

# **Small Molecule Orbitrap Customer Applications**

Ken Matuszak

Sr. CMD Product Specialist



The world leader in serving science



## Mass Spectrometry at Winfield United

Dustyn Sawall
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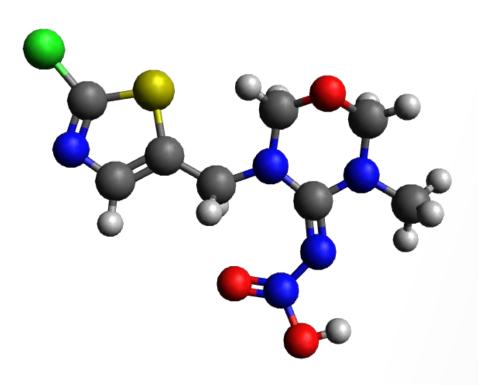
## **Analysis in Conjunction with Other Labs**

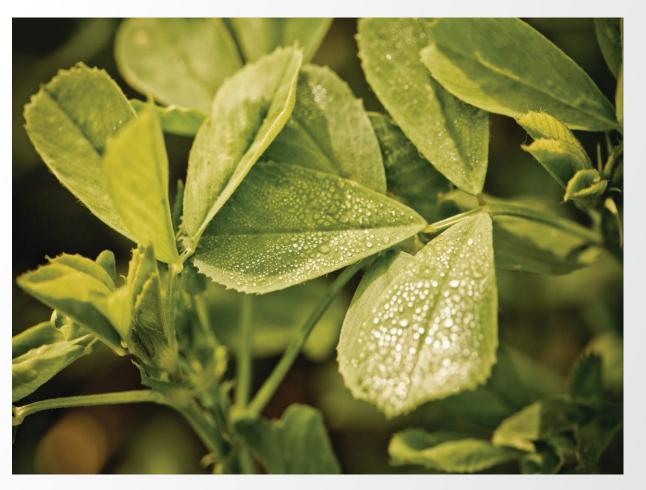
- Real world type applications in conjunction with analysis
  - Spray studies with deposition vs. drift compositions
    - Does the Herbicide go where it is put?
    - What part of it moves
- Green house testing
  - Where does the active go under growth conditions?
  - Time resolved experiments



## **Primary Use for Q-Exactive:**

- Looking for Trace amounts of Herbicides in Complex Matrices
- Time resolved residues on/in leaf surfaces
- Herbicide Degradation studies





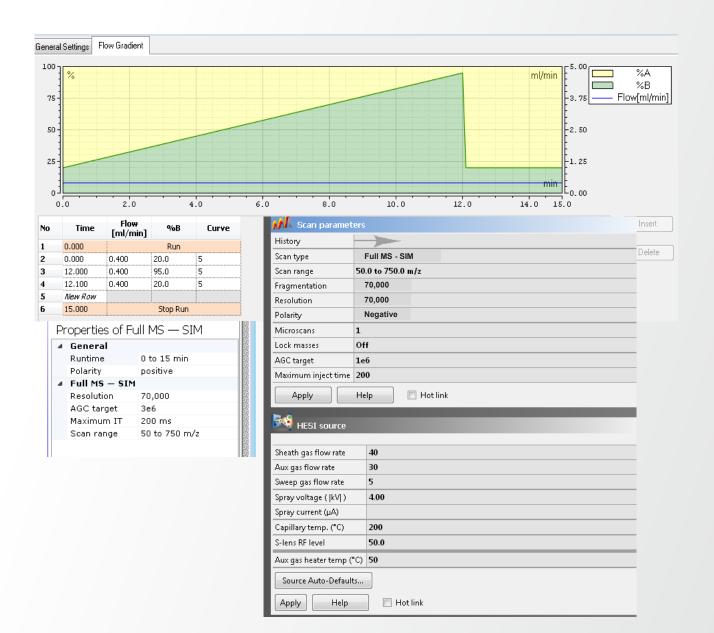


#### **Dicamba Analysis**

- A Major problem in the Agriculture industry currently is the translocation of the herbicide Dicamba beyond the desired application site.
- Experimental set up included standard filter papers attached to discrete particle size classifiers at various distances in a field that was sprayed with an Agricultural Mix including Dicamba.
- This experiment was set up in triplicate with four distinct size bins across three distances.
- The filter papers were extracted with MeOH and run on UHPLC-Orbitrap-HRMS to measure the amount of Dicamba captured at various distances from the spray event.

#### **Analysis Conditions**

- UHPLC
  - C18 Column
    - 50mm x 2.1mm, 3um Particle
    - 40C
  - 15min run
  - Mobile Phase A 0.1% Formic Acid
  - Mobile Phase B Acetonitrile
- HRMS
  - Full MS-SIM
  - (-) mode
  - 70,000 Resolution



#### **Conclusions**

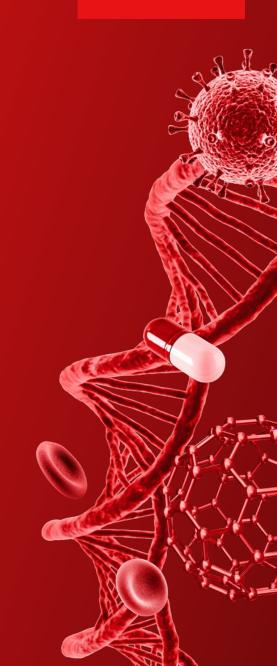


- Ion Extracted Chromatograms were generated from the SIM's.
- Enough data was collected to prove proof of concept, as Dicamba was detected in the nearest collectors.
- Unfortunately, the signals were within the LOD for the prep, they were below LOQ for this method.
- Follow up experiments are underway to concentrate the samples prior to analysis. This should bring the results into an interpretable range.
- Very few studies of this nature have even detected the Dicamba in the environment, so this is a strong step towards quantitation.



# Treated Seed Pesticide Analysis by UHPLC-HRMS

Lianna Bestvater
Canadian Grain Commission
Grain Research Laboratory – Grain Safety group
Trace Organics and Trace Elements





## **Treated Seed Analysis by UHPLC-HRMS**

- Treated seed is grain that has been treated with a pesticide product (typically fungicides) prior to planting
- In Canada, treated seed must be dyed to make seeds distinguishable from other grain
- These seeds may be inadvertently present in bulk grain if grain handling equipment is not cleaned well, and may lead to points of pesticide residues in bulk grain
- The Grain Research Laboratory routinely monitors grain using UHPLC-HRMS, and has the capacity to analyze samples for the presence of the 42 seed treatment pesticides that are active ingredients in products registered for use in Canada on grains



#### **Extraction Procedure**





Add Acetonitrile

Vortex, sonicate

Sample Dilutions

Add Internal Standard Syringe Filter

LC-HRMS







#### **UHPLC Conditions**

Column: Waters Acquity UPLC BEH C18, 1.7µm, 2.1x 100mm

Column flow: 0.4 mL/min

Column temperature: 40°C

Injection volume: 10 µL

Autosampler tray temperature: 10°C

•	Gradient:	Time (min)	%A	%B
		Initial	90	10
		0.25	90	10
		7.75	0	100
		8.50	0	100
		8.51	90	10
		12.00	90	10

#### **HRMS Parameters**

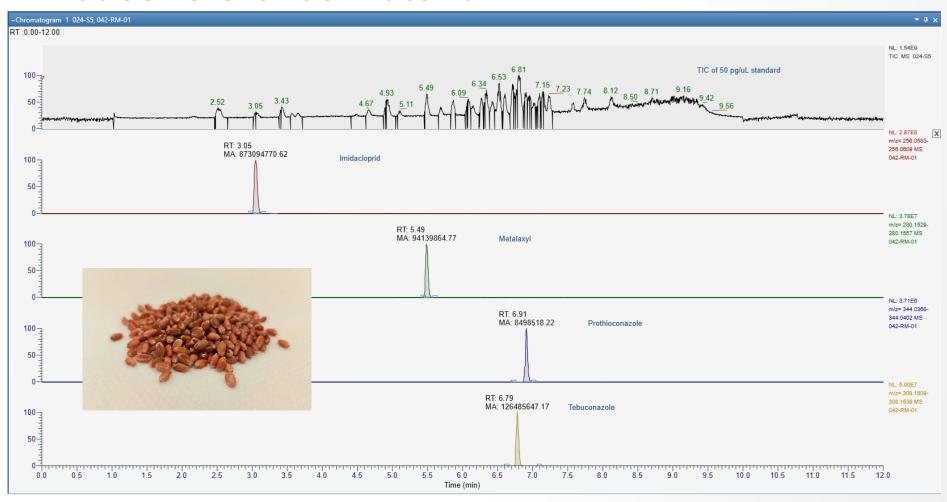
HESI source parameter	S
Sheath gas flow rate	50
Auxiliary gas flow rate	13
Capillary temperature	300°C
Sweep gas flow rate	3
Spray voltage	3.5 kV
S-lens RF level	55
Aux gas heater temperature	425°C

Experiment type	Full scan/ddMS <sup>2</sup>
Ionization mode	ESI +
Scan Range	70 to 680 m/z
Full MS	
Resolution	70,000
AGC target	1e6
Maximum IT	100 ms
ddMS2	
Resolution	17,500
AGC target	1e5
Maximum IT	50 ms
NCE stepped	10, 20, 40

#### **42 Pesticides**

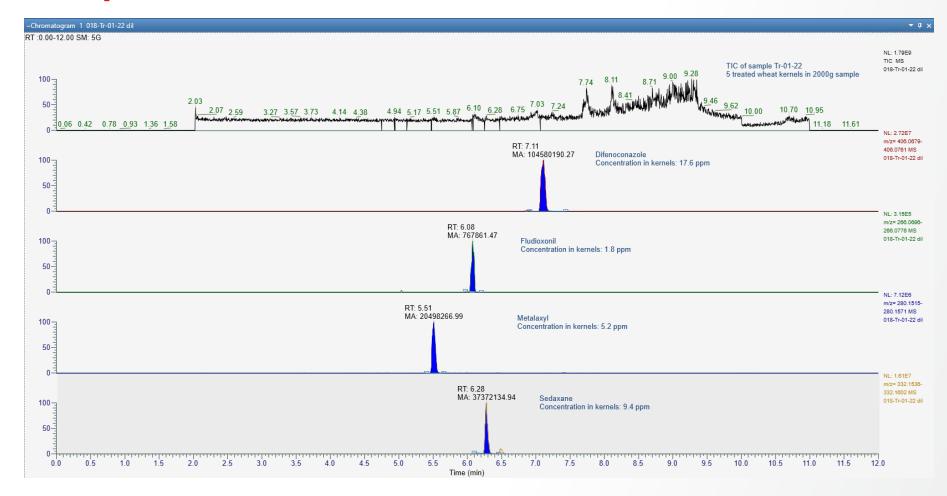
Pesticide	m/z	RT (min)	Pesticide	m/z	RT (min)
Acetamiprid	223.0745	3.40	Metalaxyl	280.1543	5.50
Azoxystrobin	404.1241	5.90	Metconazole	320.1524	7.00
Boscalid	343.0399	6.13	Oxathiapiprolin	540.1487	6.08
Broflanilide	663.0136	7.08	Penflufen	318.1976	6.70
Captan	321.9234	5.74	Penthiopyrad	360.1352	6.76
Carboxin	236.0740	5.00	Picarbutrazox	410.1935	6.61
Chlorantraniliprole	481.9781	5.70	Picoxystrobin	368.1104	6.60
Clothianidin	250.0160	3.10	Prothioconazole	344.0386	6.90
Cyantraniliprole	473.0123	5.15	Pydiflumetofen	426.0349	7.08
Diazinon	305.1083	6.81	Pyraclostrobin	388.1059	6.87
Difenoconazole	406.0720	7.20	Sedaxane	332.1569	6.30
Ethaboxam	321.0838	5.05	Sulfoxaflor	278.0569	3.50
Fludioxonil	266.0736	6.07	Tebuconazole	308.1524	4.79
Fluopyram	397.0537	6.32	Thiabendazole	202.0433	2.54
Fluoxastrobin	459.0866	6.40	Thiamethoxam	292.0266	2.60
Flupyradifurone	289.0550	3.44	Thiophanate methyl	343.0529	4.66
Fluxapyroxad	382.0973	6.19	Thiram	240.9956	4.50
Imidacloprid	256.0596	3.10	Tioxazafen	229.0430	7.00
Inpyrfluxam	334.1726	6.52	Triadimenol	294.1004	6.33
Ipconazole	334.1681	7.30	Trifloxystrobin	409.1370	7.20
Iprodione	330.0407	6.60	Triticonazole	318.1368	6.50

#### In-house reference material



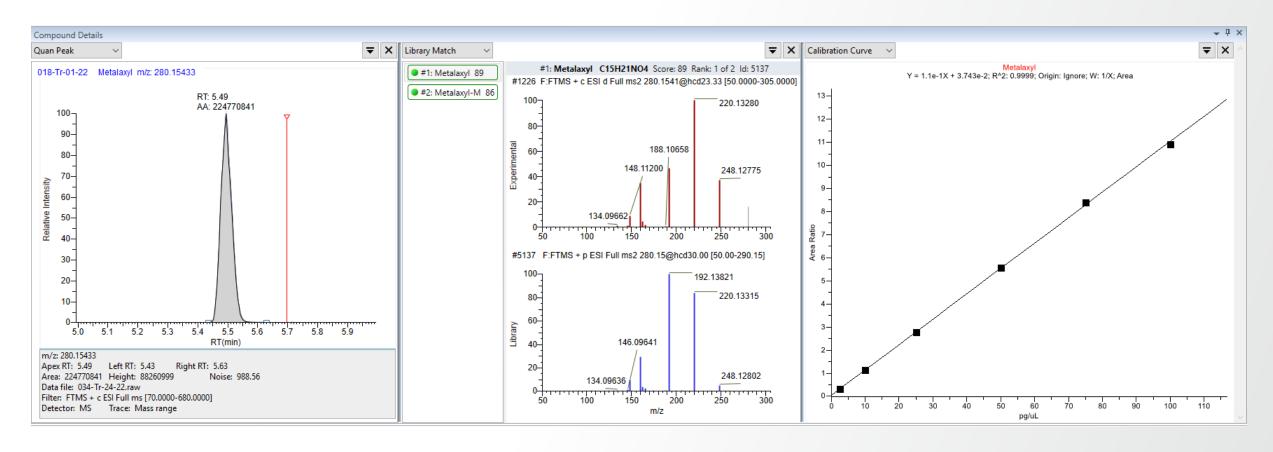
Raxil-Pro Shield seed treatment contains 4 pesticides: imidacloprid, metalaxyl, prothioconazole, tebuconazole

#### **Sample Tr-01-22**



5 treated wheat kernels in 2000g sample

#### Metalaxyl



Quantitation peak

Library match

Calibration curve

#### **Analytical Report**



Canadian Grain Commission canadienne des grains



**Grain Research Laboratory** 

1404-303 Main Street Winnipeg, Manitoba Canada R3C 3G8

2022-01-21

#### SUSPECTED TREATED SEED ANALYTICAL REPORT

Trace Organics Analysis Sample Number: Tr-01-22 Date submitted to Trace Organics Analysis:

Sample Information:

Submitted by: CGC IS file number:

Sample identification:

Bin:

Tonnes:

Type of Grain:

Wheat Sample size:

5 seed(s) from 2000 g sample. Seed(s) had a pink discoloration. Suspect condition:

Tests performed:

Trace Organic Analysis method GS-59

Results:

The overall concentration of these chemicals in the sample does not exceed Canadian maximum residue limits. Consequently, the lot is not considered to be contaminated

based on the sample provided.

More details provided on the following page.

Recommended action: No action required on food safety grounds.

Sheryl Tittlemier

Program Manager, Trace Organics and Trace Elements





Canadian Grain Commission canadienne des grains



#### Results:

Pesticide	Concentration in Stained Seed(s) (ppm)	Concentration in Total Sample, 2000 g (ppm)	Canadian Maximum Residue Limit (ppm)*	
Acetamiprid	-	-	0.1	
Azoxystobin	-	-	0.03	
Boscalid	-	-	0.2	
Broflanilide	-	-	0.01	
Captan	-	-	0.1	
Carboxin	-	-	0.2	
Chlorantraniliprole	-	-	6	
Clothianidin	-	-	0.01	
Cyantraniliprole	-	-	0.1	
Diazinon	-	-	0.1	
Difenoconazole	17.6	0.002	0.1	
Ethaboxam	-	-	0.01	
Fludioxonil	1.8	0.0002	0.02	
Fluopyram	-	-	1.5	
Fluoxastrobin	-	-	0.1	
Flupyradifurone	-	-	3	
Fluxapyroxad	-	-	0.3	
Imidacloprid	-	-	0.05	
Inpyrfluxam	-		0.01	
Ipconazole	-	-	0.01	
Iprodione	-	-	0.07	
Metalaxyl	5.2	0.0005	0.2	
Metconazole	-	-	0.15	
Oxathiapiprolin			0.1	
Penflufen	-	-	0.01	
Penthiopyrad	-	-	0.15	
Picoxystrobin	-	-	0.04	
Prothioconazole	-	-	0.35	
Pydiflumetofen	-		0.1	
Pyraclostrobin	-		0.2	
Sedaxane	9.4	0.00008	0.01	
Sulfoxaflor	-	-	0.08	
Tebuconazole	-	-	0.15	
Thiabendazole	-	-	0.1	
Thiamethoxam	-	-	0.02	
Thiophanate-methyl	-	-	0.1	
Thiram	-	-	0.1	
Tioxazafen	-	-	0.1	
Triadimenol	-		0.05	
Trifloxystrobin	-	-	0.05	
Triticonazole	+		0.01	

<sup>\*</sup> http://www.hc-sc.gc.ca/cps-spc/pest/part/protect-proteger/food-nourriture/mrl-lmr-eng.php

<sup>-</sup> Analyte was not detected above limit of quantitation, LOQ (LOQ: the lowest amount of analyte in a sample which can be quantitatively determined with suitable precision and accuracy

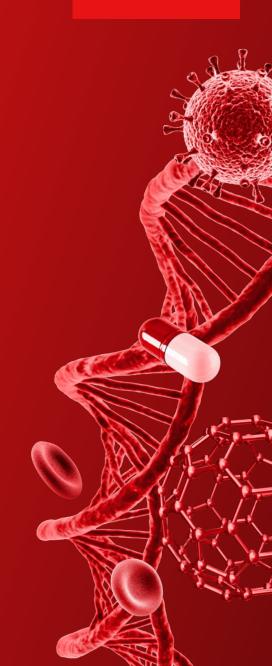






# Comprehensive Drug Screening of Whole Blood by LC-HR-MS/MS in a Forensic Laboratory

Jon Benjamin Stephenson
Toxicology Technical Leader
Georgia Bureau of Investigation Division of Forensic
Sciences

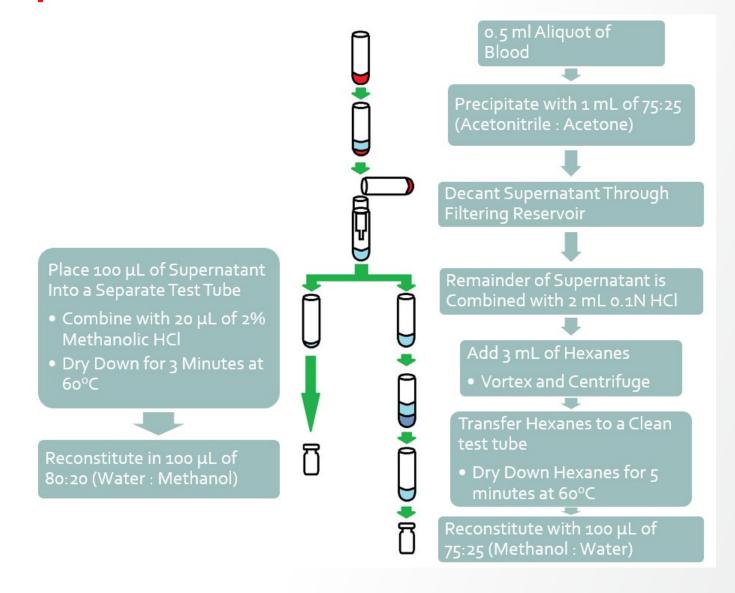


#### **Grant Funding**

This project was supported by Award No. 2016-DN-BX-K005, awarded by the National Institute
of Justice, Office of Justice Programs, U.S. Department of Justice.

The opinions, findings, and conclusions or recommendations expressed in this program are those of the author and do not necessarily reflect those of the Department of Justice.

#### **Sample Preparation**



#### **Instrument Method: LC Setup**

- ~15 millimolar Ammonium Formate
- 0.1% Formic Acid
- Optima Grade Water

#### Mobile Phase B

- ~15 millimolar Ammonium Formate
- o.1% Formic Acid
- 50 Acetonitrile : 50 Methanol

#### Syringe Wash

• 50 Methanol : 50 Water

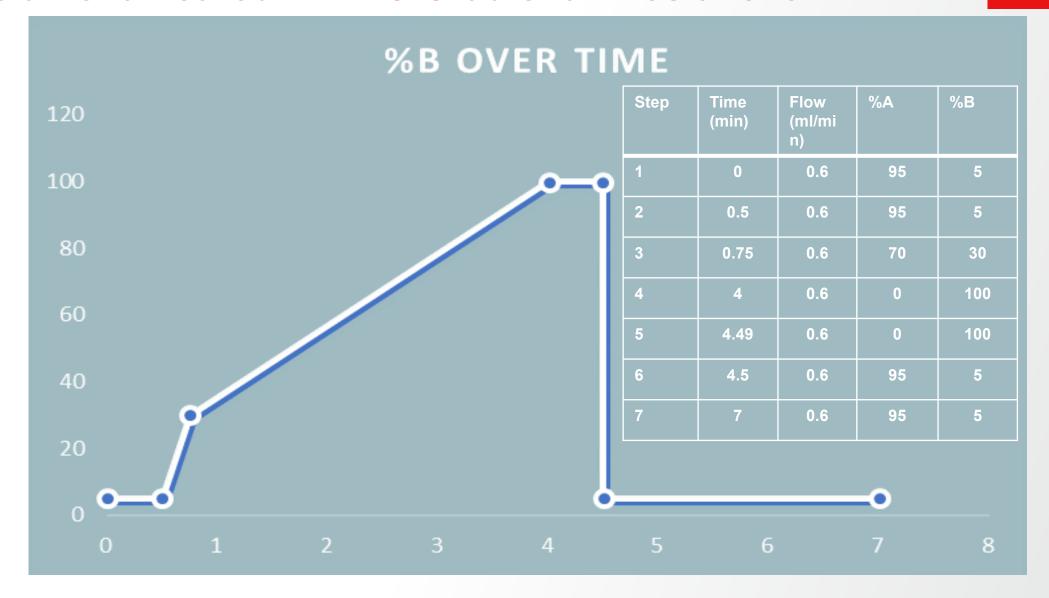
#### Rear Seal Wash

- 75 % Isopropyl Alcohol
- 25 % Water
- 0.1 % Formic Acid

- 2.1X100 mm
- 2.7 micron packing material



#### Instrument Method: HPLC Gradient - Positive Ion



#### **Instrument Method: Orbitrap Positive Mode**

- Inclusion list of compounds
  - Accurate mass and name of each compound
  - 0.4 min retention time windows
  - Polarity
  - Exclude
    - Formula
    - Species
    - Charge State
- MS1
  - 35,000 Resolution
  - Scan Range 50 to 750 m/z
- MS2
- 17,500 Resolution
- Stepped Collision Energy
  - 10, 25, 50
- Minimum AGC target
  - Positive Mode 3e3
- Dynamic Exclusions 6 s



#### Instrument Method: HPLC Gradient – Pos/Neg Ion





#### **Instrument Method: Orbitrap Pos/Neg Mode**

- Inclusion list of compounds
  - Accurate mass and name of each compound
  - 0.4 min retention time windows
  - Polarity
  - Exclude
    - Formula
    - Species
    - Charge State
- MS1
  - 35,000 Resolution
  - Scan Range 50 to 750 m/z
- MS2
- 17,500 Resolution
- Collision Energy
  - NCE 10 for Barbituates
  - NCE 30 for THC and Related compounds
- Minimum AGC target
  - Positive Mode 1e3
- Dynamic Exclusions (Auto)

#### **Mass Calibration and QC Checks**

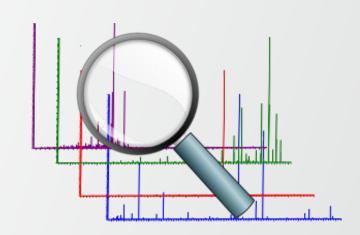
- Performed weekly
  - Standard Mass Calibration for Positive and Negative modes
  - Custom Calibration done for Positive mode to include lower mass ion 74.09643
- Daily Test mixture
  - Checks
    - Chromatography
    - Mass assignments
    - Sensitivity
  - Masses across the scan range within 5ppm
    - Amphetamine 136.1121
    - Diphenhydramine 256.1696
    - Fentanyl 337.2274
    - Buprenorphine 468.3108
    - Butalbital 223.1088
    - Butabarbital 211.1088

#### **Data Processing**

- Generally desire more drugs to pass through processing filters than accidentally exclude something of importance.
- Mass Spectrum Matching
  - Threshold 50
  - Passing value 50
- Isotopic pattern
  - 50% fit threshold (positive mode)
  - 60 ppm mass deviation allowed
  - 100% intensity deviation
  - No requirement in negative mode
- Mass Accuracy
  - 5-10 ppm accuracy
- Retention time
  - 30 second window
- Reviewing scientist evaluates the data and determines what analytes are and are not present based upon the software's pared back results

#### **Validation: LOIs**

- Generally the Positive Mode LOI for most drugs is at least 2 μg/L
  - Exceptions
    - MDA, Meprobamate, Phentermine, Zolpidem, and Zopiclone
      - 20 μg/L
    - Phenylephrine, Methylone, and Levetiracetam
      - 10 μg/L
    - Duloxetine, Carisoprodol, and Bromazepam
      - 5 μg/L
    - Fentalogs
      - Furanylfentanyl, carfentanil, and parafluorofentanyl down to 1 μg/L



#### **Validation: Matrix Effects**

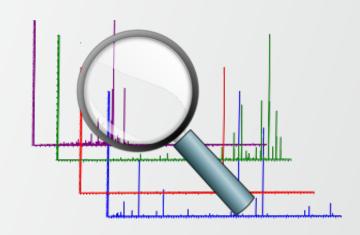
- Matrix effects were determined by fortifying three separate lots of negative blood post extraction and comparing the average area to the area of an un-extracted non-matrix containing sample.
- Ion suppression or enhancement was universally below 20% with
  - Some analytes such as morphine, nalorphine, perphenazine, norbuprenorphine, mitragynine, nifedipine, acetaminophen, and ranitidine showing ion enhancement from 500% up to more than 1000%
  - Due to the method being a qualitative screen this has no real negative effect on the results as all the reported LOIs are within an acceptable range of detection.



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#### **Validation: % Recovery**

- Percent recovery was determined by averaging the area of three separate lots of negative blood that was fortified, extracted, and compared to the area of an un-extracted non-matrix containing sample.
- Percent recovery was typically around 100% or greater (mostly due to ion enhancement)
  - Some analytes showed recovery levels below 50% such as meclizine, flubromazolam,
     lurasidone, ziprasidone, and sertraline
  - Even with some lowered % recovery the LOIs for the analytes remained in an acceptable range.



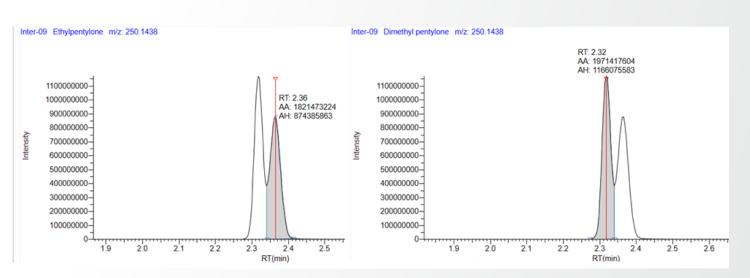
#### Validation: Interferences



- Using resolution equation the resolution between all known isomers was determined
- R= 2 x [(tB tA) / (WB+WA)]
- Resolution (R) equation, tB is the retention time of the later eluting analyte, tA is the retention time of the earlier eluting analyte, WB is the peak width of the later eluting analyte and WA is the peak width of the earlier eluting analyte.
- Resolution of 1.5 indicates two peaks are fully resolved.
- Due to this method being qualititaive only full resolution is not required for identification.
- Example: Ethylpentylone and dimethylpentylone have a resolution of 0.43 and identification is still possible

• Amitriptyline and maprotiline with a resolution of 0.35 are not sufficiently resolved to identify each compound in a single

extract



#### **Validation: Interferences**

Analyte	Mass	RT	Peak Width (min)	Analytes Used to Calculate Resolution	Resolution
Methamphetamine	150.1277	1.99	0.055	methmphetmine/phentermine	1.8
Phentermine	150.1277	2.08	0.045		
bk-MDDMA	222.1125	1.9	0.09	bk-MDDMA/ethylone	0.666666667
Butylone	222.1125	2.03	0.07	bk-MDDMA/butylone	1.625
Ethylone	222.1125	1.96	0.09	butylone/ethylone	0.875
Metaxalone	222.1125	3.22	0.075	metaxalone/butylone	16.4137931
Methylphenidate	234.1489	2.29	0.07	methylphenidate/normeperidine	2
Normeperidine	234.1489	2.43	0.07		
bk-DMBDB (Dibutylone)	236.1281	2.06	0.08	bk-DMDB/eutylone	0.75
Eutylone	236.1281	2.12	0.08	pentylone/eutylone	2
Pentylone	236.1281	2.27	0.07		
Diphenhydramine	256.1696	2.83	0.085	diphenhydramine/atomoxetine	2.875
Atomoxetine	256.1696	3.06	0.075		
Propranolol	260.1645	2.75	0.085	propranolol/ramelteon	7.5
Ramelteon	260.1645	3.35	0.075		
EMDP	264.1747	3.58	0.085	EMDP/notriptyline	2.818181818
Nortriptyline	264.1747	3.27	0.135		
Norvenlafaxine	264.1958	2.16	0.07	norvenlafaxine/tramadol	1.62962963
Tramadol	264.1958	2.27	0.065		
EDDP	278.1903	2.93	0.095	EDDP/maprotiline	3.428571429
Amitriptyline	278.1903	3.26	0.09	maprotiline/amitriptyline	0.352941176
Maprotiline	278.1903	3.23	0.08	, , , , , , , , , , , , , , , , , , , ,	
Imipramine	281.2012	3.19	0.085	imipramine/4-ANPP	5.529411765
4-ANPP	281.2012	2.72	0.085		
Hydromorphone	286.1438	1.69	0.095	morphine/hydromorphone	1.828571429
morphine	286.1438	1.53	0.08	hydromorphone/norcodeine	1.485714286
Norcodeine	286.1438	1.82	0.08	.,,,	
Hydrocodone	300.1594	1.92	0.09	hydrocodone/codeine	1.142857143
Codeine	300.1594	1.82	0.085		
Temazepam	301.0738	3.48	0.07	temazepam/clobazam	0.533333333
clobazam	301.0738	3.44	0.08		
Noroxycodone	302.1387	1.88	0.07	noroxycodone/oxymorphone	3.294117647
Oxymorphone	302.1387	1.6	0.1		
Scopolamine	304.1543	1.91	0.065	scopolamine/cocaine	9
Cocaine	304.1543	2.54	0.075		
Norpropoxyphene*	308.2009	3.11	0.065	norpropoxyphene/benztropine	1.866666667
Benztropine	308.2009	3.25	0.085		2.55555557
AH-7921	329.1182	2.91	0.085	AH-7921/U-47700	1.222222222
U-47700	329.1182	2.8	0.095		
Crotonyl fentanyl	349.2274	2.83	0.000	crotonyl fentanyl/cyclopropylfentanyl	unresolved
Cyclopropylfentanyl	349.2274	2.83			
FIBF	369.2337	2.97		FIBF/parafluorobuytyrlfentanyl	unresolved
Parafluorobuytyrlfentanyl	369.2337	3	<del>                                     </del>	· · · · · · · · · · · · · · · · · · ·	
Dimethylpentylone	250.1438	2.3	0.095	dimethylpentylone/ethylpentylone	0.432432432
ethylpentylone	250.1438	2.34	0.09		
Oxcarbazepine**	253.0972	2.8	0.095	oxcarbazepine/phenytoin	2.171428571
Phenytoin**	253.0972	2.99	0.08	-near seaching buerthough	2.272.20371
Oxcarbazepine ***	253.0972	4.98	0.21	oxcarbazepine/phenytoin	1.58974359
Phenytoin***	253.0972	5.29	0.18		

<sup>\*</sup> Mass after in-source loss of water.

<sup>\*\*\*</sup> positive/negative switching method



# Orbitrap-based Veterinary Toxicology Screening

David J. Borts, PhD

**Clinical Professor** 

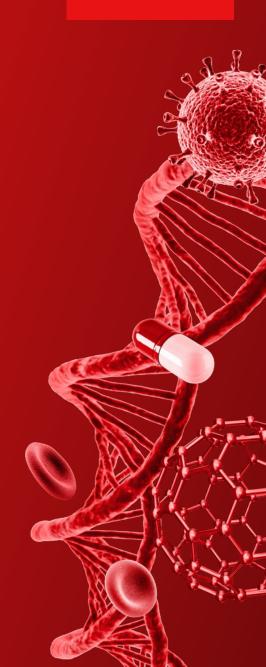
**Analytical Chemistry Services** 

**Veterinary Diagnostic Laboratory** 

Department of Veterinary Diagnostic and Production Animal Medicine

**College of Veterinary Medicine** 

**Iowa State University** 



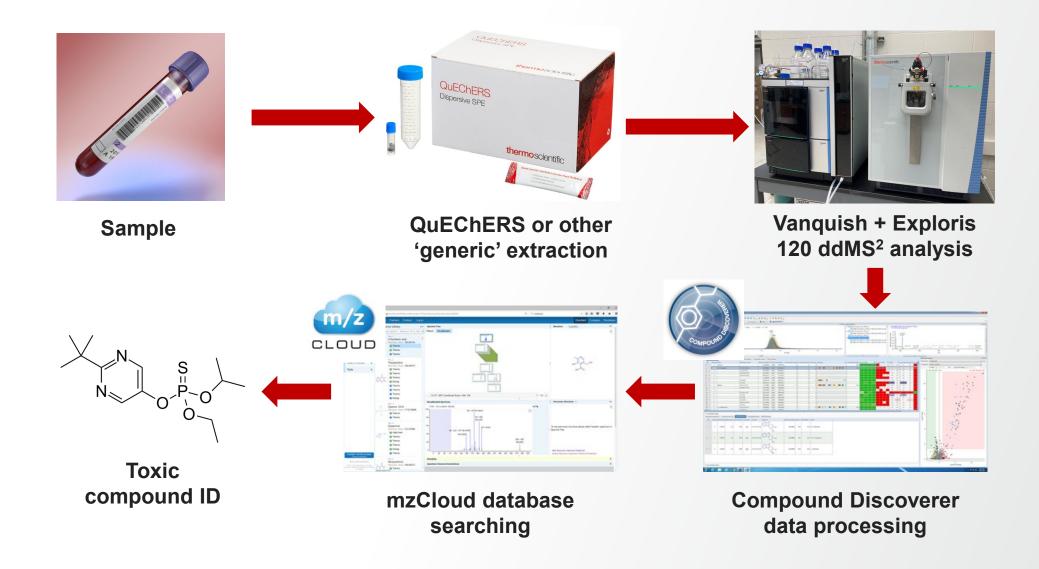
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#### **Orbitrap-based Veterinary Toxicology Screening**

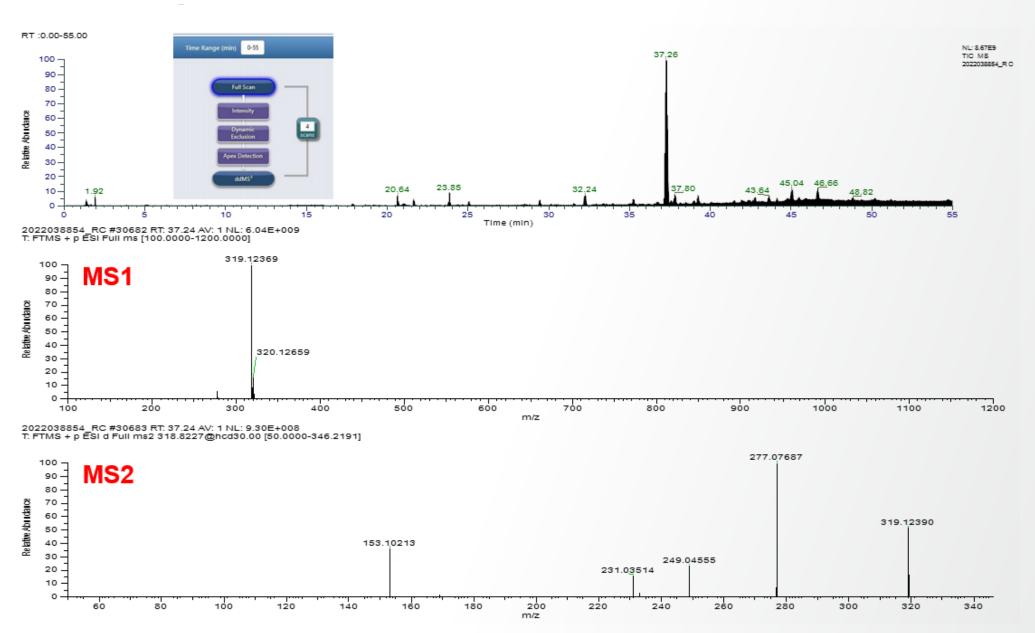
- Case history
  - Midwest US cow-calf farm
  - Calves sudden ataxia, tremors, and death
  - Cows same observations ~ 96 hours later
  - Cow rumen (stomach) contents submitted initially
  - Cow milk and calf rumen contents subsequently analyzed
  - Orbitrap-based LC/MS general unknown toxicology screening performed



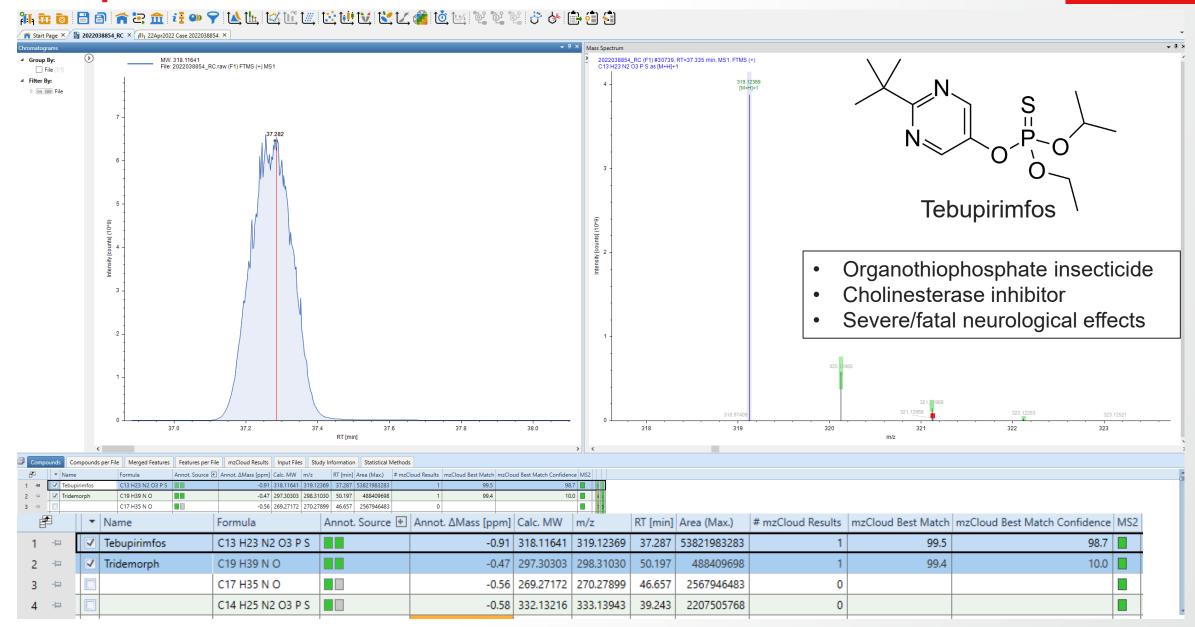
## General Unknown Tox Screening – Orbitrap-based Workflow



#### Rumen contents extract

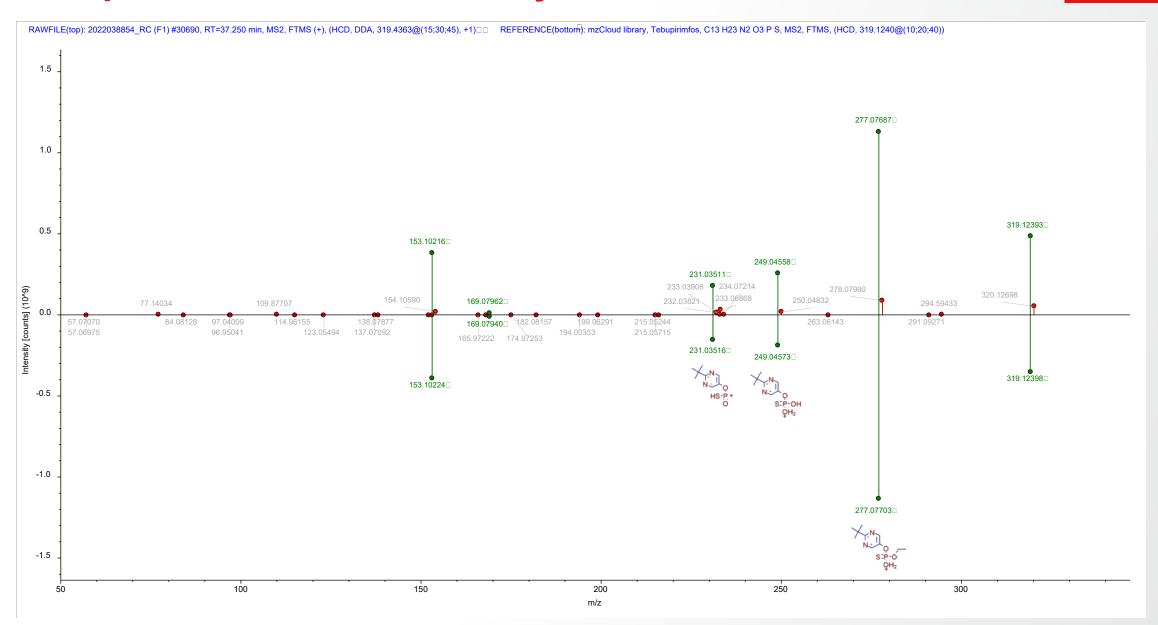


#### **Compound Discoverer**





## **Tebupirimfos mzCloud Library Match Score = 99.5**



### **Orbitrap-based Veterinary Toxicology Screening**

- Case conclusions
  - Tebupirimfos identified in cow and calf rumen and cow milk
  - On-farm investigation revealed farmhand had inadvertently mixed a tebupirimfos-containing insecticide into cow feed (had similar appearance to a commonly used mineral supplement)
  - Only cows had access to contaminated feed
  - Nursing calves received toxic dose of tebupirimfos through cow milk
  - Feeding equipment cleaned and contaminated feed replaced
  - No further symptoms or fatalities were observed in herd

#### Thermo Fisher SCIENTIFIC

#### **Acknowledgements**

- Iowa State University Veterinary Diagnostic Laboratory
  - Analytical Chemistry Services Section



David Borts,
PhD
Clinical
Professor



Dwayne Schrunk, MS Lab Manager



Scott Radke, DVM, MS,
DABVT
Clinical Assistant Professor



Laura Burns, MS Lead LC/MS Chemist

#### Summary



- Orbitraps are very cool and powerful devices
- Orbitraps are very versatile and can be used in qualitative or quantitative environments across a wide range of industries.



## Thank you!