

MULTIRESIDUE PESTICIDE ANALYSIS IN FRUIT AND VEGETABLE COMMODITIES USING BOTH UPLC AND APGC ON A SINGLE MASS SPECTROMETER PLATFORM

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INTRODUCTION

Hundreds of pesticides are commercially available and approved for use on various crops making pesticide residue analysis an important component of ensuring food safety. Maximum Residue Levels (MRLs) are set at the highest level of pesticide that would be expected to be found in that crop when it has been treated in line with good agricultural practice. National authorities control and enforce MRLs by testing samples for pesticide residue levels using analytical surveillance programs. These programs check for compliance with MRLs, assess dietary exposure, and check for use of unauthorized pesticides. The food industry also carries out its own due diligence analyses. With the large amount of pesticides available for use, it is important to be able to rapidly and reliably analyze samples for many pesticides in a single method. Mass spectrometry coupled with both gas (GC) and liquid chromatography (LC) is needed to provide comprehensive analysis of a wide range of pesticide residues with sufficient sensitivity to meet global MRL regulations. The use of Quick, Easy, Cheap, Efficient, Rugged and Safe (QuEChERS) sample extraction and clean up has streamlined analytical efficiencies for multi residue analyses.¹ The advantage of ultra performance liquid chromatography (UPLC) coupled with tandem quadrupole mass spectrometry (MS/MS) for multi residue pesticide analysis is widely reported.² More recently the use of GC-MS/MS operated at atmospheric pressure (APGC) has been shown to offer significant improvements in performance over electron impact (EI) for challenging pesticides, in terms of selectivity, specificity and speed of analysis.^{3,4} For this analysis, a single workflow for multi residue analysis of pesticides is demonstrated on a variety of fruit and vegetable samples on the same tandem quadrupole MS instrument, with less than 30 minutes needed to switch between chromatographic inlets. The performance of the method is highlighted in terms of sensitivity, repeatability, and linearity for both LC and GC in compliance with the SANTE guidelines (11945/2015) for pesticide analysis.⁵



Figure 1. Xevo TQ-S micro with the Universal Source allowing for UPLC and APGC analysis on the same mass spectrometer.

RESULTS AND DISCUSSION

Fast Scanning Speeds of Xevo TQ-S micro

Each analysis method contained approximately 200 pesticide residues acquiring at least 2 MRM transitions per compound. The mass spectrometer must be able to rapidly scan through all of the MRM channels to acquire the required data (Figure 2).

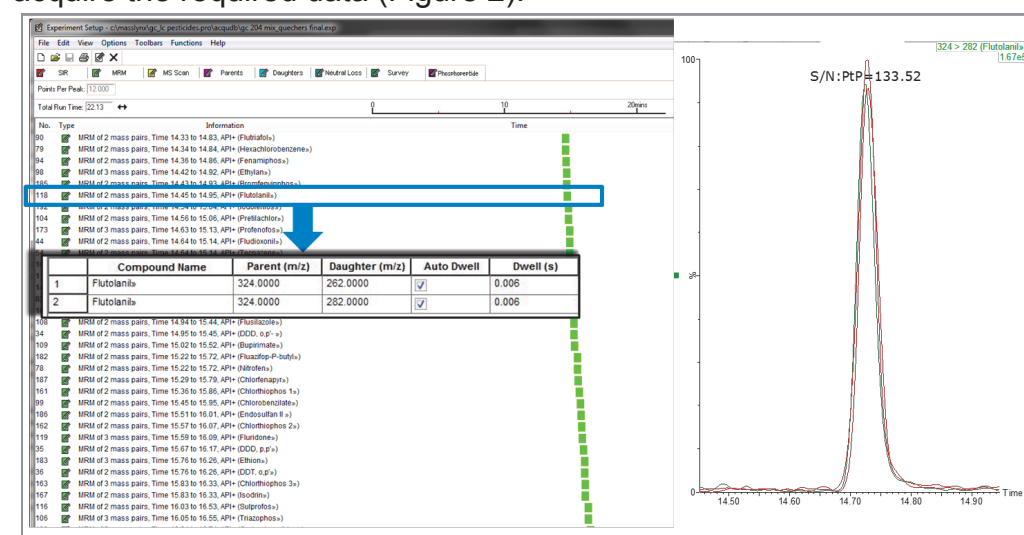


Figure 2. The fast scanning of the Xevo TQ-S micro collects the required data points (minimum 12 points per peak) while maintaining peak quality.

Pesticides in Matrix

Matrix matched standards were prepared in matrix over a range of 0.001 - 0.050 mg/kg. Replicate injections made using the UPLC and APGC methods. In accordance with the SANTE/11945/2015 guidelines, the data were fitted with the best fit calibration; for the UPLC data, the response was shown to be linear whereas the APGC response was best fitted with a quadratic calibration.⁵ (Figure 3). A majority of the compounds had correlation coefficient (R^2) values of 0.995 or greater with residuals from triplicate injections within $\pm 20\%$.

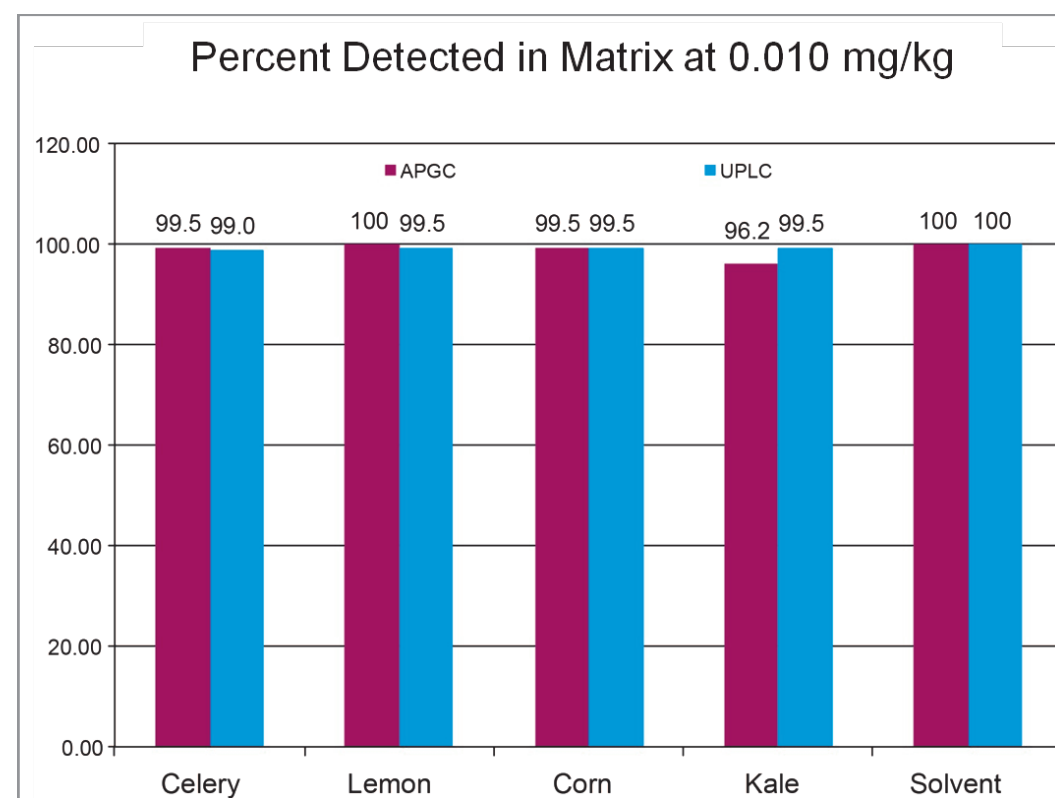


Figure 4. Percentage of pesticides detected in the 0.010 mg/kg standard for each matrix using both APGC and UPLC.

Figure 4 demonstrates the percentage of pesticides in each method detected in the spiked matrices at 0.010 mg/kg, but many pesticides could also be detected at 0.001 mg/kg. The precision of the measurements was excellent with > 90% of the detected pesticides exhibiting RSDs less than 10% (n=3), except the APGC kale analysis which had > 80% of pesticides exhibiting RSDs less than 10% (Figure 5). Ion ratios in matrix were also shown to be within 30% tolerance of the reference values.

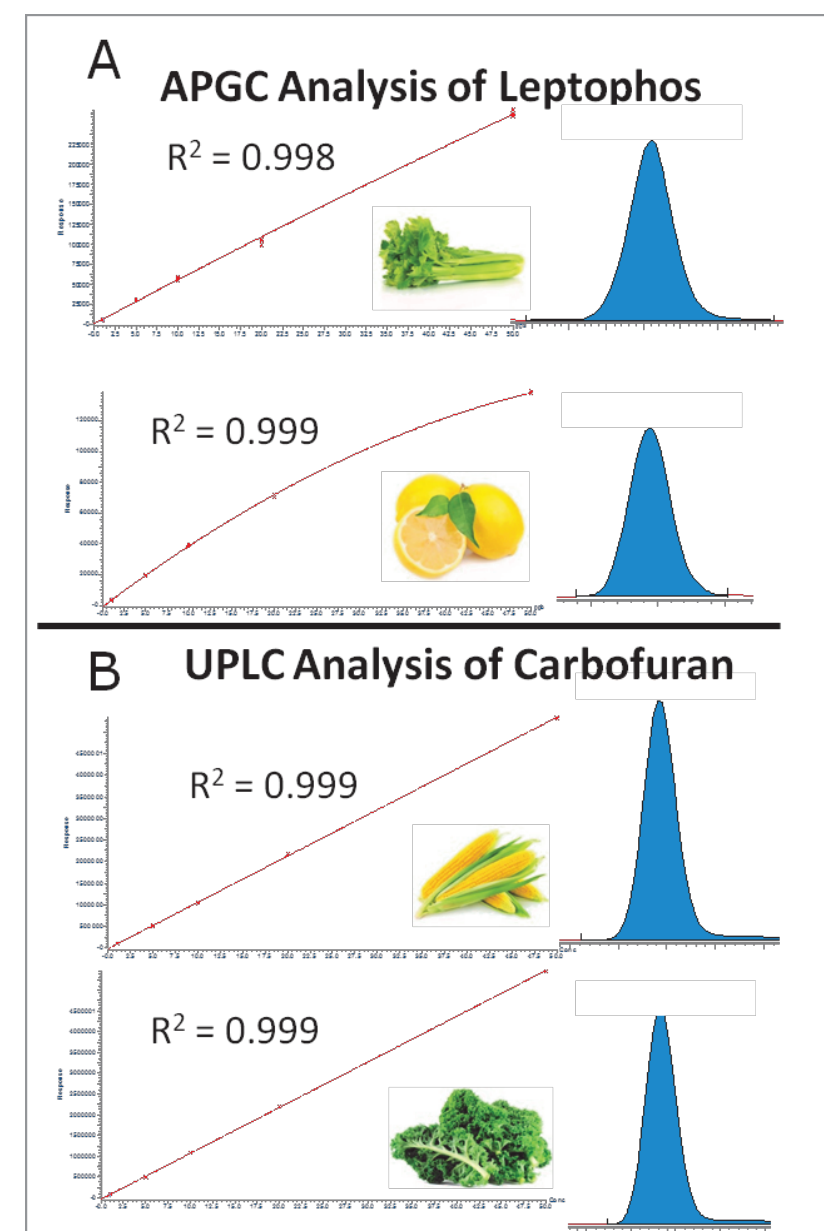


Figure 3. Matrix matched calibration curves and chromatograms for standards at 0.001 mg/kg. (A) APGC analysis of leptophos in celery and lemon, (B) UPLC analysis of carbofuran in corn and kale.

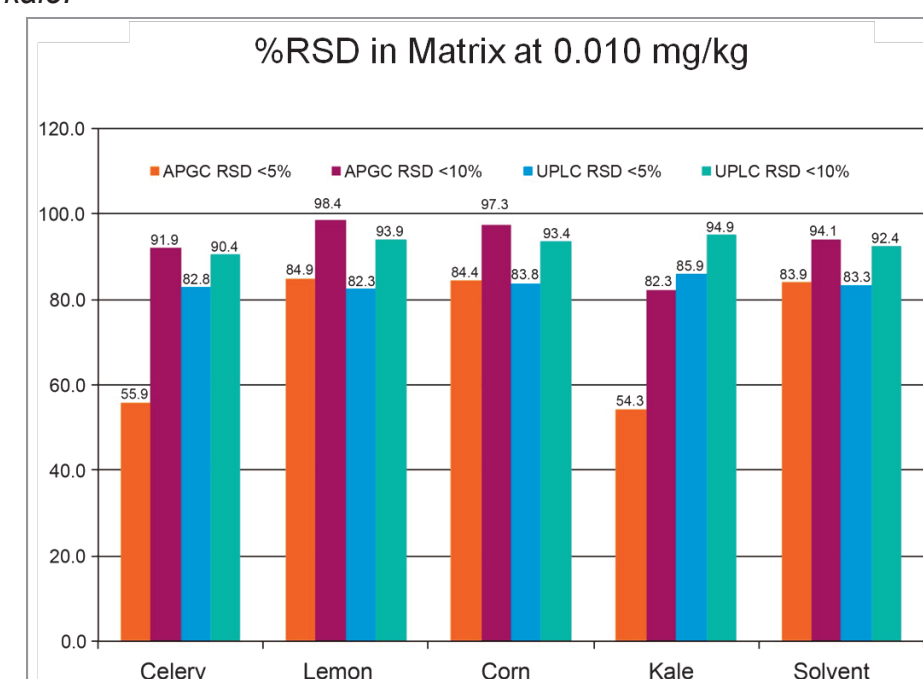


Figure 5. Percentage of compounds detected at 0.010 mg/kg at respective RSDs.

METHODS

Commodities Studied

Four different classes of food commodities were studied:



Lemon - high acid content



Celery - high water content



Kale - high water content



Corn - high starch/protein content

Pesticide Residue Classes

Each analysis method (APGC and UPLC) contained approximately 200 pesticide residues from various classes.

UPLC pesticide classes:

- Organophosphorus
- Carbamates
- Organonitrogen

APGC pesticide classes:

- Organophosphorus
- Organochlorine
- Organonitrogen
- Pyrethroids
- Herbicide Methyl Esters

Method Generation and Management using QuanPedia

MS Methods, TargetLynx Processing Methods, and UPLC Methods were generated and maintained using QuanPedia databases.

For full method details, see Waters application note 720006013en

UPLC-MS/MS Parameters

LC System: ACQUITY UPLC H Class

Column: ACQUITY BEH C18 2.1 x 100 mm, 1.7µm

Mobile Phase A: Water + 10 mM ammonium acetate

Mobile Phase B: Methanol + 10 mM ammonium acetate

MS System: Xevo TQ-S micro

Ionization Mode: ESI+

Sample Extraction and Cleanup

Weigh 15 g of homogenised sample

Add 15 mL acidified acetonitrile (1% acetic acid)

Add pouch of AOAC DisQuE salts

Shake 1 min, centrifuge for 5 min at 4000 rpm

Perform DisQuE cleanup according to Table 1

Shake for 1 min, centrifuge for 5 min at 6000 rpm

For LC analysis: 100 µL of supernatant + 400 µL water

For GC analysis: Evaporate 100 µL supernatant and reconstitute to 100 µL in hexane

Sample	MgSO ₄	PSA	GCB	Volume
Celery	150 mg	25 mg	7.5 mg	1 mL
Lemon	150 mg	25 mg	-	1 mL
Corn	150 mg	25 mg	-	1 mL
Kale	900 mg	150 mg	150 mg	6 mL

Table 1. dSPE cleanup conditions used for each sample matrix.

APGC-MS/MS Parameters

GC System: 7890A

Autosampler: CTC PAL RTC

Column: 30 m x 0.25 mm x 0.25 µm Rxi-5MS

MS System: Xevo TQ-S micro

Ionization mode: API+

Ionization mechanism: Proton transfer (3 vials of water in source)

CONCLUSIONS

- Complex multi residue pesticide analysis was demonstrated using both UPLC and APGC analysis on the same tandem quadrupole instrument
- Instrument methods were generated and maintained using QuanPedia databases making method generation and maintenance fast and simple.
- The reliable scanning speed of the TQ-S micro produced accurate and precise measurements for large multi residue methods.
- The performance of the analysis complied with the SANTE guidelines for pesticide residue analysis.
- Detection at the default maximum residue limit of 0.010 mg/kg was easily achieved for > 99% of pesticides analyzed with good precision (RSDs < 10%) for most analytes in the food samples.
- The flexibility of the Universal Source architecture provides access to both UPLC-MS/MS and GC-MS/MS on the same instrument, allowing for an increase of laboratory efficiency, while maintaining required sensitivity and repeatedly.

References

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