

**Title: Determination of Selected Pesticides Collected on XAD-4 Resin by High Performance Liquid Chromatography Ion Trap Mass Spectrometry and Gas Chromatography Mass Spectrometry**

1. Scope

This section method (SM) is for the analysis of the selected pesticides trapped in XAD-4 resin and extracted with ethyl acetate. These pesticides are azinphos-methyl, chlorothalonil, chlorpyrifos, cypermethrin, DEF, diazinon, dicofol, dimethoate, diuron, endosulfan, endosulfan sulfate, EPTC, malathion, metolachlor, molinate, DDVP, norflurazon, oryzalin, oxyfluorfen, permethrin, phosmet, propanil, propargite, simazine, thiobencarb, trifluralin, and the oxygen analogs of chlorpyrifos, diazinon, dimethoate and malathion. It is to be followed by all authorized section personnel. The reporting limits vary from 0.25  $\mu\text{g}$  to 2.0  $\mu\text{g}$  depending on the instrument sensitivity for individual compounds.

2. Principle:

Residues of the selected pesticides are extracted from XAD-4 resin using ethyl acetates. Nineteen compounds are determined by the injection of sample extract into an HPLC equipped with a C-18 column and a mass spectrometer (LC-MS). The remaining compounds, which cannot be confidently analyzed with LC-MS, are determined by the injection of sample extract into a GC equipped with a mass selective detector (GC-MSD). The confirmation of compound identity on LC-MS is achieved simultaneously with collision-induced dissociation to produce a product ion for each of the analytes. The confirmation of compound identity with GC-MSD is achieved by the ratio of selected ions.

3. Safety:

- 3.1 All general laboratory safety rules for sample preparation and analysis shall be followed.
- 3.2 All solvents should be handled with care in a ventilated area.

4. Interferences:

On LC-MS it appears that the extract enhances the response of phosmet and azinphos methyl most of the time. On GC-MS it appears that the extract deteriorates the analytical column quickly and reduces the response of some compounds. The response and peak shape of some compounds in matrix extracts versus in plain

solvent shows great differences. To be consistent we have prepared standards in matrix extract to be used in GC-MS and LC-MS analysis.

5. Apparatus and Equipment:

- 5.1 Rotary evaporator (Büchi/Brinkman or equivalent)
- 5.2 Nitrogen evaporator (Meyer N-EVAP Organomation Model # 112 or equivalent)
- 5.3 Vortex-vibrating mixer
- 5.4 Separatory funnel, 250 mL
- 5.5 Conical tube with glass stopper, 15-mL graduated
- 5.6 Boiling flask, 500-mL
- 5.7 Funnel, 15 cm diameter
- 5.8 Disposable Pasteur pipettes, and other laboratory ware as needed
- 5.9 Liquid chromatograph (Waters Model 2695 HPLC) equipped with a Thermo Finnigan Ion Trap Mass detector.
- 5.10 Gas chromatograph (Agilent Model 6890) equipped with a mass spectrometer (Agilent model 5972)

6. Reagents and Supplies: (All reagents shall meet the minimum requirement in HPLC, residue and pesticide analysis)

- 6.1 Acetic acid,
- 6.2 Ammonium formate, (Aldrich, reagent grade or equivalent)
- 6.3 Ethyl acetate, (Fisher, reagent grade or equivalent)
- 6.4 Formic acid, HPLC grade (Fisher #A35-500 or equivalent)
- 6.3 Methanol, (Burdick & Jackson, MS grade, or equivalent)
- 6.4 Nitrogen, refrigerated liquid or nitrogen generator with capacity of delivering 20 liters per minute
- 6.5 Standards: The individual 1.0 mg/mL stock standards of each compound were obtained from the CDFCA/CAC Standard Repository.

Azinphos-methyl	CAS Number 86-50-0
Chlorothalonil	CAS Number 1897-45-6
Chlorpyrifos	CAS Number 2921-88-2
Cypermethrin	CAS Number 52315-07-8
DEF	CAS Number 78-48-8
Diazinon	CAS Number 333-41-5
Dicofol	CAS Number 115-32-2
Dimethoate	CAS Number 60-51-5
Diuron	CAS Number 330-54-1
Endosulfan $\alpha$	CAS Number 959-98-8

Endosulfan $\beta$	CAS Number 332-13-65-9
Endosulfan sulfate	CAS Number 1031-07-08
EPTC	CAS Number 759-94-4
Malathion	CAS Number 121-75-5
Metolachlor	CAS Number 51218-45-2
Molinate	CAS Number 2212-67-1
DDVP	CAS Number 62-73-7
Norflurazon	CAS Number 27314-13-2
Oryzalin	CAS Number 19044-88-3
Oxyfluorfen	CAS Number 42874-03-3
Permethrin	CAS Number 52645-53-1
Phosmet	CAS Number 732-11-6
Propanil	CAS Number 709-98-8
Propargite	CAS Number 2312-35-8
Simazine	CAS Number 122-34-9
Thiobencarb	CAS Number 28249-77-6
Trifluralin	CAS Number 1582-09-8
Chlorpyrifos OA	CAS Number 0000-00-00
Diazinon OA	CAS Number 962-58-3
Dimethoate OA	CAS Number 1113-02-6
Malathion OA	CAS Number 0000-00-00

- 6.6 Water, MS grade, Burdick & Jackson or equivalent
- 6.7 Analytical column: Waters SymmetryShield RP<sub>18</sub> 5  $\mu$ m, 3.9 x 150 mm column (part number, 186000108) or equivalent
- 6.8 Guard column: Waters Symmetryshield RP 18 5  $\mu$ m, 3.9 x 20 mm cartridge (part number,
- 6.9 Guard column cartridge holder: Waters Sentry guard holder universal (P/N wat064610)

## 7. Standards Preparation:

- 7.1 Dilute the 1.0 mg/mL standards, obtained from the CDFA/CAC Standards Repository, with methanol. The concentration of each individual standard is 20 ng/ $\mu$ L or 10 ng/ $\mu$ L.
- 7.2 The CDFA/CAC Standards Repository prepares the combination standards. One contains 26 compounds without the oxygen analogies (A). The other contains the 4 oxygen analogies (B). Each component of the combinations is 100  $\mu$ g/mL prepared in methanol.

- 7.3 A combined stock solution is prepared by mixing equal amount of (A) and (B). This combined stock solution is for preparing working standards and QC spiking.
- 7.4 Working standards are prepared in sample matrix solution. The dilution ratios are:

Concentration (ng/ $\mu$ L)	Amount of stock solution ( $\mu$ L)	Amount of matrix extract ( $\mu$ L)	Final volume (mL)
0.10	4	1996	2.0
0.50	20	1980	2.0
1.0	40	1960	2.0
2.0	80	1920	2.0
5.0	200	1800	2.0

Working standards should be kept no longer than a week.

8. Sample Preservation and Storage:

All samples and sample extracts shall be stored in the refrigerator (0-5 °C).

9. Test Sample Preparation:

9.1 Sample Preparation

- 9.1.1 Remove samples from refrigerator and allow them to reach ambient temperature.
- 9.1.2 Clip the sample tube on a laboratory rack in a hood, remove the stopper from the ends, put a 250 mL round flask under it, and place the bottom of the sample tube just inside the flask.
- 9.1.3 Set a 250 mL separatory funnel as a solvent reservoir on a ring and place the tip of the funnel just inside of the sample tube.
- 9.1.4 Pour 100 mL ethyl acetate into the separatory funnel. Open the stopcock and adjust the flow to a rate that the top of the resin bed maintains about 0.5 inch of solvent.
- 9.1.5 Allow the entire 100mLs of solvent to run through the sample tube.

- 9.1.6 Evaporate the solvent to about 10 mL on a rotary vacuum evaporator at 45°C and 24 inches vacuum.
- 9.1.7 Quantitatively transfer the solution to a 15 mL conical centrifuge tube and evaporate on a N-evap at 40°C to about 1 mL.
- 9.1.8 Perform solvent exchange by adding 4 mL methanol and evaporate to about 1 mL.
- 9.1.9 Adjust final volume to 2.0 mL with methanol. Vortex for 20 seconds and transfer to three auto-sampler vials, two with insert for analysis and one without insert for storage for in case of re-analysis.

## 10 Instrument Calibration:

- 10.1 The calibration standard curves consist of six levels. The lowest level must be at or below the corresponding reporting limits. (The current working standard levels are 0.02, 0.1, 0.5, 1.0, 2.0, and 5.0 ng/μL as prepared in 7.3.
- 10.2 The calibration curves for the LC-MS and GC-MS are generally obtained using linear regression. Quadratic fit may be used if the response of certain compounds exhibited quadratic behavior.

## 11 Analysis:

### 11.1 Injection Scheme

Both LC-MS and GC-MS may need to be conditioned with a matrix standard or a sample extract 2 to 5 runs before running the following sequence: A set of calibration standards in matrix, a matrix blank, a matrix spike, a set of up to 10 test samples, then a set of standards in matrix, etc.

### 11.2 HPLC-MS Instrumentation

- 11.2.1 Waters model 2695 HPLC and auto-sampler with column heater and remote control through Thermo Finnigan Xcalibur system.
- 11.2.2 Column: Waters SymmetryShield RP<sub>18</sub>, 5 μm, 3.9 x 150 mm column (part no. 186000108)

11.2.3 Guard column cartridge Holder and cartridge: Waters Sentry guard holder universal (P/N wat064610); Waters SymmetryShield RP 18 5 $\mu$ m, 3.9 x 20mm.

11.2.4 Column Temperature: 40 °C

11.2.5 Mobile Phase: Gradient  
Solvent C: 3762 mL water, 200 mL methanol, 38 mL 1 M ammonium formate and 4.0 mL formic acid  
Solvent D: 3600 mL methanol, 360 mL water, 36 mL 1.0 M ammonium formate, 4 mL formic acid  
Flow rate: 0.75 mL/min  
Gradient:

Time(min)	Flow rate (mL/min)	A	B	C	D
0	0.75	0	0	85	15
3	0.75	0	0	85	15
4	0.75	0	0	50	50
10	0.75	0	0	50	50
21	0.75	0	0	25	75
22	0.75	0	0	5	95
27	0.75	0	0	5	95
30	0.75	0	0	85	15
34	0.75	0	0	85	15

11.2.6 Injection Volume: 10  $\mu$ L

11.2.8 Liquid Chromatograph Mass spectrometer (LC-MS) and Operating Parameters

Model: Finnigan Model DECA ion trap MS  
Ion Source Type: Atmospheric pressure Ionization (APCI)  
Source Polarity: Positive  
APCI Vaporizer Temp: 500°C  
Capillary Temperature: 200°C  
Sheath Gas: 44  
Auxiliary Gas: 3  
Mode of Operation: MS/MS  
Retention time, molecular mass, ion filter range and product ions are listed below:

Compound Name	Retention Time (min)	Molecular Masses	Ion Filter Range	Product Ions
Dimethoate OA	3.16	213	214±1.5	<u>183</u>
Dimethoate	7.40	229	230±1.5	<u>199</u>
Malathion OA	10.04	314	315±1.5	126.9, 173, 268.8
Simazine	11.55	201, 203	203±1.5	124, 132, 174
Diazinon OA	15.06	288	289±1.5	153, 261, 289
Norflurazon	16.23	303, 305	305±2	284, 286, 302, 304, 316, 317
Diuron	17.06	232, 234	235±2.5	72
Phosmet	17.40	317	318±1	<u>160</u> , 285.7
Azinphos-methyl	17.18	317	318±1	160, 171, <u>261</u>
Molinate	18.05	187	188±1	<u>126</u>
Malathion	18.56	330	334±6	<u>285</u>
Metolachlor	19.59	283, 285	284±2	<u>252</u>
Propanil	19.63	217, 219	218±2.5	<u>162</u>
Chlorpyrifos OA	20.35	333, 335, 337	334±6	306, 308, 310
EPTC	20.55	189	190±1	128
Oryzalin	21.53	346	347±1	247, 288, 305
Diazinon	22.39	304	305±1.5	153, 169
Thiobencarb	24.43	257, 259	258±2.5	100, 125, 258
DEF	27.00	314	315±2.5	200.9, 257, 259

Note: The real retention times are expected within 30 seconds of that stated above when the column is new. The column conditions, temperature, mobile phase, etc. may slightly shift retention time.

11.2.9 Operating parameter detail and Tune method are listed in Appendix 1 and Appendix 2

11.3 GC-MSD Instrumentation:

11.3.1 Agilent GC-MSD model HP6890 with auto sampler. Operating software is Enviroquant ChemStation version G1701B.01.00

11.3.2 Column: HP-5MS 30m x 0.25mm x 0.25µm

11.3.3 Pre-column: not used

11.3.4 Temperature program

Injector Temperature: 250 °C

Oven Temperature:

Oven Ramp	Program (°C/min)	Temperature (°C)	Hold (min)
initial		70	2
Ramp 1	15	190	5
Ramp 2	15	250	5
Ramp 3	15	270	8

### 11.3.5 Retention times and ions selected for SIM acquisition:

Compound name	Retention time	Selected ions	Starting time
DDVP	7.25	109,145,185	5.0
EPTC*	8.35	128,189,86	8.10
Molinate*	10.01	126,187,98	9.80
Trifluralin	11.02	264, 306,264, 335	10.8
Dimethoate*	11.86	87, 125, 229	11.80
Diazinon*	12.58	179, 304, 137	12.50
Chlorothalonil	12.76	266, 229, 109	12.50
Propanil*	14.07	161,163,217	14.00
Malathion*	15.83	173, 125, 93	15.75
Metolachlor*	15.89	162, 238, 146	15.75
Chlorpyrifos	15.98	314, 197, 258	15.75
Thiobencarb *	16.04	100, 257, 125	15.75
Dicofol p, p'	16.46	139, 250, 111	16.28
Endosulfan 1	18.02	195, 339, 241, 261	17.80
DEF*	18.61	202, 169, 147	18.50
Oxyfluorfen	18.68	252, 300, 361	18.50
Endosulfan Sulfate	20.04	272, 387, 229, 422	19.90
Propargite	20.5-20.6	135, 173, 350	20.45
Phosmet	21.31	160, 77, 93	20.60
Azinphos Methyl	22.56	160, 77, 132	20.60
Permethrin	24.8,25.09	183, 163, 127	24.80
Cypermethrin	26.41	163, 161,209	26.00

\*Can be analyzed by LCMS confidently

## 12. Quality Control:

- 12.1 Each set of samples shall have a matrix blank and minimum of one matrix spike sample. Each set contains up to 12 samples.



- 12.2 The matrix blank shall be free of target compounds.
- 12.3 The recoveries of the matrix spike should be within the control limits.
- 12.4 The retention time shall be within  $\pm 20$  seconds of that of the standard.
- 12.5 To maintain the instrument performance, trim the front 10 inches off the GC column every 30 hours (about 50 injections). The response of the 0.02 ng/uL combination standards is a good measure of the instrument performance.
- 12.6 The sample extract shall be diluted if results fall outside the linear range of the standard curve.
- 12.7 Add additional levels if there is a need to extend the standard curve range.
- 12.8 Method Detection Limits (MDL)

The method detection limit refers to the lowest concentration of analyte that a method can detect reliably. To determine the MDL, 7 replicate XAD-4 sample tubes are spiked at 0.2  $\mu\text{g}$  or 0.4  $\mu\text{g}$ . The standard deviation from the spiked sample recoveries are used to calculate the MDL for each analyte using the follow equation:

$$\text{MDL} = tS$$

Where t is the Student t test value for the 99% confidence level with n-1 degrees of freedom and S denotes the standard deviation obtained from n replicate analyses. For the n=7 replicate used to determine the MDL, t=3.143.

- 12.9 Reporting limit (RL):

The reporting limit (RL) refers to the level at which reliable quantitative results may be obtained. The MDL is used as a guide to determine the RL. In general, the RL is chosen in a range 1-5 times the MDL. The response reproducibility of each compound is also considered to determine the RL

MDL data and the RL are tabulated in Appendix 3 and 4.

- 12.10 Method Validation Recovery Data and Control Limits:

12.10.1 The method validation consisted of five sample sets. Each set included 4 levels of fortification (0.5, 1.0, 5.0 and 10.0 µg) and a method blank. All spikes, method blank and samples were processed through the entire analytical method.

12.10.2 Upper and Lower warning and control limits are set at  $\pm 2$  and  $\pm 3$  standard deviations of the average % recovery, respectively.

12.10.3 Method validation results and control limits are tabulated in appendix 4.

### 13. Calculations:

13.1 The quantification is based on the sum of area counts of the product ion and the precursor of the compound analyzed. The calculation is based on external standard (ESTD).

13.2 The software LCQuan in the Xcalibur system and the software Enviroquant are used to establish the standard curve and to calculate the analytes in the samples. The correlation coefficient, slope, intercept of the linear regression line are calculated once the calibration standards are defined. The equation for calculating analytes is as follows:

$$y = mx + b$$

Where: y = peak response  
m = slope  
b = intercept  
x = concentration of compound

When the unit and the dilution factor are entered correctly in the analysis sequence, the software will then correctly generate the results.

13.3 Results can be manually calculated by a single point standard. The unit is µg/sample. This calculation is to verify the results derived from the software

The general equation is as follows:

$$\mu\text{g} = \frac{(\text{sample peak area}) (\text{std. conc. ng}/\mu\text{L}) (\text{std. vol. Injected}(\mu\text{L})) (\text{sample final vol.}, (\text{mL}))}{(\text{std. peak area}) (\text{sample vol. Injected } (\mu\text{L}))}$$

#### 14. Reporting Procedure:

##### 14.1 Perform Quantification with LCQuan:

###### 14.1.1 Create a new Processing method

- Open a raw file
- Select calibration options
- Identify components
- Define calibration settings. Or

###### 14.1.2 Open an existing quantification method and save to an appropriate sub-directory with a new name

###### 14.1.3 Open the sequence and review the sequence. Or

###### 14.1.4 Go to the appropriate sub-directory and select the raw files to be used as standards. Place the standard raw files to the appropriate calibration levels. Select the unknown raw files to be calculated.

###### 14.1.5 Review the calibration

###### 14.1.6 Review all calculated results

###### 14.1.7 Create, review, and print peak integration report, calibration report, and summary report

##### 14.2 Perform Quantification with EnviroQuant:

###### 14.2.1 Load a standard data file

- Integrate the data file
- Edit compounds based on retention time and identity
- Review the window range of each compound and adjust it as needed.
- Reintegrate the data file based on the new method
- Update levels
- View the calibration curves
- Save as a new method

- 14.2.2 Load a sample data file
  - Do quantification with this new method with new calibration curves
  - Review each compound and do integration correction
  - Save this reviewed file
  - Print this reviewed data file

#### 14.3 Acceptance Criteria:

- 14.3.1 Peak retention time between standards, QC spikes and unknowns shall be within 20 seconds. If there is a known reason for retention time shifting, an explanation memo shall be included.
- 14.3.2 Peak response shall be within the calibration range
- 14.3.3 The  $R^2$  of calibration curve or overlay calibration curves shall be 0.990 or better.
- 14.3.4 Recoveries of spike QC shall be within the established control range, otherwise a rerun of the entire set shall be performed. If problems remain, an explanation memo shall be included.
- 14.3.5 The ratio of product ion and precursor ion between standard and unknown shall be consistent and the variation of the ratio between standard and unknown shall be within  $\pm 20\%$ .
- 14.3.6 Manual single point calculation result is acceptable with explanation

#### 14.4 Reporting:

- 14.4.1 Sample results are reported out according to the client's analytical laboratory specification sheet.
- 14.4.2 Fill out COC, QC sheet, and control chart.
- 14.4.3 Prepare data package. Peer review. Report.

### 15 Discussion

- 15.1 It was a challenging task to analyze chlorothalonil by GC. Since the beginning of this project, we were able to analyze it by GC for a period of 8 months. The

sensitivity was getting better gradually. Later, we injected air sample extracts of the quartz and glass fiber filters. Immediately after this event, the response of chlorothalonil was getting worse. Eventually we were unable to meet the method reporting limit. A lot of attempts failed to restore the response of chlorothalonil by GC-MS method. Finally, a LC-MS method on negative mode was developed. The chlorothalonil molecular mass is 264 (266, 268, etc). During the infusion of chlorothalonil into the LC-MS we observed the ions  $245^-$ ,  $247^-$ , and  $249^-$ . It appears the chlorothalonil molecule fragmented to ions  $245^-$ ,  $247^-$ , and  $249^-$ . Loss of 19 is likely due to a chlorine on the aromatic ring being replaced by a hydroxyl group. Since we have this method we successfully reanalyzed all those samples that did not meet the reporting limit.

15.2 The GC column maintenance: About every 80-100 injections we trim off 15 inches of column from the front end. To compensate the retention time change due to a shorter column we add 10-15 seconds to the initial temperature holding time. Then check the retention time for each compound and make necessary changes.

15.3 Excluding ammonium salt from the mobile phase greatly increase the response of propanil by LC-MS

16. References:

## APPENDIX I Operating parameters

Creator: PAUL LEE

MS Run Time (min): 35.00

Divert Valve: in use during run

Divert Time (min)	Valve State
0.00	To Waste
1.99	To Source
32.41	To Waste

Contact Closure: not used during run

MS Detector Settings:

Acquisition Start Delay (min): 2.00

Real-time modifications to method disabled

Segment 1 Information

Duration (min): 10.76

Number of Scan Events: 3

Tune Method: oryzalin

Scan Event Details:

- 1: Pos ·(230.0)->o(60.0-250.0)  
MS/MS: Amp. 22.0% Q 0.250 Time 30.000 IsoW 3.0
- 2: Pos ·(214.0)->o(55.0-250.0)  
MS/MS: Amp. 25.0% Q 0.250 Time 30.000 IsoW 3.0
- 3: Pos ·(315.0)->o(85.0-350.0)  
MS/MS: Amp. 23.0% Q 0.250 Time 30.000 IsoW 3.0

Segment 2 Information

Duration (min): 2.26

Number of Scan Events: 1

Tune Method: oryzalin

Scan Event Details:

- 1: Pos ·(203.0)->o(55.0-250.0)  
MS/MS: Amp. 36.0% Q 0.250 Time 30.000 IsoW 3.0

Segment 3 Information

Duration (min): 3.18  
Number of Scan Events: 2  
Tune Method: oryzalin

Scan Event Details:

- 1: Pos ·(305.0)->o(80.0-350.0)  
MS/MS: Amp. 36.0% Q 0.250 Time 30.000 IsoW 4.0
- 2: Pos ·(289.0)->o(75.0-350.0)  
MS/MS: Amp. 30.0% Q 0.250 Time 30.000 IsoW 3.0

Segment 4 Information

Duration (min): 1.77  
Number of Scan Events: 4  
Tune Method: oryzalin

Scan Event Details:

- 1: Pos ·(305.0)->o(80.0-350.0)  
MS/MS: Amp. 36.0% Q 0.250 Time 30.000 IsoW 4.0
- 2: Pos ·(235.0)->o(60.0-300.0)  
MS/MS: Amp. 35.0% Q 0.250 Time 30.000 IsoW 5.0
- 3: Pos ·(318.0)->o(85.0-350.0)  
MS/MS: Amp. 20.0% Q 0.250 Time 30.000 IsoW 2.0
- 4: Pos ·(188.0)->o(50.0-250.0)  
MS/MS: Amp. 27.0% Q 0.250 Time 30.000 IsoW 2.0

Segment 5 Information

Duration (min): 0.94  
Number of Scan Events: 3  
Tune Method: oryzalin

Scan Event Details:

- 1: Pos ·(188.0)->o(50.0-250.0)  
MS/MS: Amp. 27.0% Q 0.250 Time 30.000 IsoW 2.0
- 2: Pos ·(334.0)->o(90.0-400.0)  
MS/MS: Amp. 25.0% Q 0.250 Time 30.000 IsoW 12.0
- 3: Pos ·(318.0)->o(85.0-350.0)

MS/MS: Amp. 20.0% Q 0.250 Time 30.000 IsoW 2.0

#### Segment 6 Information

Duration (min): 1.55  
Number of Scan Events: 4  
Tune Method: oryzalin

#### Scan Event Details:

- 1: Pos ·(284.0)->o(75.0-350.0)  
MS/MS: Amp. 36.0% Q 0.250 Time 30.000 IsoW 4.0
- 2: Pos ·(218.0)->o(60.0-250.0)  
MS/MS: Amp. 35.0% Q 0.250 Time 30.000 IsoW 5.0
- 3: Pos ·(334.0)->o(90.0-400.0)  
MS/MS: Amp. 25.0% Q 0.250 Time 30.000 IsoW 12.0
- 4: Pos ·(190.0)->o(50.0-250.0)  
MS/MS: Amp. 27.0% Q 0.250 Time 30.000 IsoW 2.0

#### Segment 7 Information

Duration (min): 2.77  
Number of Scan Events: 4  
Tune Method: oryzalin

#### Scan Event Details:

- 1: Pos ·(347.0)->o(95.0-400.0)  
MS/MS: Amp. 30.0% Q 0.250 Time 30.000 IsoW 2.0
- 2: Pos ·(305.0)->o(80.0-350.0)  
MS/MS: Amp. 30.0% Q 0.250 Time 30.000 IsoW 3.0
- 3: Pos ·(190.0)->o(50.0-250.0)  
MS/MS: Amp. 27.0% Q 0.250 Time 30.000 IsoW 2.0
- 4: Pos ·(334.0)->o(90.0-400.0)  
MS/MS: Amp. 25.0% Q 0.250 Time 30.000 IsoW 12.0

#### Segment 8 Information

Duration (min): 1.57  
Number of Scan Events: 1  
Tune Method: oryzalin



Scan Event Details:

1: Pos ·(258.0)->o(70.0-300.0)  
MS/MS: Amp. 28.0% Q 0.250 Time 30.000 IsoW 5.0

Segment 9 Information

Duration (min): 1.60  
Number of Scan Events: 4  
Tune Method: oryzalin

Scan Event Details:

1: Neg ·(332.0)->o(90.0-400.0)  
MS/MS: Amp. 32.0% Q 0.250 Time 30.000 IsoW 4.0

2: Pos ·(350.0)->o(95.0-400.0)  
MS/MS: Amp. 28.0% Q 0.250 Time 30.000 IsoW 5.0

3: Neg ·(360.0)->o(95.0-380.0)  
MS/MS: Amp. 38.0% Q 0.250 Time 30.000 IsoW 4.0

4: Pos ·(335.0)->o(90.0-400.0)  
MS/MS: Amp. 35.0% Q 0.250 Time 30.000 IsoW 3.0

Segment 10 Information

Duration (min): 3.79  
Number of Scan Events: 2  
Tune Method: oryzalin

Scan Event Details:

1: Pos ·(315.0)->o(85.0-400.0)  
MS/MS: Amp. 32.0% Q 0.250 Time 30.000 IsoW 3.0

2: Pos ·(350.0)->o(95.0-400.0)  
MS/MS: Amp. 28.0% Q 0.250 Time 30.000 IsoW 5.0

Segment 11 Information

Duration (min): 4.81  
Number of Scan Events: 1  
Tune Method: oryzalin

Scan Event Details:

1: Pos ·(350.0)->o(95.0-400.0)  
MS/MS: Amp. 28.0% Q 0.250 Time 30.000 IsoW 5.0

Custom Dependent Data Settings:  
Not enabled

## APPENDIX 2 Tune Method

Capillary Temp (C):	200
APCI Vaporizer Temp (C):	500
Ion Time (ms):	5
Sheath Gas Flow (l):	44
Aux Gas Flow (l):	3
Source Type:	APCI
Injection Waveforms:	Off
AGC:	On

### POSITIVE POLARITY

Source Voltage (kV):	6
Source Current (uA):	5
Capillary Voltage (V):	3
Tube Lens Offset (V):	15
Octapole RF Amplifier (Vp-p):	290
Octapole 1 Offset (V):	-7.75
Octapole 2 Offset (V):	-10
Entrance Lens (V):	-26
InterOctapole Lens Voltage (V):	-50
Trap DC Offset Voltage (V):	-10
Zoom Micro Scans:	5
Zoom AGC Target:	20000000
Zoom Max Ion Time (ms):	50
Full Micro Scans:	3
Full AGC Target:	50000000
Full Max Ion Time (ms):	300
SIM Micro Scans:	3
SIM AGC Target:	20000000
SIM Max Ion Time (ms):	100
MSn Micro Scans:	2
MSn AGC Target:	20000000
MSn Max Ion Time (ms):	1000

### NEGATIVE POLARITY

Source Voltage (kV):	5
Source Current (uA):	80
Capillary Voltage (V):	-47
Tube Lens Offset (V):	-35
Octapole RF Amplifier (Vp-p):	120

Octapole 1 Offset (V): 6.25  
Octapole 2 Offset (V): 12  
InterOctapole Lens Voltage (V): 68  
Entrance Lens (V): 52  
Trap DC Offset Voltage (V): 10  
Zoom Micro Scans: 5  
Zoom AGC Target: 20000000  
Zoom Max Ion Time (ms): 0  
Full Micro Scans: 3  
Full AGC Target: 50000000  
Full Max Ion Time (ms): 200  
SIM Micro Scans: 5  
SIM AGC Target: 20000000  
SIM Max Ion Time (ms): 200  
MSn Micro Scans: 2  
MSn AGC Target: 20000000  
MSn Max Ion Time (ms): 800

### Appendix 3

#### Method Detection Limit (MDL) data

	Spiked (ug)	Replicates							Average	Stdev	MDL=STDEV* 3.143	RL (suggest)
		1	2	3	4	5	6	7				
DimethoateOA	0.2	0.195	0.192	0.209	0.191	0.203	0.214	0.213	0.199	0.013	0.042	0.250
Dimethoate	0.2	0.160	0.173	0.208	0.192	0.200	0.175	0.189	0.186	0.016	0.050	0.250
Malathion OA	0.2	0.185	0.186	0.191	0.186	0.192	0.179	0.183	0.189	0.009	0.028	0.250
Simazine	0.2	0.181	0.177	0.179	0.190	0.182	0.183	0.199	0.186	0.008	0.026	0.250
Diazinon OA	0.2	0.213	0.190	0.219	0.227	0.229	0.233	0.225	0.218	0.014	0.045	0.250
Norflurazon	0.2	0.184	0.175	0.144	0.213	0.211	0.178	0.218	0.187	0.026	0.081	0.250
Diuron	0.2	0.175	0.203	0.132	0.205	0.178	0.137	0.223	0.184	0.035	0.111	0.250
Phosmet	0.4	NO Data										
Azinphos Methyl	0.4	0.377	0.364	0.278	0.372	0.349	0.257	0.412	0.345	0.052	0.164	1.000

Molinate	0.2	0.151	0.167	0.179	0.171	0.148	0.176	0.150	0.163	0.012	0.039	0.250
Malathion	0.2	0.152	0.141	0.131	0.118	0.130	0.157	0.155	0.141	0.015	0.047	0.250
Metolachlor	0.2	0.166	0.177	0.204	0.184	0.173	0.175	0.176	0.184	0.019	0.059	0.250
Propanil	0.2	0.178	0.182	0.158	0.209	0.189	0.186	0.199	0.184	0.016	0.050	0.250
Chlorpyrifos OA	0.2	0.199	0.225	0.198	0.174	0.171	0.182	0.171	0.189	0.020	0.063	0.250
EPTC	0.2	0.162	0.157	0.176	0.177	0.167	0.143	0.173	0.165	0.011	0.036	0.250
Oryzalin	0.2	0.216	0.229	0.228	0.232	0.227	0.220	0.220	0.227	0.010	0.030	0.250
Diazinon	0.2	0.185	0.176	0.196	0.192	0.191	0.189	0.185	0.190	0.008	0.025	0.250
Thioben carb	0.2	0.229	0.162	0.269	0.198	0.181	0.156	0.226	0.200	0.038	0.121	0.500
DEF	0.2	0.191	0.190	0.190	0.204	0.193	0.199	0.226	0.198	0.012	0.038	0.250

	Spiked (ug)	MDL-1	MDL-2	MDL-3	MDL-4	MDL-5	MDL-6	MDL-7	Average	STDEV	MDL =3.143 x stdev	RL
Dichlorvos	0.2	0.19	0.2	0.18	0.2	0.23	0.21	0.16	0.196	0.022	0.070	1.0
Trifluralin	0.2	0.26	0.26	0.26	0.26	0.29	0.26	0.26	0.264	0.011	0.036	0.5
Chlorothalonil	0.4	0.544	0.624	0.504	0.63	0.674	0.4	0.514	0.556	0.094	0.295	2.0
Chlorpyrifos	0.2	0.136	0.201	0.175	0.193	0.252	0.201	0.194	0.193	0.035	0.109	1.0
Dicifol	0.2	0.192	0.214	0.205	0.205	0.239	0.208	0.2199	0.212	0.015	0.046	1.0
Endosulfan	0.2	0.172	0.209	0.199	0.187	0.241	0.204	0.185	0.200	0.022	0.070	1.0
Oxyfluorfen	0.4	0.354	0.408	0.357	0.394	0.477	0.367	0.424	0.397	0.044	0.138	1.0
Endosulfan sulfate	0.2	0.216	0.242	0.185	0.186	0.256	0.172	0.191	0.207	0.032	0.100	1.0
Propargite	0.4	0.36	0.41	0.35	0.36	0.4	0.34	0.36	0.369	0.026	0.082	1.0
Permethrin	0.4	0.402	0.472	0.431	0.463	0.555	0.434	0.489	0.464	0.050	0.156	1.0
Cypermethrin	0.4	0.45	0.49	0.5	0.51	0.53	0.44	0.49	0.487	0.032	0.101	1.0

## Appendix 4 Method Validation Data

	Spike Level	Set 1		Set 2		Set 3		Set 4		Set 5	
		Result	Recovery	Result	Recovery	Result	Recovery	Result	Recovery	Result	Recovery
	(ug)	(ug)	(%)	(ug)	(%)	(ug)	(%)	(ug)	(%)	(ug)	(%)
DimethoateOA	0.5	0.599	120	0.53	106	0.524	105	0.445	89	0.469	94
	1	1.11	111	1.13	113	0.993	99	0.921	92	0.924	92
	5	5.26	105	5.33	107	4.73	95	4.15	83	4.63	93
	10	10.33	103	10.78	108	9.62	96	9.75	97	9.47	95
Dimethoate	0.5	0.411	82	0.505	101	0.498	100	0.51	102	0.415	83
	1	0.914	91	0.864	86	0.959	96	1.15	115	0.792	79
	5	4.02	80	4.47	89	5.00	100	4.21	84	4.6	92
	10	8.71	87	9.81	98	9.03	90	10.7	107	8.7	87

Malathion OA	0.5	0.546	109	0.603	121	0.521	104	0.44	88	0.477	95
	1	1.14	114	1.03	103	1.06	106	0.947	95	1.03	103
	5	5.09	102	5.02	100	5.46	109	4.85	97	4.73	95
	10	10.57	106	10.23	102	10.0	100	9.39	94	10.6	106
Simazine	0.5	0.463	93	0.506	101	0.439	88	0.538	108	0.442	88
	1	1.06	106	0.963	96	0.815	82	1.09	109	0.917	92
	5	4.51	90	4.66	93	4.61	92	5.06	101	4.61	92
	10	9.49	95	9.88	99	8.66	87	11.1	111	9.01	90
Diazinon OA	0.5	0.516	103	0.651	130	0.509	102	0.28	94	0.494	99
	1	1.05	105	1.18	118	0.948	95	0.576	97	0.992	99
	5	5.03	101	5.75	115	4.96	99	2.73	92	4.01	80
	10	10.07	101	11.09	111	9.66	97	6.21	105	10.3	103
Norflurazon	0.5	0.45	90	0.563	113	0.349	70	0.503	101	0.554	111
	1	1.06	106	1.1	110	0.864	86	1.06	106	0.837	84
	5	4.85	97	5.3	106	4.25	85	5.12	102	4.1	82
	10	9.59	96	9.44	94	8.08	81	10.9	109	8.71	87
Diuron	0.5	0.493	99	0.424	85	0.438	88	0.499	100	0.61	122
	1	1.1	110	1.05	105	0.915	91	1.09	109	1.15	115
	5	5.87	117	4.8	96	5.06	101	4.56	91	4.07	81
	10	9.76	98	11.37	114	9.75	98	10.1	101	7.26	73
Phosmet	1.0/0.5	1.48	148	1.235	123	1.30	130	0.392	78.4	0.437	87
	2.0/1.0	3.64	182	2	100	2.73	136	0.884	88.4	1.02	102
	10/5.0	14.4	144	10.1	101	11.58	116	0.460	92	5.28	106
	20/10.	33.13	166	15.88	79	20.6	103	1.09	109	11.8	118
Azinphos Methyl	1.0/0.5	0.553	55	0.691	69	0.713	71	0.476	95	0.297	59
	2.0/1.0	1.79	89	1.73	86	2.11	105	1.00	100	0.647	65
	10/5.0	8.08	81	8.15	81	9.63	96	4.85	97	3.55	71
	20.0/10.0	16.55	83	15.88	79	19.2	96	11.3	113	8.34	83
Molinate	0.5	0.304	61	0.328	66	0.390	78	0.448	90	0.389	78
	1	0.636	64	0.62	62	0.774	77	0.907	91	0.821	82
	5	2.98	60	3.35	67	5.01	100	4.54	91	4.28	86
	10	7.84	78	8.32	83	9.48	95	10.2	102	9.34	93
Malathion	0.5	0.369	74	0.496	99	0.342	68	0.577	115	0.335	67
	1	0.835	84	0.84	84	0.928	93	1.08	108	0.68	68
	5	4.01	80	4.02	80	5.29	106	6.15	123	4.19	84
	10	8.2	82	7.57	76	8.53	85	11.9	119	8.82	88
Metolachlor	0.5	0.391	78	0.451	90	0.354	71	0.556	111	0.449	90
	1	0.938	94	0.938	94	0.935	93	1.09	109	0.887	89
	5	4.02	80	3.96	79	4.97	99	5.77	115	5.13	103
	10	9.07	91	8.5	85	6.69	67	10.9	109	10.6	106
Propanil	0.5	0.407	81	0.504	101	0.488	98	0.484	97	0.53	106
	1	0.841	84	1.02	102	0.865	86	0.969	97	0.904	90
	5	3.5	70	4.44	89	4.76	95	4.75	95	4.87	97
	10	7.91	79	9.63	96	9.52	95	10.6	106	11	110

Chlorpyrifos OA	0.5	0.546	109	0.577	115	0.474	95	0.723	145	0.46	92
	1	1.25	125	1.09	109	1.02	102	1.3	130	0.969	97
	5	4.17	83	4.56	91	4.90	98	4.94	99	5.24	105
	10	9.83	98	9.3	93	6.48	65	10.2	102	10.9	109
EPTC	0.5	0.27	54	0.343	69	0.415	83	0.446	89	0.394	79
	1	0.344	34	0.689	69	0.676	68	0.928	93	0.773	77
	5	2.2	44	3.04	61	4.08	82	4.68	94	4.08	82
	10	7.42	74	8.12	81	8.12	81	10.4	104	8.58	86
Oryzalin	0.5	0.544	109	0.519	104	0.439	88	0.605	121	0.554	111
	1	0.989	99	0.949	95	0.955	95	1.08	108	1.02	102
	5	4.49	90	3.76	75	4.80	96	5.73	115	5.26	105
	10	10.99	110	8.32	83	8.45	84	11.1	111	11.6	116
Diazinon	0.5	0.418	84	0.407	81	0.387	77	0.401	80	0.474	95
	1	0.884	88	0.772	77	0.938	94	0.899	90	0.959	96
	5	3.97	79	3.72	74	5.07	101	5.05	101	4.66	93
	10	8.47	85	8.13	81	8.82	88	11.4	114	9.86	99
Thioben carb	0.5	0.435	87	0.517	103	0.511	102	0.427	85	0.463	93
	1	0.947	95	0.815	81	0.879	88	0.926	93	0.943	94
	5	4.31	86	4.00	80	5.22	104	4.8	96	4.79	96
	10	8.41	84	8.83	88	10.1	101	9.69	97	9.87	99
DEF	0.5	0.434	87	0.463	93	0.496	99	0.488	98	0.478	96
	1	0.915	91	0.922	92	0.951	95	0.991	99	0.948	95
	5	4.01	80	4.38	88	5.06	101	4.98	100	4.8	96
	10	9.11	91	9.03	90	9.30	93	10.8	108	9.97	100

Analyte	Spike Level (ug)	SET 1		SET 2		SET 3		SET 4		SET 5	
		Result (ug)	Recovery (%)	Result (ug)	Recovery (%)	Result (ug)	Recovery (%)	Result (ug)	Recovery (%)	Result (ug)	Recovery (%)
1) DICHLORVOS	0.5	0.227	45	0.576	115	0.550	0.5	0.570	114	0.680	136
	1.0	0.499	50	1.04	104	0.960	96	1.09	109	1.01	101
	5	3.36	67	3.34	67	4.68	94	4.09	82	5.24	105
	10	6.58	66	9.70	97	12.6	126	10.8	108	9.48	95
2) TRIFLURALIN	0.5	0.486	97	0.506	101	0.530	106	0.600	120	0.630	126
	1.0	0.955	96	1.05	105	0.840	84	1.08	108	0.970	97
	5.0	4.78	96	4.25	85	4.68	94	4.23	85	5.15	103
	10	9.86	99	9.42	94	11.8	118	12.4	124	11.0	110
3) CHLOROTHA LONIL	1.0/0.5	1.20	120	1.17	117	1.45	145	0.749	150	1.31	262
	2.0/1.0	1.91	95	2.14	107	2.08	104	1.03	103	1.49	149
	10/5.0	8.62	86	9.26	93	9.59	96	4.64	93	5.38	108
	20.0/10.0	23.6	118	9.55	48	23.4	117	15.8	158	12.2	122
4)	0.5	0.470	94	0.459	92	0.640	128	0.510	102	0.530	106

CHLORPYRIFOS											
	1.0	0.877	88	1.05	105	0.920	92	1.19	119	0.900	90
	5	4.73	95	4.10	82	2.88	58	5.29	106	5.34	107
	10.0	7.80	78	9.36	94	10.4	104	12.4	124	9.92	99
5) DICOFOLOP, P'	0.5	0.498	100	0.484	97	0.500	100	0.380	76	0.430	86
	1.0	0.941	94	1.05	105	0.850	85	1.02	102	0.830	83
	5.0	5.09	102	4.20	84	4.29	86	4.78	96	5.23	105
	10	8.32	83	9.72	97	11.3	113	9.94	99	11.2	112
6) ENDOSULFAN (Thiodan 1)	0.5	0.516	103	0.419	84	0.440	88	0.650	130	0.53	106
	1.0	1.05	105	1.17	117	0.800	80	1.08	108	0.91	91
	5.0	5.08	102	4.84	97	4.78	96	4.67	93	5.55	111
	10	10.8	108	9.7	97	12.0	120	10.7	107	10.6	106
7) OXYFLUORFEN	1.0/0.5	1.03	103	1.03	103	1.13	113.0	0.440	88	0.78	156
	2.0/1.0	1.84	92	2.03	102	1.88	94	1.06	106	1.06	106
	10/5.0	9.70	97	8.93	89	9.94	99	4.78	96	5.52	110
	20.0/10.0	18.9	95	19.8	99	23.2	116	10.5	105	11.1	111
8) ENDOSULFAN SULFATE	0.5	0.534	107	0.555	111	0.670	134	0.510	102	0.620	124
	1.0	0.882	88	0.966	97	1.05	105	1.11	111	0.930	93
	5	4.49	90	4.88	98	5.23	105	4.93	99	5.46	109
	10.0	12.0	120	9.00	90	12.3	123	11.4	114	11.0	110
9) PROPARGITE	1.0/0.5	0.972	97	0.801	80	1.36	136	0.390	78	0.550	110
	2.0/1.0	2.13	107	2.11	106	2.03	102	1.11	111	1.10	110
	10/5.0	12.3	123	8.41	84	8.65	87	3.83	77	4.97	99
	20.0/10.0	18.9	95	16.8	84	21.4	107	8.65	87	9.36	94
10) PERMETHRIN	1.0/0.5	0.960	96	1.00	100	1.10	110.0	0.470	94	0.500	100
	2.0/1.0	1.93	97	2.25	112	1.90	95	1.02	102	0.870	87
	10/5.0	9.44	94	9.60	96	10.6	106	5.00	100	5.52	110
	20.0/10.0	19.0	95	19.7	99	24.9	124	10.5	105	11.2	112
11) CYPERMETHRIN	1.0/0.5	0.974	97	1.03	103	1.33	133	0.650	130	0.540	108
	2.0/1.0	1.77	89	2.11	106	2.06	103	1.20	120	0.900	90
	10.0/5.0	9.20	92	9.34	93	10.0	100	4.36	87	5.29	106
	20.0/10.0	19.6	98	16.5	83	24.0	120	11.5	115	11.5	115

## Appendix 5 Sample Storage Data

### Storage Recovery LC-MS

		Spiked (µg)	Sample 1		Sample 2		Sample 3	
			Found(µg)	Recovery (%)	Found (µg)	Recovery (%)	Found (µg))	Recovery (%)
DimethoateOA	Day 0	none	0.415		trace		0.398	
	Day 7	none	trace		0.39		0.403	
	Day14	none	0.513		0.481		0.578	
	Day 28	none	trace		trace		trace	
Dimethoate	Day 0	10.0	10.4	104	8.55	86	9.78	98
	Day 7	10.0	9.56	96	9.54	95	9.33	93
	Day14	10.0	9.68	97	10.5	105	8.04	80
	Day 28	10.0	10.7	107	9.86	99	9.75	97
Malathion OA	Day 0	none	0.194		trace		0.217	
	Day 7	none	trace		trace		0.209	
	Day14	none	0.270		0.271		0.301	
	Day 28	none	0.051		trace		trace	
Simazine	Day 0	10.0	9.99	100	9.52	95	10.3	103
	Day 7	10.0	9.63	96	9.503	95	10.7	107
	Day14	10.0	9.78	98	10.3	103	10.1	101
	Day 28	10.0	10.6	106	10.0	100	10.5	105
Diazinon OA	Day 0	none	0.405		trace		0.423	
	Day 7	none	trace		0.458		0.435	
	Day14	none	0.548		0.519		0.649	
	Day 28	none	trace		trace		trace	
Norflurazon	Day 0	10.0	10.4	104	9.38	94	10.6	106
	Day 7	10.0	9.28	93	9.341	93	9.84	98
	Day14	10.0	10.4	104	10.4	104	10.7	107
	Day 28	10.0	11.2	112	10.5	105	10.2	102
Diuron	Day 0	10.0	8.91	89	9.26	93	9.99	100
	Day 7	10.0	11.3	113	10.1	101	9.49	95
	Day14	10.0	11.5	115	10.7	107	10.4	104
	Day 28	10.0	9.54	95	10.0	100	8.98	90
Phosmet	Day 0	10.0	10.8	108	10.3	103	9.10	91
	Day 7	10.0	12.0	120	13.9	139	11.5	115
	Day14	10.0	16.6	166	18.1	181	16.4*	164
	Day 28	10.0	12.4*	124	11.5	115	11.7	117
Azinphos Methyl	Day 0	10.0	10.1	101	9.98	100	11.0	110
	Day 7	10.0	11.5	115	11.460	115	11.466	115
	Day14	10.0	9.23	92	10.8	108	9.30	93
	Day 28	10.0	10.6	106	9.99	100	9.59	96
Molinate	Day 0	10.0	8.29	83	8.77	88	7.62	76
	Day 7	10.0	8.90	89	8.212	82	7.702	77



	Day14	10.0	8.49	85	8.40	84	7.91	79
	Day 28	10.0	10.5	105	9.41	94	9.70	97
Malathion	Day 0	10.0	8.49	85	8.03	80	7.36	74
	Day 7	10.0	7.80	78	7.748	77	7.320	73
	Day14	10.0	10.2	102	9.23	92	7.42	74
	Day 28	10.0	10.4	104	9.91	99	9.72	97
Metolachlor	Day 0	10.0	11.3	113	9.62	96	11.7	117
	Day 7	10.0	9.95	100	9.274	93	9.484	95
	Day14	10.0	9.14	91	10.1	101	9.62	96
	Day 28	10.0	10.2	102	9.81	98	9.90	99
Propanil	Day 0	10.0	10.9	109	9.75	97	10.8	108
	Day 7	10.0	9.43	94	10.181	102	10.563	106
	Day14	10.0	9.47	95	10.8	108	10.0	100
	Day 28	10.0	9.14	91	9.00	90	8.93	89
Chlorpyrifos OA	Day 0	none	0.403		0.36		0.398	
	Day 7	none	0.273		0.315		0.300	
	Day14	none	trace		trace		trace	
	Day 28	none	0.303		0.322		ND	
EPTC	Day 0	10.0	8.18	82	9.13	91	7.82	78
	Day 7	10.0	8.54	85	7.836	78	7.625	76
	Day14	10.0	7.40	74	7.60	76	7.38	74
	Day 28	10.0	9.48	95	9.57	96	8.69	87
Oryzalin	Day 0	10.0	9.83	98	8.71	87	9.76	98
	Day 7	10.0	10.8	108	10.4	104	10.1	101
	Day14	10.0	10.4	104	10.6	106	10.5	105
	Day 28	10.0	11.9	119	11.7	117	11.3	113
Diazinon	Day 0	10.0	9.25	92	9.89	99	9.15	91
	Day 7	10.0	9.93	99	9.87	99	9.22	92
	Day14	10.0	9.18	92	9.86	99	9.77	98
	Day 28	10.0	11.6	116	10.8	108	10.8	108
Thioben carb	Day 0	10.0	8.72	87	9.02	90	9.41	94
	Day 7	10.0	9.94	99	10.1	101	9.66	97
	Day14	10.0	10.2	102	11.1	111	9.32	93
	Day 28	10.0	9.77	98	9.90	99	9.97	100
DEF	Day 0	10.0	9.84	98	9.51	95	10.3	103
	Day 7	10.0	9.84	98	9.93	99	10.31	103
	Day14	10.0	10.5	105	11.1	111	10.3	103
	Day 28	10.0	10.3	103	9.65	97	9.63	96

**Storage Recovery GC-MS**

		Sample 1			Sample 2		Sample 3	
		Spiked (ug)	Found (ug)	% recovery	Found (ug)	% recovery	Found (ug)	% recovery
1) DICHLORVOS	Day 0	10	10.8	108	11.4	114	10.1	101
	Day 7	10	7.38	74	10.6	106	8.78	88
	Day 14	10	11.7	117	8.1	81	12.6	126
	DAy28	10	7.82	78	7.9	79	6.16	62
2) TRIFLURALIN	Day 0	10	10.5	105	11.6	116	10.2	102
	Day 7	10	8.2	82	10.7	107	10.1	101
	Day 14	10	11.0	110	8.3	83	11.8	118
	DAy28	10	8.76	88	8.7	87	7.1	71
3) CHLOROTHALONIL	Day 0	10	13.0	130	13.9	139	11.7	117
	Day 7	10	8.32	83	12.4	124	11.5	115
	Day 14	10	12.2	122	8.8	88	13.5	135
	DAy28	10	12.2	122	13.0	130	11.6	116
4) CHLORPYRIFOS	Day 0	10	10.4	104	12.1	121	10.0	100
	Day 7	10	7.5	75	12.2	122	9.16	92
	Day 14	10	11.7	117	7.4	74	11.4	114
	DAy28	10	8.33	83	8.5	85	7.97	80
5) DICOFOL p,p'	Day 0	10	10.7	107	11.7	117	10.5	105
	Day 7	10	9.36	94	12.3	123	12.1	121
	Day 14	10	12.0	120	9.5	95	12.2	122
	DAy28	10	8.82	88	9.1	91	7.58	76
6) Endosulfan	Day 0	10	10.2	102	12.6	126	10.1	101
	Day 7	10	7.64	76	12.1	121	10.5	105
	Day 14	10	11.5	115	8.2	82	12.1	121
	DAy28	10	8.92	89	8.9	89	7.51	75
7) OXYFLUORFEN	Day 0	10	11.1	111	12.4	124	10.6	106
	Day 7	10	7.84	78	12.6	126	10.7	107
	Day 14	10	11.0	110	8.28	83	11.5	115
	DAy28	10	8.63	86	8.76	88	7.41	74
8) ENDOSULFAN SULFATE	Day 0	10	11.1	111	12.3	123	9.20	92
	Day 7	10	8.16	82	11.0	110	10.2	102
	Day 14	10	10.8	108	7.04	70	11.6	116
	DAy28	10	10.1	101	11.0	110	8.56	86
9) PROPARGITE	Day 0	10	7.28	73	7.92	79	9.26	93
	Day 7	10	8.28	83	10.6	106	8.88	89
	Day 14	10	11.6	116	7.9	79	10.2	102
	DAy28	10	7.16	72	6.83	68	7.11	71
10) PERMETHRIN	Day 0	10	10.6	106	11.9	119	10.9	109
	Day 7	10	8.54	85	11.5	115	11.4	114
	Day 14	10	11.9	119	8.7	87	11.7	117
	DAy28	10	9.62	96	9.97	100	7.37	74
11) CYPERMETHRIN	Day 0	10	10.9	109	13.2	132	11.1	111
	Day 7	10	8.78	88	12.1	121	10.2	102
	Day 14	10	11.4	114	8.26	83	11.8	118
	DAy28	10	7.51	75	7.57	76	7.12	71

## Appendix 6 Trapping Efficiency Data

### Trapping Efficiency

Sample ID		1	2	3	4	5	6	7	8	9
Spiked (µg)		10.0	10.0	10.0	5.0	5.0	5.0	2.0	2.0	2.0
DimethoateOA	No spike	0.803	0.646	1.00	0.490	0.503	0.582	0.051	0.25	trace
Dimethoate		8.39	8.95	7.60	4.30	4.01	4.21	1.53	1.75	2.05
Malathion OA	No spike	0.315	0.264	0.383	0.167	0.206	0.233	trace	trace	trace
Simazine		11.1	10.2	11.3	5.23	5.28	4.93	2.18	2.27	2.29
Diazinon OA	No spike	0.552	0.507	0.677	0.311	0.376	0.399	trace	trace	trace
Norflurazon		0.900	1.39	0.858	1.90	0.877	0.481	2.04	0.825	1.84
Diuron		8.27	7.65	8.87	4.02	3.17	4.17	2.14	1.97	2.08
Phosmet		10.6	9.79	10.5	3.89	4.66	4.51	2.36	1.76	2.24
Azinphos Methyl		6.67	6.65	7.24	2.68	3.06	3.18	1.77	1.58	1.74
Molinate		8.46	8.49	8.41	4.04	3.84	4.08	1.89	1.67	1.99
Malathion		6.89	8.73	10.4	3.80	3.56	4.45	2.26	1.84	1.93
Metolachlor		8.37	6.40	7.71	4.81	3.80	4.11	1.94	1.75	1.83
Propanil		10.5	9.32	10.2	5.08	5.01	4.58	1.68	1.91	2.31
Chlorpyrifos OA	No spike	0.47	0.45	0.496	0.434	0.437	0.456	trace	0.400	trace
EPTC		9.05	8.87	8.81	4.11	4.02	3.74	1.86	1.69	1.84
Oryzalin		1.25	1.73	1.34	2.26	0.95	0.53	1.74	0.473	1.65
Diazinon		10.6	8.54	8.84	4.14	4.35	4.57	1.83	1.73	1.42
Thioben carb		8.95	9.00	9.15	4.3	4.35	4.45	2.00	1.82	2.12
DEF		9.85	9.09	10.1	4.59	4.6	4.81	1.9	1.81	2.06

1) DICHLORVOS		11.2	12.8	9.88	5.2	5.58	5.86	1.36	1.62	1.70
2) TRIFLURALIN		6.74	9.30	6.86	4.62	3.86	4.20	1.24	1.52	1.58
3) CHLOROTHALONIL		11.4	8.92	6.54	4.94	3.14	3.52	1.74	1.58	1.86
4) CHLORPYRIFOS		10.1	11.0	8.52	4.60	4.40	4.78	1.08	1.30	1.24
5) DICOFOL p,p'		10.3	11.4	9.9	5.14	5.22	4.98	1.24	1.84	1.58
6) Endosulfan		10.1	11.5	9.66	5.28	5.00	5.36	1.14	1.68	1.48
7) OXYFLUORFEN		1.98	2.82	1.96	3.04	1.70	1.54	1.38	1.10	1.60

8) ENDOSULFAN SULFATE		11.3	11.3	9.5	5.72	4.80	4.82	1.30	1.78	1.70
9) PROPARGITE		9.34	9.64	7.92	3.64	3.92	4.06	1.02	1.28	1.08
10) PERMETHRIN		8.70	10.1	9.24	5.36	4.90	4.72	1.18	1.88	1.62
11) CYPERMETHRIN		9.86	10.0	9.24	6.22	5.44	5.46	2.44	2.82	2.68

## Appendix 7

### Cleaning XAD-4 resin

1. Measure 1.6 kg XAD-4 resin (about 3.0-3.5 liters) to a 6 liters Teflon (or Glass) container.
2. Wet with 2 liters of residue grade methanol. Then add 2 liters of 0.25 N HCl. Stir occasionally with a glass or Teflon rod for about 30 minutes.
3. Transfer the content to a glass cylinder (An open-end cylinder with rubber stopper tightly fit the bottom. A glass tube is inserted through the stopper and connected to a d.i water faucet.). Cover the top with a fine screen .
4. Overfill the cylinder with d.i water through the bottom at about 30 mL/min for at least 4 hours. The pH should be that of d.i water, or no Cl<sup>-</sup> reaction with a drop of 0.1 N AgNO<sub>3</sub> solution added to a 1 mL of out going water.
5. Transfer the water washed XAD-4 to the container of Soxhlet extractor. Drain the water through the stopcock. Add residue grade methanol to just cover the resin. Stir with a glass rod to help remove the air bubble.( Do not disturb the glass wool plug.). Drain the methanol through the stopcock.
6. Add about 4 liters methanol to the resin and allow the methanol to be siphoned to the solvent flask. The solvent flask should be 60% full.
7. Start the extraction and continue for at least 72 hours
8. Change extraction solvent to ethyl acetate and continue the process for at least 72 hours.
9. Dry the resin at 28 inches vacuum or higher and 40 °C for at least 72 hours until all traces of ethyl acetate is gone. During the drying process allow dry air sweep through the oven at a flow rate that can just be felt by placing finger on the air inlet tube.

Note: Use residue grade methanol and ethyl acetate to wash the resin

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