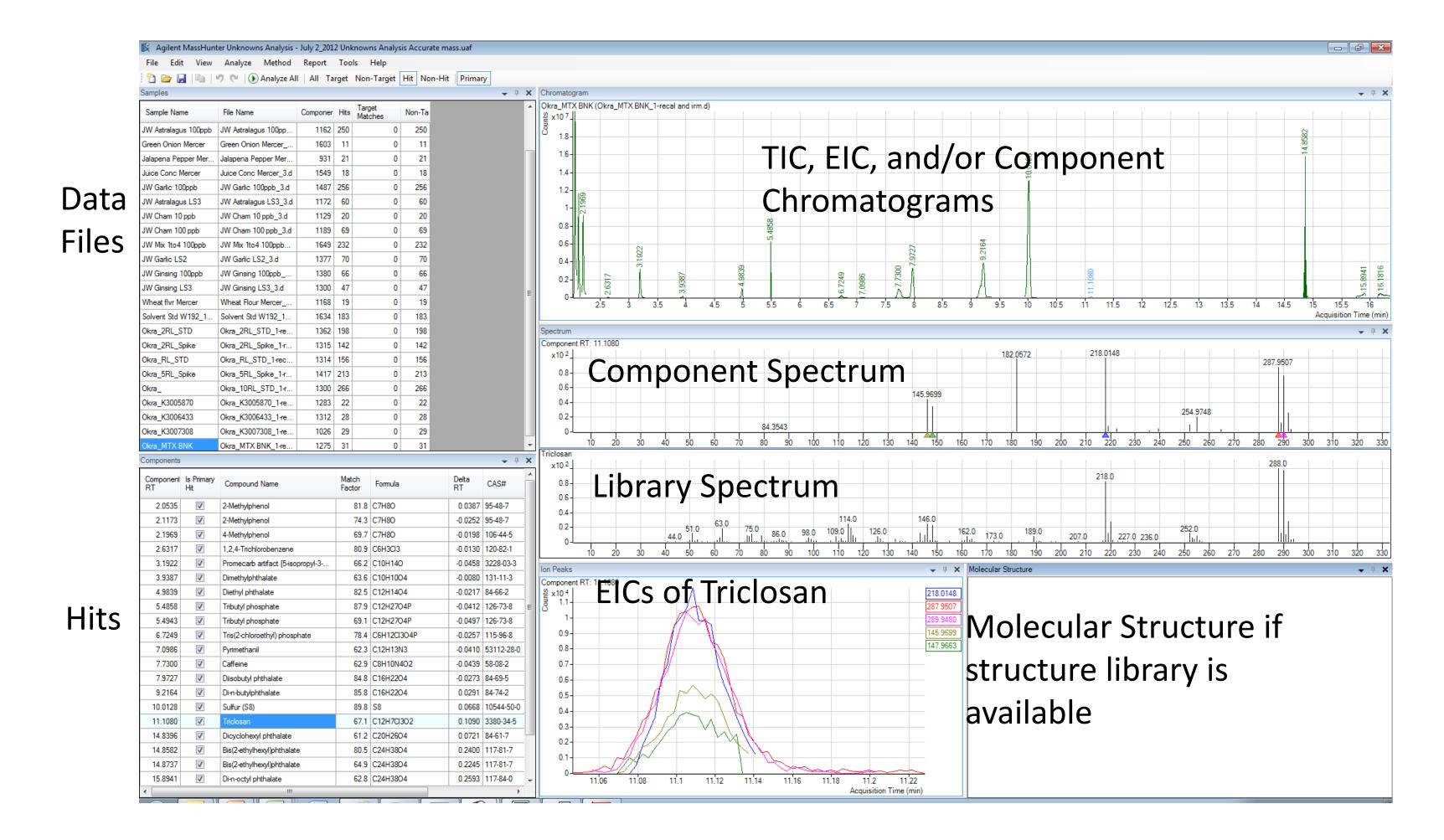
Finding Pesticides using an Agilent 7200 GC/Q-TOF and Unknowns Analysis Software <u>Philip L. Wylie¹</u> and Chris Sandy², (1) Agilent Technologies, Wilmington, DE, USA (phil_wylie@agilent.com) (2) Agilent Technologies, Wokingham, Berkshire, GB

Introduction

With increased international trade in food and food ingredients, there is even more emphasis on food safety. GC with triple quadrupole mass spectrometry (GC/QQQ) is the gold standard for the target analysis of volatile contaminants. But, if a residue is present that is not on the target list, it will be missed. This presentation describes a new tool, GC with a quadrupole time of flight (GC/Q-TOF) mass spectrometer, for the retrospective analysis of unknowns (Figure 1). The TOF provides high resolution and accurate mass measurements.

Instrumentation and Software

Identifying Triclosan in an Okra "Blank"



GC	Agilent 7890A
Mass Spectrometer	Agilent 7200 Q-TOF in TOF mode
Autosampler	Agilent 7693A
GC Column	Agilent 15m X 0.25mm X 0.25um HP-5MS UI
Oven Temp Program	70°C (1 min), 50°C/min to 150°C (0 min), 6°C/min to 200°C (0 min), 16°C/min to 280°C (5 min)
Retention Time Locking	Chlorpyrifos-methyl locked to 8.298 min (constant pressure)
Software	MassHunter Acquisition, Qualitative Analysis, Quantitative Analysis, and Unknowns Analysis

Figure 2. One of the hits is Triclosan, a disinfectant used in soaps, toothpaste, deoderants, and many other consumer products. Residues are widespread in the environment.

Figure 3. Use MassHunter's Accurate Mass Calculator to find actual monoisotopic accurate mass

lass Calculator							
Base formula (M)	-	Species		Calc m/z 🧹	Mono m/z	Diff (ppm)	Defect
C12 H7 Cl3 O2 -			M⁺+	287.9506	287.9506	-0.3	-0.04
Species to calculate							
 Positive ions Negative ions 							
Neutral							
Radical							
+H +Na	-						
⊨ +К	ш						
+ ×							
Number of charges: 1							
Number of charges: 1							
Mass comparison	_						
Comparison m/z: 287.9507	-						
III	•						

- Accurate mass
- High resolution
- MS/MS capability
- Removable ion source
- High dynamic range
- Internal reference mass
- Fast scanning



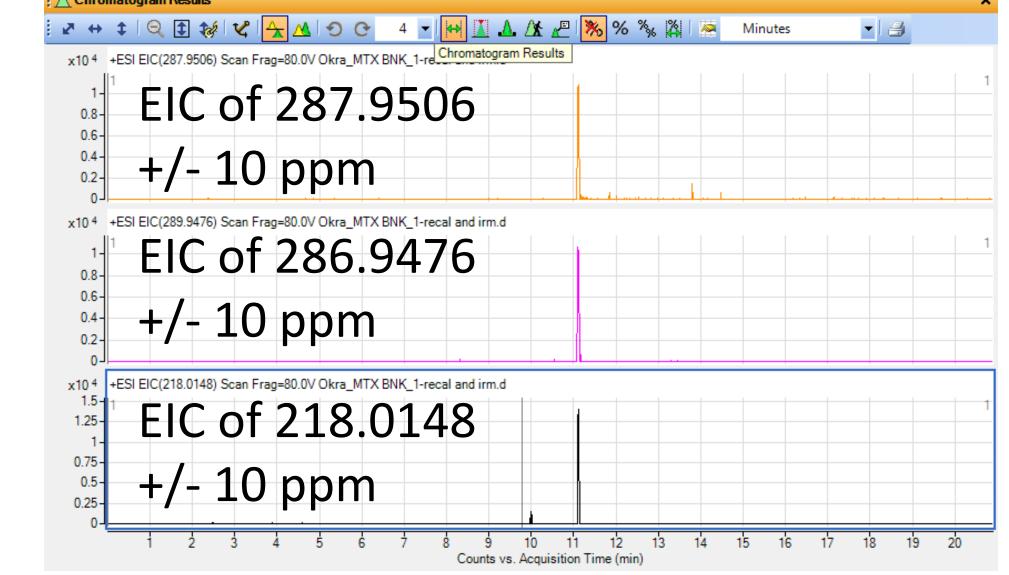
Figure 1. Agilent 7200 GC/Q-TOF

Unknowns Analysis

- 1. Analyze sample by GC/TOF
- 2. Run quant method first or go directly to Unknowns Anal.

Figure 4. EICs for the monoisotopic molecular ion and the molecular ion with one ³⁷Cl. *m/z* 218.0148 was the measured

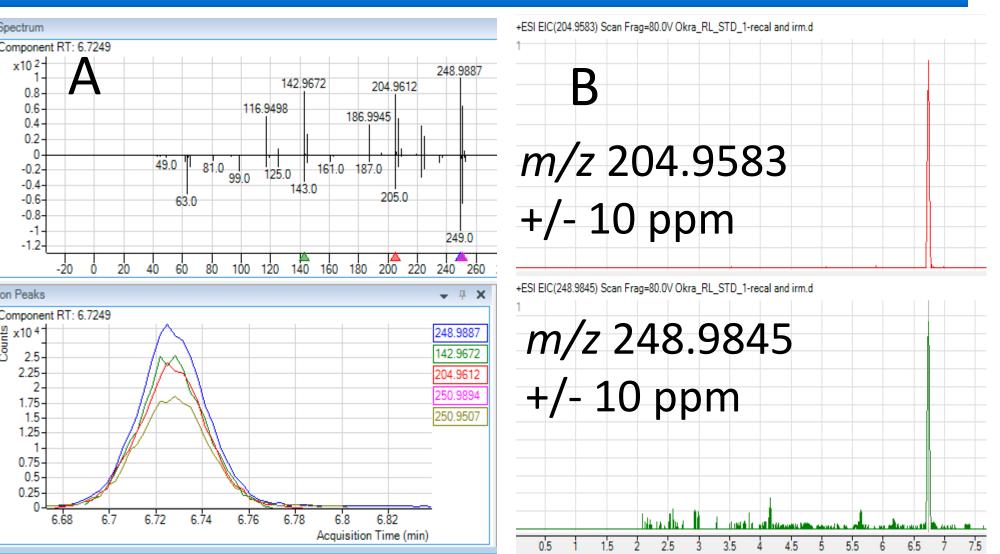
value.



Identifying a Flame Retardant in an Okra Blank

Tris(2-chloroethyl) phosphate (TCEP) was identified by the Unknowns Analysis software in the same Okra "blank." TCEP is carcinogenic and has shown broad toxicity in animal models.

Figure 5. A) Comparison of the spectrum & EICs for the decon-



- 3. Set up method parameters for deconvolution, library searching and hit matching
- 4. Analyze samples
 - 1. Deconvolute chromatogram(s)
 - 2. Library search all components
 - 1. Use Agilent RT-Locked Pesticide and Endocrine Disruptor Library
 - 2. Match library retention times to filter out false positives
 - 3. Report identified compounds
 - 4. Display results as shown in Figure 2
- 5. Review Unknowns Analysis results
- 6. Confirm in MassHunter Qual by extracting out accurate masses of identified compounds.

voluted component identified as TCEP. B) Two accurate mass EICs for TCEP



- MassHunter Unknowns Analysis software is a simple, but powerful tool to identify unknowns, i.e., compounds not on the target list but in a mass spectral library.
 Accurate mass offers high selectivity and the ability to calculate formulas for the molecular ion and fragments.
- 3. MS/MS can aid in true unknown characterization.