

Multiresidue Method for the Quantification of Pesticides in Fruits, Vegetables, Cereals and Black Tea using UPLC-MS/MS

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Abstract

In this application note we demonstrate a multi-residue UPLC-MS/MS method developed for the determination of 552 pesticides and relevant metabolites in fruits, vegetables, cereals and black tea extracts using ACQUITY UPLC I-Class coupled with Xevo TQ-XS

Benefits

- The ACQUITY UPLC I-Class coupled with Xevo TQ-XS allows for the determination of 552 pesticides in various commodities at concentrations below the default EU MRL in a single method
- In all five matrices, despite the dilution of the crude QuEChERS extract, most of the compounds were detected below the typical EU default MRL of 0.01 mg/kg
- The Gaussian peak shape of early eluting compounds is maintained by use of the post injector mixing kit

Introduction

As the population grows, demand for food consumption and global trade in the food industry has also increased. Hundreds of pesticides are routinely used for crop protection across the globe, traces of pesticides left in treated commodities are called “residues”. Regulations are in place for Maximum Residue Levels (MRL), that are legally tolerated in or on food and feed when pesticides are applied correctly in accordance with Good Agricultural Practice. A growing target list of pesticides in complex matrices, and the need for low limits of detection, bring various challenges for multi-residue methods.

A multi-residue method for 552 pesticides and relevant metabolites was developed for various food commodities. Extracts from representative commodities, including high-water content (spinach), high acid and highwater content (strawberry), high oil and very low-water content (soybean), high protein and low-water and fat content (wheat flour), and difficult or unique commodities (black tea) were chosen to assess the performance of the UPLC-MS/MS method.

Experimental

Sample preparation

Representative samples from different commodity groups were chosen including high-water content (spinach), high acid and high-water content (strawberry), high oil and very low-water content (soybean), high protein and low water and fat content (wheat flour), and difficult or unique commodities (black tea). These samples were purchased from a local retail store. The samples were immediately homogenized in a food processor and frozen until extraction was performed. The samples were extracted using the QuEChERS CEN method. For high aqueous samples, such as spinach and strawberry, samples were prepared according to the European Union Reference Laboratory (EURL) fruits and vegetable method.¹ For intermediate or low water content, such as soybean and wheat flour, the EURL method for cereals and feeding stuff was used.² For low water and high carbohydrates, such as tea, the QuEChERS extraction with sample cleanup method was followed.³

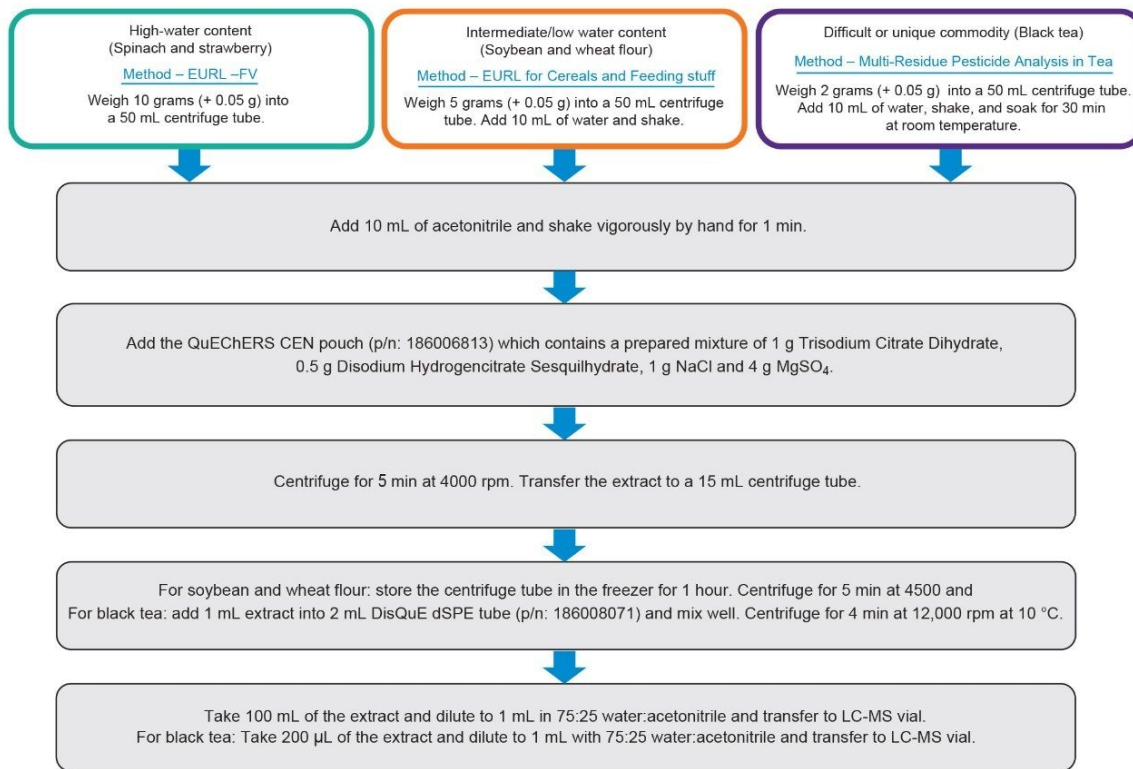


Figure 1. Sample preparation method for strawberry, spinach, soybean, wheat flour, and black tea.

The performance of the UPLC-MS/MS method was assessed by spiking a mixture of 256 pesticides (see Annex for details) into extracts of spinach, strawberry, soybean, wheat flour and black tea at eight concentrations between 0.0001 to 0.1 mg/kg. Matrix-matched calibration standards were injected in a way that simulated the analysis of a batch of samples; calibration standards bracketed a series of replicate injections (n=6) at three matrix-matched levels.

	Dilution factor from sample preparation	Target in sample concentration (mg/kg)	Equivalent in vial concentration (ng/mL)
Strawberry	10	0.01	1
Spinach	10	0.01	1
Soybean	20	0.01	0.5
Wheat flour	20	0.01	0.5
Black tea	25	0.01	0.4

Table 1. An example of the default MRL of 0.01 mg/kg in strawberry, spinach, soybean, wheat flour and black tea along with their dilution factors from QuEChERS sample preparation and their equivalent concentration in vial. For ease of discussion, all concentrations are reported in mg/kg for standards, QCs and samples discussed in this application note.

UPLC-MS/MS Conditions

UPLC system:	ACQUITY UPLC I-Class with FL Sample Manager
Column:	ACQUITY UPLC HSS T3 Column, 1.8 μ m, 2.1 \times 100 mm (186003539)
Post injector mixing kit:	50 μ L extension loop (430002012)
Mobile phase A:	5 mM ammonium formate in water + 0.1% formic acid
Mobile phase B:	5 mM ammonium formate in 50:50 MeCN: MeOH + 0.1% formic acid
Injection volume:	5 μ L (PLNO mode)
Column temp.:	40 $^{\circ}$ C

Sample temp.:	10 °C
Run time:	19 minutes
Data acquisition and processing:	MassLynx 4.2 and TargetLynx XS
MS instrument:	Xevo TQ-XS
Ionization:	Electrospray
Polarity:	Positive and negative ion mode
Capillary voltage:	+2.0/-2.0 kV
Desolvation temp.:	300 °C
Desolvation gas flow:	1000 L/Hr
Source temp.:	150 °C
Cone gas flow:	150 L/Hr
Strong needle wash (SNW):	0.1% formic acid in acetonitrile
Weak needle wash (WNW):	Mobile phase A
Seal wash:	10% methanol in water

Gradient:

Time	Flow rate	% A	% B	Curve
Initial	0.5	99.0	1.0	Initial
0.5	0.5	99.0	1.0	6
3.50	0.5	60.0	40.0	6
12.50	0.5	15.0	85.0	6
12.60	0.5	1.0	99.0	6
15.00	0.5	1.0	99.0	6
15.10	0.5	99.0	1.0	6
19.00	0.5	99.0	1.0	6

The LC-MS/MS method for 552 pesticides was created using the Quanpedia database. The Quanpedia database automatically creates LC and MS acquisition methods as well as processing methods from a compendium of compound specific MS parameters such as transitions, cone voltage, and collision energy. The MS method used for this work contains at least 2 MRM transitions per pesticide resulting in a method with more than 1000 MRMs. The auto-dwell functionality in the MS method managed the dwell time for each transition, ensuring the required number of data points (10–14) is available across each peak. Figure 2 shows an example of method creation in Quanpedia for multi-residue analysis.

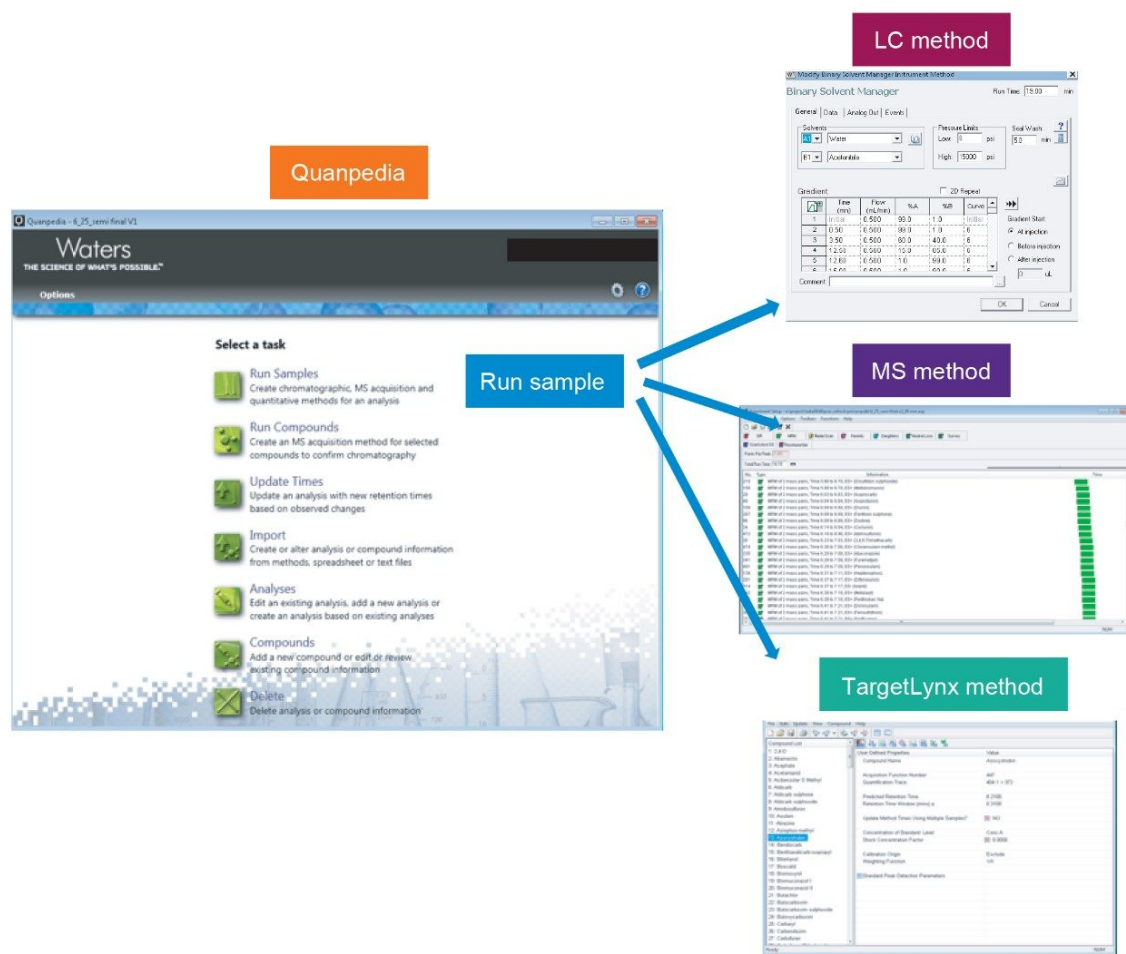


Figure 2. Creation of a multi-residue method for the determination of 552 pesticides using Quanpedia.

Results and Discussion

Chromatography

This multi-residue method was developed for a wide range of pesticides with varying chemical properties. A few of these pesticides, such as methamidophos and acephate, are very polar and elute early in the chromatogram. Injecting samples containing a moderate organic content (25%) often results in fronting and/or splitting peaks

for the early eluting compounds. Reducing organic content in the sample diluent prior to its injection onto the column may help to improve the peak shape of early eluting analytes. Installing a post injector mixing kit, between the injector port and column, allows the injection of typical QuEChERS extracts into high aqueous mobile phase without compromising peak shape. Before making the injection, the extension loop is filled with the high aqueous mobile phase, which provides more volume to aid dispersion of the sample into the aqueous solvent prior to transfer onto the column. In this way, a moderate level (25%) of organic solvent can still be used to prepare the sample, in which most of the pesticides remain soluble, whilst still providing good peak shape for the very polar analytes. Figure 3 shows the peak shape of methamidophos with and without the post injector mixing kit; good peak shape is observed with the extension loop fitted, which provided more reliable quantitation and greater sensitivity and hence a lower limit of detection (LOD).

Methamidophos

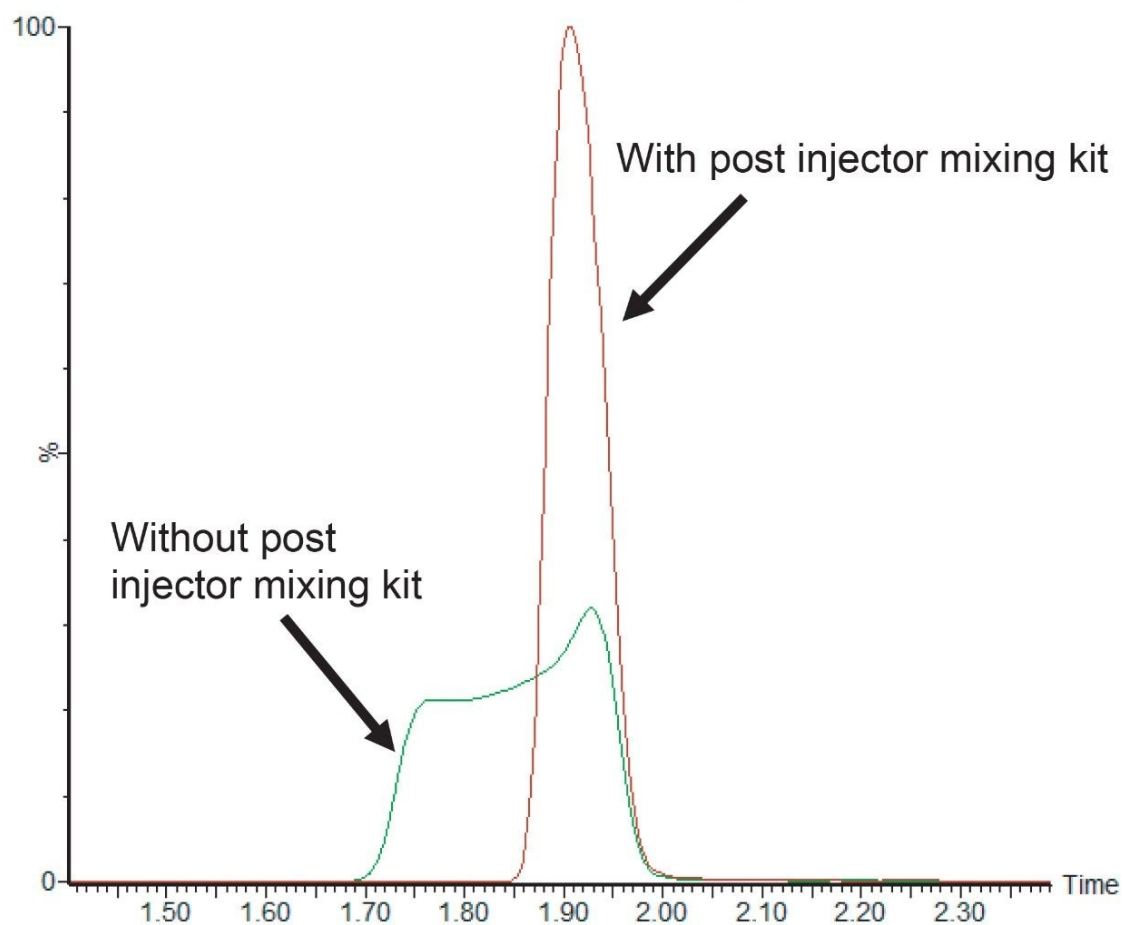


Figure 3. Chromatogram of methamidophos with (green trace) and without (red trace) the post injector mixing kit fitted.

As shown in Figure 4, the retention time of methamidophos is obtained at ~1.9 minutes with a peak width at the base line of ~7 seconds in all studied matrices. The retention time of methamidophos is greater than two times the retention time corresponding to the void volume of the column (void time 0.46 min).

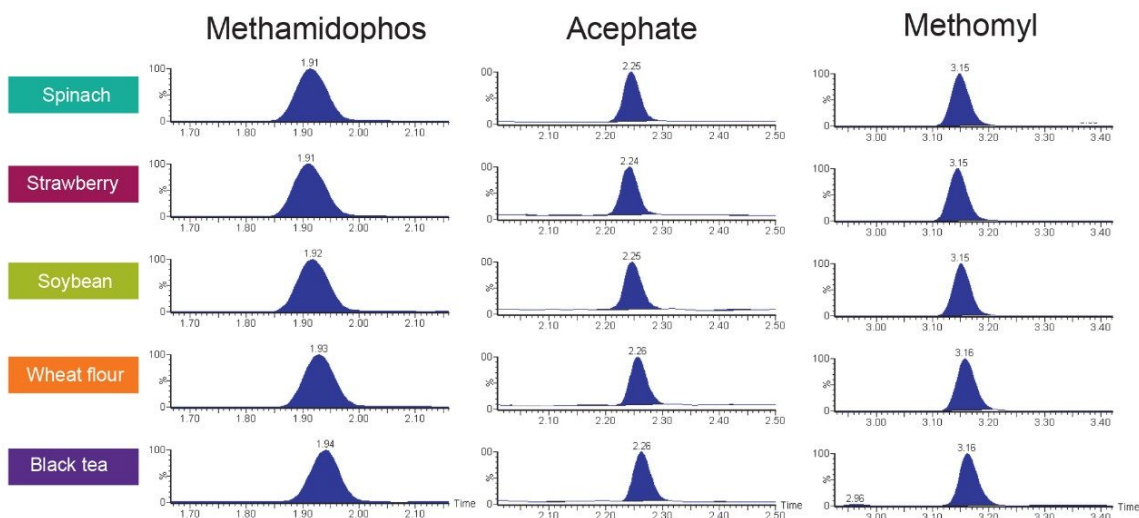


Figure 4. Chromatograms for some of the very polar analytes across matrices with the post injector mixing kit fitted.

Sensitivity, Quantification, Precision, and Matrix Effects

The calibration graphs for the majority of the 256 compounds in all the matrices showed values for coefficient of determination (R^2) greater than 0.99 and back-calculated concentrations (residuals) were all within the $\pm 20\%$ SANTE tolerance.⁴ Figure 5 shows matrix matched calibration graphs for metoxuron, a representative analyte, in five studied matrices.

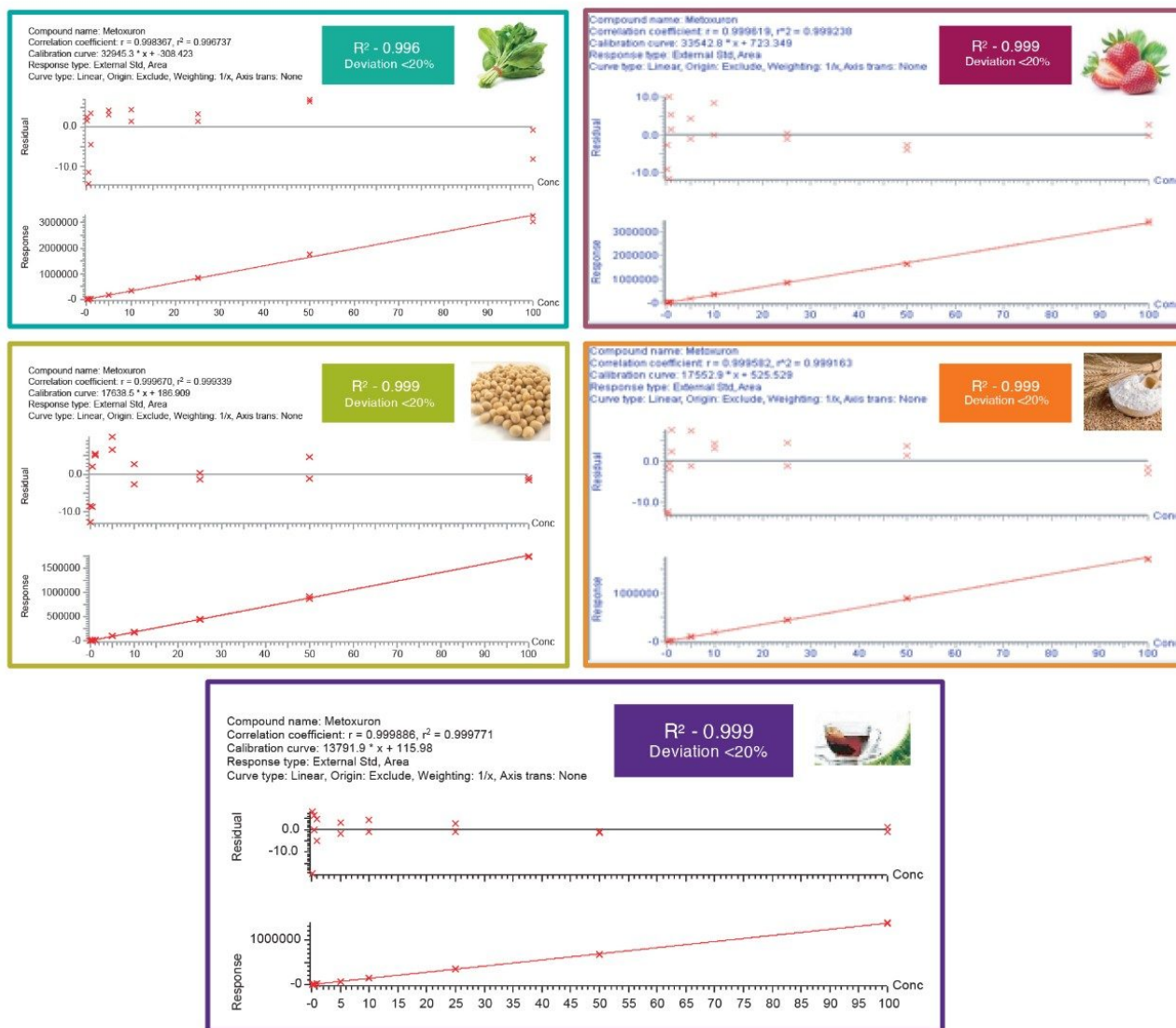


Figure 5. Matrix-matched calibration graphs from 0.0001 to 0.1 mg/kg (equivalent to 0.1 to 100 ng/mL) for metoxuron in spinach, strawberry, soybean, wheat flour and black tea.

The precision of the LC-MS/MS measurements was calculated for the 256 representative pesticides from the replicate (n=6) determination of matrix-matched standards at three concentrations (0.005, 0.01, and 0.05 mg/kg). The precision of the measurement was good with more than 85% of the detected pesticides exhibiting RSDs for peak area of less than 10% (see Figure 6).

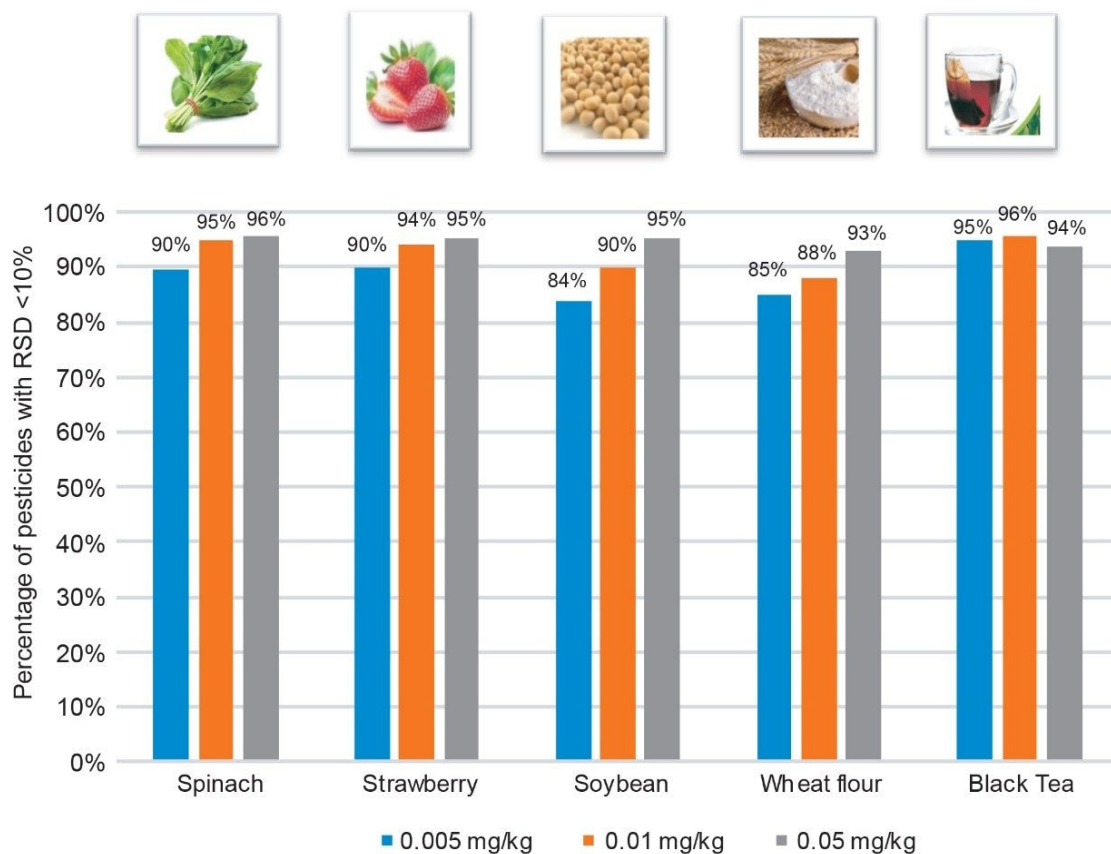


Figure 6. The percentage of pesticides with RSDs <10% in various matrices at 0.005 mg/kg, 0.01 mg/kg, and 0.05 mg/kg.

Matrix effects for the 256 compounds in the various matrices were calculated by comparing the ratio of the slope of the matrix-matched calibration graph for each commodity to that of the solvent calibration graph. Figure 7 summarizes the range of matrix effects observed for each commodity. All commodities show some degree of matrix suppression (response suppressed by >20%) and enhancement (response of the analyte increased by >20%) so matrix-matched calibration is recommended. Procedural calibration or standard addition are alternative approaches which compensate for matrix effects and recovery losses.

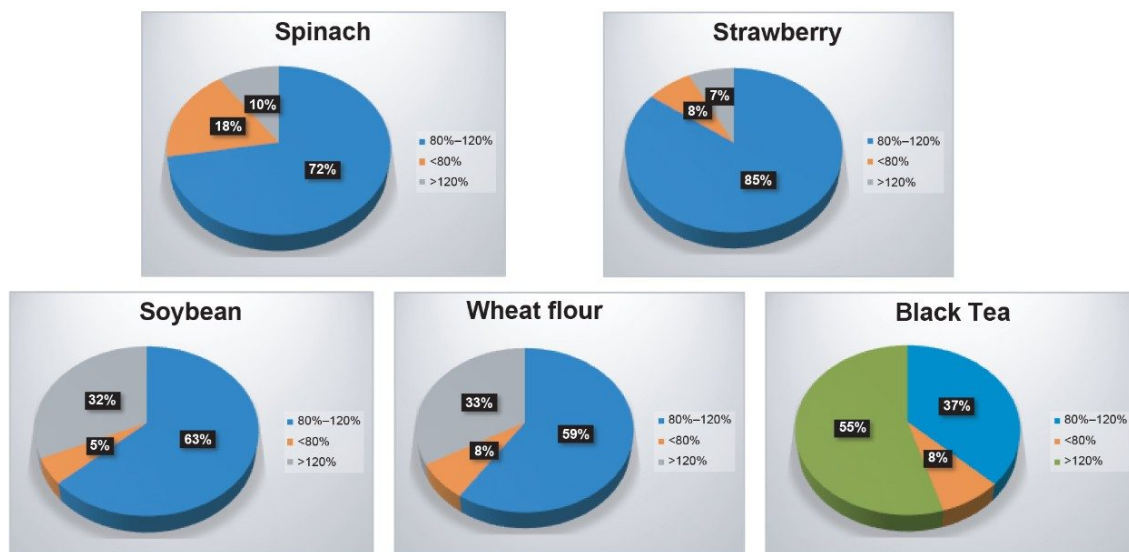


Figure 7. Percentage of pesticides that exhibited significant matrix effects for spinach, strawberry, soybean, wheat flour and black tea.

In order to investigate matrix effects further, soybean samples were prepared using QuEChERS and the extracts diluted (5x and 10x). Matrix-matched calibrants were prepared to 0.02 mg/kg in both crude and diluted extracts and analyzed. To investigate the impact of co-extractives, full scan RADAR acquisition was enabled alongside the MRM transitions for the 552 compounds of interest. Figure 8 shows an overlay of the total ion current (TIC) chromatograms from RADAR for the various soybean extracts. There is a significant peak observed at 3.9 minute in the crude extract trace (blue). One compound of interest, carbofuran 3-hydroxy, also elutes at 3.9 minute in the chromatogram based on this LC gradient.

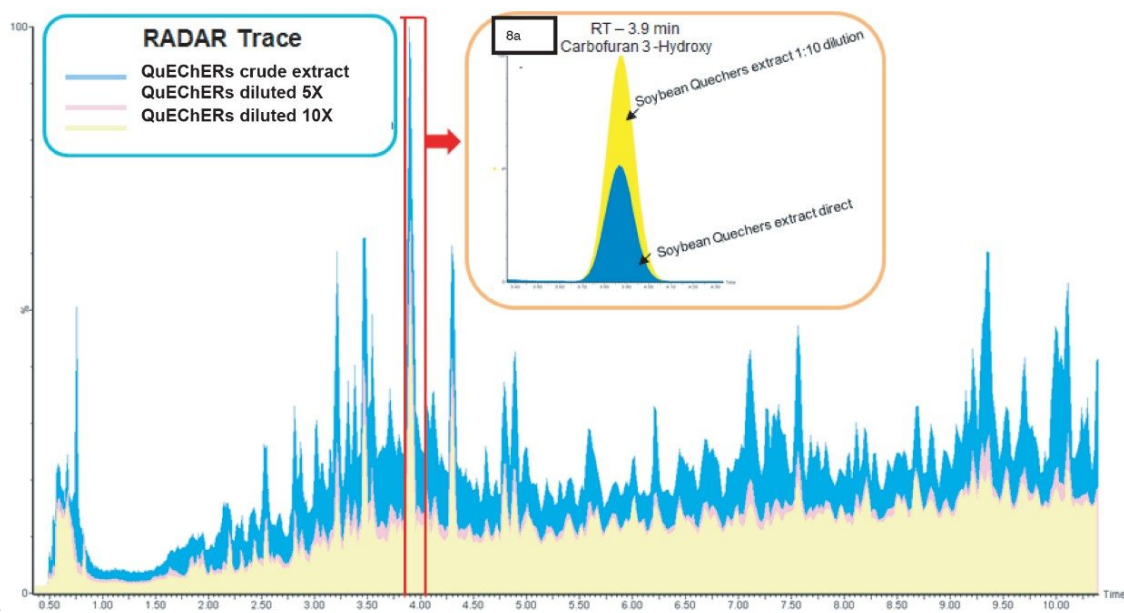


Figure 8. RADAR trace of soybean QuEChERS crude extract (blue), extract diluted 5 times (pink) and extract diluted 10 times (yellow). 8a shows chromatogram of carbofuran 3-hydroxy in QuEChERS crude extract (blue) and extract diluted 10 times (yellow).

The presence of co-eluting matrix co-extractives can cause ion enhancement/suppression or isobaric interference. As can be seen in inset 8a, the peak area of carbofuran 3-hydroxy is almost doubled when the QuEChERS crude extract of soybean is diluted 10 times (yellow) compared to the crude QuEChERS extract (blue). The dilution of these QuEChERS crude extracts (as shown in Figure 8) reduces the matrix effect while still achieving the required LOD, without the need for sample cleanup. Dilution also reduces the loading of co-extractives into the system, increasing UPLC column longevity, decreasing the frequency of routine instrument maintenance and improving analytical productivity.

Identification Criteria

Identification criteria, retention times and ion ratios, were calculated and flagged using TargetLynx XS. The retention time and ion ratio of each analyte detected in the sample should correspond to that of the calibration standard reference.⁴ The retention times of all 256 representative pesticides, in all the matrices studied, were found to be within the tolerance of ± 0.1 min. The ion ratios from the analysis of matrix-matched calibrants were within $\pm 30\%$ of the reference values for all 256 compounds.

Conclusion

- A multi-residue UPLC-MS/MS method has been developed for the determination of 552 pesticides and relevant metabolites for pesticide residue analysis.
- Quanpedia is an extendable and searchable database, from which chromatographic, MS and TargetLynx processing methods can be generated for the determination of 552 pesticides in an efficient manner. Quanpedia offers fast and simple ways for method generation and maintenance.
- The performance of the UPLC-MS/MS method was evaluated for the determination of 256 representative analytes in extracts from various commodities.
- The post injector mixing kit enables the reliable injection of typical QuEChERS extracts into high aqueous mobile phase without compromising peak shape of early eluting analytes.
- The dilution of the QuEChERS crude extracts reduces the loading of co-extractives into the system and decreases the frequency of routine instrument maintenance.
- Detection of most compounds in matrix-matched calibrants at concentrations below the typical EU default MRL of 0.01 mg/kg was achieved in the five matrices.

References

1. EURL-FV Multiresidue Method using QuEChERS followed by GC-QqQ/MS/MS and LC-QqQ/ MS/MS for Fruits and Vegetables. https://www.eurl-pesticides.eu/library/docs/fv/CRLFV_Multiresidue_methods.pdf <
https://www.eurl-pesticides.eu/library/docs/fv/CRLFV_Multiresidue_methods.pdf>
2. EURL for cereals and feeding stuff: Determination of pesticide residues in wheat, rye, oat and rice by LC-MS/MS and GC-MS/MS (QuEChERS method). [https://www.eurl-pesticides.eu/userfiles/file/\(23B\)%20Appendix7_2017_dec_Validation%20report%2023B%20samstik%20H_cereals.pdf](https://www.eurl-pesticides.eu/userfiles/file/(23B)%20Appendix7_2017_dec_Validation%20report%2023B%20samstik%20H_cereals.pdf) <
[https://www.eurl-pesticides.eu/userfiles/file/\(23B\)%20Appendix7_2017_dec_Validation%20report%2023B%20samstik%20H_cereals.pdf](https://www.eurl-pesticides.eu/userfiles/file/(23B)%20Appendix7_2017_dec_Validation%20report%2023B%20samstik%20H_cereals.pdf)>

3. Multi-Residue Pesticide Analysis in Tea: Optimized Cleanup After QuEChERS Extraction for UPLC-MS/MS and GC-MS/MS Analysis. https://www.waters.com/waters/library.htm?cid=511436&lid=134775094&locale=en_US <
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4. SANTE/12682/2019: https://ec.europa.eu/food/sites/food/files/plant/docs/pesticides_mrl_guidelines_wrkdoc_2019-12682.pdf <
https://ec.europa.eu/food/sites/food/files/plant/docs/pesticides_mrl_guidelines_wrkdoc_2019-12682.pdf>

Annex

2,4-D	Cyproconazole I
Abamectin B1b	Cyproconazole II
Acephate	Cyprodinil
Acetamiprid	Cyromazine
Acibenzolar S Methyl	Demeton-S-methyl
Alachlor	Demeton-S-methyl sulphone
Aldicarb	Demeton-S-methyl sulphoxide (oxydemeton methyl)
Aldicarb sulphone	Desmedipham
Aldicarb sulphoxide	Diafenthiuron
Ametoctradin	Dichlorprop
Amidosulfuron	Diclobutrazol
Asulam	Diethofencarb
Azinphos-methyl	Difenoconazole
Atrazine	Diflubenzuron
Azoxystrobin	Dimethoate
Bendiocarb	Dimethomorph E
Benthiavalicarb-isopropyl	Dimethomorph Z
Bispyribac-sodium	Dimethylaminosulfanilide acid (DMSA)
Bitertanol	Dimoxystrobin
Boscalid	Diniconazole
Bromoxynil	Dinotefuran
Bromuconazol I	Disulfoton
Bromuconazol II	Disulfoton sulphone
Butachlor	Disulfoton sulphoxide
Butocarboxim	Diuron
Butocarboxim sulphoxide	Dodine
Butoxycarboxim	Emamectin benzoate B1a
Carbaryl	Emamectin benzoate B1b
Carbetamide	Epoxiconazole
Carbofuran	Ethiofencarb
Carbofuran (3-hydroxy)	Ethiofencarb sulphone
Carboxin	Ethiofencarb sulphoxide
Chlorantraniliprole	Ethirimol
Chlorfenvinphos	Etofenprox
Chloridazon	Etoxazole
Chlortoluron	Fenamidone
Chromafenozide	Fenamiphos
Clethodim	Fenamiphos sulfone
Clofentezine	Fenamiphos sulfoxide
Clothianidin	Fenarimol
Cyazofamid	Fenbutatin oxide
Cycloxydim	Fenbuconazole
Cyflufenamid	Fenhexamid
Cymoxanil	Fenoxycarb

Fenpropimorph
Fenpyroximate
Fensulfothion
Fensulfothion oxon
Fensulfothion oxon sulfone
Fensulfothion sulfone
Fenthion
Fenthion sulphone
Fenthion sulphoxide
Fipronil
Flonicamid
Fluazifop
Fluazifop-P-butyl
Fluazinam
Flubendiamide
Fludioxonil
Flufenacet
Flufenoxuron
Fluometuron
Fluopicolide
Fluoxastrobin
Fluquinconazole
Flusilazole
Flutriafol
Fluxapyroxad
Fonofos
Formetanate-HCl
Fosthiazate
Furathiocarb
Furmecyclox
Halofenozide
Halosulfuron methyl
Haloxyfop (free acid)
Heptenophos
Hexaconazole
Hexythiazox
Imazalil
Imidacloprid
Indoxacarb
Ioxynil
Iprovalicarb
Isazofos
Isocarbofos
Isofenphos
Isofenphos-methyl
Isoprocarb

Isoprothiolane
Isoproturon
Isoxaben
Isoxaflutole
Kresoxim-methyl
Lenacil
Linuron
Lufenuron
Malathion
Mandipropamid
Mecarbam
Mepanipyrim
Mesosulfuron methyl
Mepronil
Metaflumizone
Metaxyl
Metamitron
Metconazole
Methabenzthiazuron
Methamidophos
Methiocarb
Methiocarb sulfone
Methiocarb sulfoxide
Methomyl
Methoxyfenozide
Metobromuron
Metolachlor
Metolcarb
Metosulam
Metoxuron
Metrafenone
Metsulfuron-methyl
Mevinphos I
Molinate
Mevinphos II
Monocrotophos
Monuron
Myclobutanil
N-2,4-dimethylphenyl-N'-methylformamidine
Neoquassin
Nitenpyram
Nuarimol
Ofurace
Omethoate
Oxadixyl
Oxamyl

Paclbutrazol
Penconazole
Pencycuron
Phenmedipham
Phenthoate
Phorate
Phorate sulphone
Phorate sulphoxide
Phosphamidon
Phoxim
Picolinafen
Picoxystrobin
Piperonyl butoxide
Pirimicarb
Pirimicarb desmethyl
Prochloraz
Profenofos
Promecarb
Prometryn
Propamocarb (free base)
Propaquizafop
Propiconazole
Propyzamide
Propoxur
Prosulfuron
Prothioconazole-desthio
Pymetrozine
Pyraclostrobin
Pyrethrins I
Pyrethrins II
Pyrimethanil
Pyriproxyfen
Quassin
Quinmerac
Quinoxifen
Rimsulfuron
Rotenone
Spinosad A

Spinosad D
Spiromesifen
Spirotetramat
Spirotetramat BYI08330 enol-glucoside
Spirotetramat BYI08330 ketohydroxy
Spirotetramat BYI08330 monohydroxy
Spirotetramat BYI08330enol
Spiroxamine
Sulcotrione
Tebuconazole
Tebufenozide
Tebufenpyrad
Tebuthiuron
Teflubenzuron
Terbufos
Terbufos sulfone
Terbufos sulfoxide
Tetraconazole
Thiabendazole
Thiacloprid
Thiamethoxam
Thiodicarb
Thiophanate-methyl
Tolfenpyrad
Triclopyr
Triadimefon
Triadimenol
Triasulfuron
Triazamate acid
Triazophos
Tricyclazole
Trifloxystrobin
Triflumizole
Triflumuron
Triforine I
Triforine II
Triticonazole
Zoxamide

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