

**ACCURATE-MASS DATABASE
FOR SCREENING OF
PESTICIDE RESIDUES IN
FRUITS AND VEGETABLES BY
GAS CHROMATOGRAPHY
TIME-OF-FLIGHT MASS
SPECTROMETRY**



EURL-FV



In practice, target analysis approach

(analysing 100-400 of GC and LC amenable compounds)



MEANS THAT THE MAJORITY OF THE LOW-FREQUENCY OR MISUSED COMPOUNDS ARE NOT LOOKED FOR

Target methods



EURL

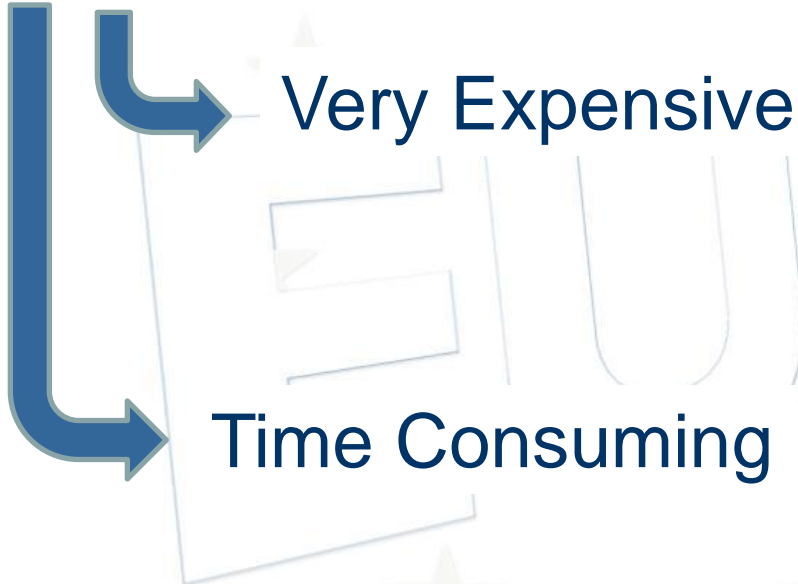
Target methods



Very Expensive



Target methods





Screening methods

- ✓ Practically all applications have been focused on LC (Time of flight)
- ✓ LC-TOF-MS (libraries containing fragments of compounds and retention time).
- ✓ Focused in GC (single quadrupole), commercial libraries.

GC-TOF/MS (full scan)

(Ability to record an unlimited number of compounds)

- ✓ GC libraries (accurate masses of molecular ions and fragments and retention times)

Experimental conditions

Injection volume: 2 μ L

Mode: Splitless MM Inlet

Injector temperature: 280°C

Agilent 7200 Q-TOF GC/MS

OVEN PROGRAM

Rate(°C/min)	Value(°C)	Hold Time(min)	Run Time(min)
	60	1	1
40	120	0	2.5
5	310	0	40.5

Experimental conditions

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Agilent 7200 Q-TOF GC/MS

OVEN PROGRAM

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	60	1	1
40	120	0	2.5
5	310	0	40.5

Post Run Time: 2 min

Post Run temperature: 310°C

**BACKFLUSHING
CAPABILITIES**

Retention time locked with constant flow (Trifluralin)

Experimental conditions

Injection volume: 2 μ L

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OVEN PROGRAM

Rate(°C/min)	Value(°C)	Hold Time(min)	Run Time(min)
	60	1	1
40	120	0	2.5
5	310	0	40.5

Post Run Time: 2 min

Post Run temperature: 310°C

**BACKFLUSHING
CAPABILITIES**

Retention time locked with constant flow (Trifluralin)

Electron Ionization

Acquisition in full scan at 2 GHz

Ion source temperature: 280°C

Transfer line temperature: 280°C

BACKFLUSH CONFIGURATION

2 columns: HP5MSI (350°C) 15mx250µmx0.25µm

Constant Flow

Column 1 = 1.225ml/min

Column 2 = 1.425ml/min

Front Inlet Flowpath

Backflush technique

Post-Column Backflush

Post-Column Details

Backflush Start Time:

Postrun Duration:

Oven Temperature:

Restrictor Temperature:

Void Volumes Backflushed:

Inlet Pressure	Column Flow	Aux Pressure	Column Flow	Detector Pressure
<input type="text" value="1 psi"/>	<input type="text" value="-5.851 mL/min"/>	<input type="text" value="50 psi"/>	<input type="text" value="6.217 mL/min"/>	<input type="text" value="0 psi"/>

Internal Reference Mass

Scan data acquired at 1:07:08 PM

File and Reports | Autotune | TOF Mass Calibration | Manual Tune | Vacuum Control | Removable Ion Source

Run

Tune Masses Profile Display

Enabled	Mass (m/z)	Window
<input checked="" type="checkbox"/>	68.99466	1
<input checked="" type="checkbox"/>	130.99147	1
<input checked="" type="checkbox"/>	168.988266	1
<input checked="" type="checkbox"/>	218.985077	1
<input checked="" type="checkbox"/>	263.986572	1
<input checked="" type="checkbox"/>	413.976959	1
<input checked="" type="checkbox"/>	463.9738	1
<input checked="" type="checkbox"/>	501.970581	1
<input checked="" type="checkbox"/>	613.9642	1
<input type="checkbox"/>		

Ion Source Quadrupole | Collision Cell | Transfer Optics | TOF | Detector | Acquisition

Source Type : EI+

Source Temp. °C

Filament 1 2

Emission μA

Energy eV

Repeller V

Source Body V

Extractor V

Ion Focus V

Entrance Lens V

EI Cal Valve Off

CI Cal Valve

IRM Off Off

Gas Control

Gas Flow %

Gas is turned off and isolation valve is closed

Range from --- to --- step ---

Ramp From To Step Dwell Time

New Sample(s)

	Name	Vial	Method File	Data File	Type	Level	Dil.	Comment	Keyword	Method Path
1			Metod...L.ei.m		Keyword				MassCal	D:\...methods
2	Solvent_2	1	Metod...L.ei.m	Solvent_2	Sample					D:\...methods
3			Metod...L.ei.m		Keyword				MassCal	D:\...methods
4	Blanco_Tomate_1	2	Metod...L.ei.m	Blanco_Tomate_1	Sample					D:\...methods
5			Metod...L.ei.m		Keyword				MassCal	D:\...methods
6	Blanco_Tomate_2	2	Metod...L.ei.m	Blanco_Tomate_2	Sample					D:\...methods

Internal Reference Mass

Scan data acquired at 1:07:08 PM

File and Reports | Autotune | TOF Mass Calibration | Manual Tune | Vacuum Control | Removable Ion Source

Run

Tune Masses Profile Display

Enabled	Mass (m/z)	Window
<input checked="" type="checkbox"/>	68.99466	1
<input checked="" type="checkbox"/>	130.99147	1
<input checked="" type="checkbox"/>	168.988266	1
<input checked="" type="checkbox"/>	218.985077	1
<input checked="" type="checkbox"/>	263.986572	1
<input checked="" type="checkbox"/>	413.976959	1
<input checked="" type="checkbox"/>	463.9738	1
<input checked="" type="checkbox"/>	501.970581	1
<input checked="" type="checkbox"/>	613.9642	1
<input type="checkbox"/>		

Ion Source: Quadrupole | Collision Cell | Transfer Optics | TOF | Detector | Acquisition

Source Type: EI+

Source Temp. 280 °C

Filament 1

Emission 4.6 μA

Energy 70.0 eV

Repeller 42.0 V

Source Body 46.0 V

Extractor 40.5 V

Ion Focus -144.0 V

Entrance Lens 9.0 V

Gas Control: Gas Flow 10 %

Get Parameter Default

Ramp: From N/A To N/A Step N/A Dwell Time 25

Close Help

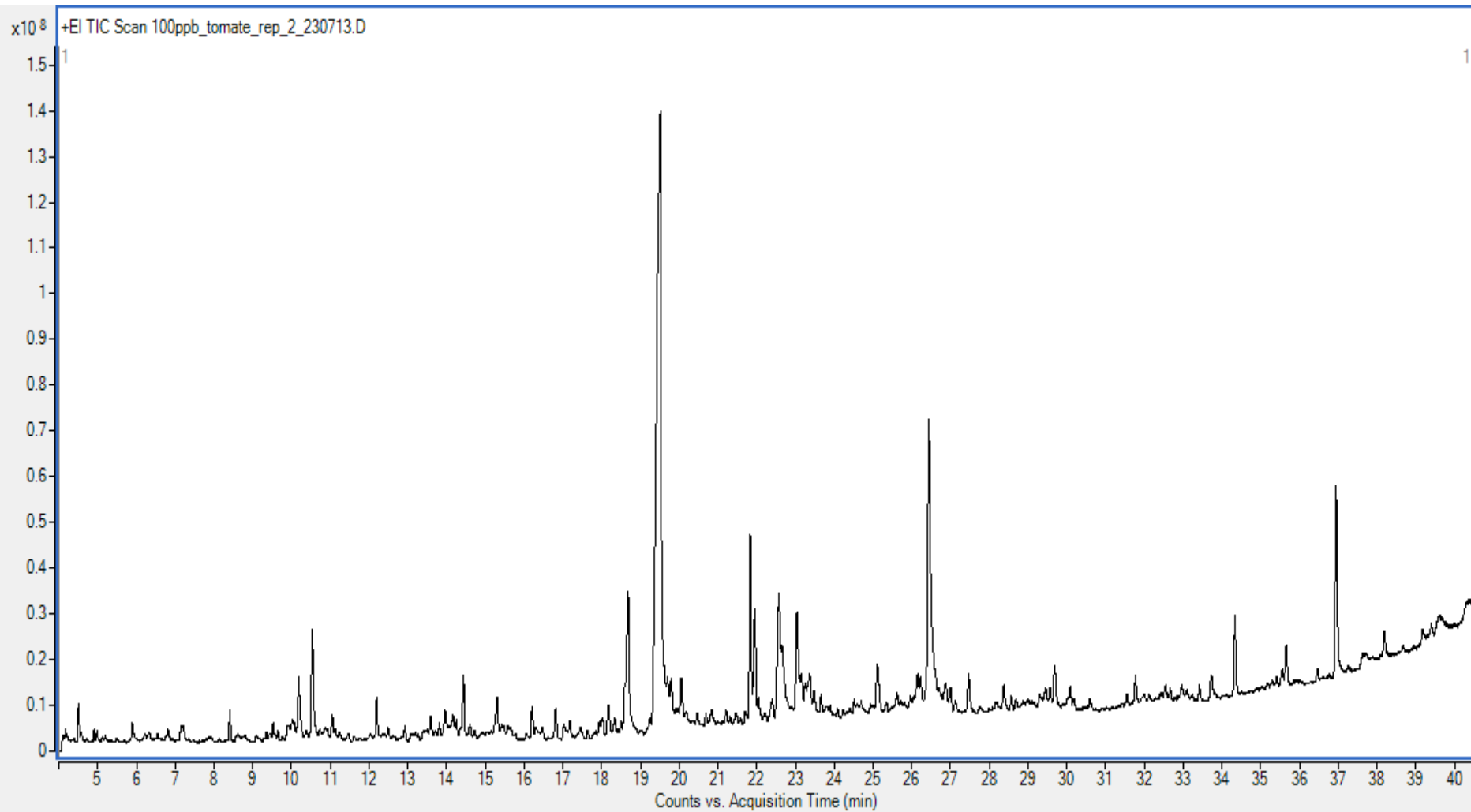
New Sample(s) | Tools

Name	Vial	Method File	Data File	Type	Level	Dil.	Comment	Keyword	Method Path
1		Metod_L.ei.m		Keyword				MassCal	D:_methods
2	Solvent_2	Metod_L.ei.m	Solvent_2	Sample					D:_methods
3		Metod_L.ei.m		Keyword				MassCal	D:_methods
4	Blanco_Tomate_1	Metod_L.ei.m	Blanco_Tomate_1	Sample					D:_methods
5		Metod_L.ei.m		Keyword				MassCal	D:_methods
6	Blanco_Tomate_2	Metod_L.ei.m	Blanco_Tomate_2	Sample					D:_methods

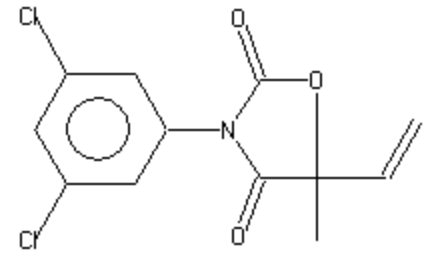
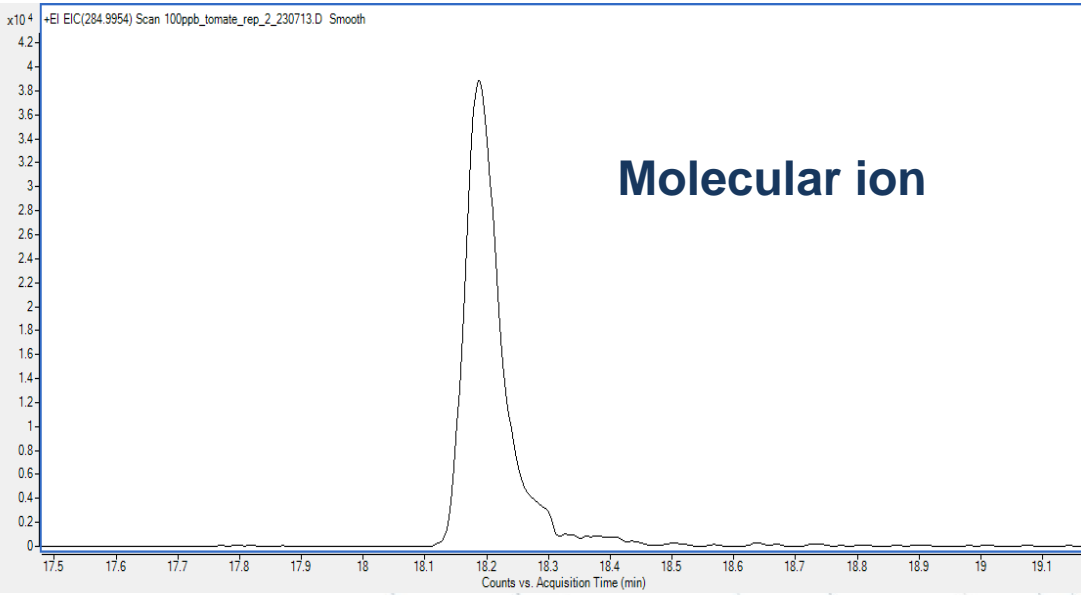
GC-Q-TOF DATABASE

- **Many ions in the mass spectra.**
- **The molecular ion is not always present in the spectra.**

100ppb Tomato TIC

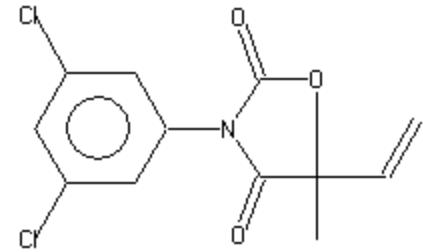
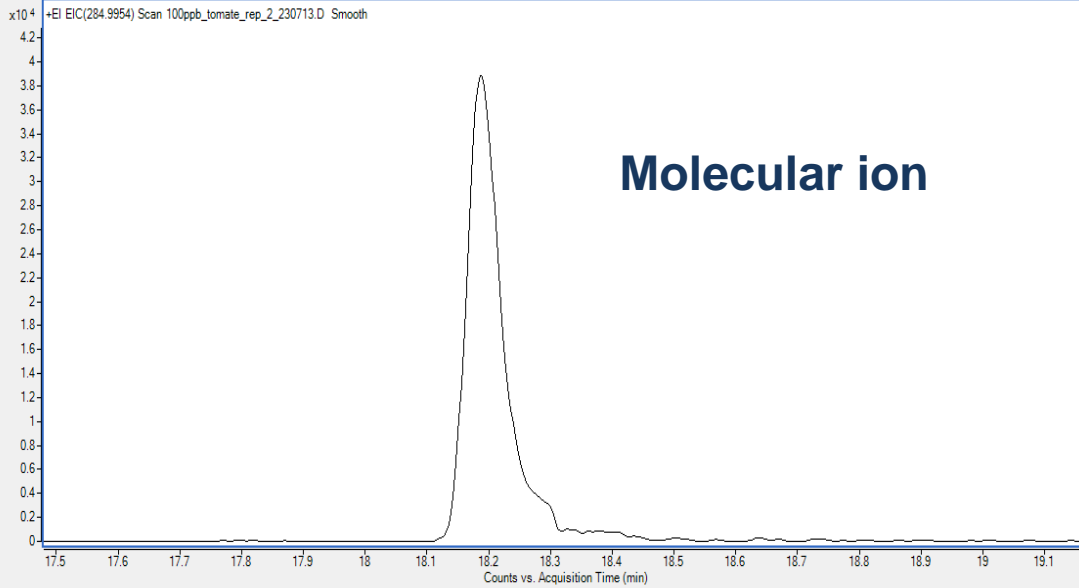


100ppb Tomate EIC (284.9954_Vinclozolin Parent Ion)



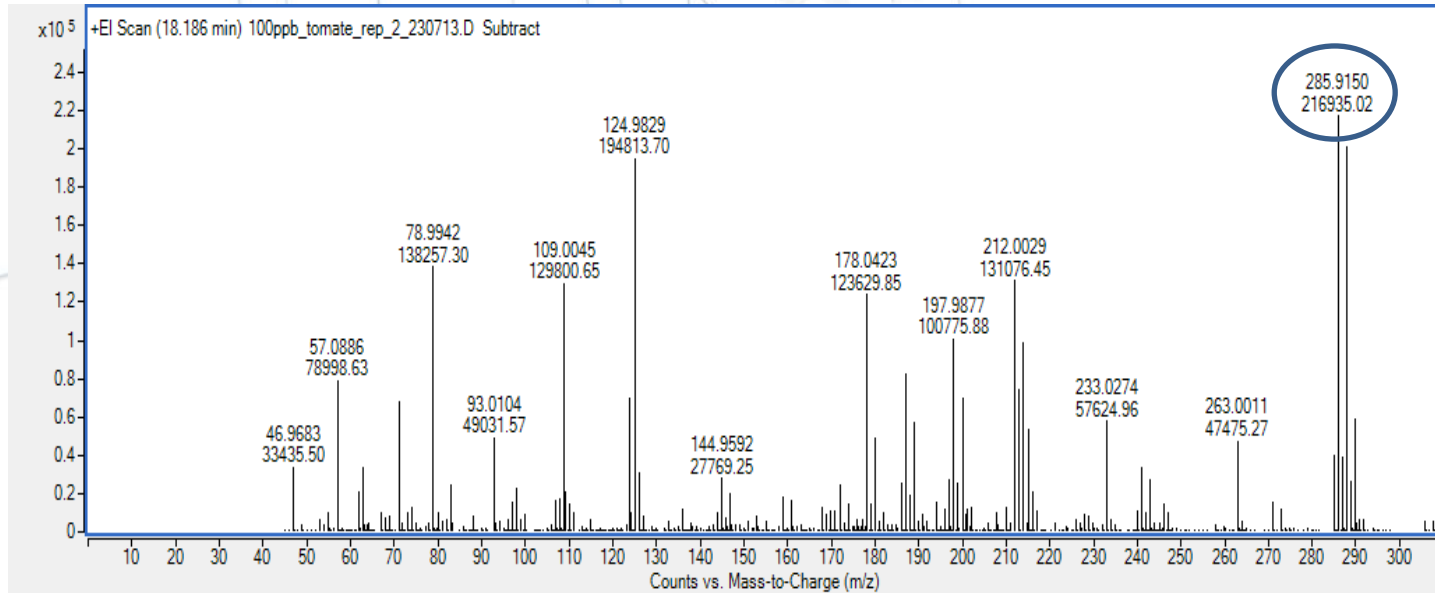
Vinclozolin

100ppb Tomato EIC (284.9954_Vinclozolin Parent Ion)

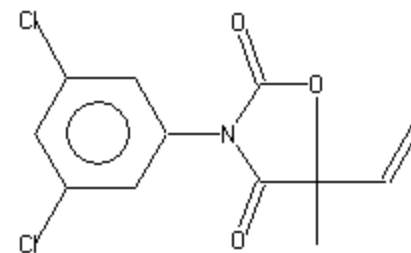
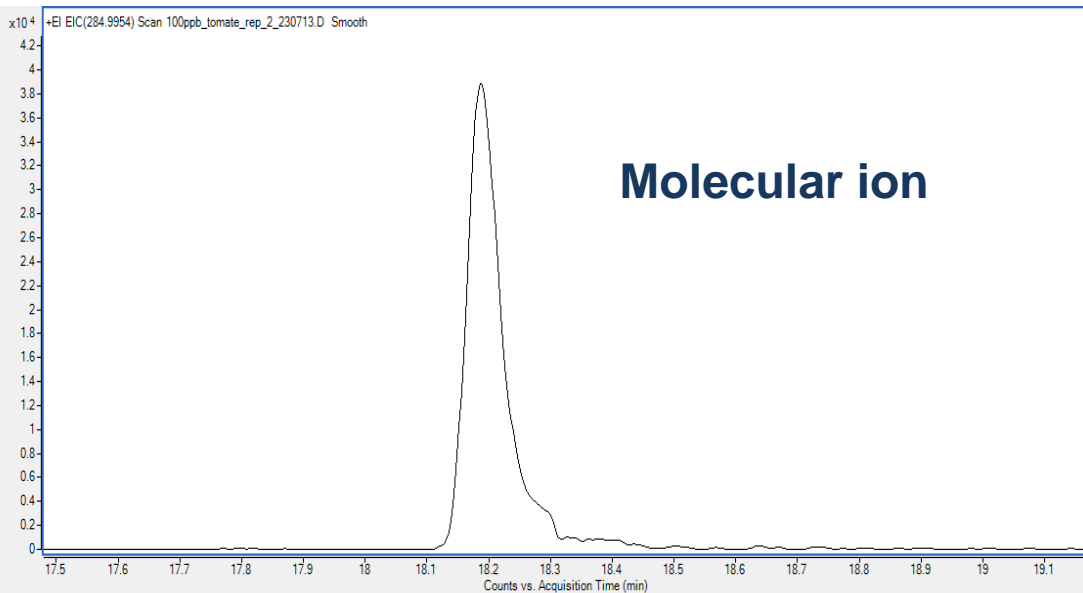


Vinclozolin

100ppb Tomato Mass Spectrum (tr = 18.186 min)



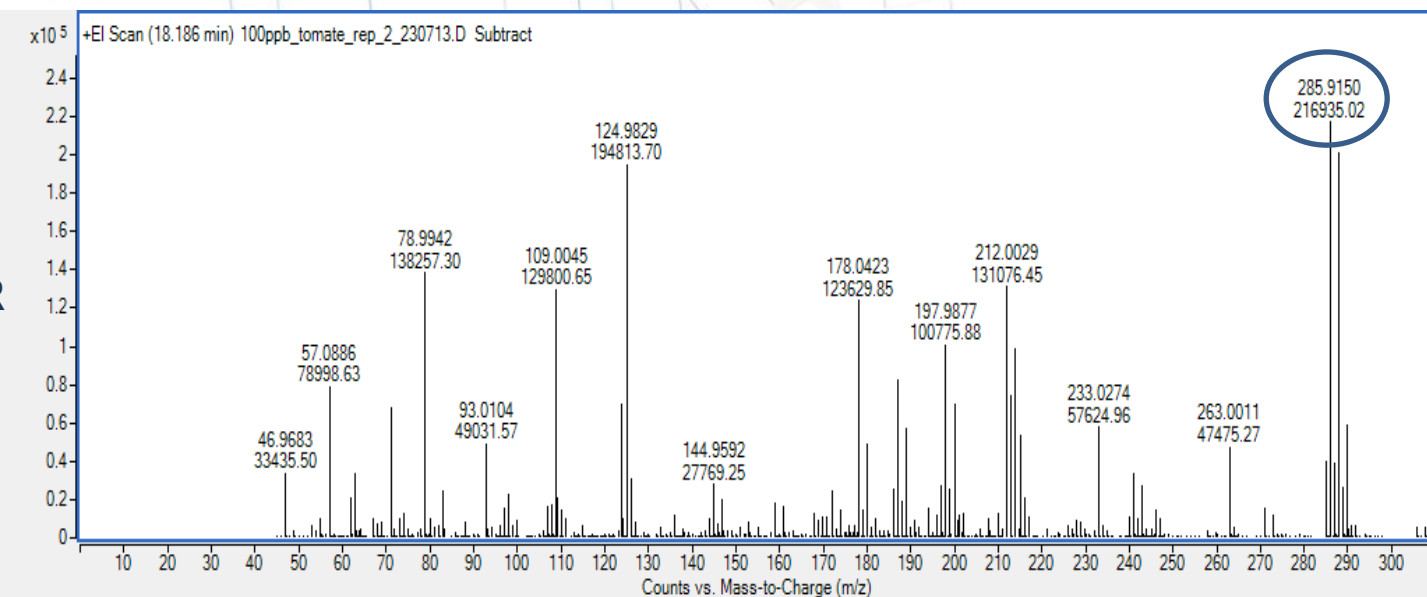
100ppb Tomato EIC (284.9954_Vinclozolin Parent Ion)



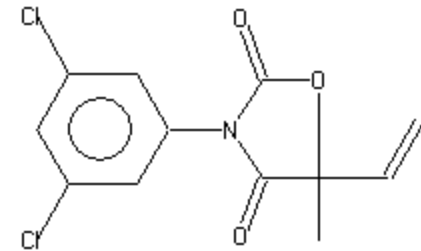
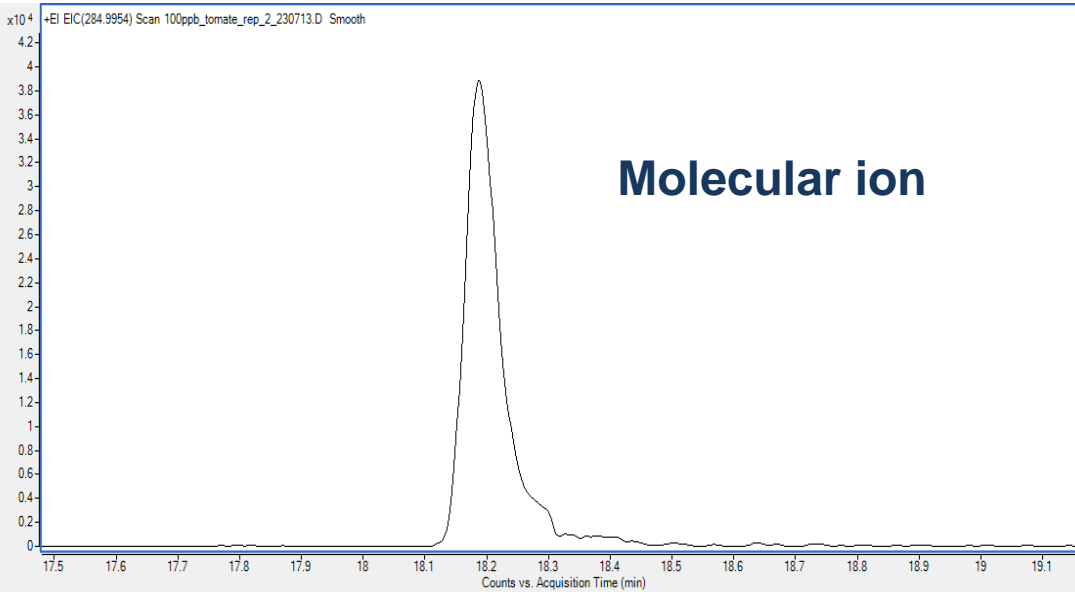
Vinclozolin

100ppb Tomato Mass Spectrum (tr = 18.186 min)

**NIST
MASS INTERPRETER**

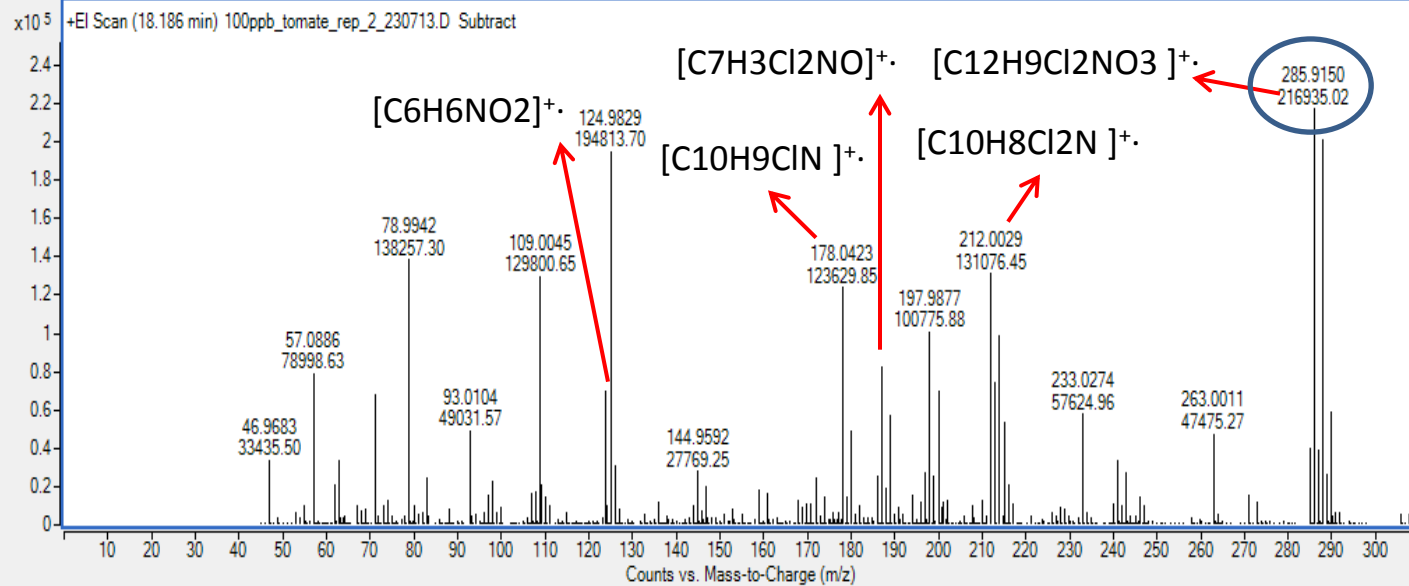


100ppb Tomato EIC (284.9954_Vinclozolin Parent Ion)

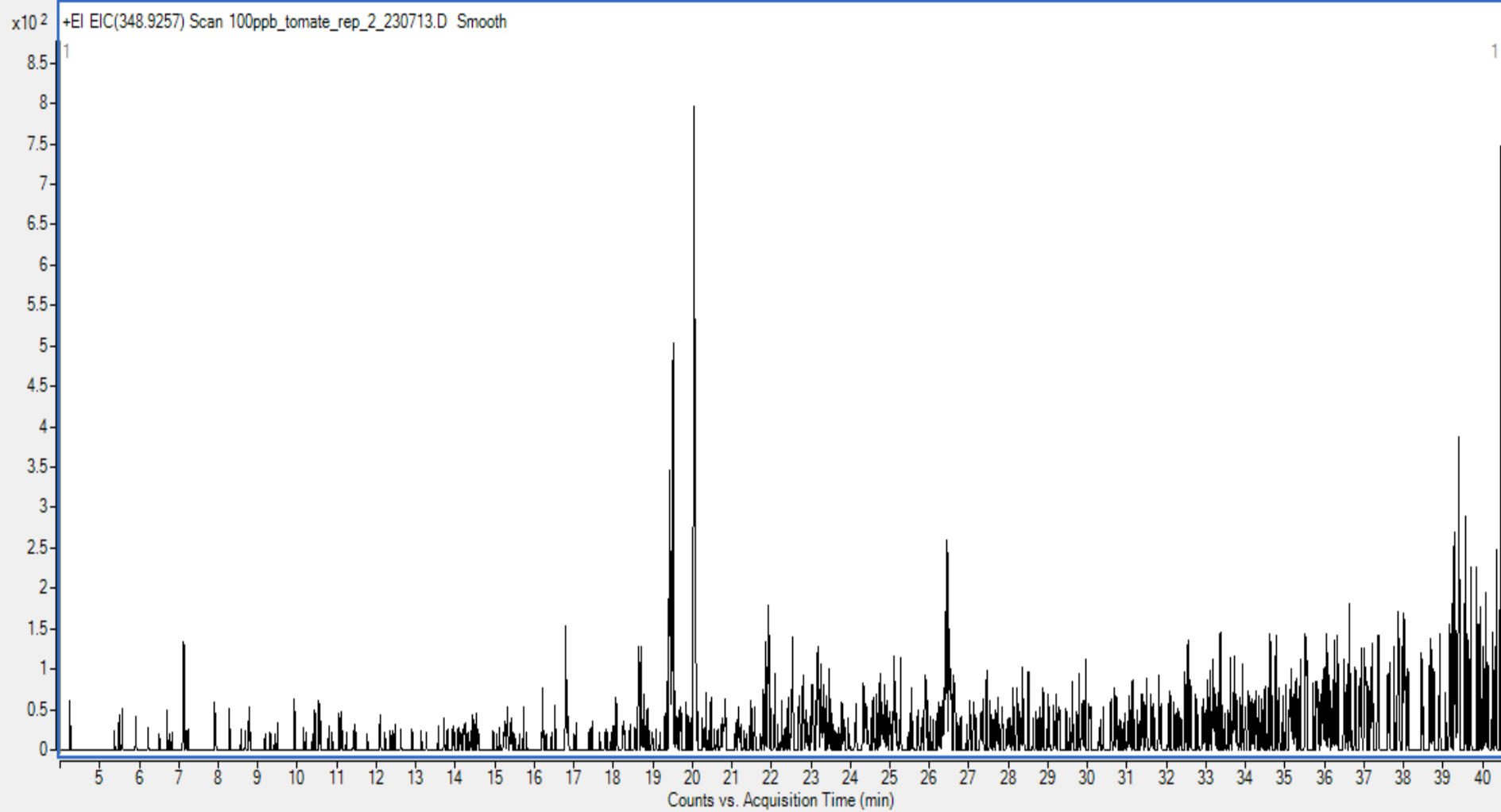
**Vinclozolin**

100ppb Tomato Mass Spectrum (tr = 18.186 min)

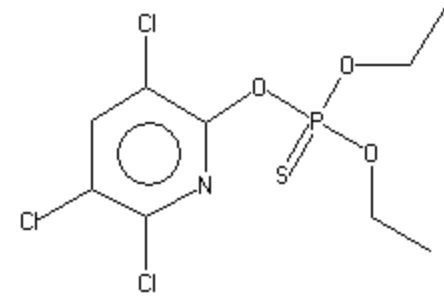
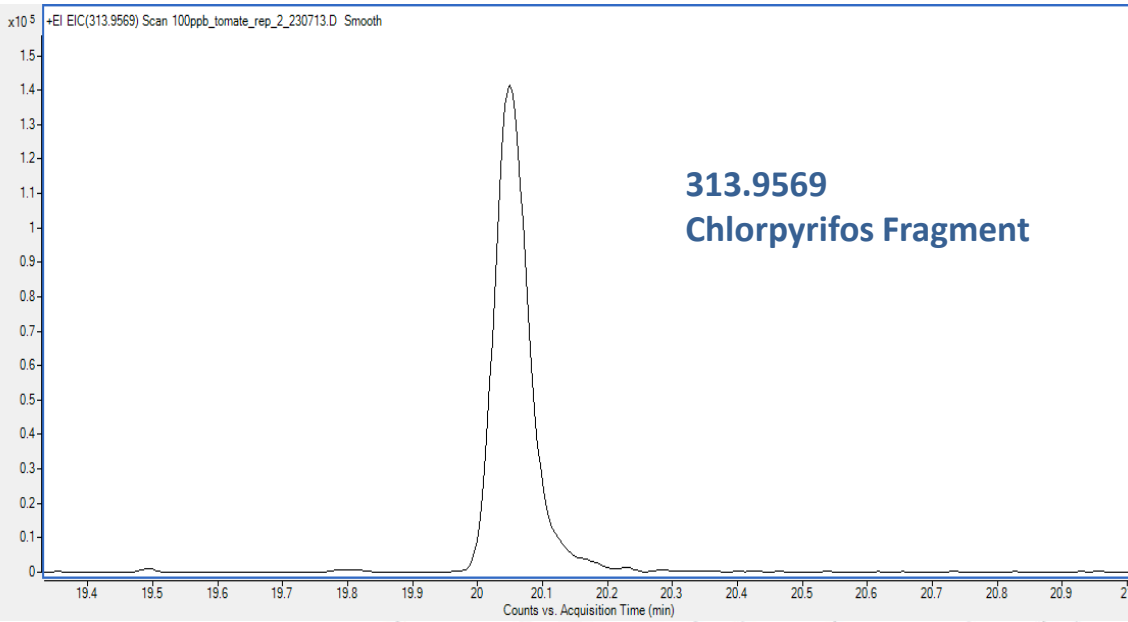
NIST
MASS INTERPRETER



100ppb Tomate EIC (348.9257_Chlorpyrifos Parent Ion)

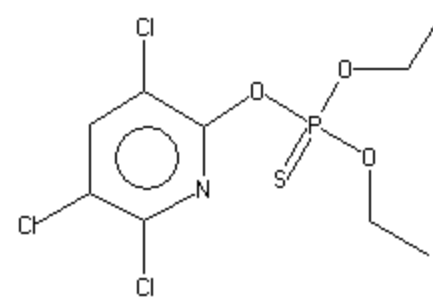


100ppb Tomato EIC (313.9569_Chlorpyrifos Fragment)

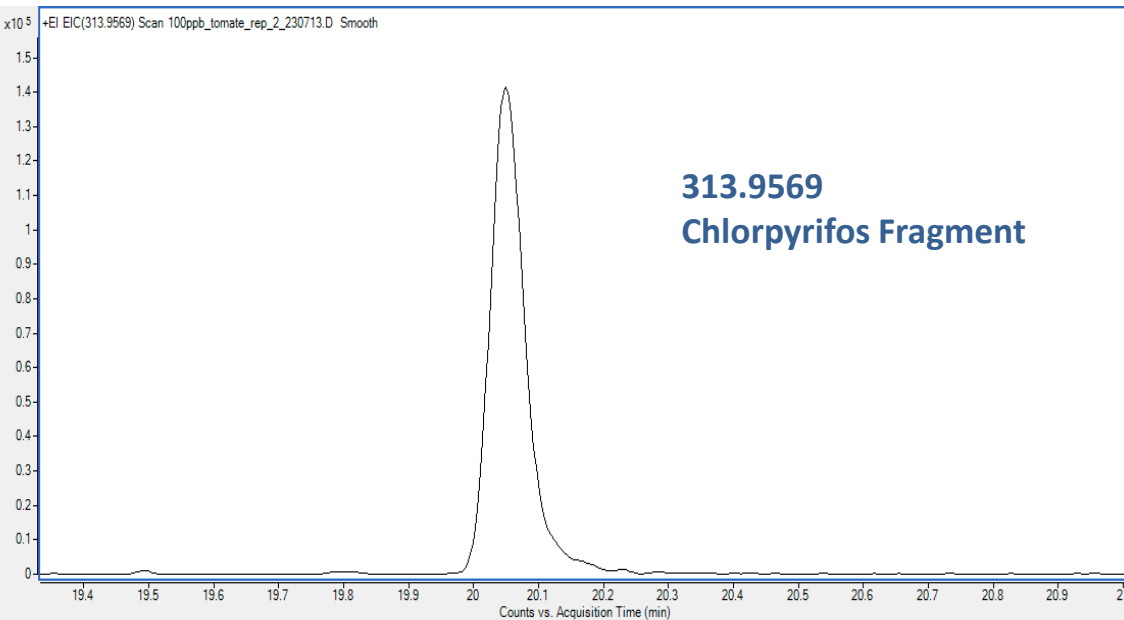


Chorpyrifos

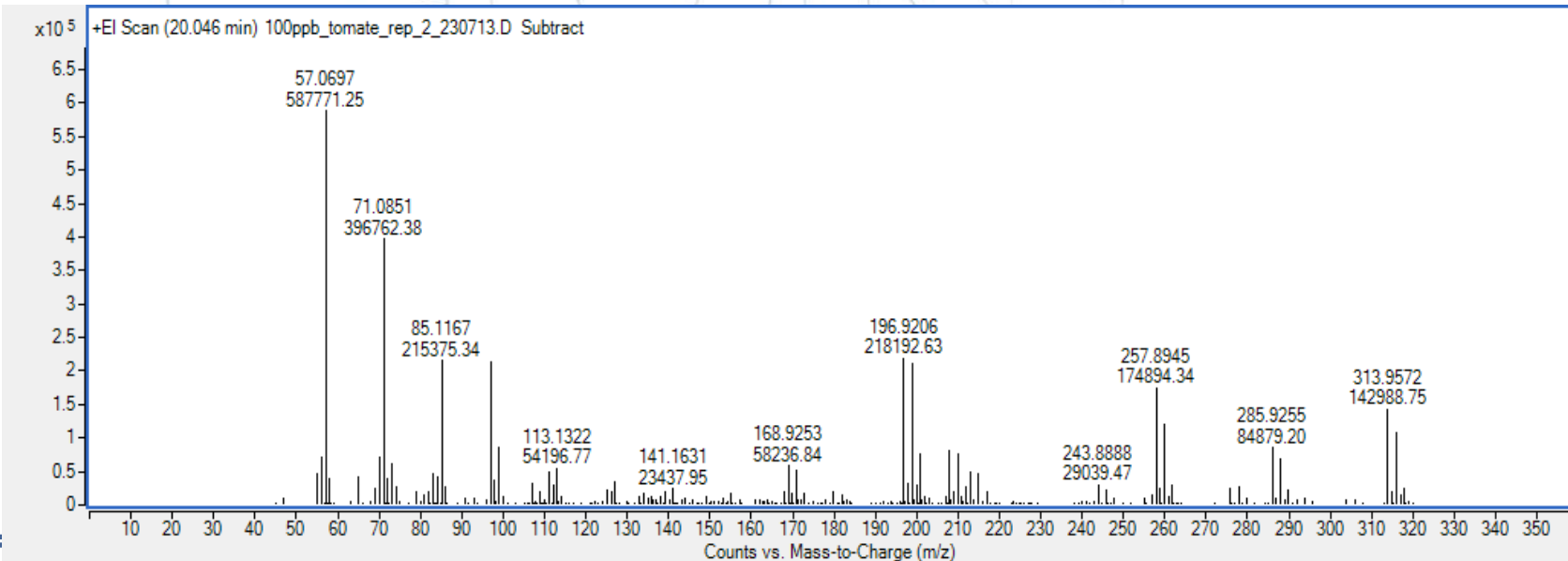
100ppb Tomato EIC (313.9569_Chlorpyrifos Fragment)



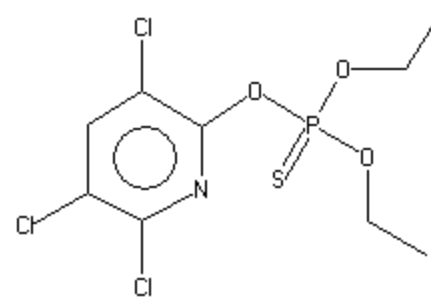
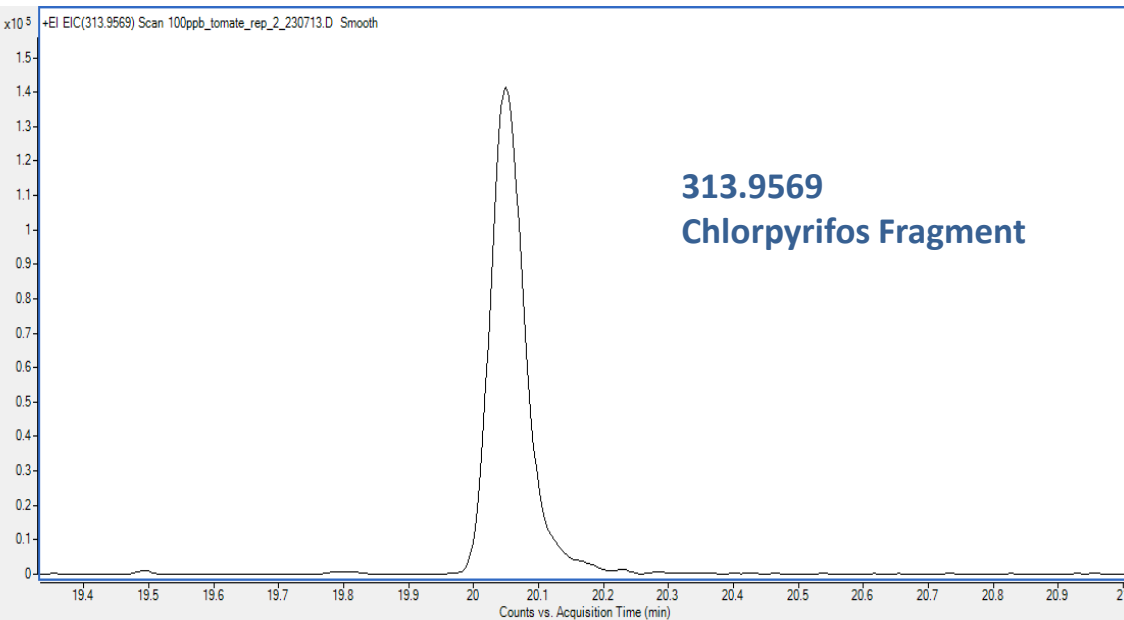
Chlorpyrifos



100ppb Tomato Mass Spectrum (tr = 20.046 min)

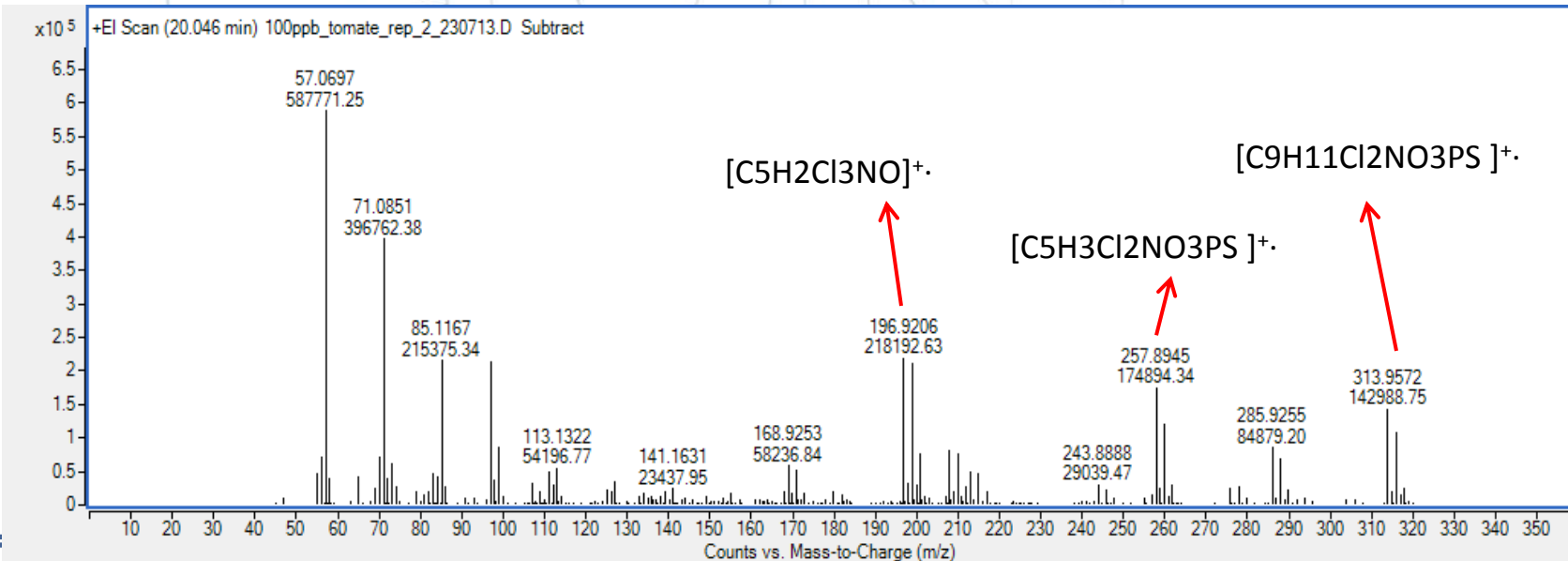


100ppb Tomato EIC (313.9569_Chlorpyrifos Fragment)



Chlorpyrifos

100ppb Tomato Mass Spectrum (tr = 20.046 min)



ACCURATE MASS DATABASE

### Formula	Retention Time	Mass	Compound name
# Formula	RT	Mass	Cpd
C17H16Br2O3	28,12	425,9466	Bromopropylate
C13H9Br2O	28,12	338,9020	Bromopropylate F1
C7H6BrO	28,12	184,9602	Bromopropylate F2
C6H4Br	28,12	154,9496	Bromopropylate F3
C13H12BrCl2N3O	27,92	374,9541	Bromuconazole
C10H8BrCl2O	27,92	292,9136	Bromuconazole F1
C13H12Cl2N3O	27,92	296,0357	Bromuconazole F2
C10H9Cl2O	27,92	215,0030	Bromuconazole F3
C7H3Cl2O	27,92	172,9561	Bromuconazole F4
C8H7Cl2	27,92	172,9925	Bromuconazole F5
C13H24N4O3S	24,01	316,1569	Bupirimate
C11H18N3O3S	24,01	272,1069	Bupirimate F1
C11H18N3O	24,01	208,1450	Bupirimate F2
C5H4N2O	24,01	108,0324	Bupirimate F3
C16H23N3OS	23,85	305,1562	Buprofezin
C9H9N2S	23,85	177,0486	Buprofezin F1
C7H5NO	23,85	119,0371	Buprofezin F2
C7H8N	23,85	106,0657	Buprofezin F3
C7H11N2OS	23,85	171,0592	Buprofezin F4
C8H16N2S	23,85	172,1034	Buprofezin F5
Continue

ACCURATE MASS DATABASE

### Formula	Retention Time	Mass	Compound name
# Formula	RT	Mass	Cpd
C17H16Br2O3	28,12	425,9466	Bromopropylate
C13H9Br2O	28,12	338,9020	Bromopropylate F1
C7H6BrO	28,12	184,9602	Bromopropylate F2
C6H4Br	28,12	154,9496	Bromopropylate F3
C13H12BrCl2N3O	27,92	374,9541	Bromuconazole
C10H8BrCl2O	27,92	292,9136	Bromuconazole F1
C:	87 PESTICIDES		
C:	2 FRAGMENTS AT LEAST FOR EACH PESTICIDE		
C:	RETENTION TIMES OF EACH PESTICIDE		
C15H24N4O3S	24,01	310,1509	Bupirimate
C11H18N3O3S	24,01	272,1069	Bupirimate F1
C11H18N3O	24,01	208,1450	Bupirimate F2
C5H4N2O	24,01	108,0324	Bupirimate F3
C16H23N3OS	23,85	305,1562	Buprofezin
C9H9N2S	23,85	177,0486	Buprofezin F1
C7H5NO	23,85	119,0371	Buprofezin F2
C7H8N	23,85	106,0657	Buprofezin F3
C7H11N2OS	23,85	171,0592	Buprofezin F4
C8H16N2S	23,85	172,1034	Buprofezin F5
Continue

Database_LAPRW_221113_melectron (2).csv [Sólo lectura] - Microsoft Excel

Archivo Inicio Insertar Diseño de página Fórmulas Datos Revisar Vista

Calibri 11 Fuente Alineación Ajustar texto Combinar y centrar General Número Estilos

A1 # Compound database,,,,

	A	B	C	D	E	F	G	H	I	J	K	L	M
19	# Fields are separated by comma. Use quotes around a field that contains a comma.,,,,												
20	#,,,,												
21	# The first two lines have to be comments which start with the '#' character.,,,,												
22	# First line is 'Agilent TOF Formula data store',,,,												
23	# Second line is 'Version:' followed by a version number.,,,,												
24	#,,,,												
25	# Additionally, comments (such as these) may be inserted on individual,,,												
26	# lines by specifying a '#' character at the beginning of the line.,,,,												
27	#,,,,												
28	#,,,,												
29	### Formula, Retention Time, Mass, Compound name, Description												
30	# Formula, RT, Mass, Cpd, Comments												
31	C9H17N5S,18.46917,227.1199174,Ametryn,												
32	C8H14N5S,18.46917,212.0964423,Ametryn F1,												
33	C6H9N5S,18.46917,183.0573171,Ametryn F2,												
34	C6H10N4S,18.46917,170.0620681,Ametryn F3,												
35	C5H8N5S,18.46917,170.0495,Ametryn F4,												
36	C10H12N3O3PS2,29.38219,317.0052203,Azinphos Methyl,												
37	C8H6N3O,29.38219,160.0505379,Azinphos Methyl F1,												
38	C7H6N3,29.38219,132.0556233,Azinphos Methyl F2,												
39	C7H4N2O,29.38219,132.0318139,Azinphos Methyl F3,												
40	C8H6NO,29.38219,132.0443899,Azinphos Methyl F4,												
41	C7H4O,29.38219,104.0256659,Azinphos Methyl F5,												
42	C20H23NO3,26.00279,325.1672447,Benalaxyl,												
43	C18H20NO,26.00279,266.1539404,Benalaxyl F1,												

Database LAPRW_221113_melectron

Listo

Data base in Excel file (csv format)

Method Editor: Find Compounds by Formula - Options

Find Compounds by Formula | Method Items

Formula Source | Formula Matching | Positive Ions | Negative Ions | Results | Result Filters | Fragment Confirmation

Source of formulas to confirm

- These formulas:
(type a comma-separated list of formulas, e.g., "C6H6, CH4")
- Compound exchange file (.CEF):
- Database / Library ▲
D:\MassHunter\PCDL\default_GCMIA_2.csv ▲
- Worklist

Matches per formula

Maximum number of matches: 1

Automatically increase for isomeric compounds

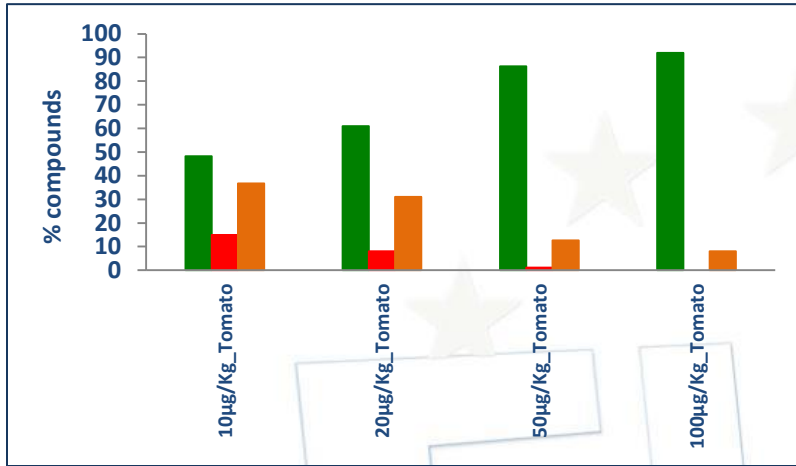
Values to match

Retention time tolerance: 0,5 min
Mass error tolerance 10 ppm

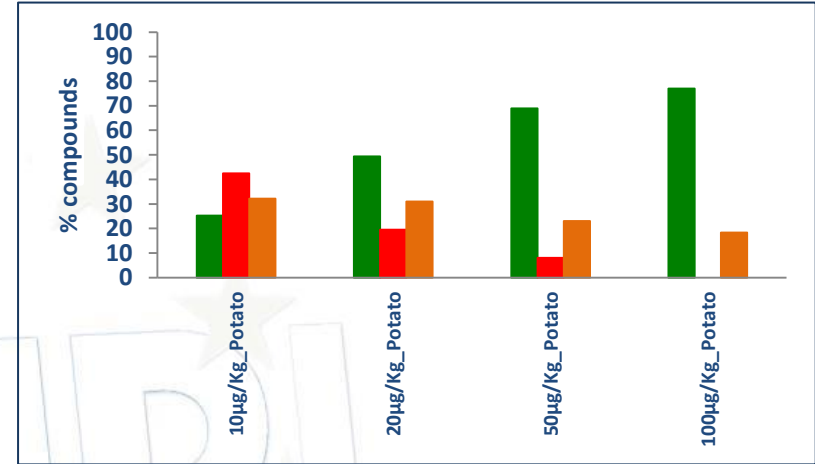
Data base in Excel file (csv format)

AUTOMATIC IDENTIFICATION

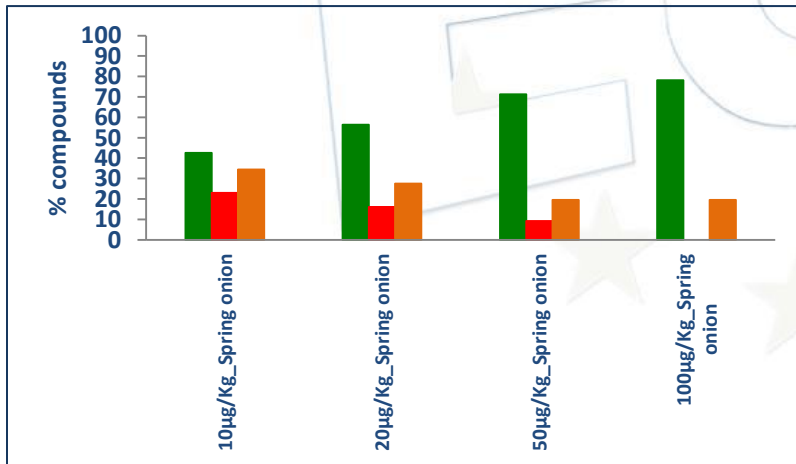
TOMATO



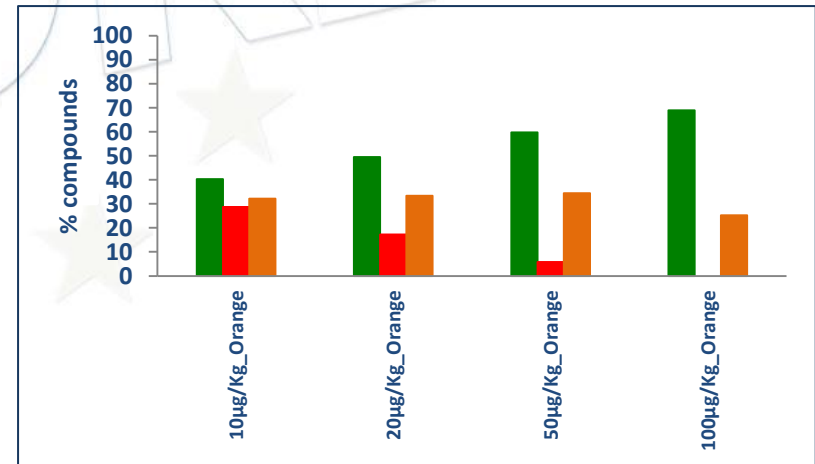
POTATO



SPRING ONION



ORANGE



■ % DETECTED

■ % NOT DETECTED

■ % ONE FRAGMENT DETECTED

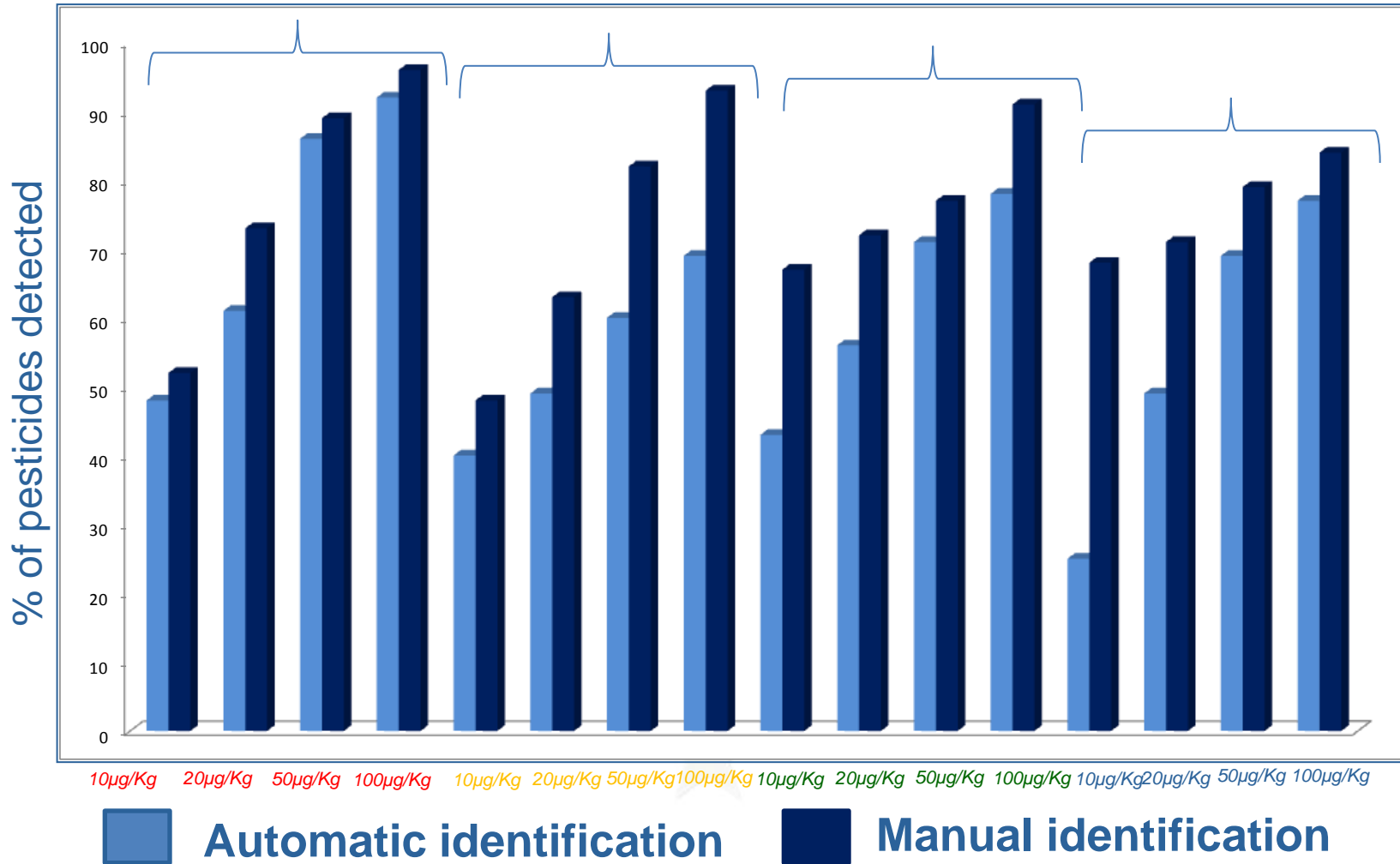
AUTOMATIC IDENTIFICATION VS MANUAL IDENTIFICATION

TOMATO

ORANGE

SPRING ONION

POTATO

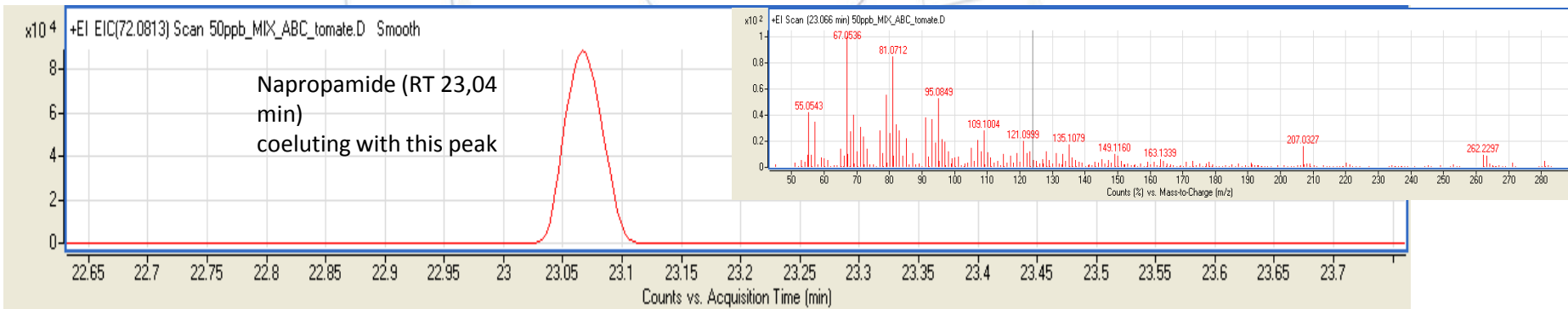
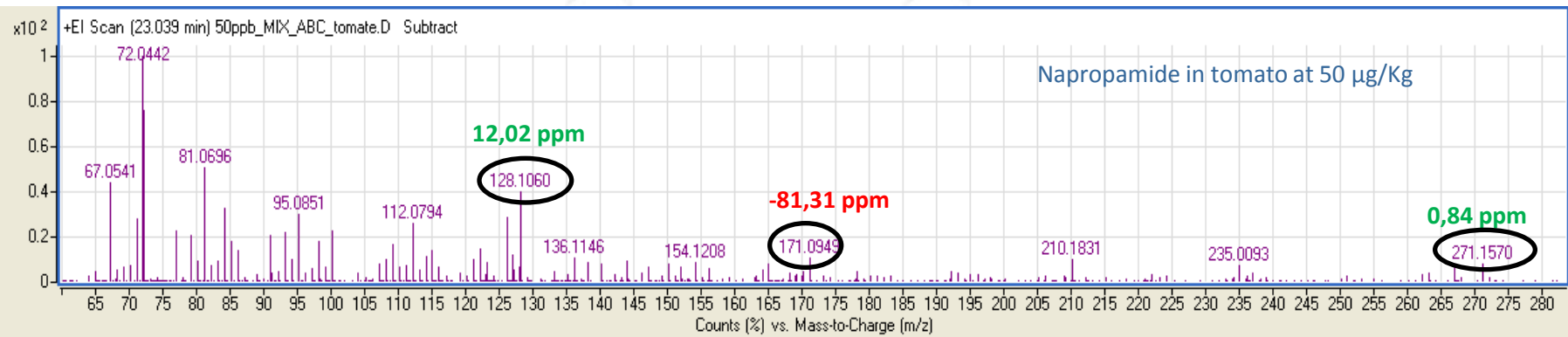


NOT AUTOMATICALLY IDENTIFIED

One ion is automatically detected

Cpd	Label	Name	Formula	Score	Mass	Mass (DB)	Diff (Tgt, ppm)	Diff (Tgt, mDa)
106	Cpd 106: Napropamide	Napropamide	C17 H21 N O2	81,36	271,1563	271,1572	-3,6	-0,98

C17H21NO2	22,97	271,1572	Napropamide
C12H11O	22,97	171,081	Napropamide F1
C7H14NO	22,97	128,1075	Napropamide F3



A large, faint watermark of the EURL logo, consisting of the letters 'EURL' in a bold, sans-serif font, surrounded by a circle of twelve yellow stars, similar to the European Union flag.

QUANTIFICATION

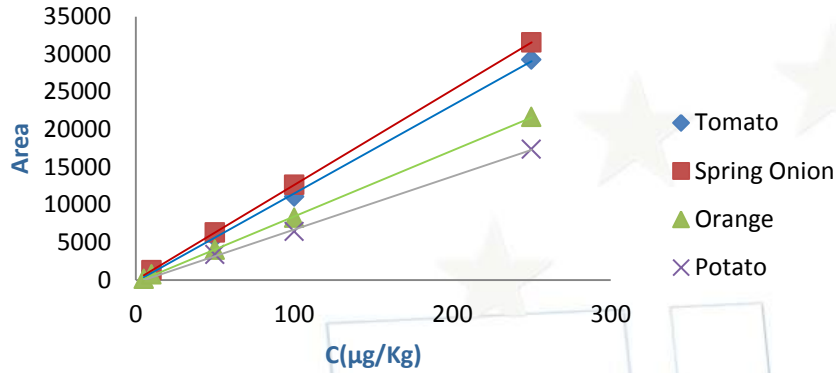
A large, faint watermark of the EURL logo, consisting of the letters 'EURL' in a stylized font surrounded by twelve yellow stars, is centered in the background.

MATRIX EFFECT

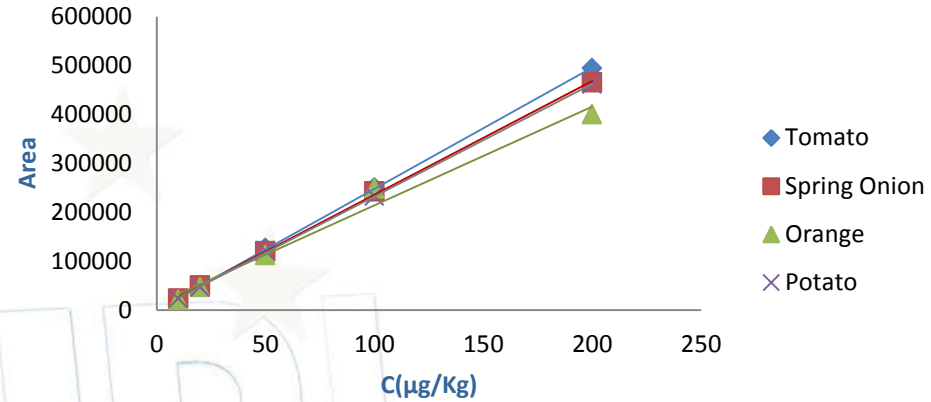
GC-QqQ/MS

GC-QTOF/MS

Chlozolate



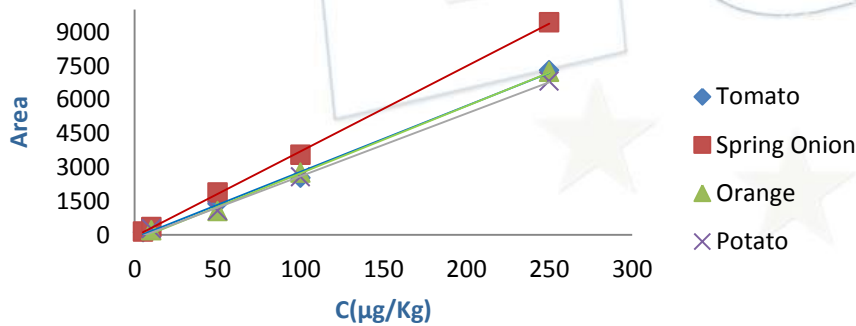
Chlozolate



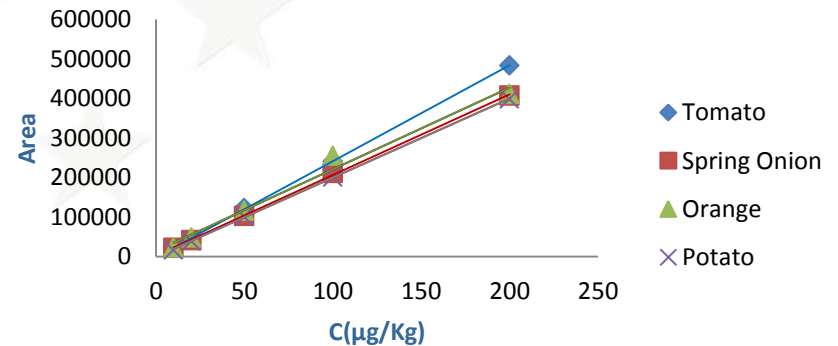
5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Endosulphan sulfate



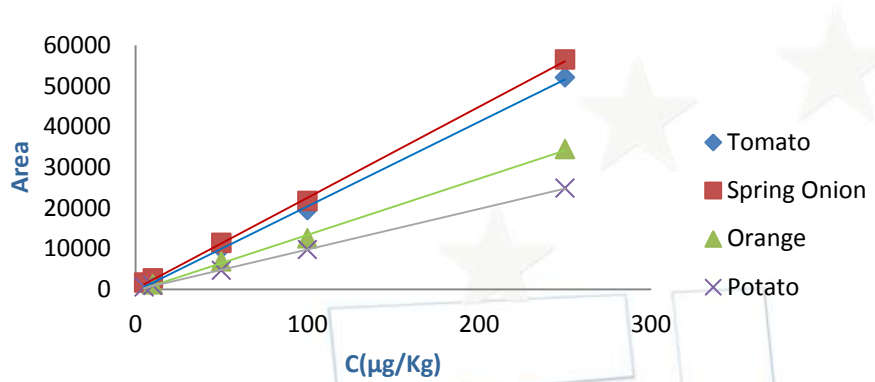
Endosulphan sulfate



GC-QqQ/MS

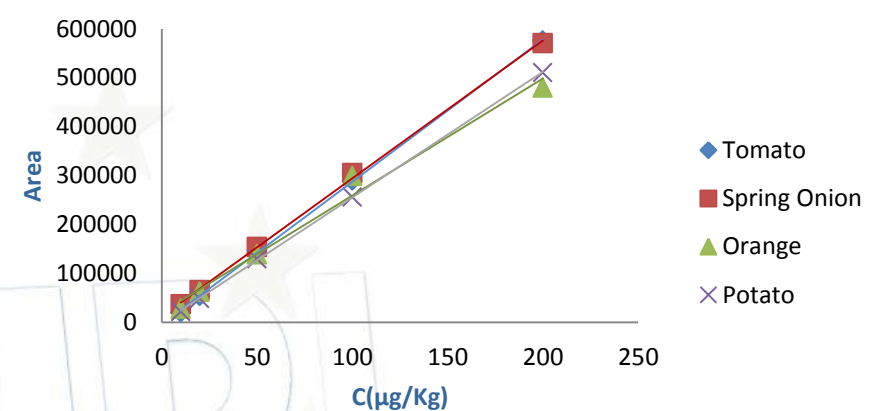
GC-QTOF/MS

Metalaxyl



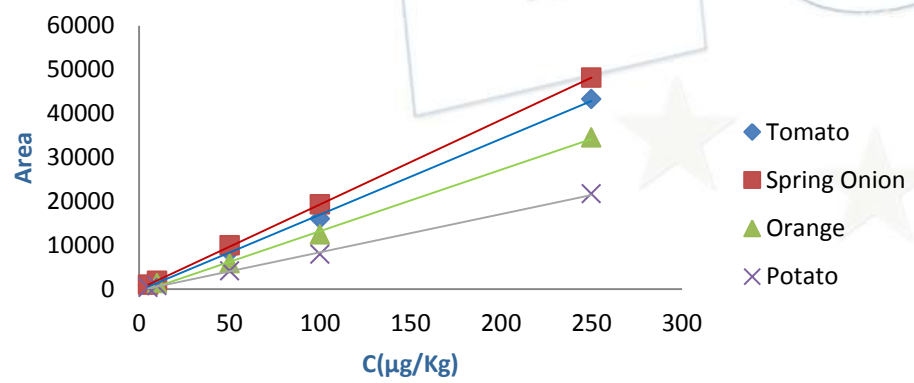
5-10-50-100-250µg/Kg

Metalaxyl

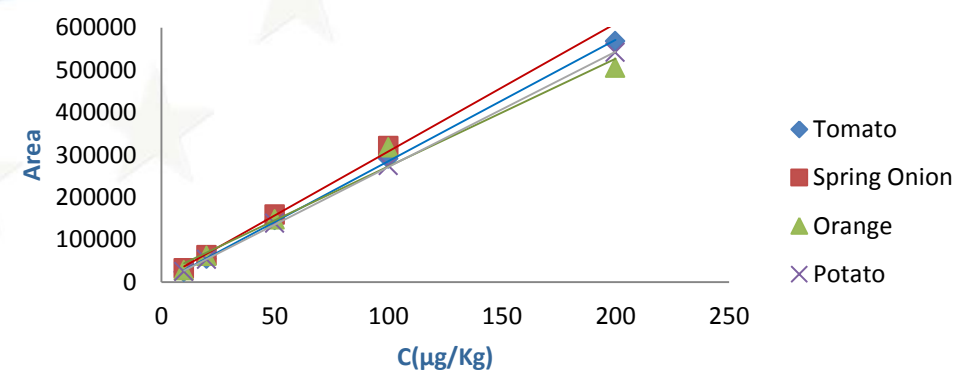


10-20-50-100-200µg/Kg

Vinclozolin



Vinclozolin

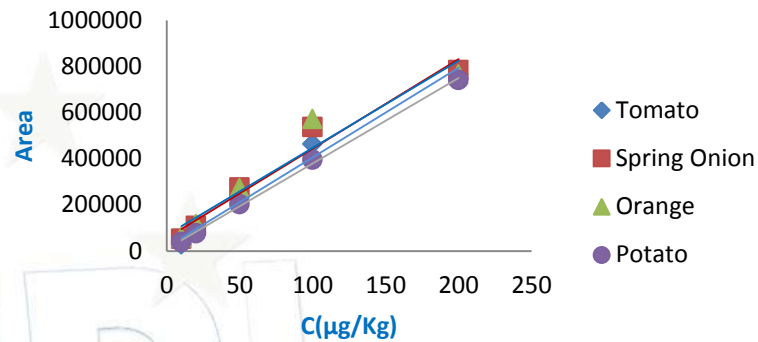
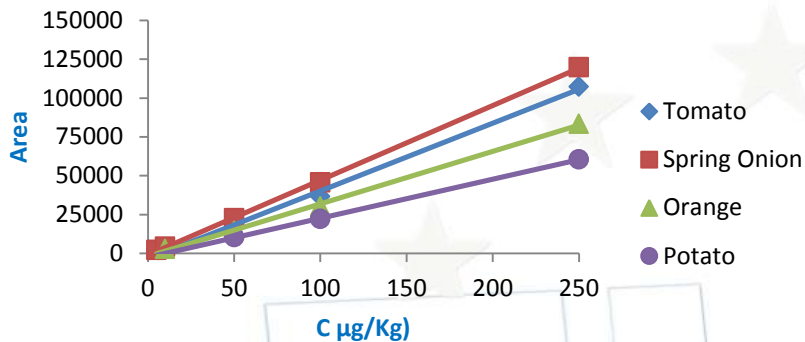


GC-QqQ/MS

GC-QTOF/MS

Ametryn

Ametryn

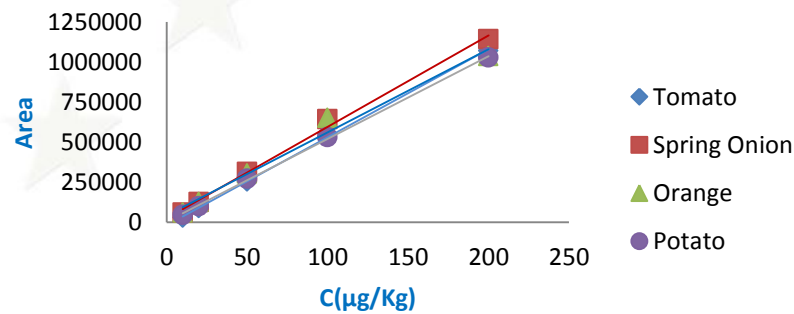
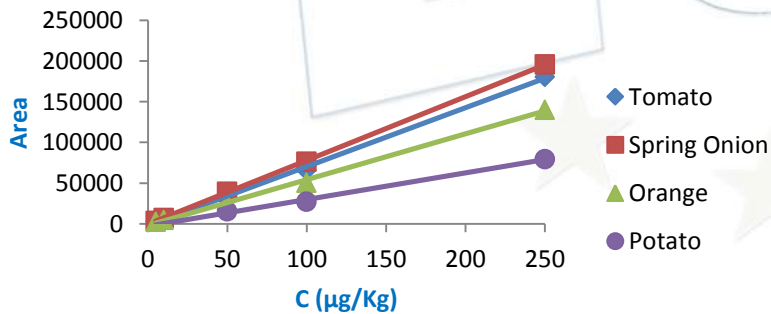


5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Cadusafos

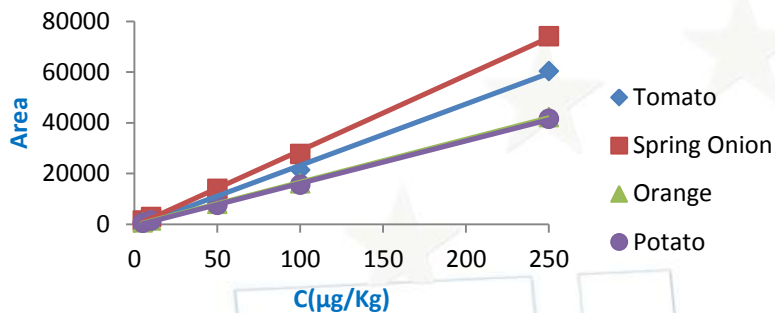
Cadusafos



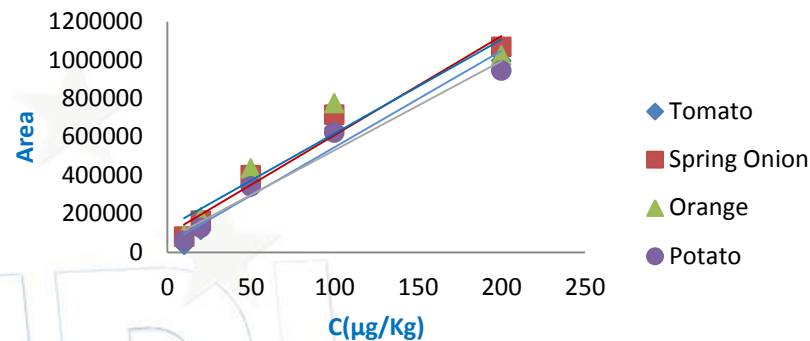
GC-QqQ/MS

GC-QTOF/MS

Chlorpyrifos-Methyl



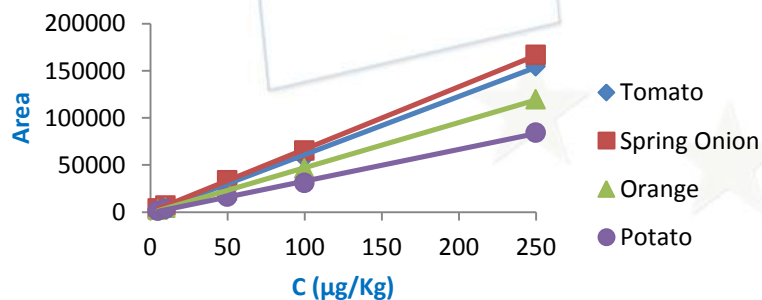
Chlorpyrifos_Methyl



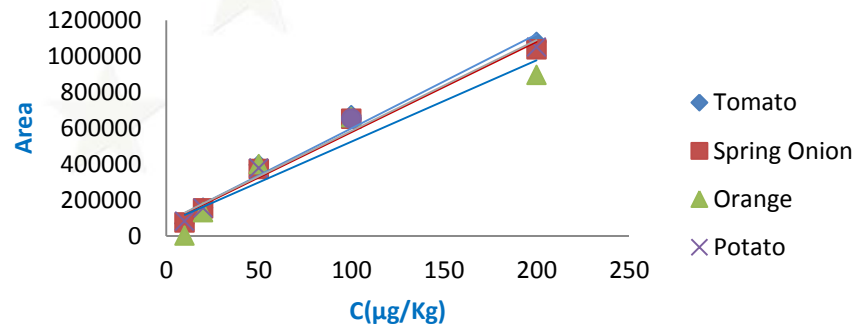
5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Ethofumesate



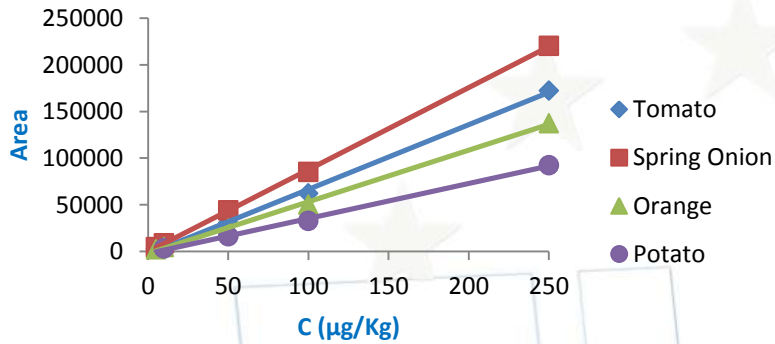
Ethofumesate



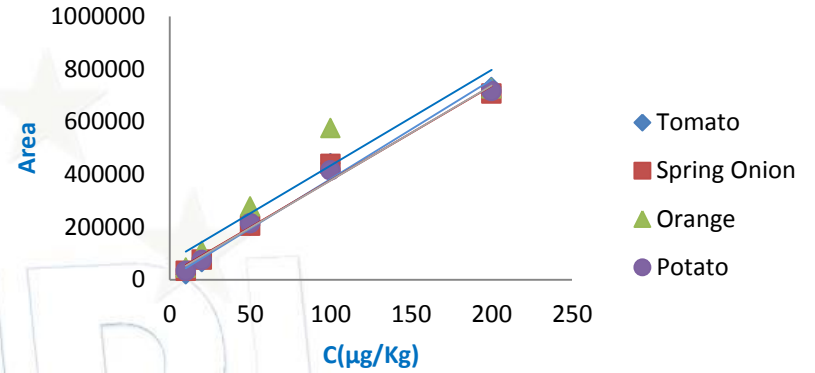
GC-QqQ/MS

GC-QTOF/MS

Malathion



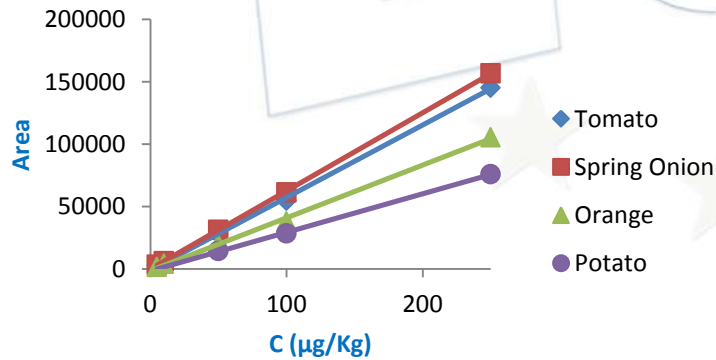
Malathion



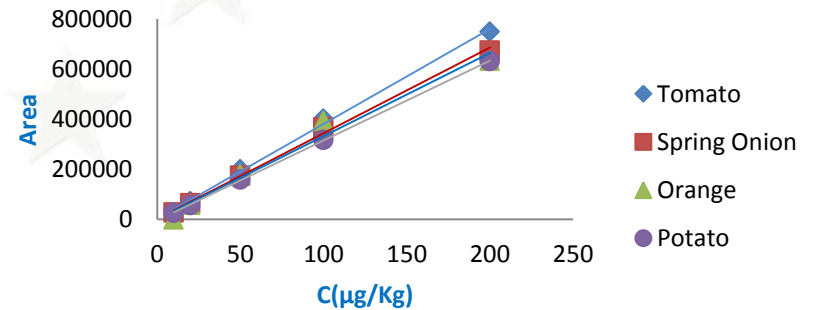
5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Nuarimol



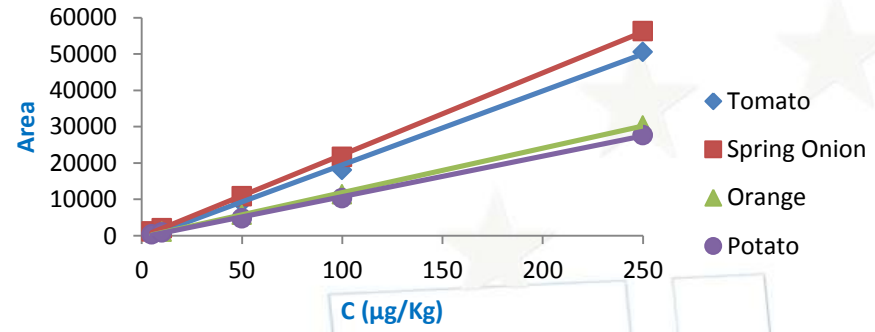
Nuarimol



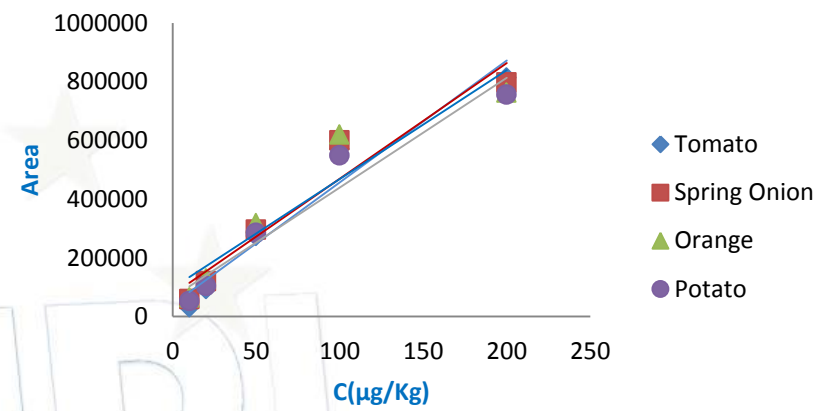
GC-QqQ/MS

GC-QTOF/MS

Pirimiphos-Methyl



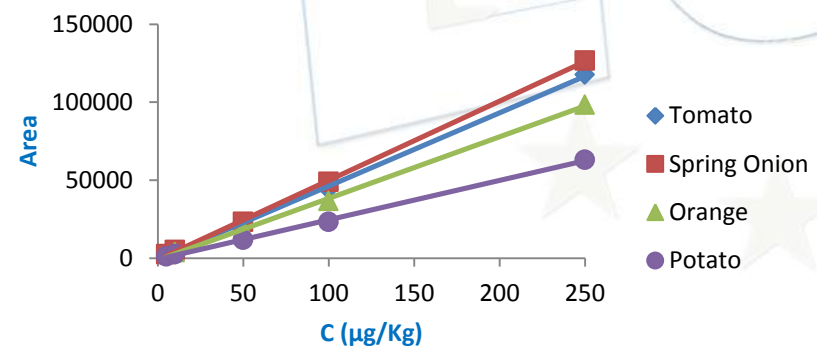
Pirimiphos-Methyl



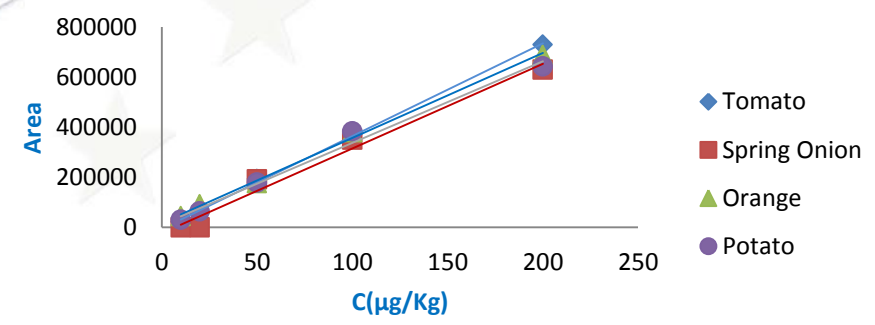
5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Triadimefon



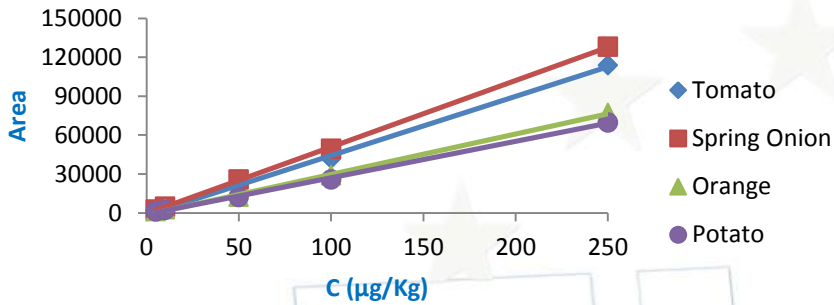
Triadimefon



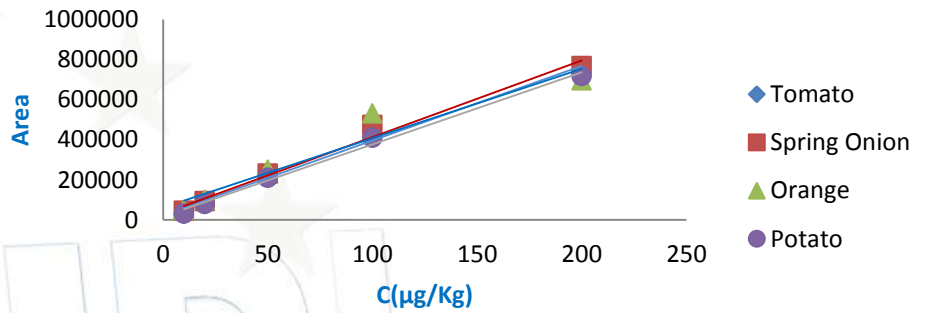
GC-QqQ/MS

GC-QTOF/MS

Prothiophos



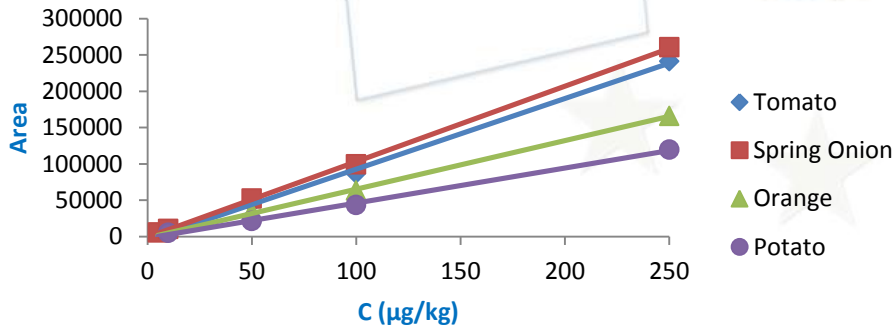
Prothiophos



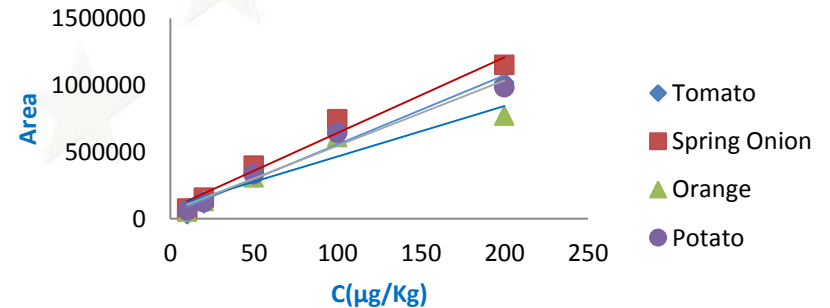
5-10-50-100-250µg/Kg

10-20-50-100-200µg/Kg

Terbumeton



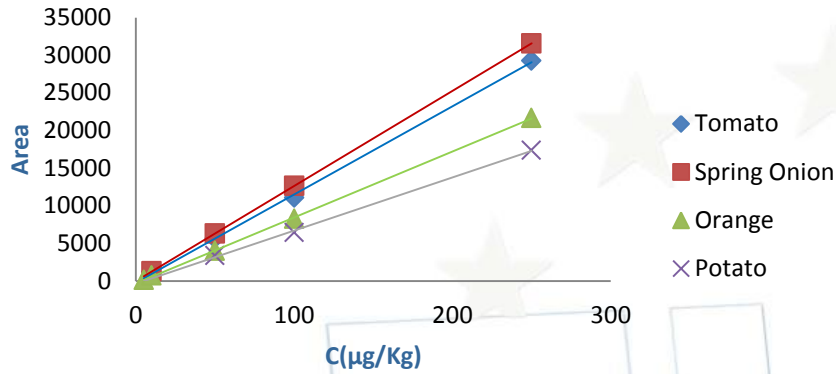
Terbumeton



GC-QqQ/MS

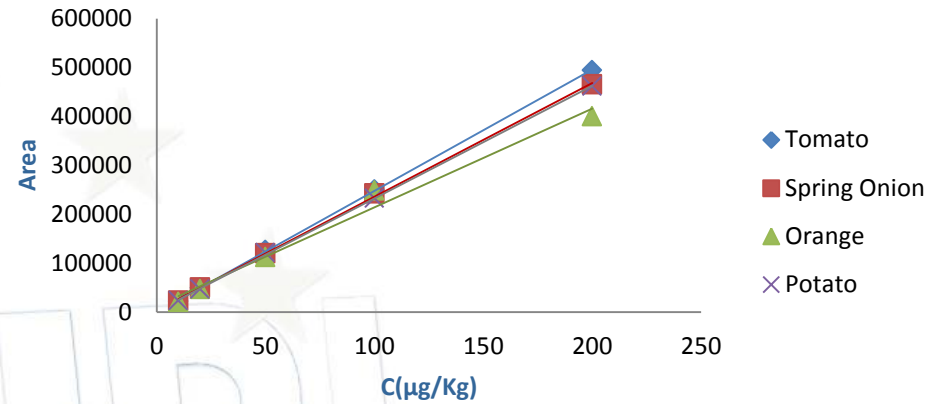
GC-QTOF/MS

Chlozolate



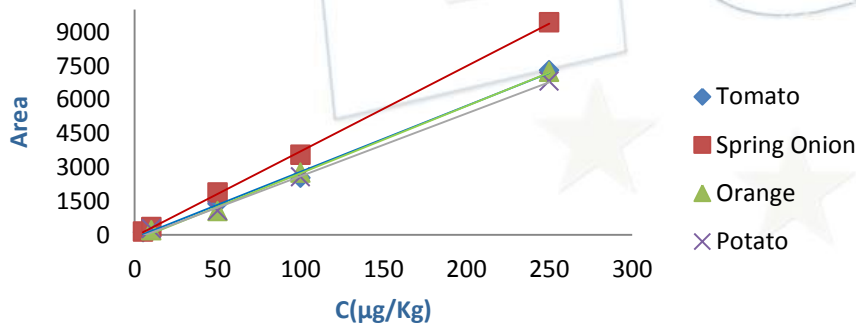
5-10-50-100-250µg/Kg

Chlozolate

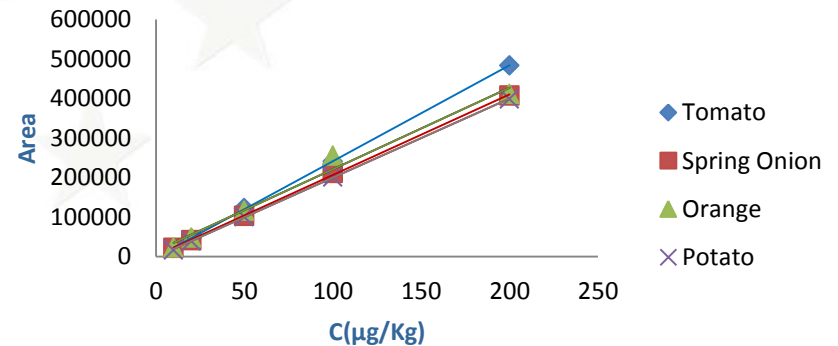


10-20-50-100-200µg/Kg

Endosulphan sulfate



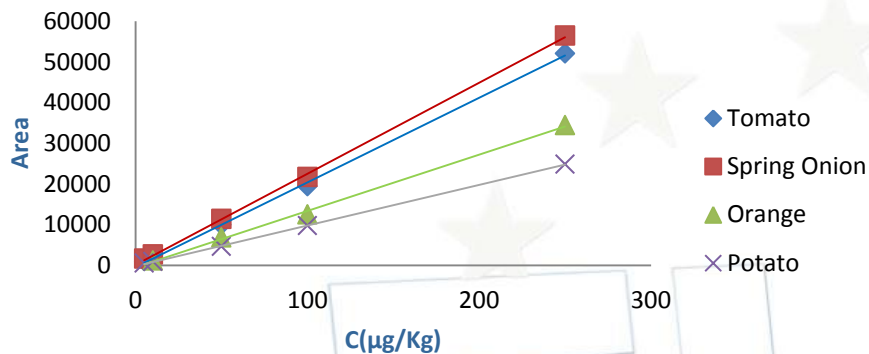
Endosulphan sulfate



GC-QqQ/MS

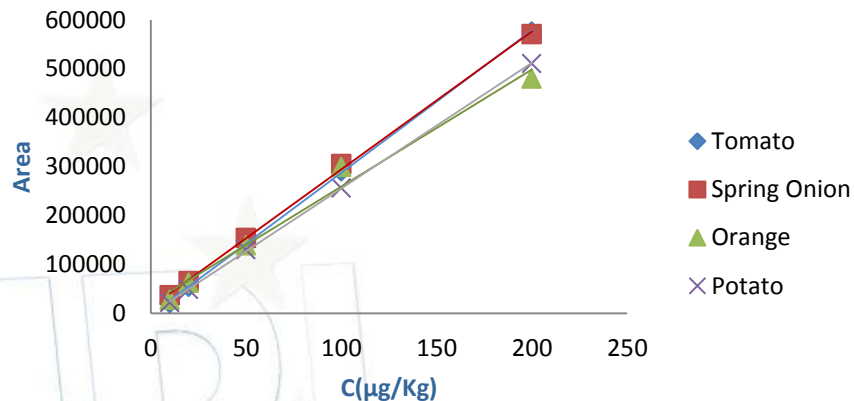
GC-QTOF/MS

Metalaxyl



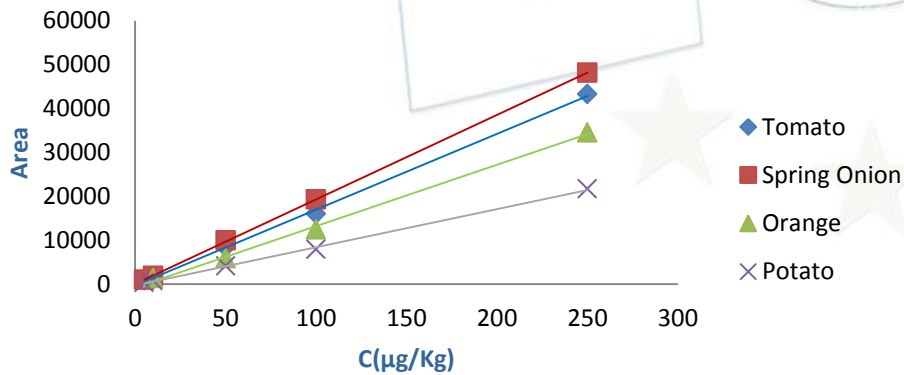
5-10-50-100-250µg/Kg

Metalaxyl

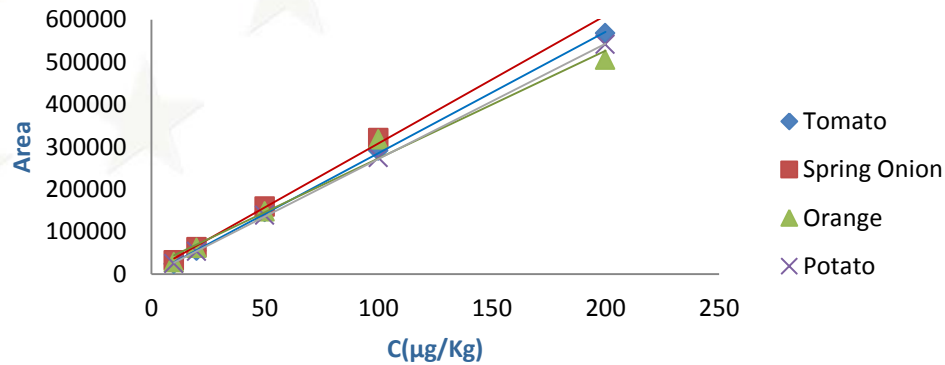


10-20-50-100-200µg/Kg

Vinclozolin

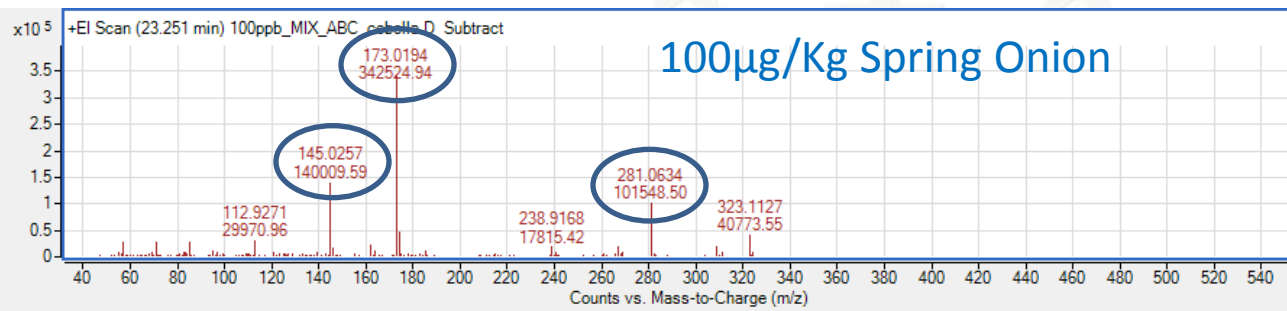
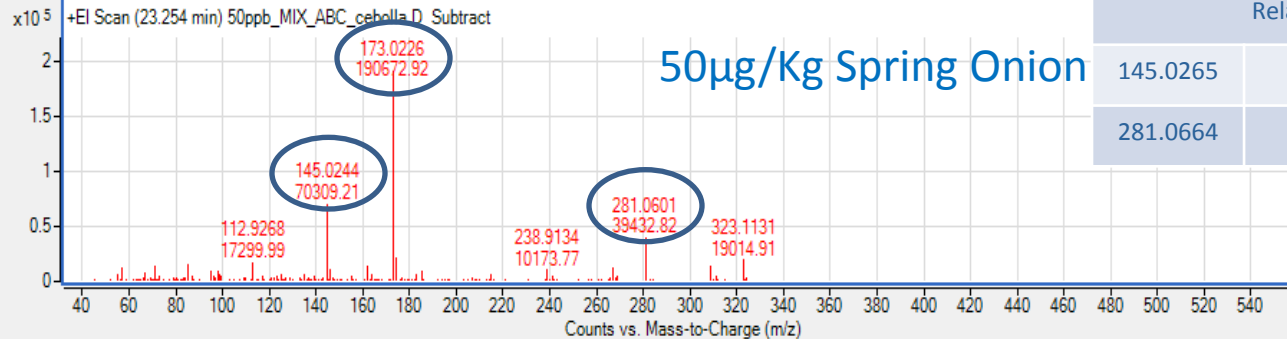
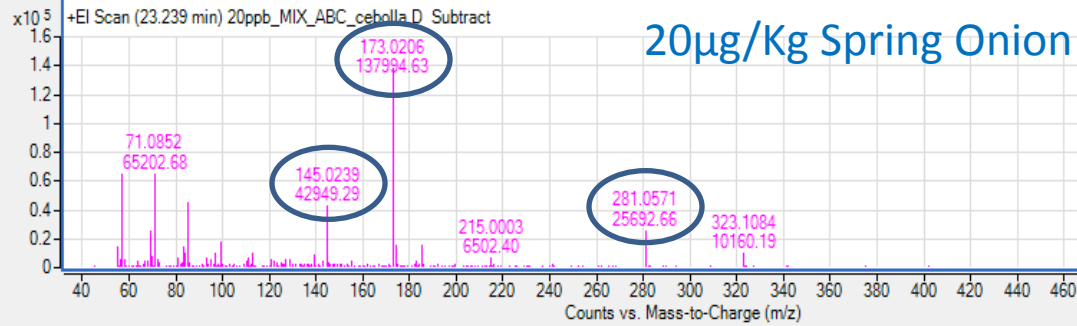
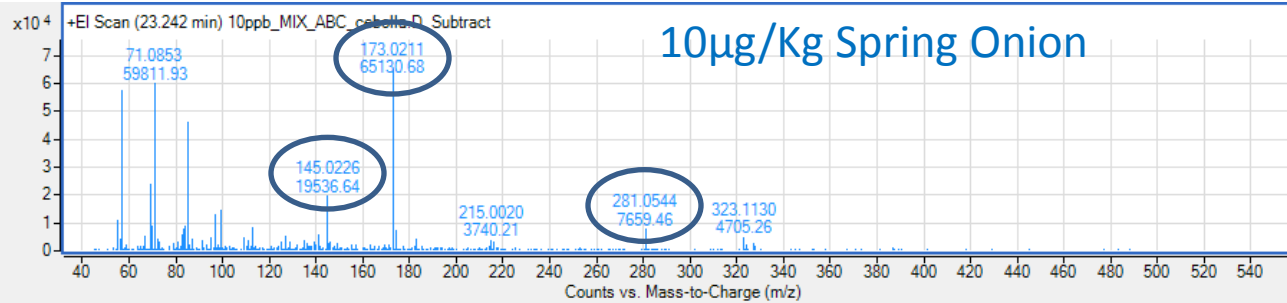


Vinclozolin

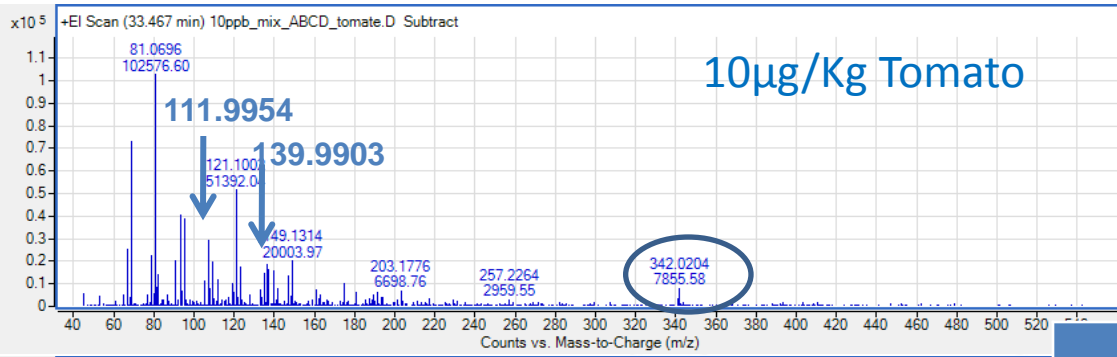


RELATIVE ABUNDANCES AT DIFFERENT CONCENTRATIONS

FLUTOLANIL

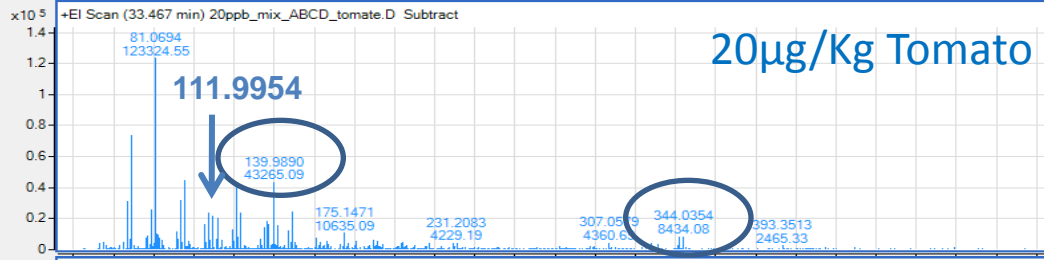


	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
173.0214	65130.68	137994.63	190672.92	342524.94
145.0265	19536.64	42949.29	70309.21	140009.59
281.0664	7659.46	25692.66	39432.82	101548.50
Relative Abundances to base peak 173.0214				
145.0265	30%	31%	37%	41%
281.0664	12%	19%	21%	30%

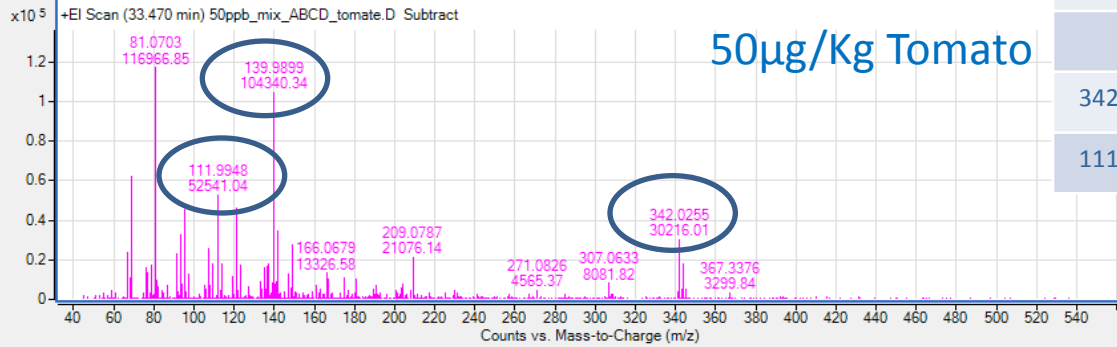


10µg/Kg Tomato

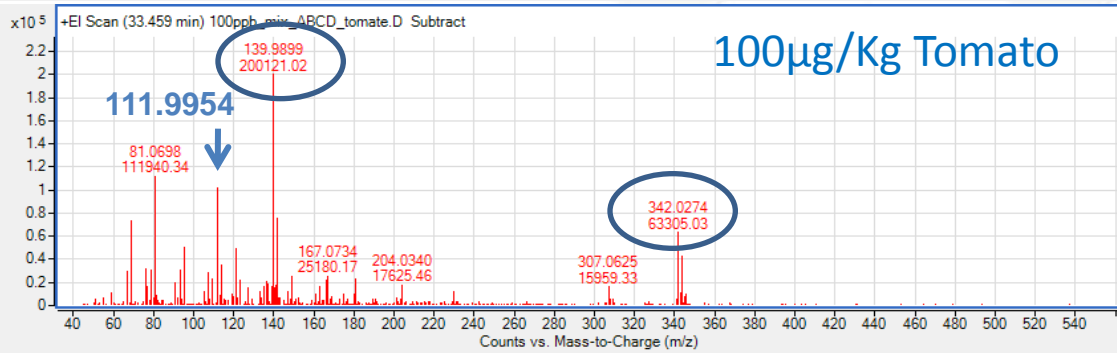
BOSCALID



20µg/Kg Tomato



50µg/Kg Tomato

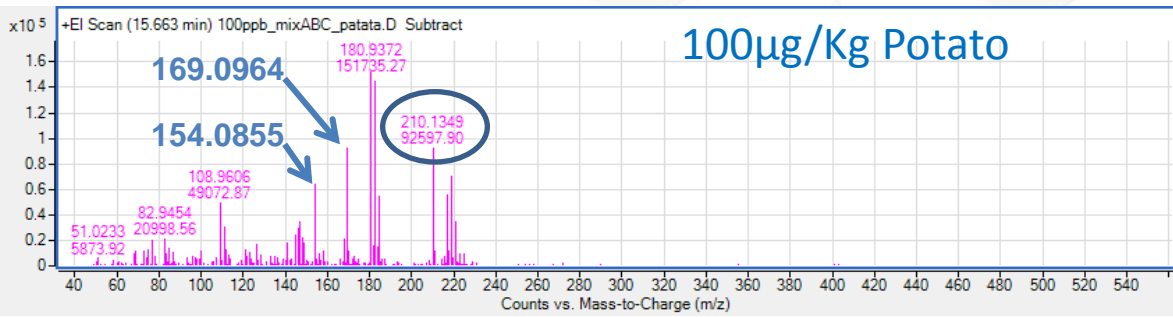
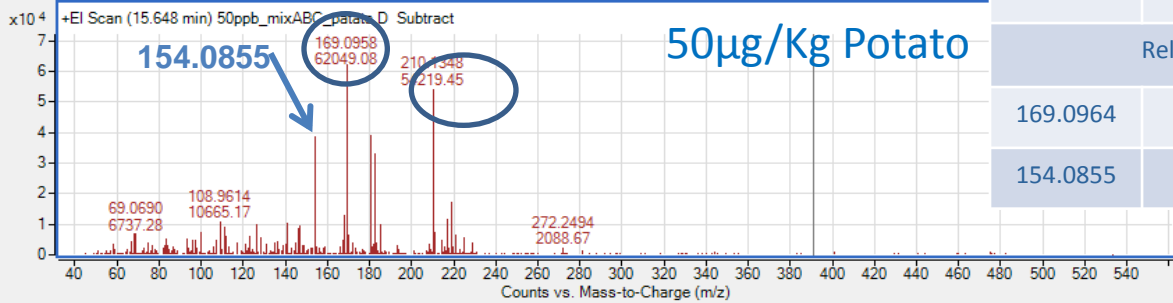
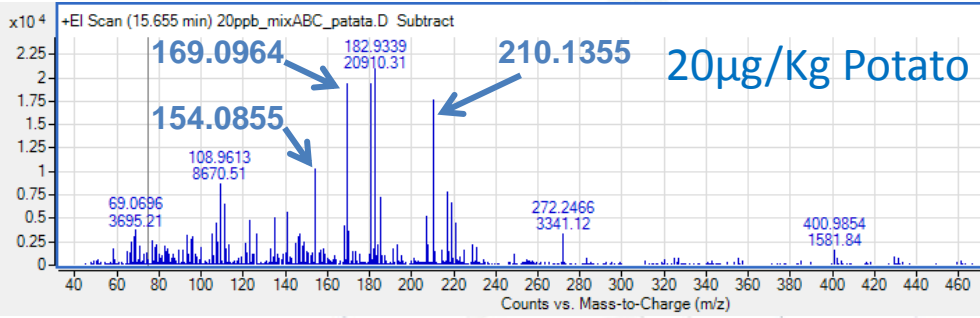
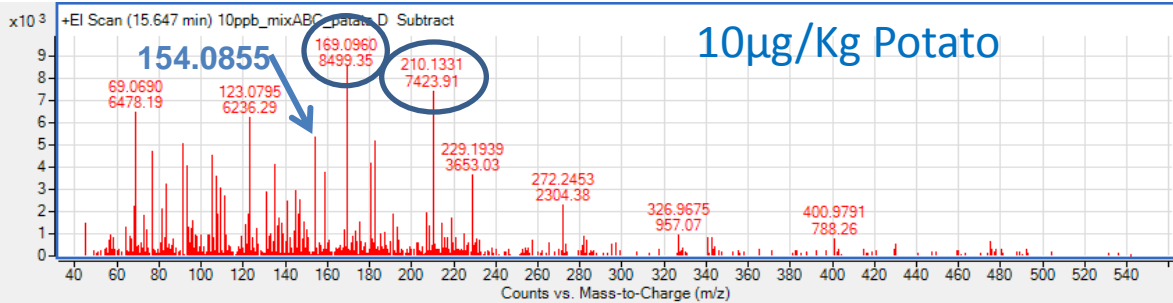


100µg/Kg Tomato

	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
139.9903	15757.57	43265.09	104340	200121
342.0327	7855.58	8177.92	30216	63305.03
111.9954	11613.48	20109.98	52541	102291
Relative Abundances to base peak 139.9903				
342.0327	44%	19%	30%	32%
111.9954	74%	46%	50%	51%



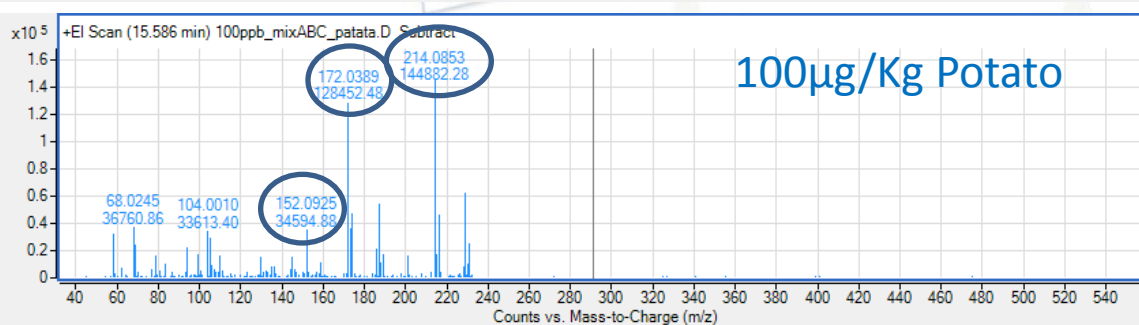
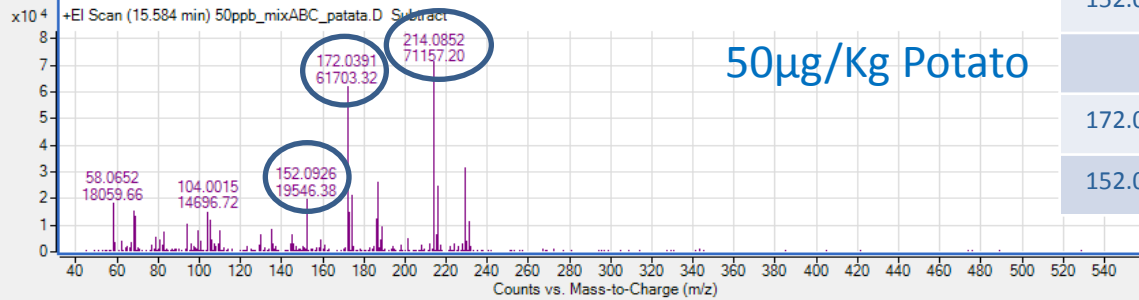
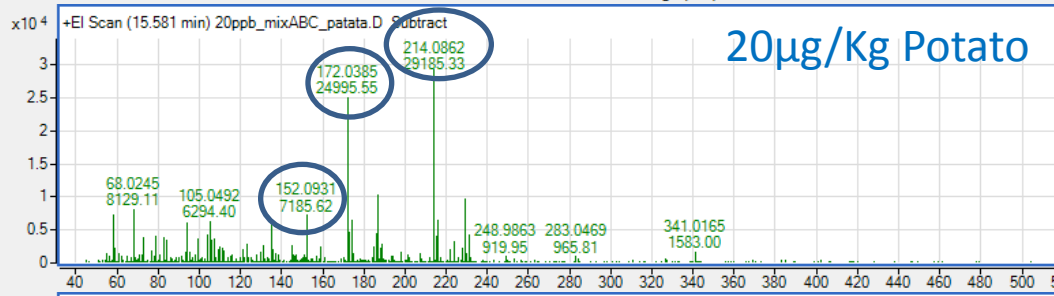
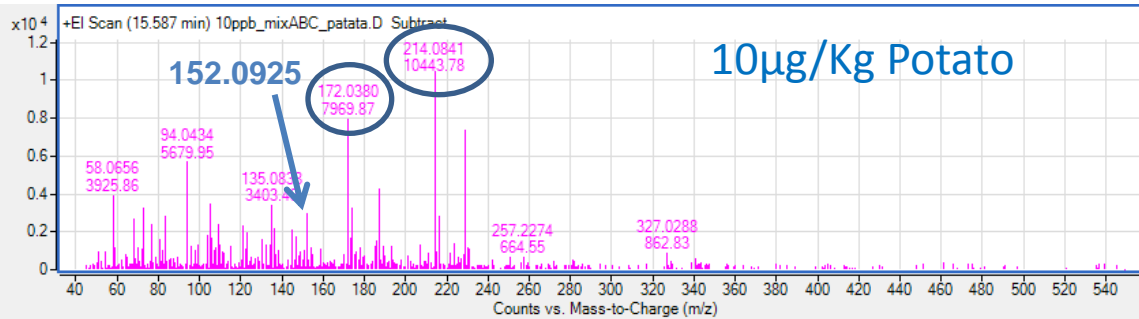
TERBUMETON



	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
210.1355	7423.91	17674.94	54219.45	92597.90
169.0964	8499.35	19325.86	62049.08	91847.98
154.0855	5352.16	10216.07	38487.03	63805.87
Relative Abundances to base peak 210.1355				
169.0964	114%	109%	114%	99%
154.0855	72%	58%	71%	69%

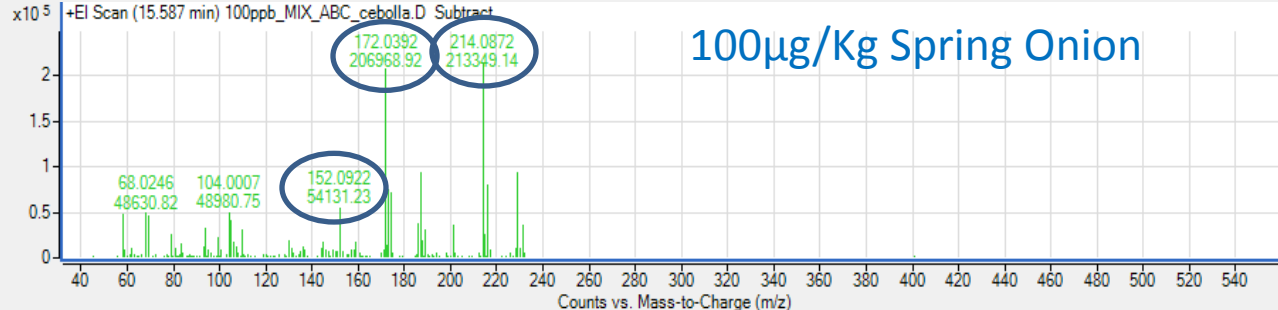
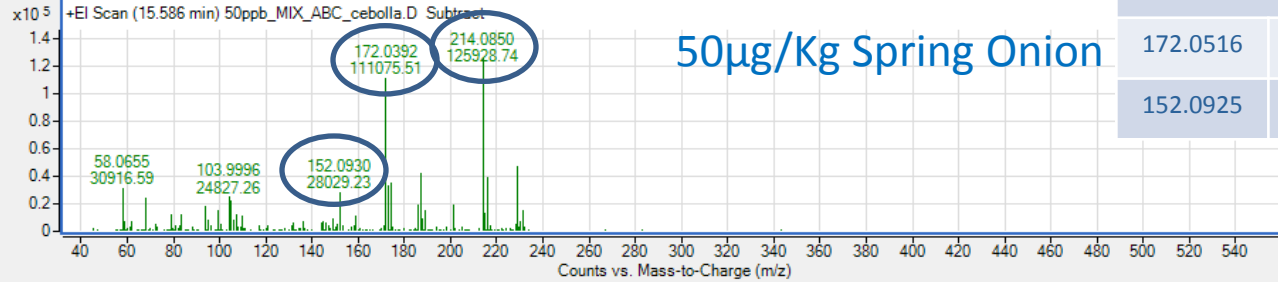
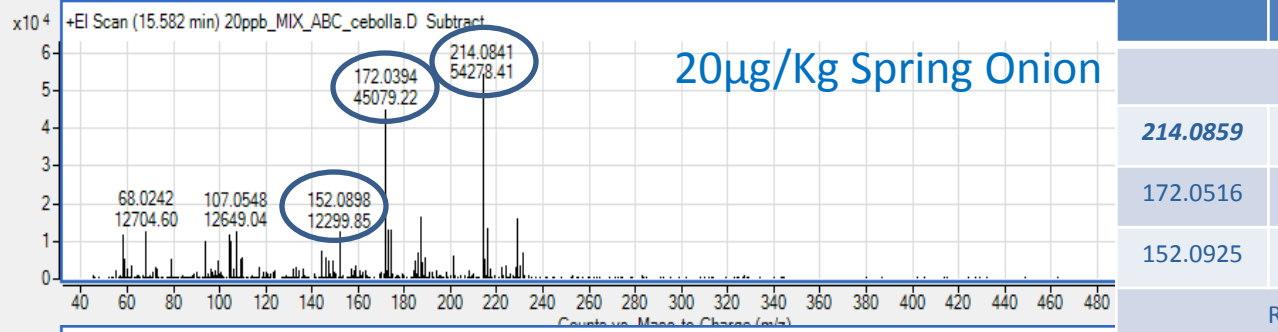
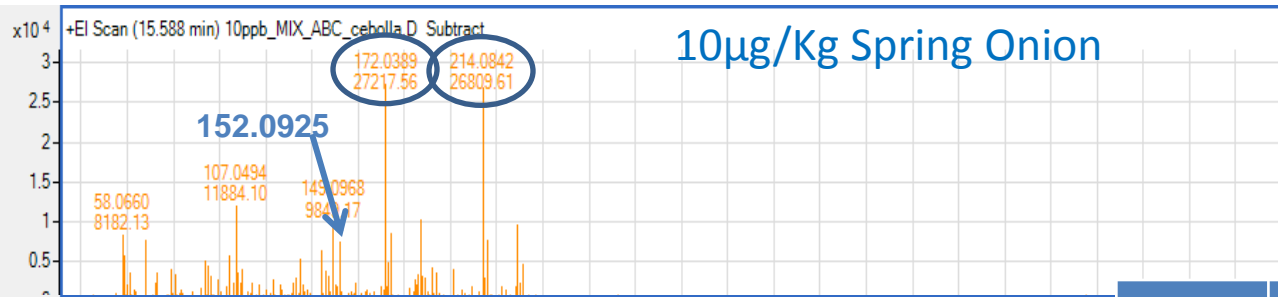
RELATIVE ABUNDANCES IN DIFFERENT MATRICES

PROPAZINE



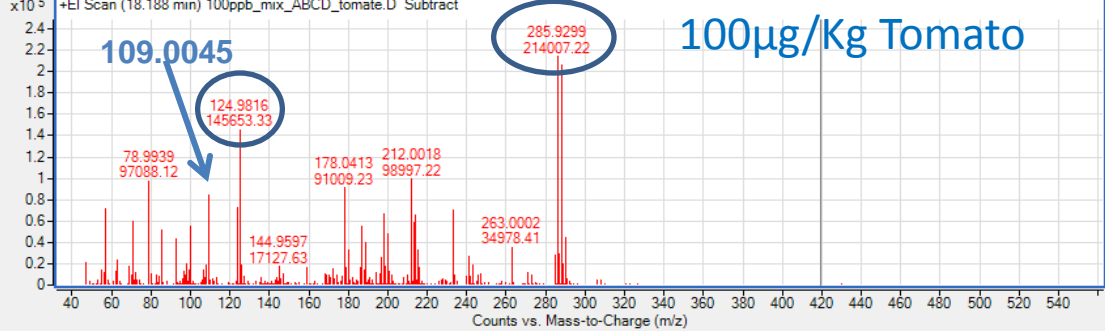
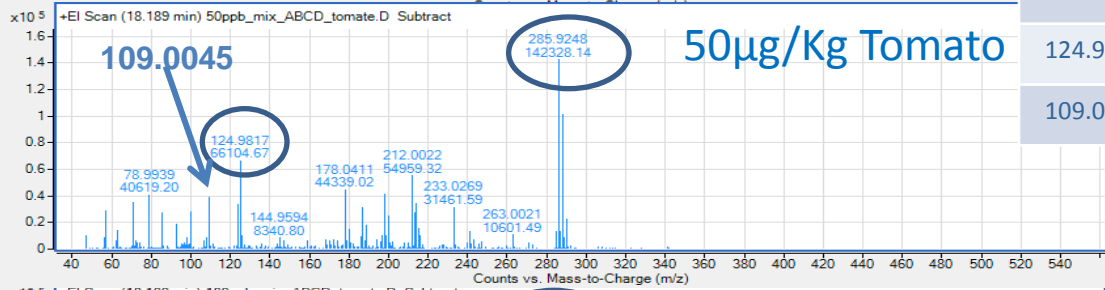
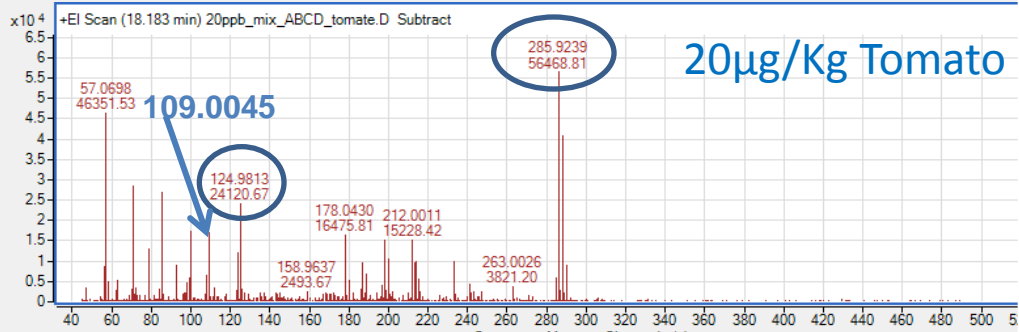
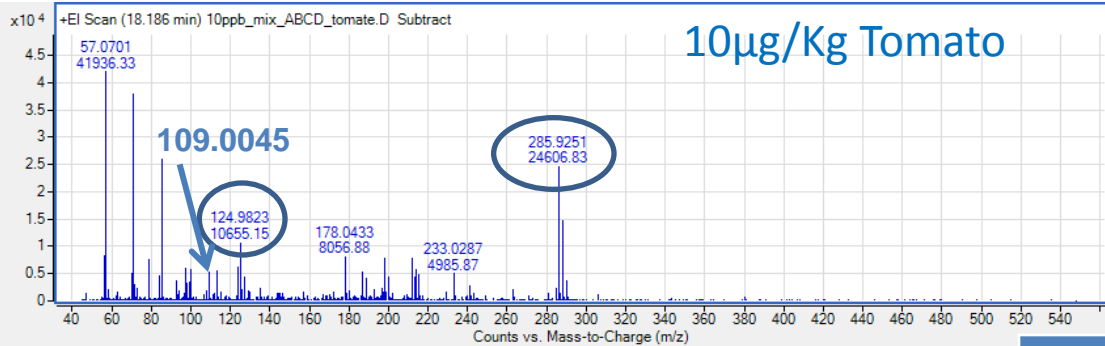
	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
214.0859	10443.78	29185.33	71157.20	144882.28
172.0516	7969.87	24995.55	61703.32	128452.48
152.0925	2947.45	7185.62	19546.38	34594.88
Relative Abundances to base peak 214.0859				
172.0516	76%	86%	87%	89%
152.0925	28%	25%	27%	24%

PROPAZINE



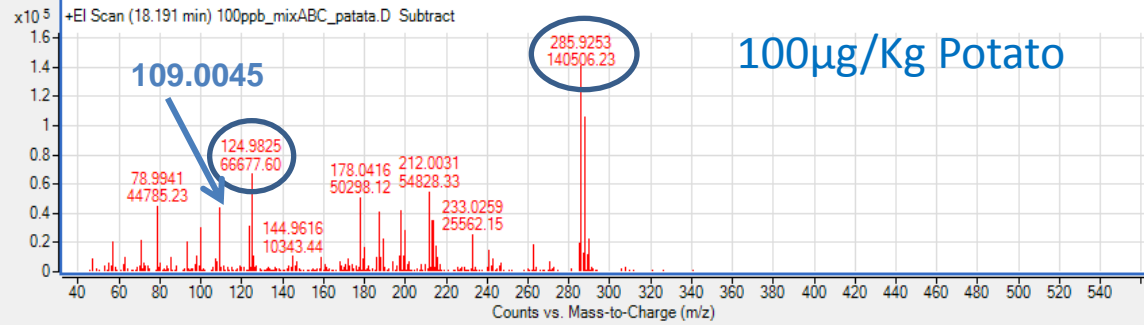
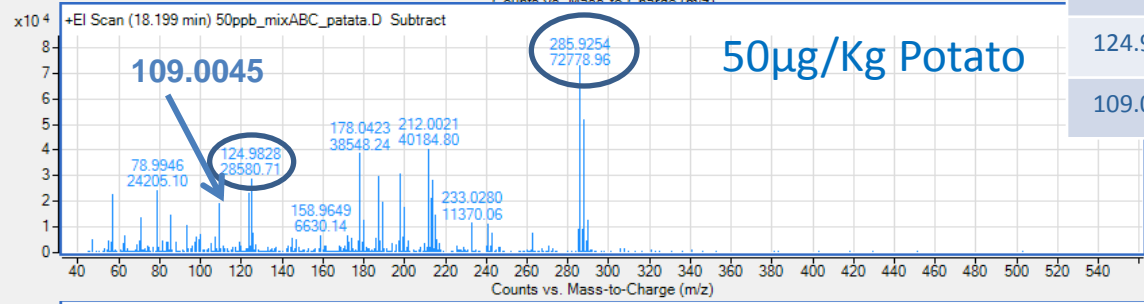
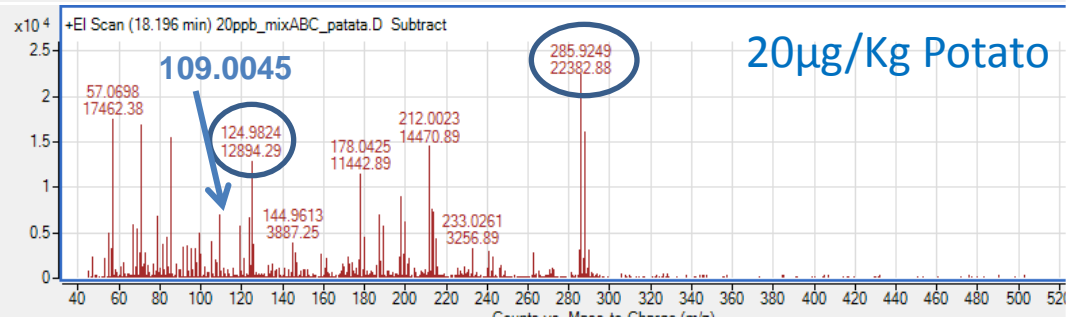
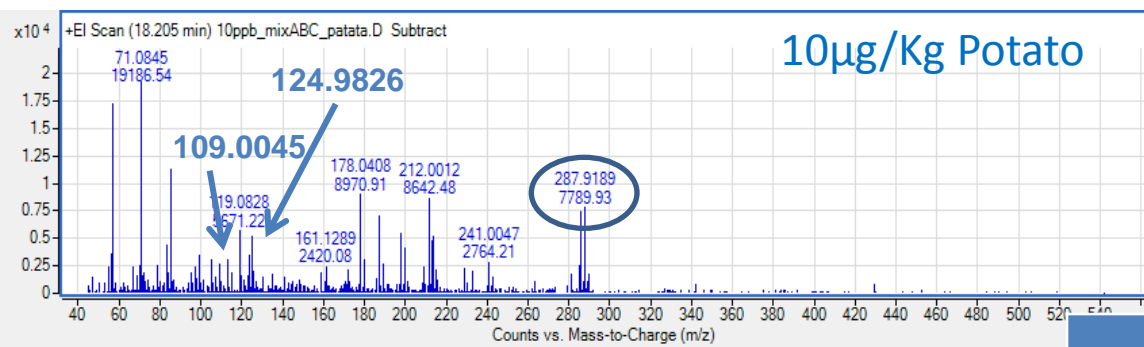
	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
214.0859	26809.61	54278.41	125928.74	213349.14
172.0516	27217.56	45079.22	111075.51	206968.92
152.0925	7313.01	12299.85	28029.23	54131.23
Relative Abundances to base peak 214.0859				
172.0516	102%	83%	88%	97%
152.0925	27%	23%	22%	25%

CHLORPYRIFOS METHYL



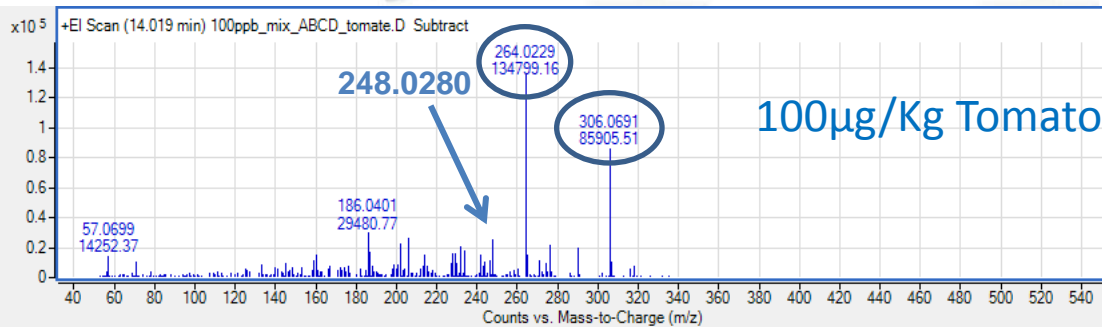
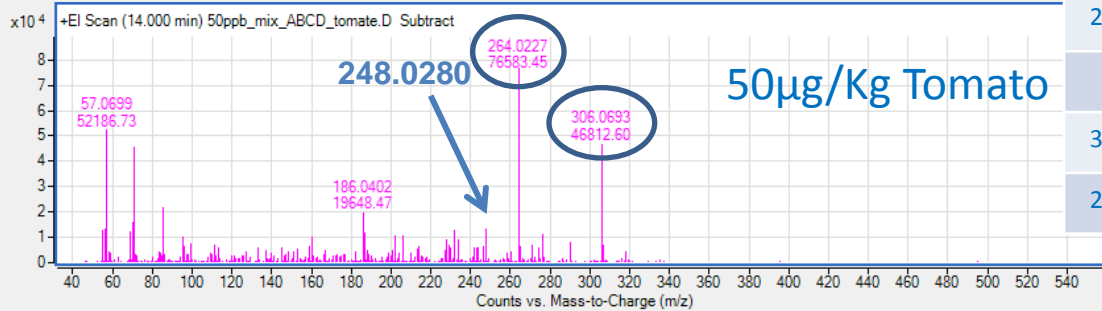
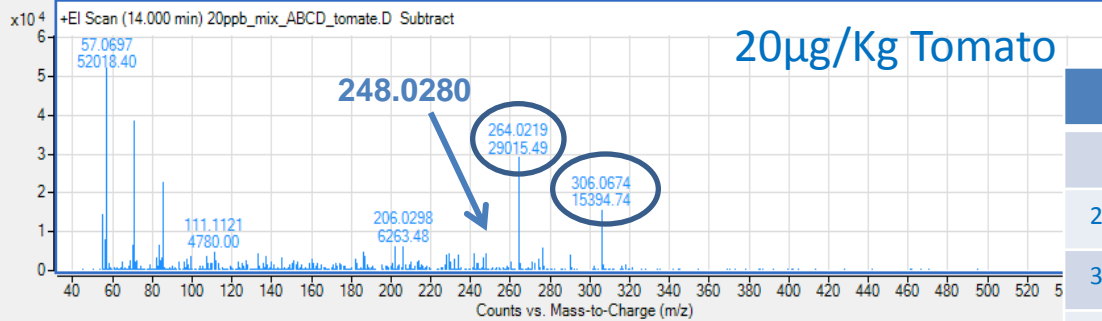
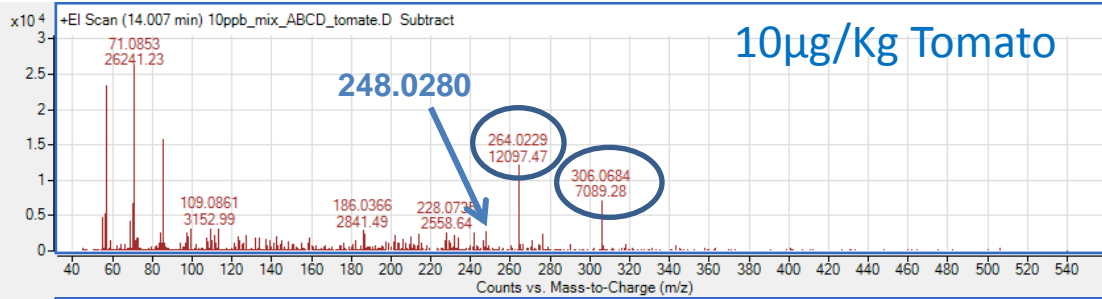
	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
285.9261	24606.83	56468.81	142328	214007.22
124.9826	10655.15	24120.67	66104.67	145653.33
109.0045	5369.61	16857.85	39130.32	84063.63
Relative Abundances to base peak 285.9261				
124.9826	43%	43%	46%	68%
109.0045	22%	30%	27%	39%

CHLORPYRIFOS METHYL



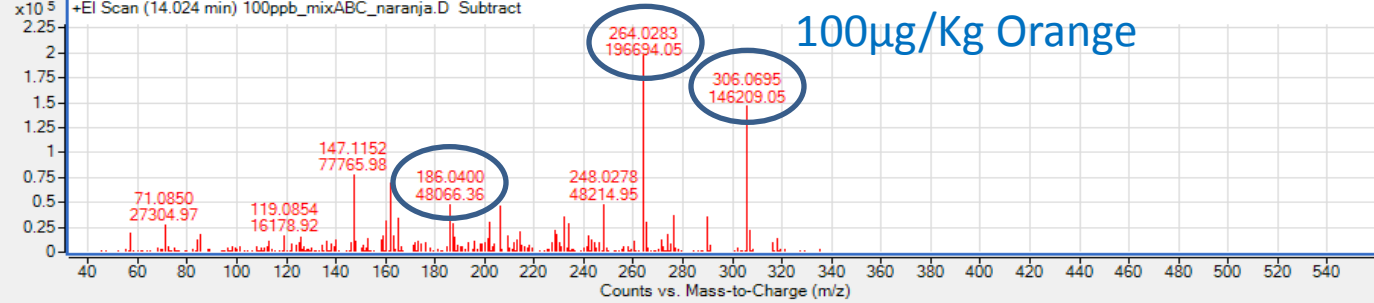
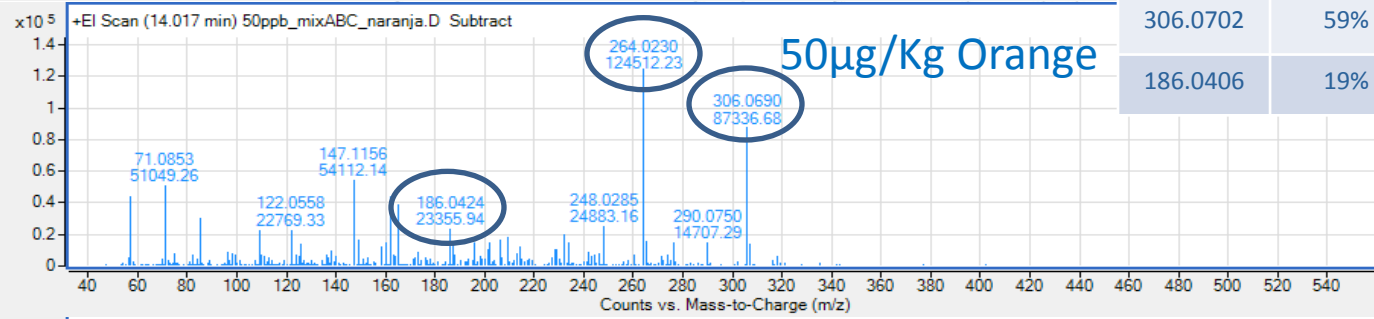
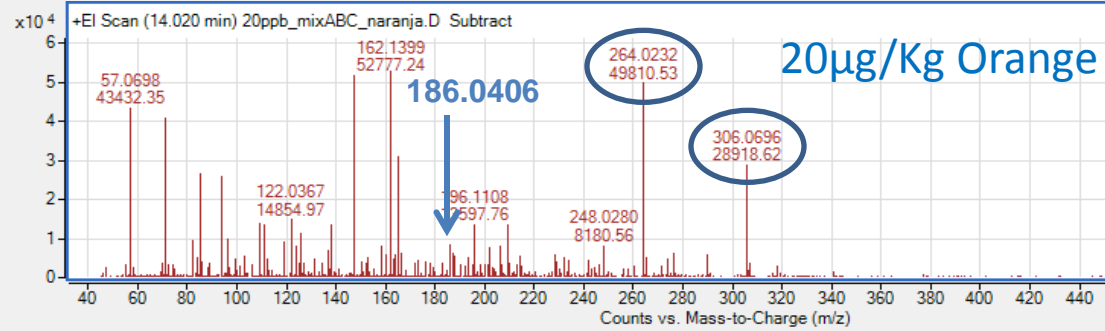
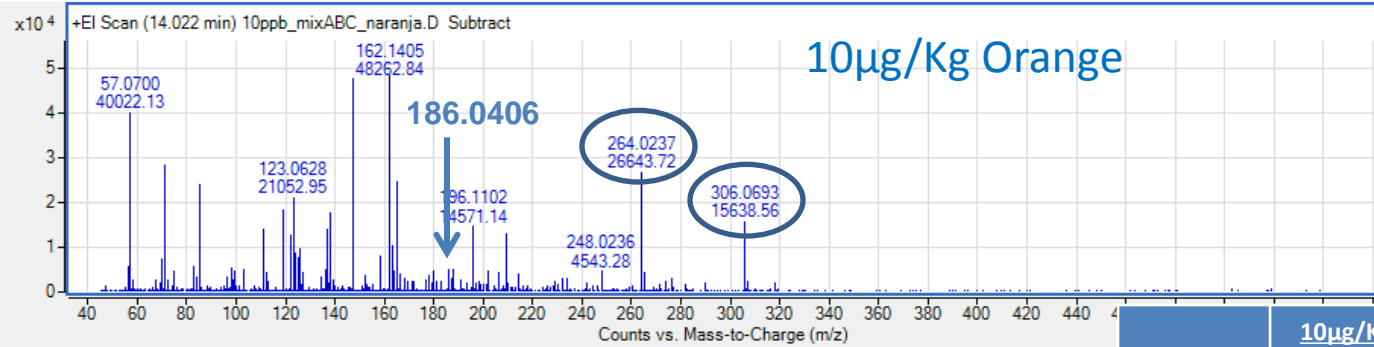
	<u>10µg/Kg</u>	<u>20µg/Kg</u>	<u>50µg/Kg</u>	<u>100µg/Kg</u>
Absolute Abundances				
285.9261	7434.42	22382.88	72778.96	140506.23
124.9826	5132.83	12894.29	28580.71	66677.60
109.0045	2606.15	6930.35	19063.65	44127.69
Relative Abundances to base peak 285.9261				
124.9826	69%	58%	39%	47%
109.0045	35%	31%	26%	31%

TRIFLURALIN



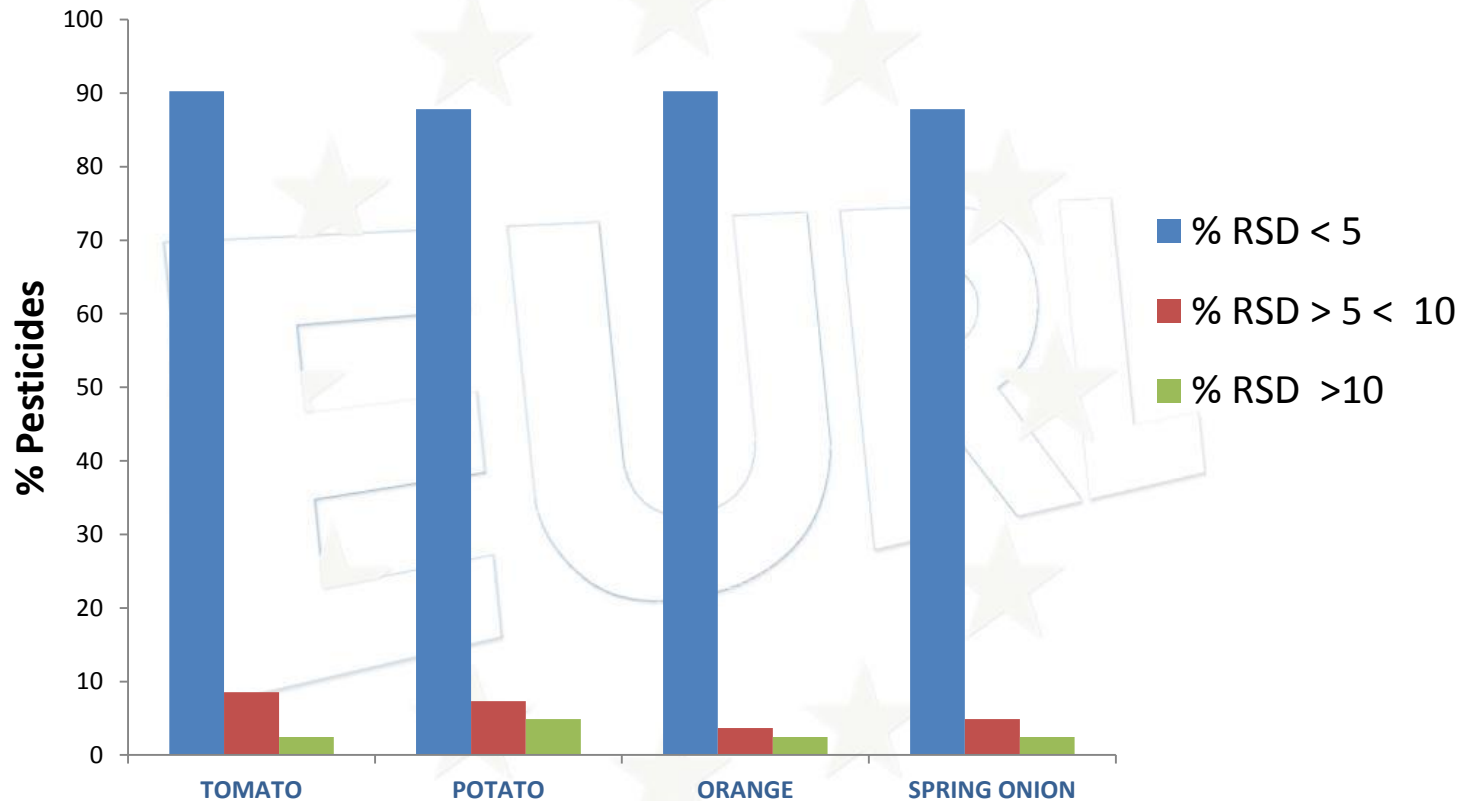
	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
264.0232	12097.47	29015.49	76583.45	134799.16
306.0702	7089.28	15394.74	46812.60	85905.51
248.0280	2802.65	4210.54	13306	24920.55
Relative Abundances to base peak 264.0232				
306.0702	59%	53%	61%	64%
248.0280	23%	15%	17%	18%

TRIFLURALIN



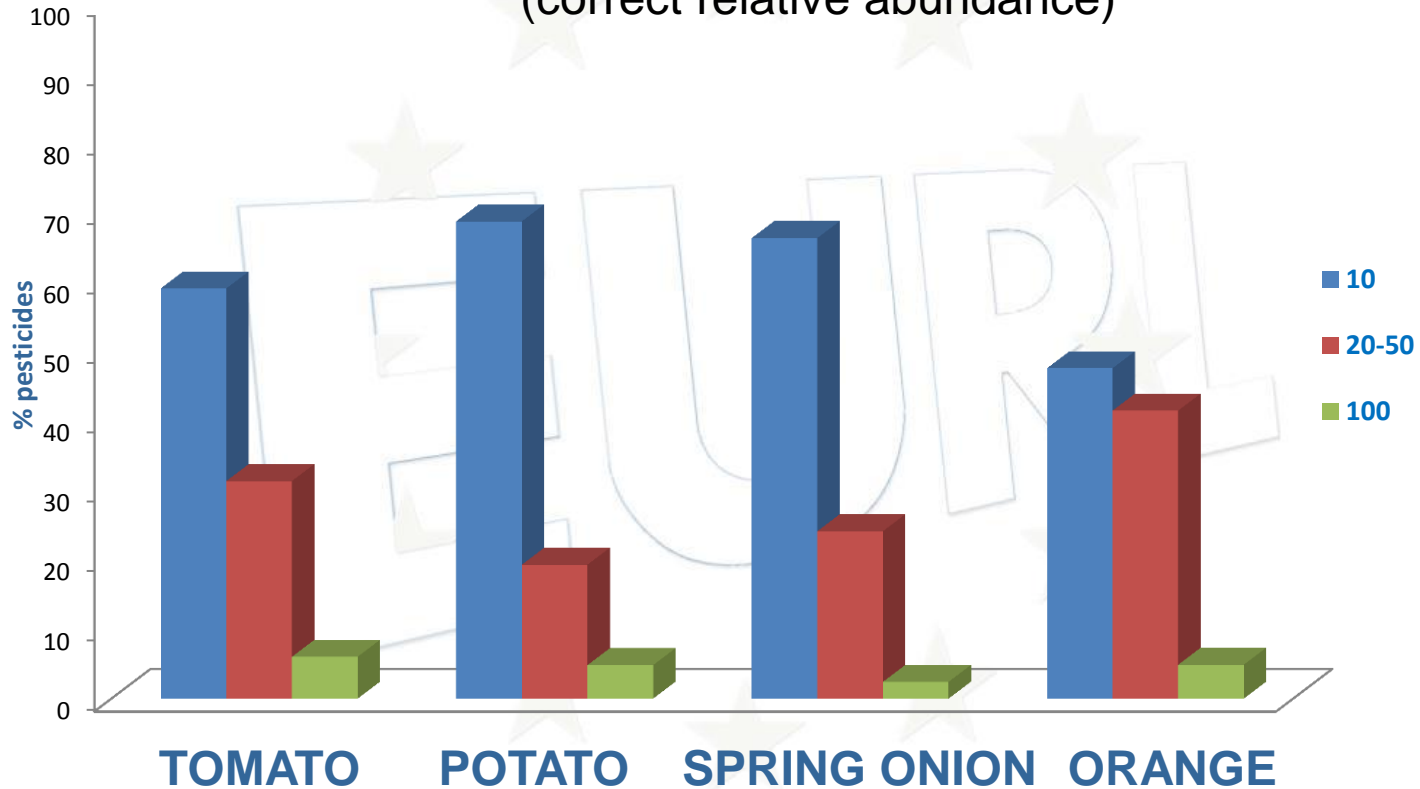
	10µg/Kg	20µg/Kg	50µg/Kg	100µg/Kg
Absolute Abundances				
264.0232	26643.72	49810.53	124512.23	196694.05
306.0702	15638.56	28918.62	87336.68	146209.05
186.0406	5121.20	8316.86	23355.94	48066.36
Relative Abundances to base peak 264.0232				
306.0702	59%	58%	70%	74%
186.0406	19%	17%	19%	24%

ACCURACY (%RSD, n =10)



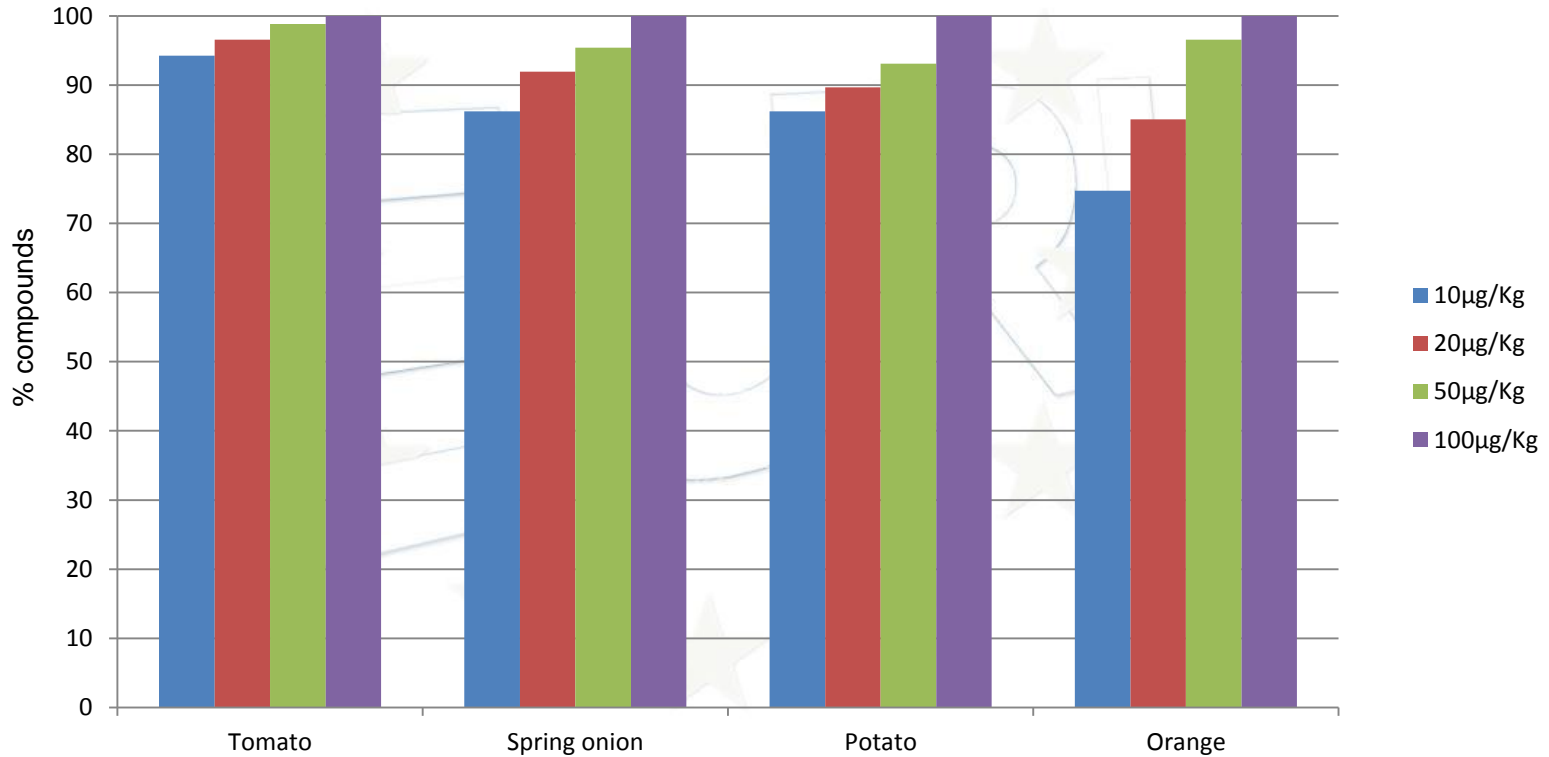
LIMITS OF DETECTION

Retention time and at least **two fragments** with error lower than 5 ppm
(correct relative abundance)



LIMITS OF DETECTION

Retention time and at least **one fragment** with error lower than 5 ppm (other fragments with error higher than 5 ppm but with the correct relative abundance)



REAL SAMPLES

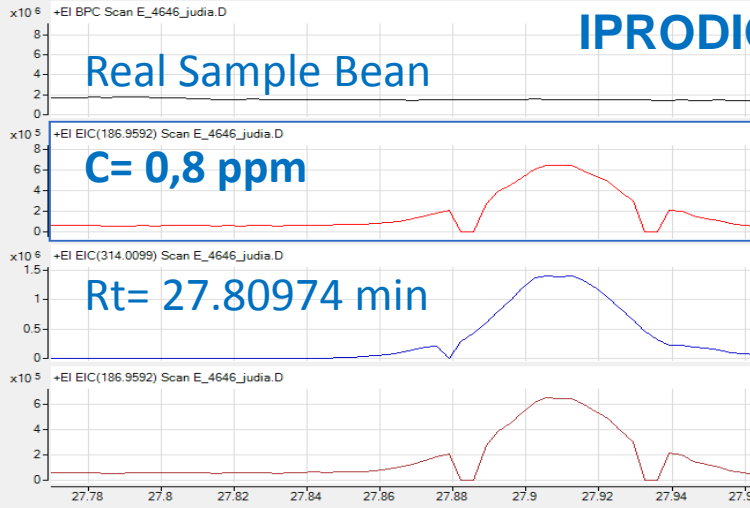
Muestra	GC-TOF automatico	GCQQMS	GC-TOF manual
Pepper	Boscalid (0.25mg/Kg)	Boscalid(0.23mg/Kg)	
Tomato_1	Iprodione (0.20mg/Kg)	Iprodione(0.27mg/Kg)	
	Pyriproxifen (0.054mg/Kg)	Pyriproxifen (0.062mg/Kg)	
Tomato_2	Bifenthrin 1F(0,12mg/Kg)	Bifenthrin (0.13mg/Kg)	Ion 166.099 error higher than10 ppm
Melon_1	Bupirimate (0.15mg/Kg)	Bupirimate(0.13mg/Kg)	
Tomato_3	Buprofezin(0.49mg/Kg)	Buprofezin(0.16mg/Kg)	
	Metalaxyl(0.060mg/Kg)	Metalaxyl(0.068mg/Kg)	
Zucchini	Pirimicarb(0.33mg/Kg)	Pirimicarb(0.24mg/Kg)	
	Bupirimate(0.11mg/Kg)	Bupirimate(0.048mg/Kg)	
	Penconazole(0.091mg/Kg)	Penconazole(0.074mg/Kg)	
Brussels sprouts	Boscalid(0.21mg/Kg)	Boscalid(0.12mg/Kg)	
Tomato_4	Bifenthrin (0.10mg/Kg)	Bifenthrin (0.125mg/Kg)	Ion 166.099 error higher than10 ppm
Bean_1*	Iprodione 1F(0.830mg/Kg)	Iprodione(1.02mg/Kg)	Saturation*
Bean_2	Azoxystrobin ND	Azoxystrobin(0.337mg/Kg)	Coelution
	Iprodione(0.36mg/Kg)	Iprodione(0.25mg/Kg)	
Bean_3	Azoxystrobin ND	Azoxystrobin(0.11mg/Kg)	Coelution
	Iprodione(0.19mg/Kg)	Iprodione(0.078mg/Kg)	
Melon_2	Azoxystrobin ND	Azoxystrobin(0.16mg/Kg)	Coelution
	Bupirimate(0.11mg/Kg)	Bupirimate(0.08mg/Kg)	
Bean_4*	Iprodione ND(1.82mg/Kg)	Iprodione(1.70mg/Kg)	Saturation*
Bean_5*	Iprodione ND(1.27mg/Kg)	Iprodione(1.70mg/Kg)	Saturation*
Grape	Iprodione(0.15mg/Kg)	Iprodione(0.11mg/Kg)	
	Myclobutanil 0.041mg/Kg	Myclobutanil(0.041mg/Kg)	

*These samples were injecting diluted 1:2 to obtain a correct identification and quantification and the mass spectrum show at least 2 fragments ions.

IPRODIONE

Real Sample Bean

NOT DETECTED AUTOMATICALLY



C= 0,8 ppm

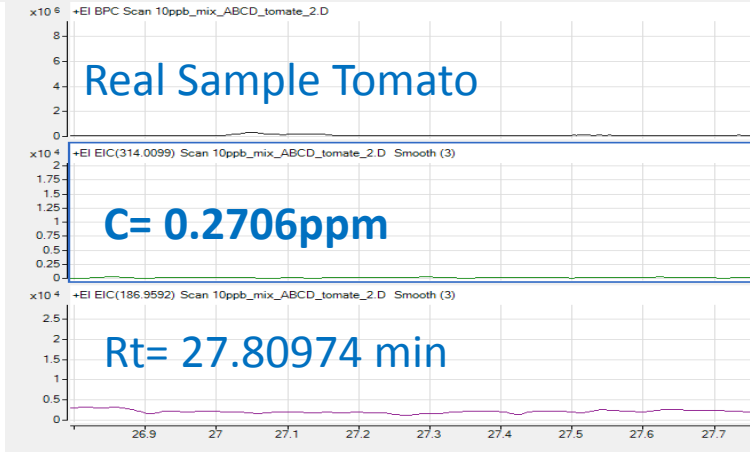
Rt= 27.80974 min

SATURATION

### Formula	Retention Time	Mass	Compound name	Description
# Formula	RT	Mass	Cpd	Comments
C13H13Cl2N3O3	27,81	329,0334	Iprodione	
C12H10Cl2N3O3	27,81	314,0099	Iprodione F1	
C7H3Cl2NO	27,81	186,9592	Iprodione F3	

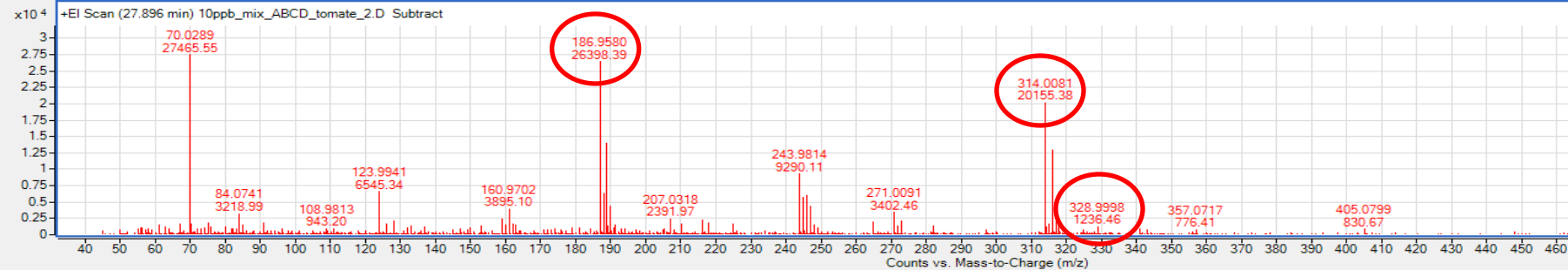
Real Sample Tomato

DETECTED AUTOMATICALLY



C= 0.2706ppm

Rt= 27.80974 min

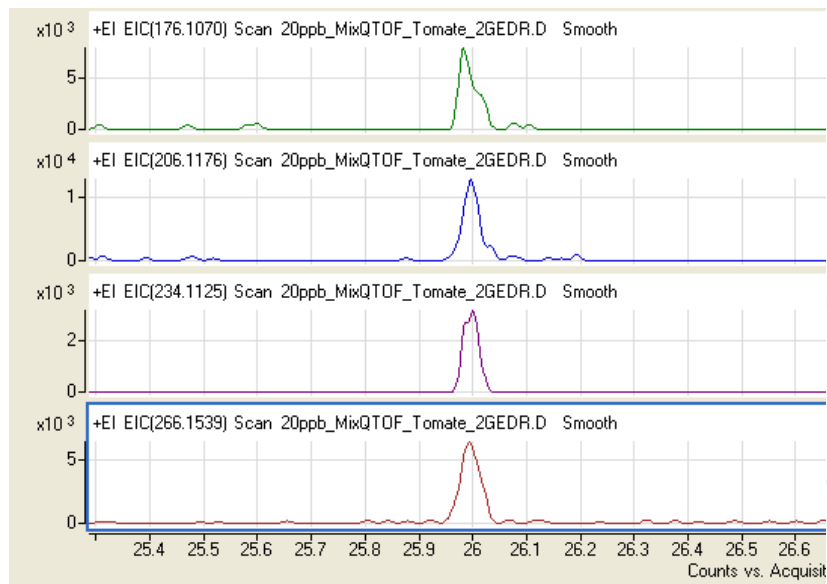


The system has 2 modes of operation :

- **2GHz Extended Dynamic Range (EDR) mode**
- **4Ghz High Resolution (HR) mode**

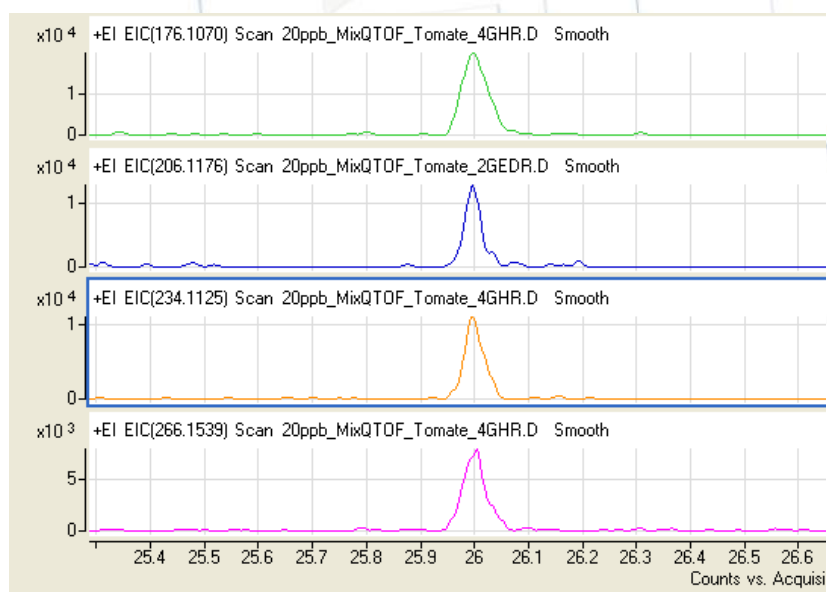
Benalaxyl

2GHz



Fragment	Molecular formula	Experimental mass	Theoretical mass	Error (ppm)
Benalaxyl F1	C18 H20 N O	266,1542	266,1539	0,83

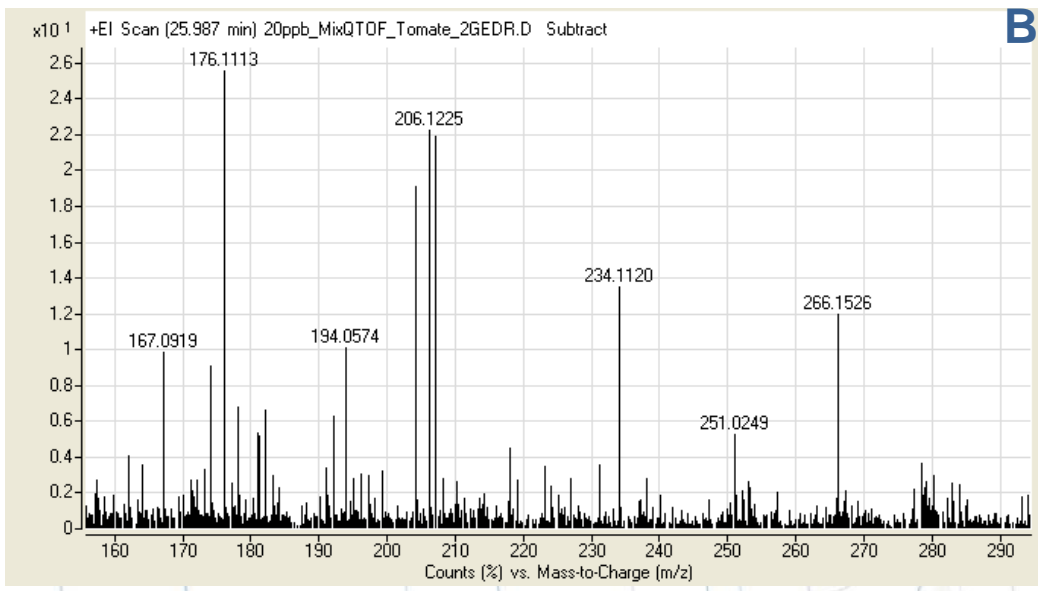
4GHz



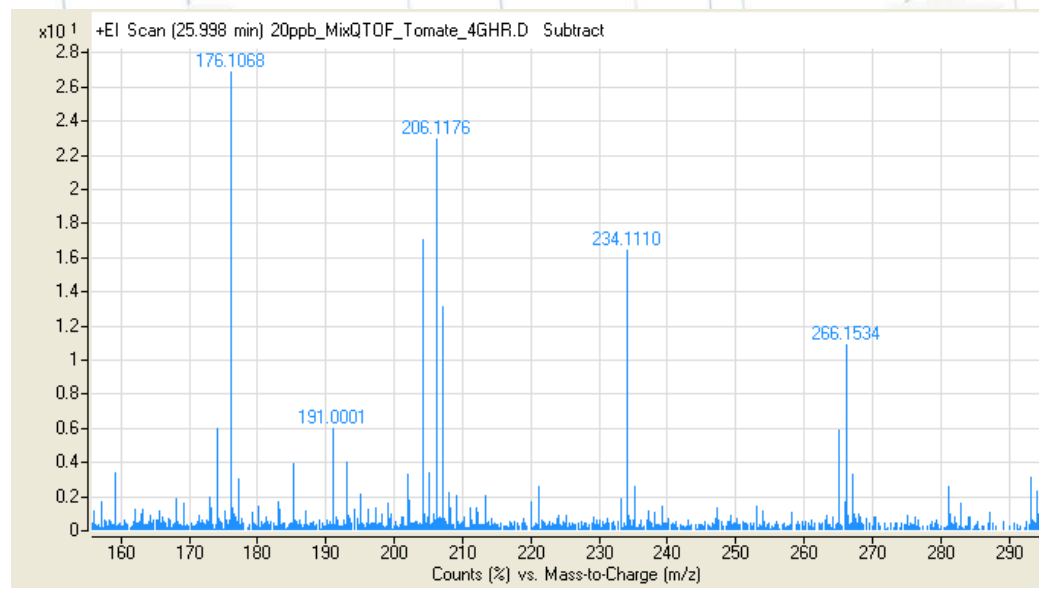
Fragment	Molecular formula	Experimental mass	Theoretical mass	Error (ppm)
Benalaxyl F1	C18 H20 N O	266,152	266,1539	0,83
Benalaxyl F2	C13 H16 N O3	234,112	234,1125	-2,19
Benalaxyl F4	C12 H16 N O2	206,1174	206,1176	-0,81
Benalaxyl F5	C11 H14 N O	176,1073	176,107	1,67

Benalaxyl

2GHz

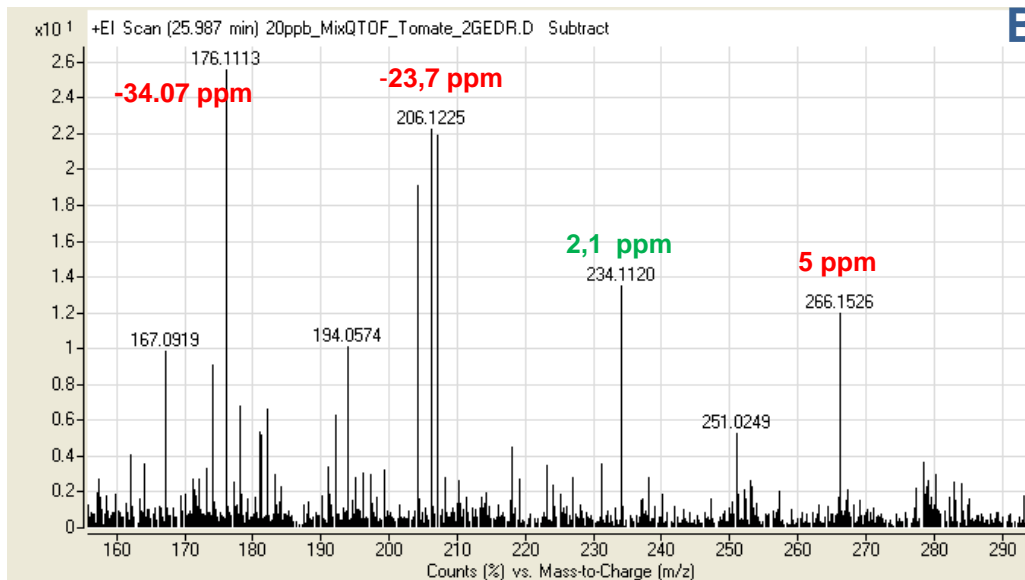


4GHz

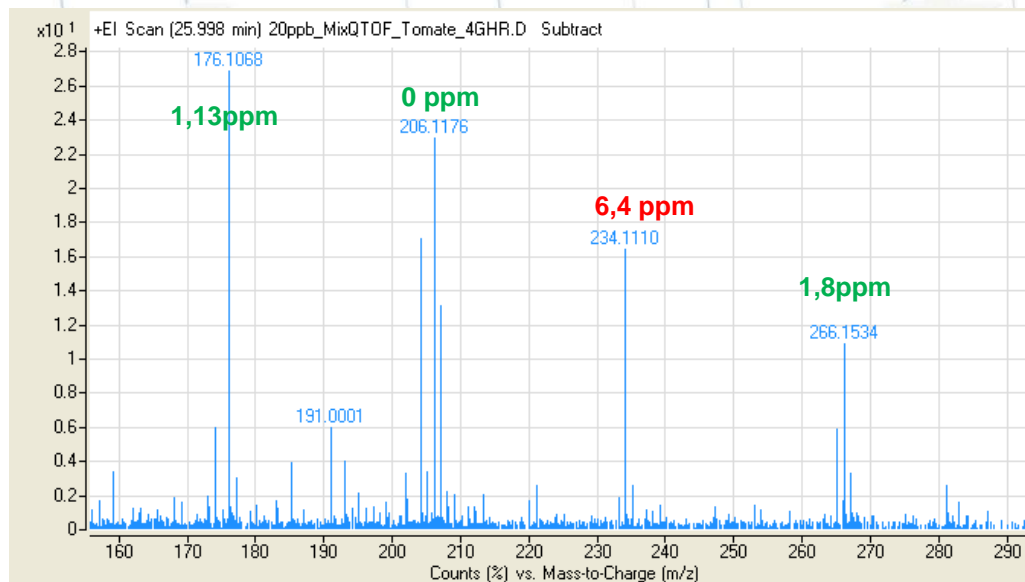


Benalaxyl

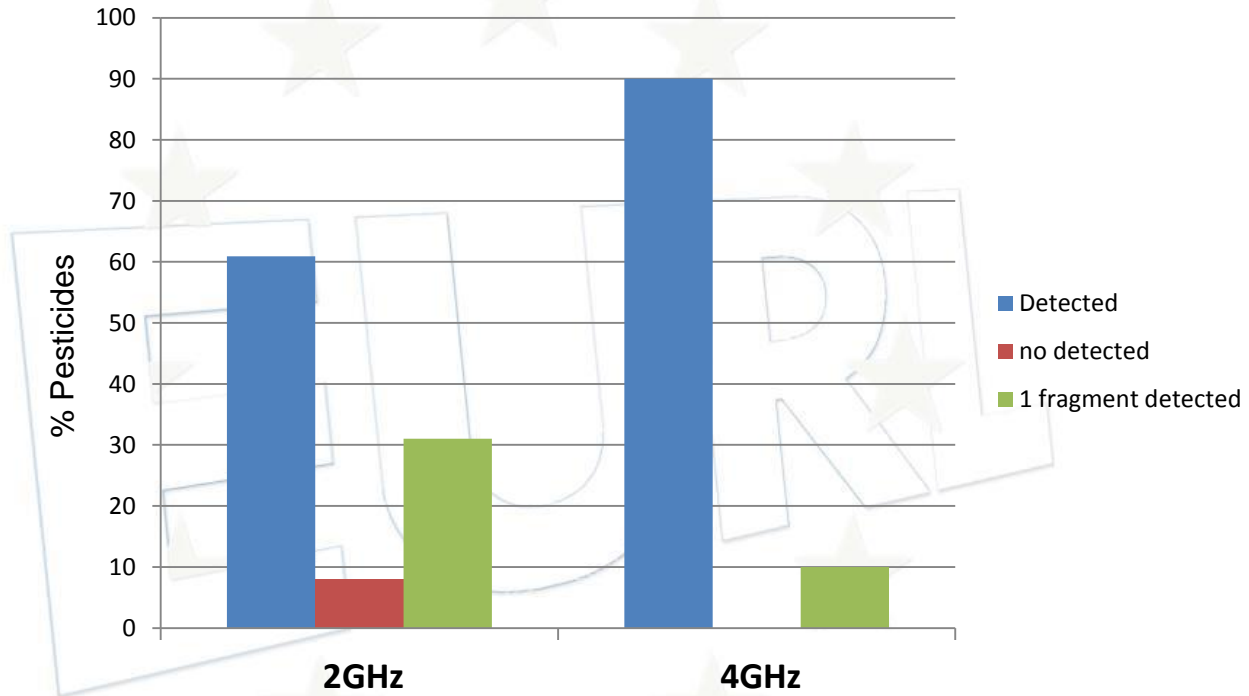
2GHz



4GHz



20 ppb in Tomato Automatic identification



A large, faint background graphic consisting of the letters "EURL-FV" in a light blue outline font, surrounded by a circle of twelve light yellow stars, similar to the European Union flag.

CONCLUSIONS

- 15 real samples were processed with the developed method, results were compared with those obtained by GC-QQQ-MS, from a total of 22 positives 14 were correctly identified automatically, and three pesticides were not identified correctly by GCQTOF because the high concentration saturate the detector, five pesticide were not correctly identified because the some of the ions showed errors higher than 10 ppm.
- This work is a first approach to the accurate mass database for automatic identification and quantification in GC-TOF-MS. It can be considered as a starting point for a large database and for an improved methodology for automatic identification.
- High Resolution (HR) mode can be considered as a alternative to improve the effectiveness of the automatic identification.