Full MS - vDIA

General

Polarity positive

Full MS

Resolution 70,000

Scan range 120 to 1000 m/z

variable DIA

Resolution 17,500

vDIA segments 8

vDIA isolation rai 50 to 150 m/z

vDIA isolation rai 140 to 240 m/z

vDIA isolation rai 230 to 330 m/z

vDIA isolation rai 320 to 420 m/z

vDIA isolation rai 410 to 510 m/z

vDIA isolation rai 500 to 600 m/z

vDIA isolation rai 590 to 690 m/z

vDIA isolation rai 680 to 780 m/z

CE / stepped CE 30, 50, 70

Fixed first mass —

AGC target 5e5

Spectrum data ty Centroid

Alternative Recommended MS Method Settings

The figure consists of three side-by-side screenshots of the Thermo Fisher MassLynx software interface, each showing the 'Properties of the method' window for a different mass spectrometry mode. A large orange arrow points downwards from each screenshot towards a small blue box labeled 'Full MS'.

Left Screenshot (vDIA mode):

- Global Settings:** User Role: Advanced; Use lock masses: off; Lock mass injecti: –; Chrom. peak wid: 12 s.
- Time:** Method duration: 17.00 min.
- Customized Tolerances (+/-):** (checkboxes)
- 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 type: Profile.
 - variable DIA:** Resolution: 17,500; # vDIA segment: 5; vDIA isolation rai: 120 to 205 m/z, 195 to 305 m/z, 295 to 405 m/z, 395 to 505 m/z, 495 to 1000 m/z; CE / stepped CE: 30, 50, 70; Fixed first mass: –; AGC target: 5e5; Maximum IT: auto; Microscans: 1; Spectrum data type: Centroid.

Middle Screenshot (Full MS mode):

- Global Settings:** User Role: Advanced; Use lock masses: off; Lock mass injecti: –; Chrom. peak wid: 12 s.
- Time:** Method duration: 17.00 min.
- Customized Tolerances (+/-):** (checkboxes)
- Properties of Full MS:**
 - General:** Polarity: positive; dd-MS²: 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 type: Profile.
 - dd-MS² Confirmation:** Resolution: 17,500; Isolation window: 3.0 m/z; Isolation offset: –; CE / stepped CE: 15, 30, 50; Fixed first mass: –; Default charge st: 1; AGC target: 5e4; Maximum IT: auto; Loop count: 1; Underfill ratio: 1.0 %; Intensity thresho: auto; Apex trigger: –; Dynamic exclusi: auto; Spectrum data type: Centroid.

Right Screenshot (PRM mode):

- Global Settings:** User Role: Advanced; Use lock masses: off; Lock mass injecti: –; Chrom. peak wid: 12 s.
- Time:** Method duration: 17.00 min.
- Customized Tolerances (+/-):** (checkboxes)
- Properties of PRM:**
 - General:** In-source CID: –.
 - Targeted-MS²:** Resolution: 70,000; Isolation window: 3.0 m/z; Isolation offset: –; CE / stepped CE: 15, 30, 50; Fixed first mass: –; Default charge st: 1; AGC target: 1e5; Maximum IT: auto; Microscans: 1; Spectrum data type: Profile.

*

Statistically identical results with symmetrical and asymmetrical vDIA settings

Targeted 330 Compounds – vDIA Screening Method

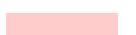
Acephate	Bromacil	Cumyluron	Dinotefuran	Fenthion-sulfone	Heptenophos	Methabenzthiazuron	Penconazole	Pyridaben	Thiacloprid
Acetamiprid	Bromoconazole	Cyanazine	Dioxacarb H	Fenthion-sulfoxide	Hexaconazole	Methamidophos	Pencycuron	Pyridate e	Thiamethoxam
Acibenzolar-S-methyl	Bupirimimate	Cyazofamid	Disulfoton	Fenuron	Hexaflumuron	Methidathion	Permethrin	Pyrimethanil	Thiazopyr H
Aclonifen	Buprofezin T	Cycloate	Dithiopyr	Fipronil	Hexazinone	Methiocarb	Phenmedipharm	Pyroquilon	Thidiazuron T
Alachlor	Butachlor	Cycluron	Diuron	Flazasulfuron	Hexythiazox T	Methiocarb sulfoxide	Phenthionate	Pyroxulam	Thiobencarb
Alanyncarb	Butafenacil	Cyflufenamid	Dodemorph	Flonicamid T	Imazalil	Methiocarb-sulfone	Phoxim	Quinoxifen	Thiodicarb
Aldicarb	Butocarboxim	Cymoxanil	Epoxiconazole	Florasulam	Imazaquin	Metholcarb	Picoxystrobin	Quinalofop T	Thifanox
Aldicarb sulfone	Butoxycarboxim	Cypermethrin T	Esprocarb	Fluazifop	Imazethapyr	Methomyl	Piperonyl butoxide	Quinalofop-p-ethyl	Thionazin
Aldicarb sulfoxide	Carbaryl	Cyproconazole	Etaconazole	Flufenacet	Imidacloprid	Methoprottryne	Piperophos	Resmethylrin	Tolfenpyrad
Allethrin	Carbendazim	Cyprodinil	Ethiofencarb	Flufenoxuron	Indoxacarb	Methoxyfenozide	Pirimicarb	Rimsulfuron	Tralkoxydim
Ametryn	Carbetamide	Cyromazine	Ethiofencarb_sulfoxide	Flumetsulam	Iprovalicarb	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	Isopenphos	Metosulam	Pretilachlor	Sethoxydim	Triazophos
Anilofos	Carbosulfan	Desmedipharm	Ethirimol	Fluopicolide	Isoprocarb	Metoxuron	Primsulfuron-methyl	Simeconazole T	Trichlorfon
Aramite H	Carboxin	Desmethyl-pirimicarb	Ethofumesate	Fluopyram	Isoprothiolane	Metrafenone	Prochloraz	Simetryn	Tricyclazole
Atrazine	Carfentrazone-ethyl	Desmetryn	Ethoxyquin	Fluoxastrobin	Isoproturon	Metribuzin	Profenos	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	Trifloxytrobin
Azinphos-ethyl	Chlорbromuron	Diclobutrazol	Etrimfos	Fluoroypyrr	Isoxaflutole	Mexacarbate	Prometryn	Sulfotep	Triflumizole
Azinphos-methyl	Chlorfenvinphos	Dicrotophos	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	Dimethylchlor	Fenhexamid	Formothion	Mefenacet	Nitenpyram	Propoxur	Terbufos	Bentazone (neg)
Bensulfuron-methyl	Clethodim	Dimethametryn	Fenobucarb	Fosthiazate	Mepanipyrim	Nuarimol	Propyzamide	Terbumeton	Bromoxynil (neg)
Benzoximate	Clofentezine	Dimethenamide	Fenoxyanil	Fuberidazole	Mepronil	Ofurace	Prosulfocarb	Terbutylazine	DNOC (neg)
Benzoylprop-ethyl	Clomazone	Dimethoate	Fenoxy carb	Furathiocarb	Mesotrione	Omethoate	Pymetrozine	Terbutryn	Fluazinam (neg) H
Bifenazate	Clopyralid	Dimethomorph	Fenpiclonil	Griseofulvin	Metalaxyd	Oxadixyl	Pyraclostrobin	Tetrachlorvinphos H	Flubendiamide (neg)
Bitertanol	Clothianidin	Dimetilan	Fenpyroximate	Halofenozone	Metamitron	Oxamyl	Pyrazophos	Tetraconazole	MCPA (neg)
Boscalid	Coumaphos	Dimoxystrobin	Fensulfothion	Haloxypoph	Metazachlor	Oxyfluorfen	Pyrethrin I	Tetramethrin	Tepraloxydim (neg)
Brodifacoum	Crotoxyphos	Diniconazole	Fenthion	Haloxypoph-methyl	Metconazole	Pacobutrazol	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

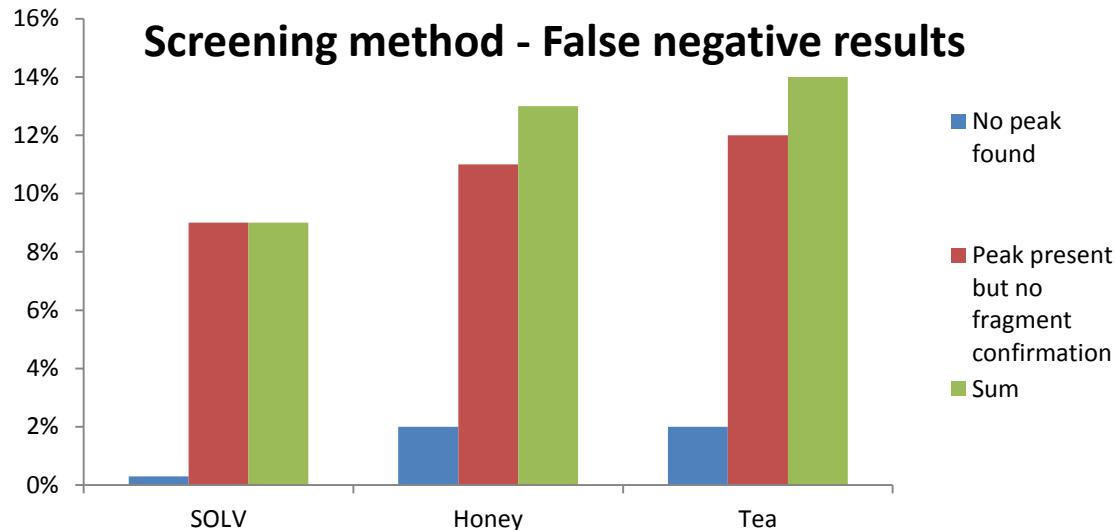


Compound H or T Missing fragment ion confirmation in one of the matrices at 10 ppb

Validation Parameters

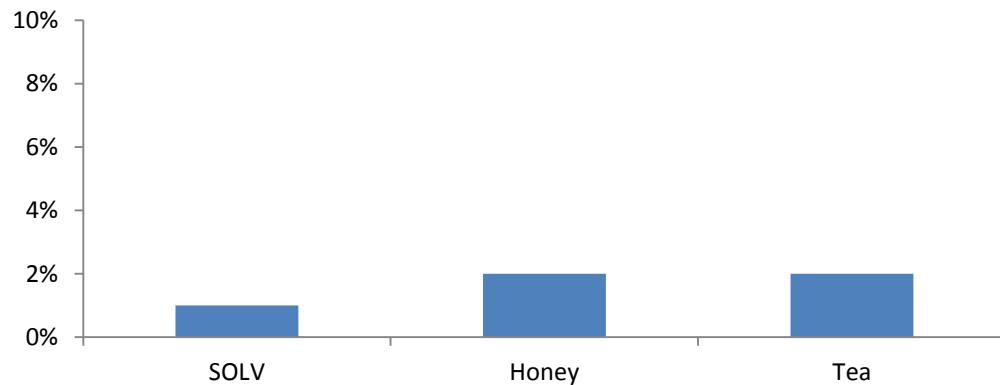
- Selectivity criteria: RT, min. detection of 1 HRAM quan and 1 HRAM qual ion with 5 ppm mass accuracy, ion ratio
- False negative and positive evaluation
- Recovery & repeatability at two concentration levels: 10 & 100 µg/kg
- Injection precision – 10 repeated injections of standard 100 µg/kg
- Screening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

Screening Method Efficiency – % of 330 Target Pesticides



Peak confirmation criteria:
 t_R , HRAM with 5 ppm accuracy and fragment ion presence

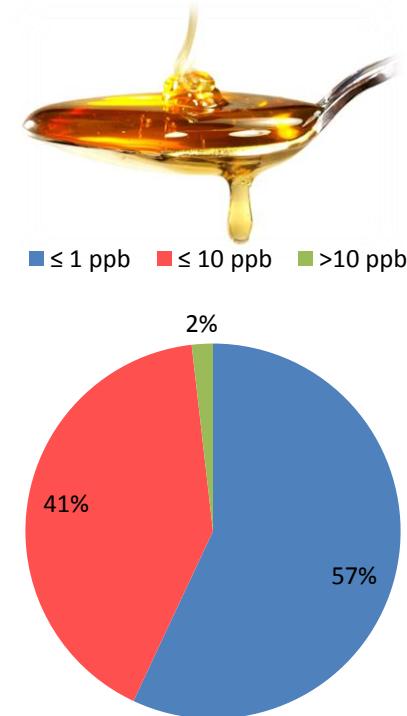
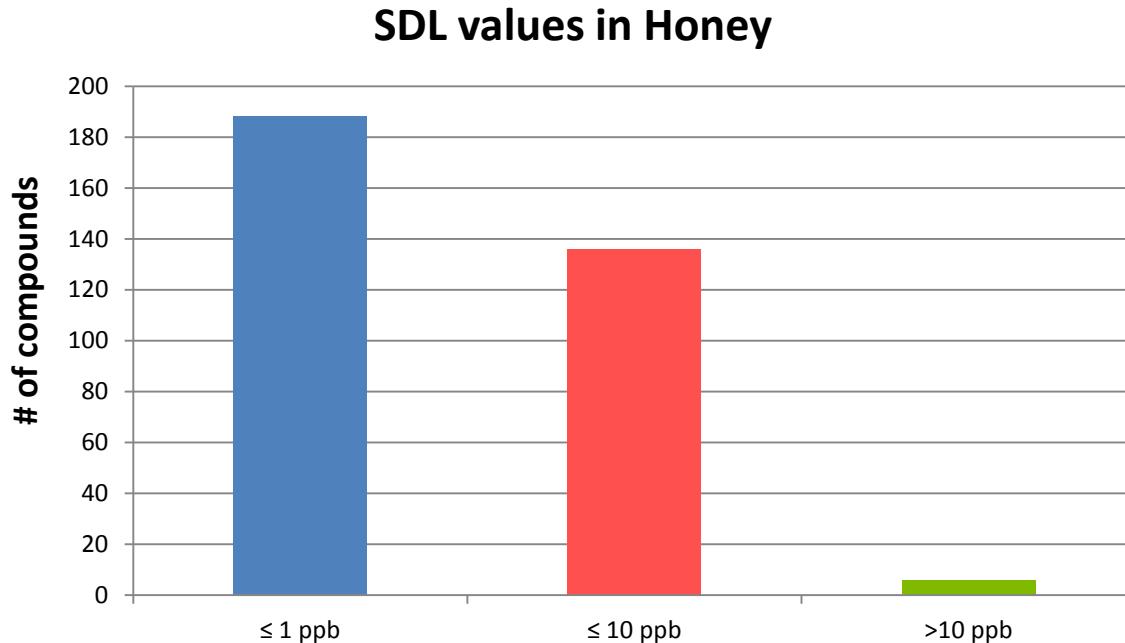
Screening method - False positive results



DG SANCO defines:
False negative < 5 %
No criteria for false positives

Results vDIA – Sensitivity Overview Honey

- Screening Detection Limits (SDL) in honey matrix

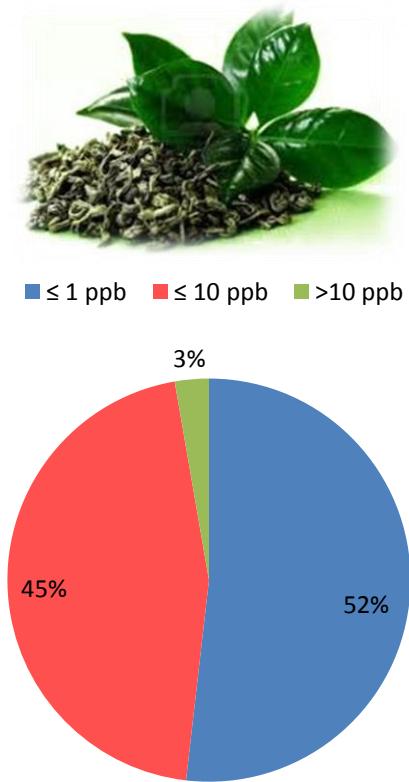
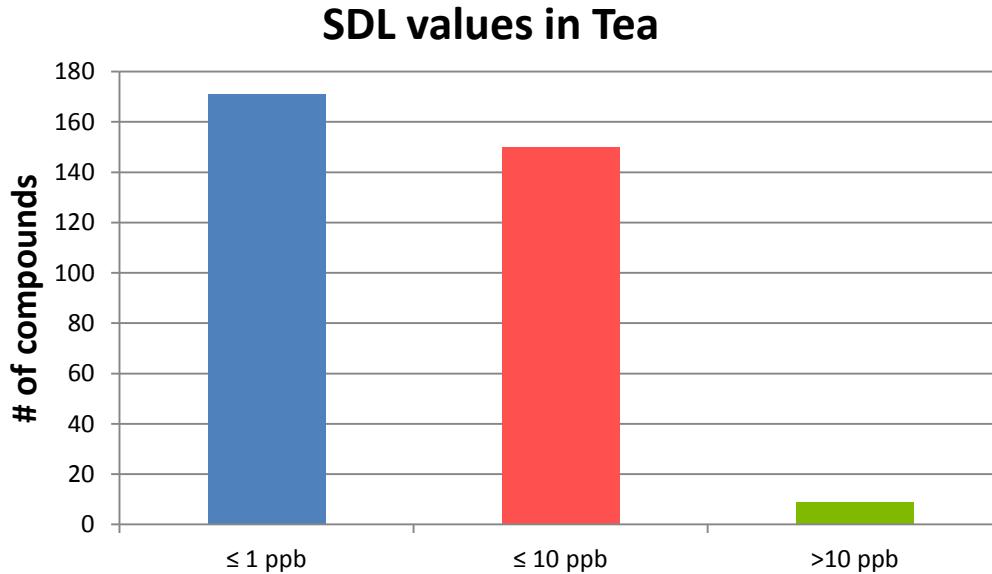


SUM = 330 compounds; 324 compounds $\text{SDL} \leq 10 \mu\text{g/kg}$

SDL determination according to SANCO12571/2013

Results vDIA – Sensitivity Overview Tea

- Screening Detection Limits – Tea matrix



SUM = 330 compounds; 321 compounds $SDL \leq 10 \mu\text{g/kg}$

SDL determination according to SANCO12571/2013

Missing Compounds at 10 ppb

SDL and MRL values for compounds not seen at 10 ppb

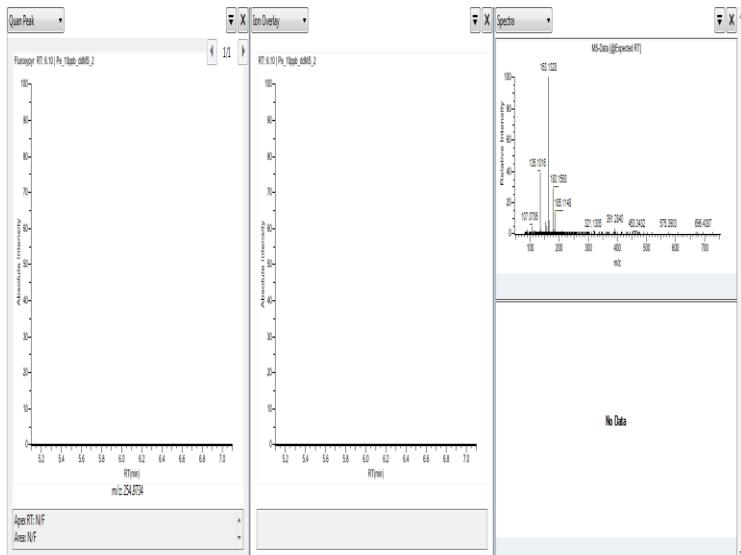
Compound name	SDL [ug/kg] in Honey	MRL for Honey [ug/kg]	SDL [ug/kg] in Tea	MRL for Tea [ug/kg]
Aramite	10	100	>100	n.d.
Bentazone	>100	50	0.5	100
Butafenacil	5	n.d.	60	n.d.
Dimethachlor	50	n.d.	40	20
Fenthion-sulfone	5	10	70	50
Hexaflumuron	3	n.d.	30	n.d.
Isoxathion	50	n.d.	50	n.d.
Mesotrione	>100	n.d.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

n.d. - not defined

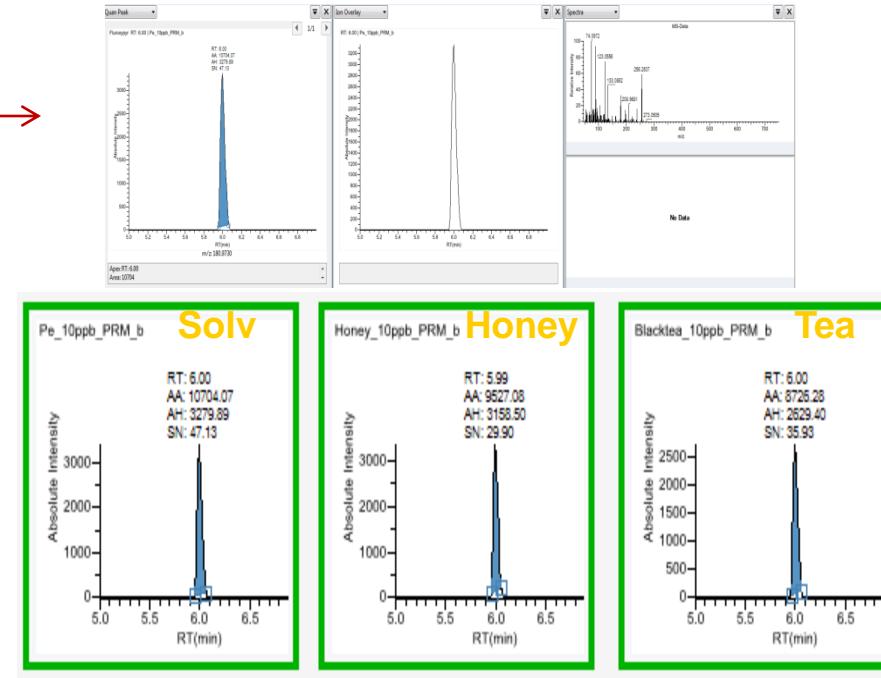
Sensitivity Improvement at MRL – ddMS vs PRM

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

Fluroxypyr 10 ppb ddMS

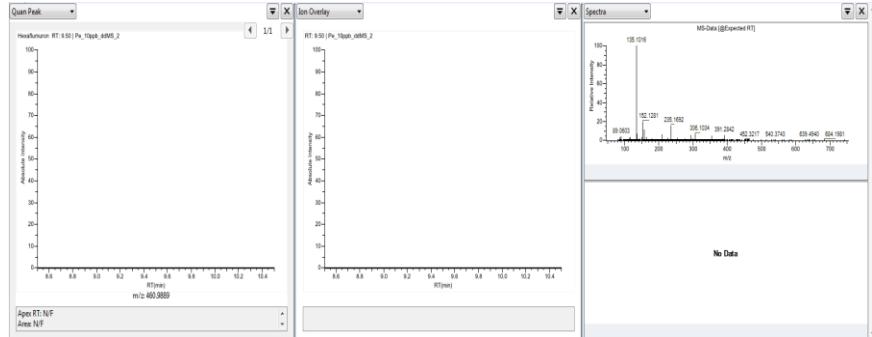


Fluroxypyr 10 ppb PRM

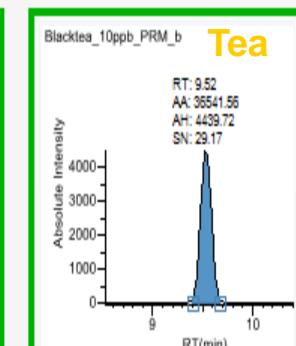
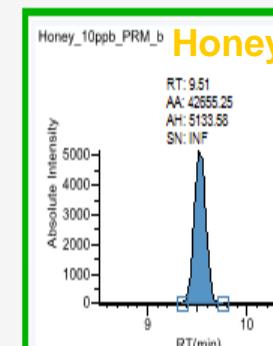
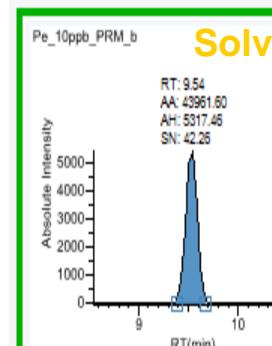
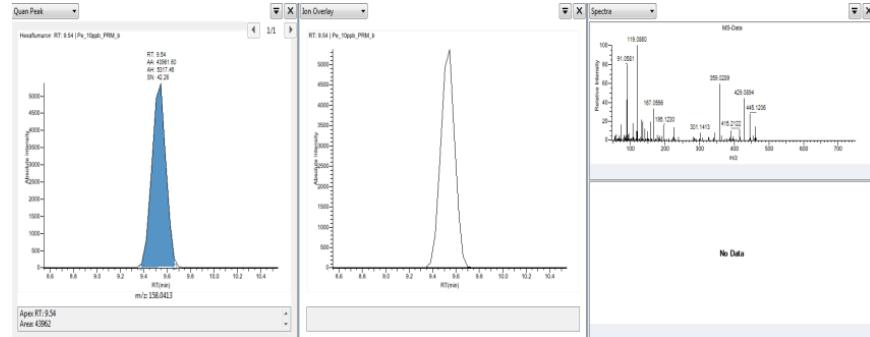


Sensitivity Improvement at MRL – ddMS vs PRM

Hexaflumuron 10 ppb ddMS

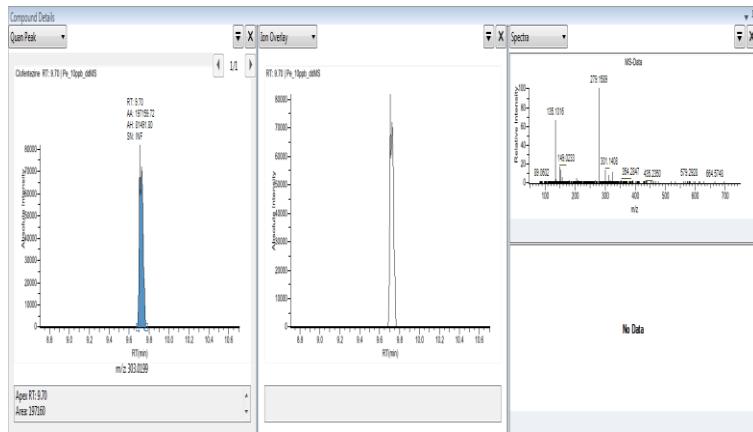


Hexaflumuron 10 ppb PRM

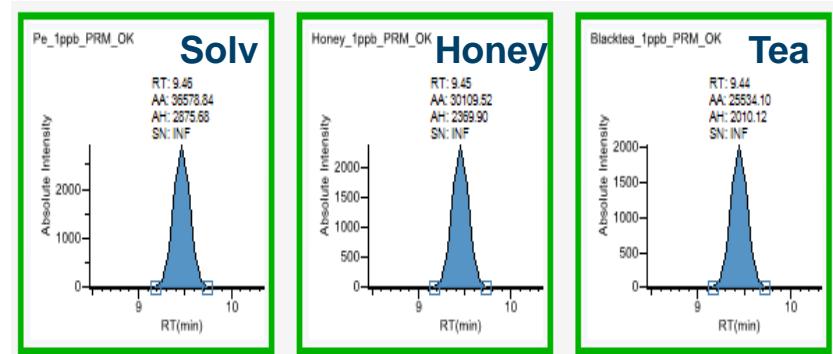
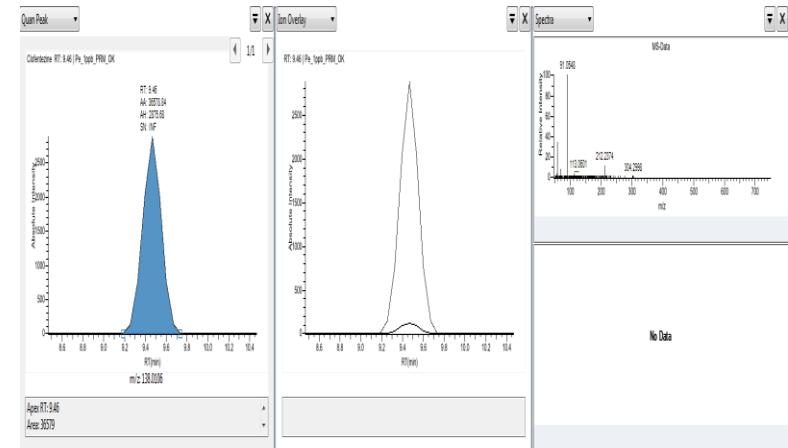


Sensitivity Improvement With Parallel Reaction Monitoring

Clofentezin 10 ppb in ddMS2

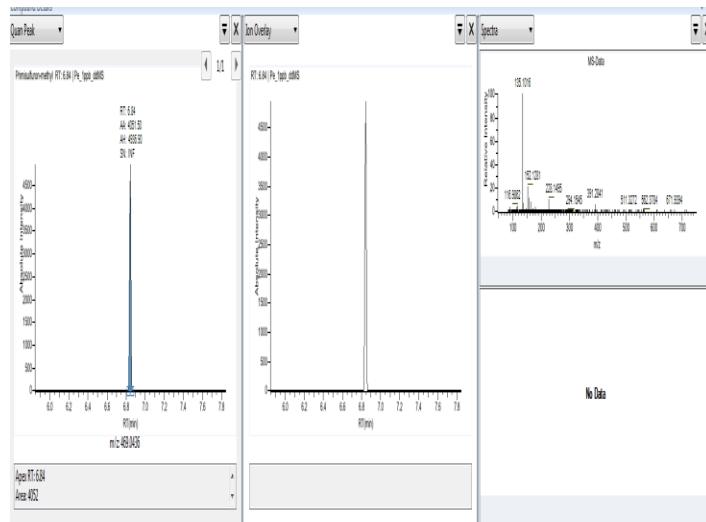


Clofentezin 1 ppb in PRM

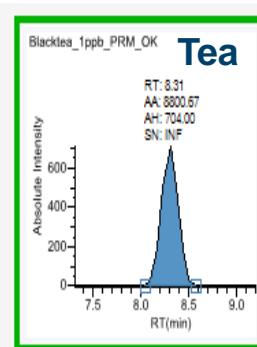
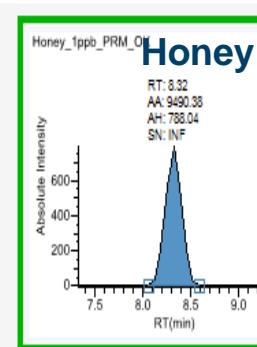
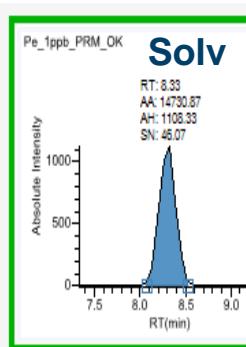
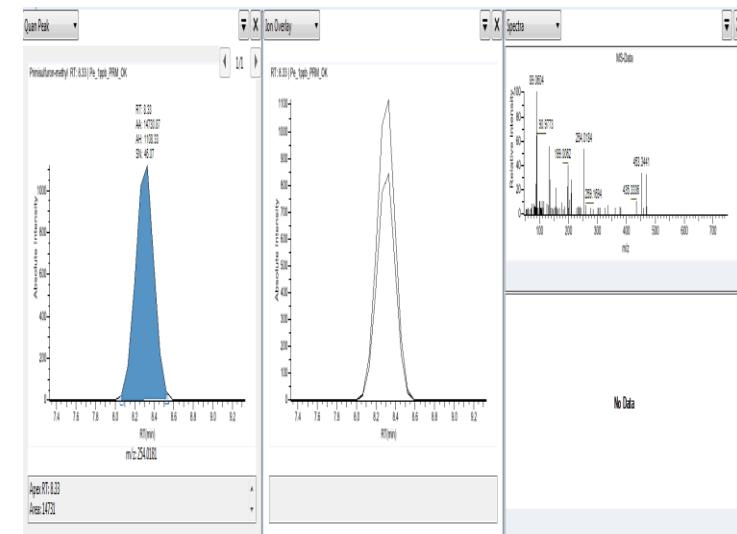


Sensitivity Improvement With Parallel Reaction Monitoring

Primisulfuron-Me 1ppb in ddMS

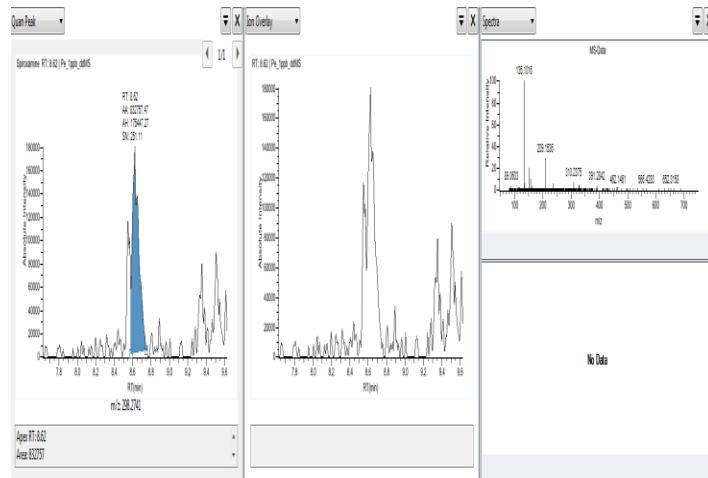


Primisulfuron-Me 1ppb in PRM

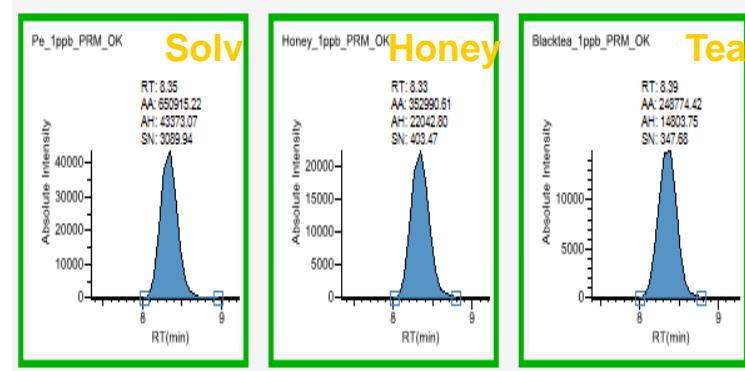
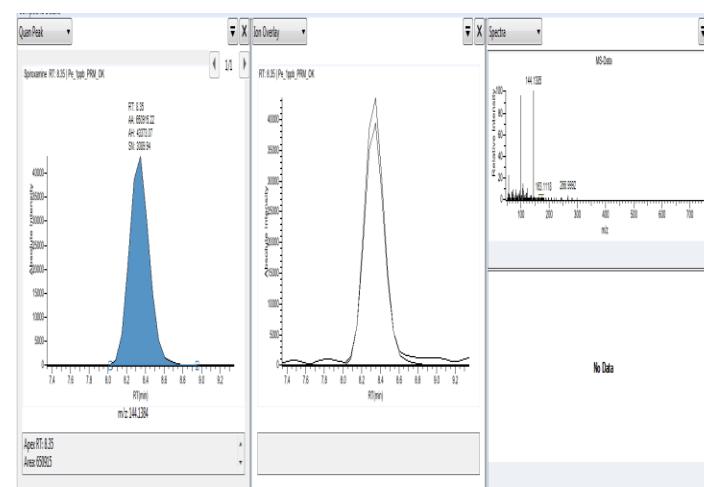


Sensitivity Improvement – With Parallel Reaction Monitoring

Spiroxamine 1ppb in ddMS



Spiroxamine 1ppb in PRM

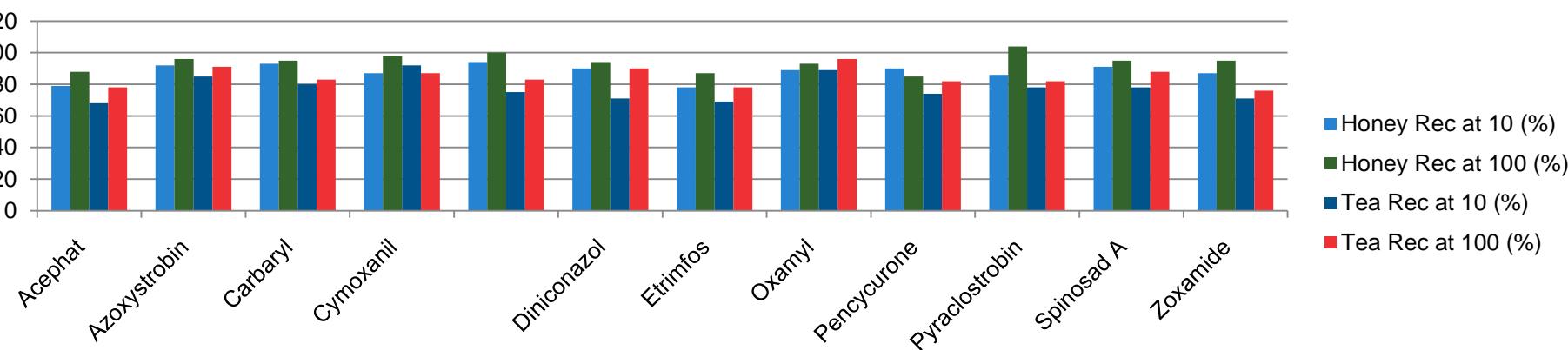


Method Sensitivity For 12 Representative Compounds - vDIA

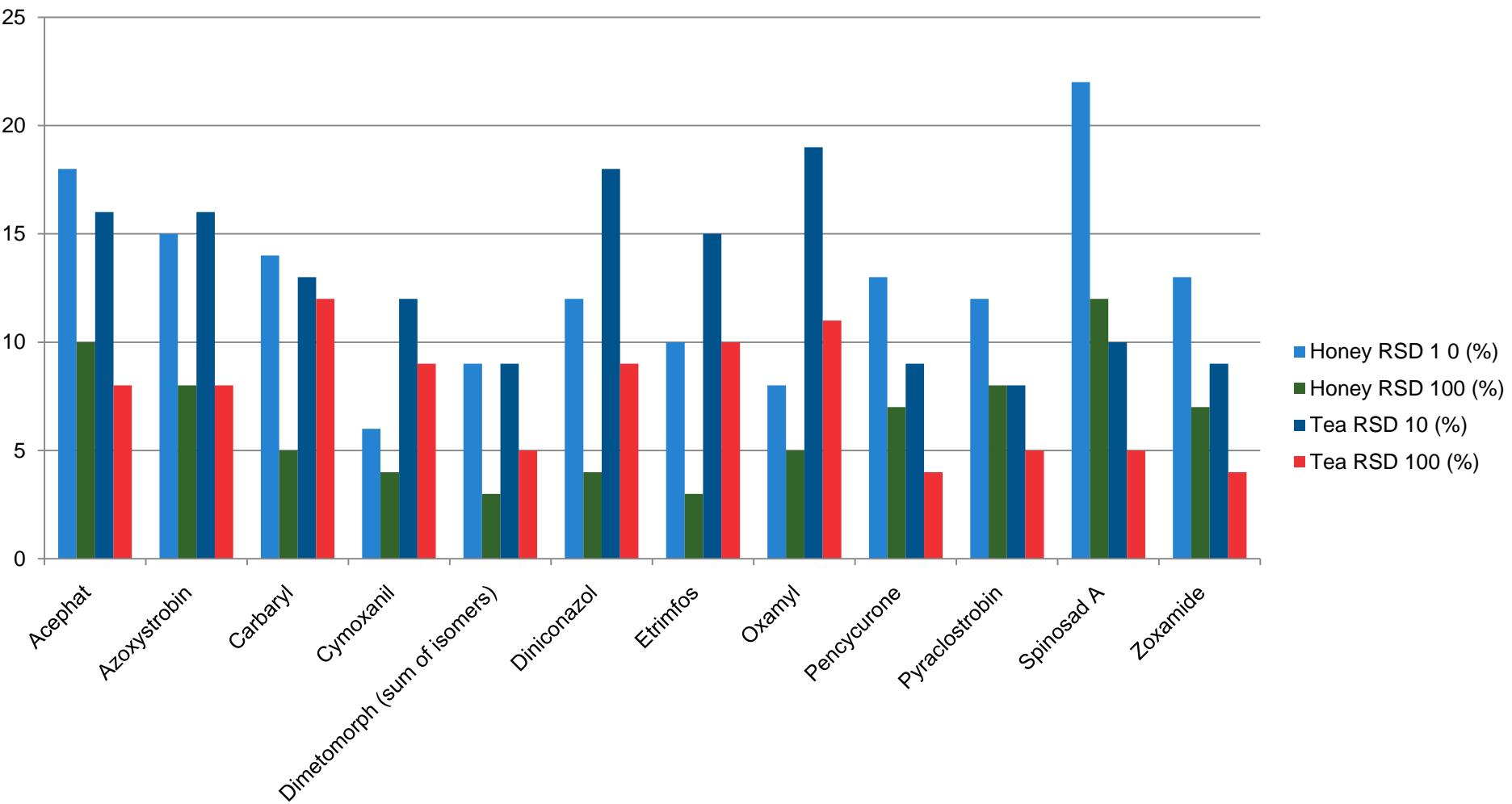
Analyte	Solvent (µg/kg)	Honey (µg/kg)	Tea (µg/kg)	MRL's (µg/kg)	
	LOQ	LOQ	LOQ	Honey	Tea
Acephate	2.5	10	10	20	50
Azoxystrobin	0.25	0.75	1.25	50	100
Carbaryl	0.25	2.5	3	50	50
Cymoxanil	0.25	1.25	5	50	50
Dimetomorph (sum of isomers)	12.5	12.5	25	50	50
Diniconazol	2	2.5	10	50	50
Etriflumuron	1.25	1.25	2.5		
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

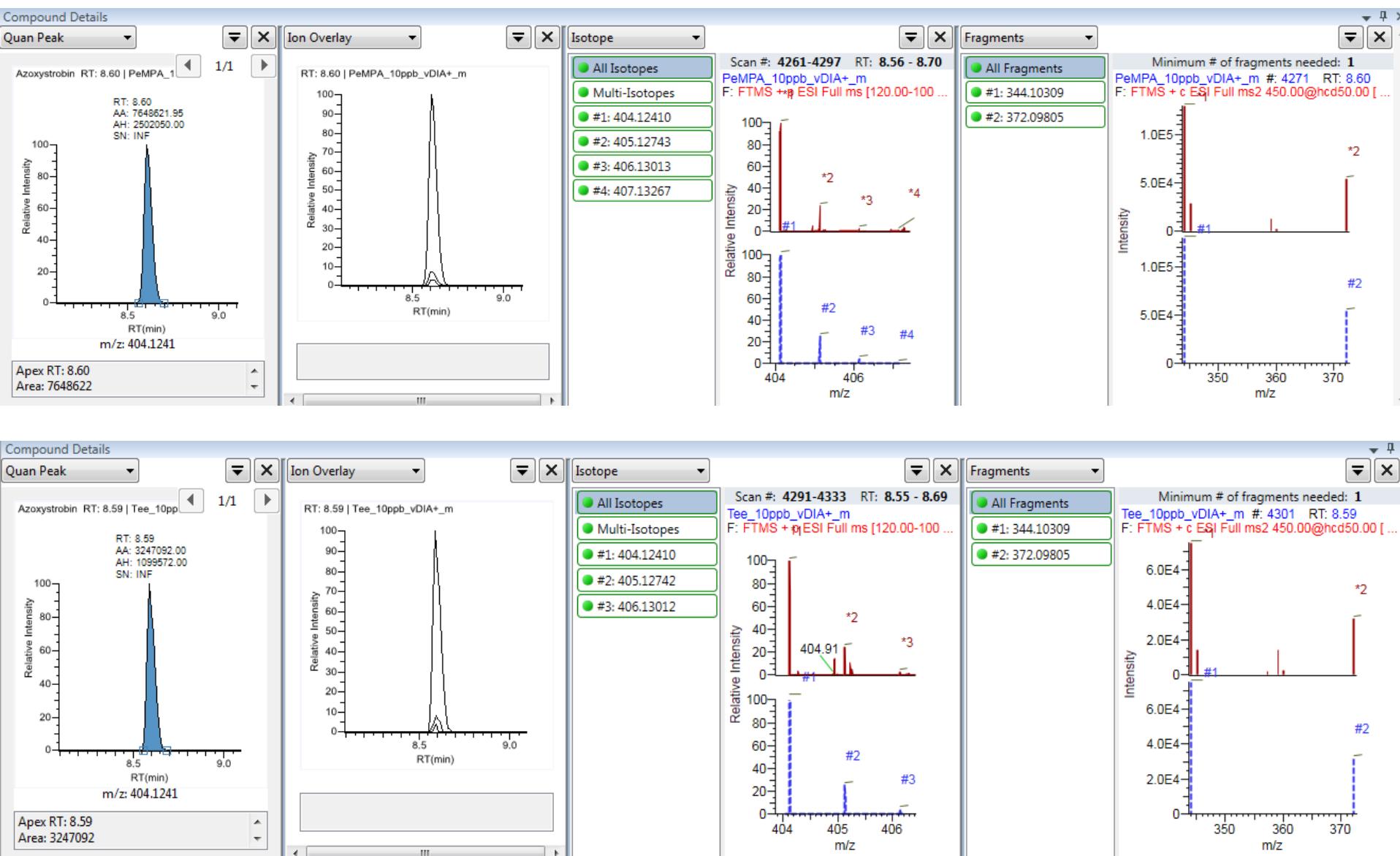
Analyte	Sp. Level 1 ($\mu\text{g/kg}$)	Sp. Level 2 ($\mu\text{g/kg}$)	Honey				Tea				Linearity in solvent
			RSD 1 (%)	RSD 2 (%)	Rec 1 (%)	Rec 2 (%)	RSD 1 (%)	RSD 2 (%)	Rec 1 (%)	Rec 2 (%)	
Acephat	10	100	18	10	79	88	16	8	68	78	0.9902
Azoxystrobin	10	100	15	8	92	96	16	8	85	91	0.9879
Carbaryl	10	100	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
Etrimfos	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



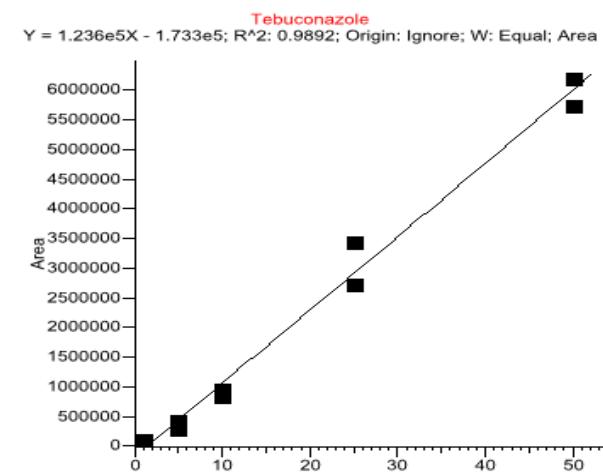
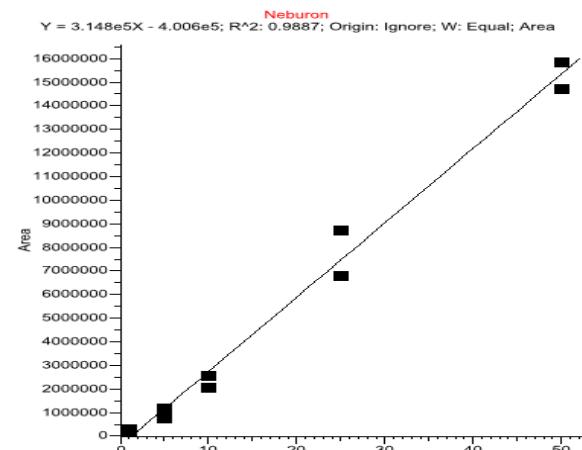
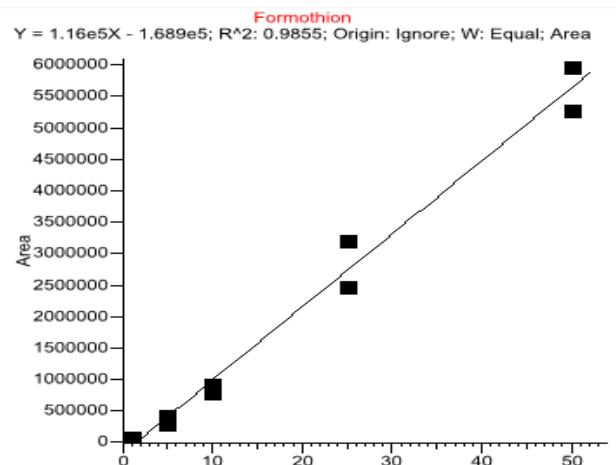
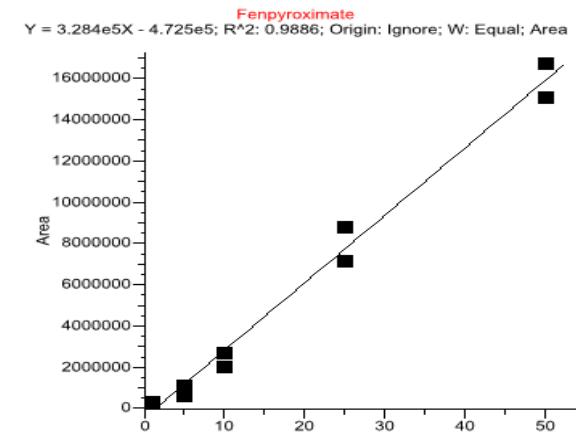
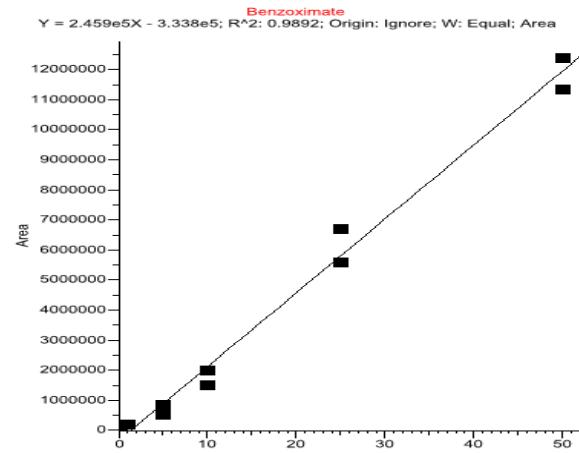
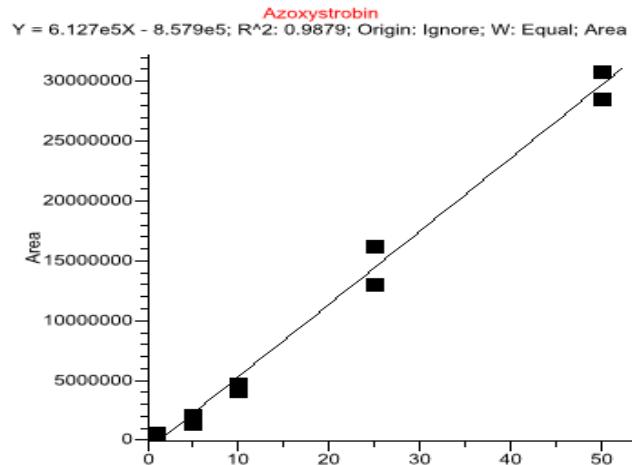
vDIA - Repeatability – For 12 Representatives



Azoxystrobin in Solvent and Tea at 10 ppb [μ g/kg]



Calibration Curves in Honey Matrix



Summary

- Complete workflow solution for multi-pesticide analysis has been evaluated
- Performance parameters of a complete routine method were tested and demonstrated that the vast majority of routinely measured LC amenable compounds can be analysed with high degree of confidence
- <10ppb (ug/kg) screening detection limit was achieved for >95 % of targeted compounds in all matrices (tea and honey)
- Excellent values under repeatability conditions
- TraceFinder software package allows streamlined method optimization, data acquisition and fast processing and reporting

Key User Benefits

Implementation

- Complete method delivered with instrument
- Full operation in <1 week from installation
- Implementation onsite by experienced application specialist

Performance

- LOQ<MRL for 95% of compounds even in heavy matrix
- Compliance with SANCO guidelines
- Robust operation due to unique source design
- High confidence in results due to HRAM data

Ease of use

- Simple method and instrument maintenance (no vent capillary exchange)
- Streamlined data management with TraceFinder

Flexibility

- Method expansion with built-in HRAM database
- Application support from Special Solutions Center Europe



ThermoFisher SCIENTIFIC

The world leader in serving science

For more details please contact:
Laszlo Hollosi,
laszlo.hollosi@thermofisher.com