

“Incredibly Powerful, Remarkably Efficient Quantitation”

Patrick M. Jeanville, Ph.D.
LC/TQ Product Manager

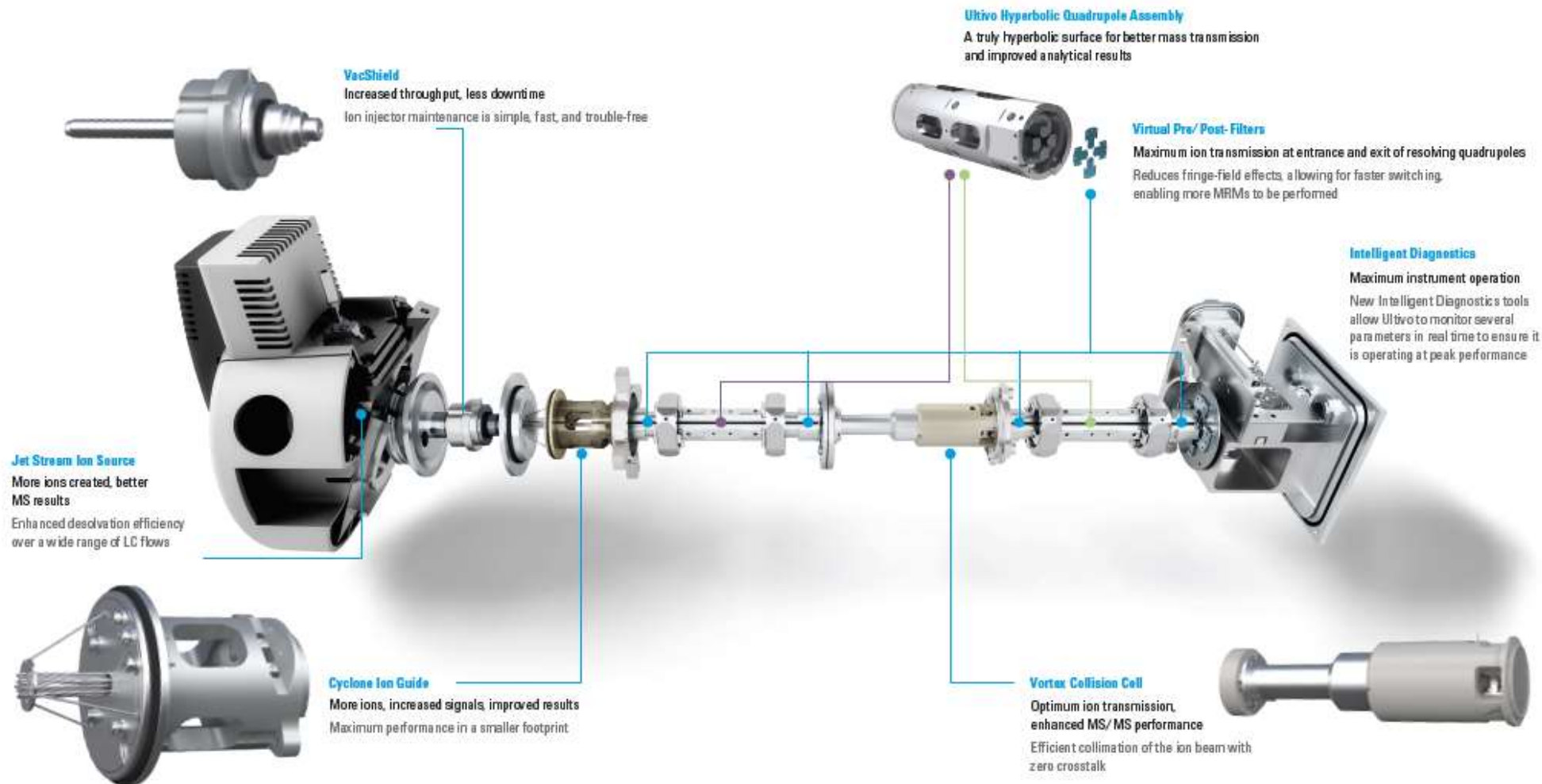
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MS Solutions Product Manager

Yanan Yang, Ph.D.
Application Scientist

Jennifer Hitchcock, Ph.D.
Application Scientist



Ultivo: Innovation with Purpose!



Trade Shows/Conferences

- HPLC 2017, Prague
- ASMS 2017
- NACRW 2017
- Taiwan MS Society
- ANZSMS
- RACI 100
- British MS Society (September)
- JASIS
- LabAfrika (Johannesburg)
- Korea MS Society (August)
- LabThailand (September)
- LabAsia (October)
- Analitica, Brazil (September)
- SETAC, Brazil (September)
- India
 - Analytika-Anacon (September)
 - Customer events (Jim Yano)
- BCEIA, China (October)
- RAFA (CZ)
- Intl POP conference Japan (sept)
- AOAC
- SETAC (No. Am)

Press Events

- HPLC 2017, Prague
- ASMS 2017
- China (Beijing)
- VIP Event, London
- Singapore (June)
- VIP Event, Mexico City
- India (Sept)

Seminars

- EMEAI – Manchester
- AFO
 - Environmental Seminar (Houston)
 - **Lunch and learn events (Aug-Oct)**
- Gr. China – Shanghai, Taiwan, Hong Kong. MS Tech seminar series throughout Gr. China
- Japan – 6X seminars (August/Sept)
- SAPK – MS Technical Seminars (4X) July
 - **Indonesia (September)**
- Korea (August)
- India (August-September)

Training Events

- EMEAI – IDO, Waldbronn AE/PS
- AFO – ASMS, WebEx
- LATAM – AE, PS, IDO
- Gr. China – PS/AE, WebEx (2), F-t-F
- Japan – PS/AE, WebEx, F-t-F
- SAPK – SMT-IDO, ANZ (TP)/WebEx
- Korea – Direct and DR training August
- India – Field training (July)
- ANZ – Software & apps (M. Tischler, T Anumol.)
- **AE Training @ SCS**
 - **Oct 2-6, Oct 9-18 (all regions ex. China)**
 - **Nov 6-10 (CN specific)**



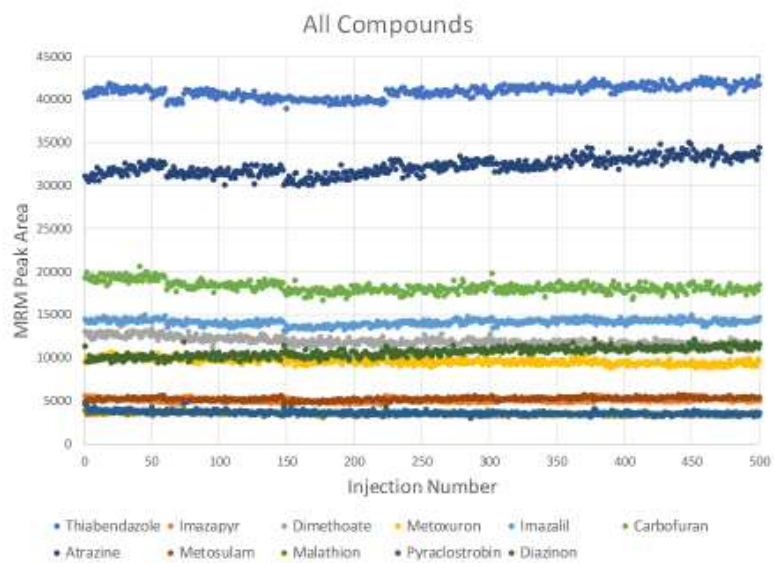
VIP Events

- EMEAI
 - HPLC 2017, Prague
 - Uppsala, Sweden
 - London, England
 - Basel, Switzerland
 - Milan, Italy
- AFO
 - Santa Clara, CA
 - Little Falls, DE
 - Chicago, IL
 - LATAM
 - Mexico City (August)
 - Sao Paulo (August)
 - Argentina (September)
 - Colombia (September)
- Gr. China
 - Beijing
- Japan
 - JASIS (September)
- SAPK
 - Singapore X2 (Monty, Mike McMullen)
 - Melbourne (RACI)
- ASMS 2017
- China (BCEIA)
- VIP Event, London
- Korea
 - Seoul (August)
- **India (September, two locations)**

Ultivo LC/TQ: Small, Robust, Easy-to-Use with EMF Feedback

Ultivo System Robustness

20 ppb Pesticide Basic Checkout Mix spiked into Avocado QuEChERS-EMR



| Compound | % RSD (n=500) |
|----------------|---------------|
| Thiabendazole | 1.8% |
| Imazapyr | 4.0% |
| Dimethoate | 4.1% |
| Metoxuron | 3.6% |
| Imazalil | 2.3% |
| Carbofuran | 3.3% |
| Atrazine | 3.0% |
| Metosulam | 3.5% |
| Malathion | 4.3% |
| Pyraclostrobin | 6.1% |
| Diazinon | 5.0% |

Data collected by Mark Sartain

Instrument Status

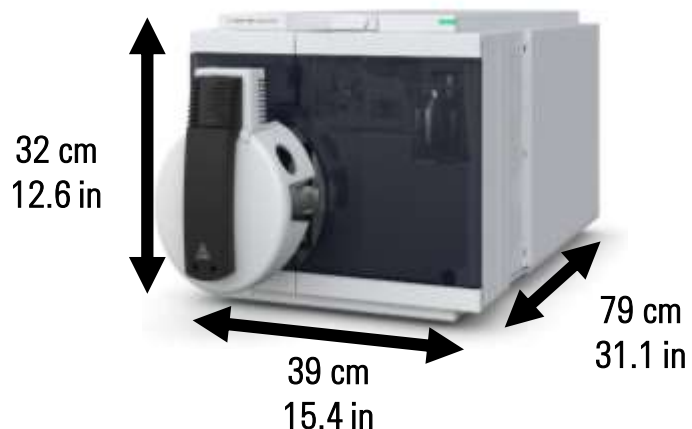
QQQ ? - =

Tune

On Off EMF ⚠

ESI

Capillary clog



Infinity II
HPLC Stack

Ultivo
Triple Quad

Incredibly Powerful, Remarkably Efficient Quantitation

MassHunter Productivity Software
Applications

Ruben Garnica, Ph. D.
MS Solutions Product Manager
Agilent Technologies, Inc.
Wilmington, DE



Ultivo LC/TQ – Reducing Pain Points



Diagnostics – Provide “real-time” information to customer concerning stability/usability of the system



Form Factor – Size/ Footprint, stackability, external appearance



Reliability – Overall (mean time to service), sample throughput, consumable components (TMP, ion injector)



Ease-of-Use - Initial set-up / post-sales support (e.g. method development), training, fit into workflow, software



Performance – Sensitivity, selectivity, throughput, positive/negative switching, mass range / resolution, matrix



Total Cost of Ownership – Initial cost, servicing cost, gas usage, training, noise reduction



MassHunter is frequently the right tool...

File Launch Edit View Find Identify Method Configuration Tools Help

Sample Table: Cal 50 ppb.d

Results Workflow

Result Summary Saved Results Method Last Run Method Workflow Target Source Filename Sample Position Acq

SS qualified (5) Default.m << Interactive >> Custom Pestl Cal 50 ppb.d Pt-A4 Custom I

Compound List: 55 found

| m/z | RT | Width | Height | Area | Score | Mining Algorithm | ID Source | Cl | File |
|-------|--------|-------|--------|---------|-------|------------------|-----------|----|------|
| 406 | 11.16 | 0.311 | 11 | 25 | 100 | Find by MRM | Acq | 15 | |
| 230 | 3.776 | 0.257 | 150784 | 568... | 100 | Find by MRM | Acq | 24 | |
| 326 | 10.974 | 0.351 | 115962 | 491... | 100 | Find by MRM | Acq | 30 | |
| 330 | 9.987 | 0.353 | 179107 | 764... | 100 | Find by MRM | Acq | 20 | |
| 265.1 | 3.431 | 0.273 | 112619 | 378... | 100 | Find by MRM | Acq | 12 | |
| 385 | 12.751 | 0.283 | 121865 | 458... | 100 | Find by MRM | Acq | 8 | |
| 243 | 8.05 | 0.323 | 96378 | 388... | 100 | Find by MRM | Acq | 16 | |
| 218 | 7.3 | 0.322 | 11887 | 4950... | 100 | Find by MRM | Acq | 24 | |
| 331 | 8.775 | 0.32 | 39532 | 170... | 100 | Find by MRM | Acq | 20 | |
| 337.1 | 9.696 | 0.351 | 86434 | 363... | 100 | Find by MRM | Acq | 22 | |
| 462.9 | 12.598 | 0.304 | 2070 | 7531 | 100 | Find by MRM | Acq | 13 | |
| 247 | 7.544 | 0.232 | 345 | 1389 | 100 | Find by MRM | Acq | 32 | |
| 266.1 | 7.547 | 0.273 | 13351 | 55867 | 100 | Find by MRM | Acq | 8 | |
| 328 | 3.417 | 0.259 | 206463 | 700... | 100 | Find by MRM | Acq | 28 | |
| 213.1 | 5.853 | 0.281 | 178987 | 669... | 100 | Find by MRM | Acq | 15 | |
| 459 | 8.587 | 0.331 | 185909 | 755... | 100 | Find by MRM | Acq | 15 | |
| 376 | 8.623 | 0.269 | 23768 | 181... | 100 | Find by MRM | Acq | 24 | |
| 116 | 8.32 | 0.256 | 148293 | 595... | 100 | Find by MRM | Acq | 15 | |

Method Editor: Workflow

Run Method Workflow

Method Automation

Workflow

Additional Chromatograms

Reports

Export

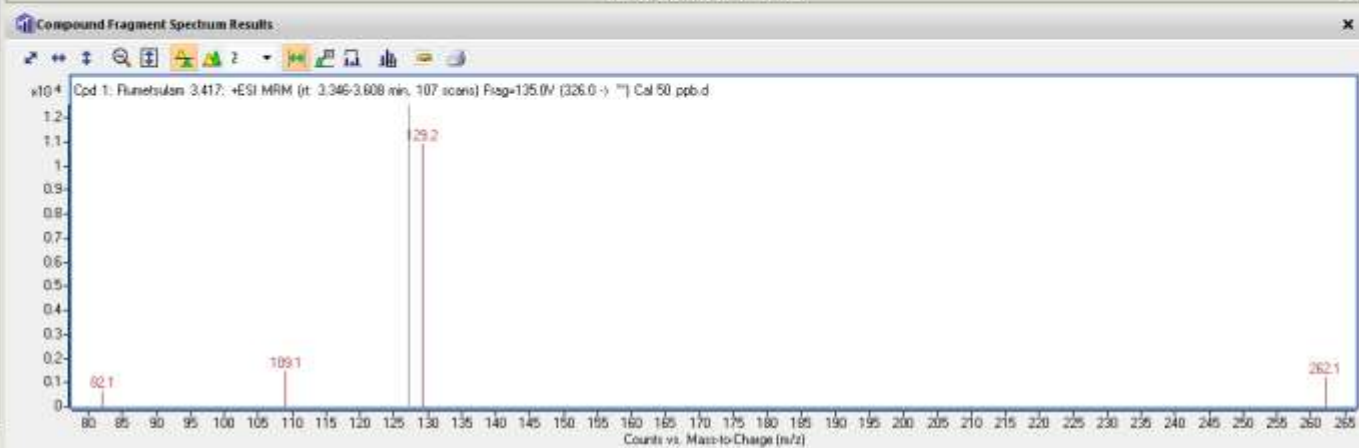
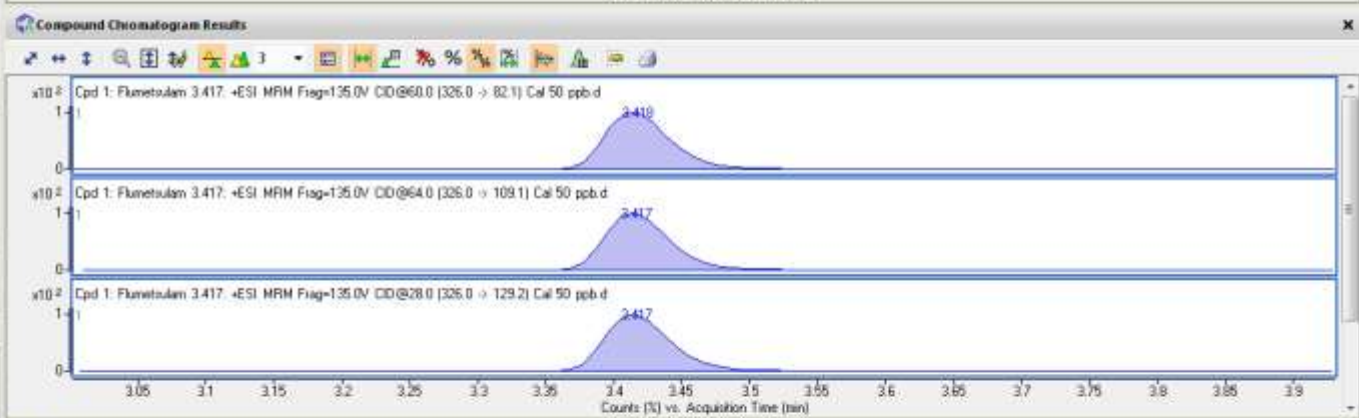
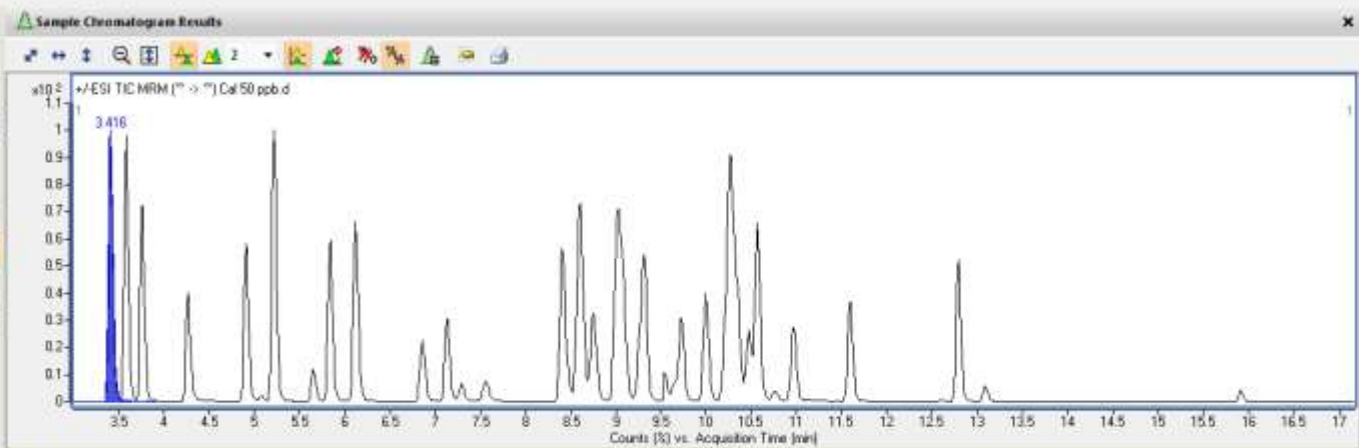
Target/Suspect Screening

MS Extraction

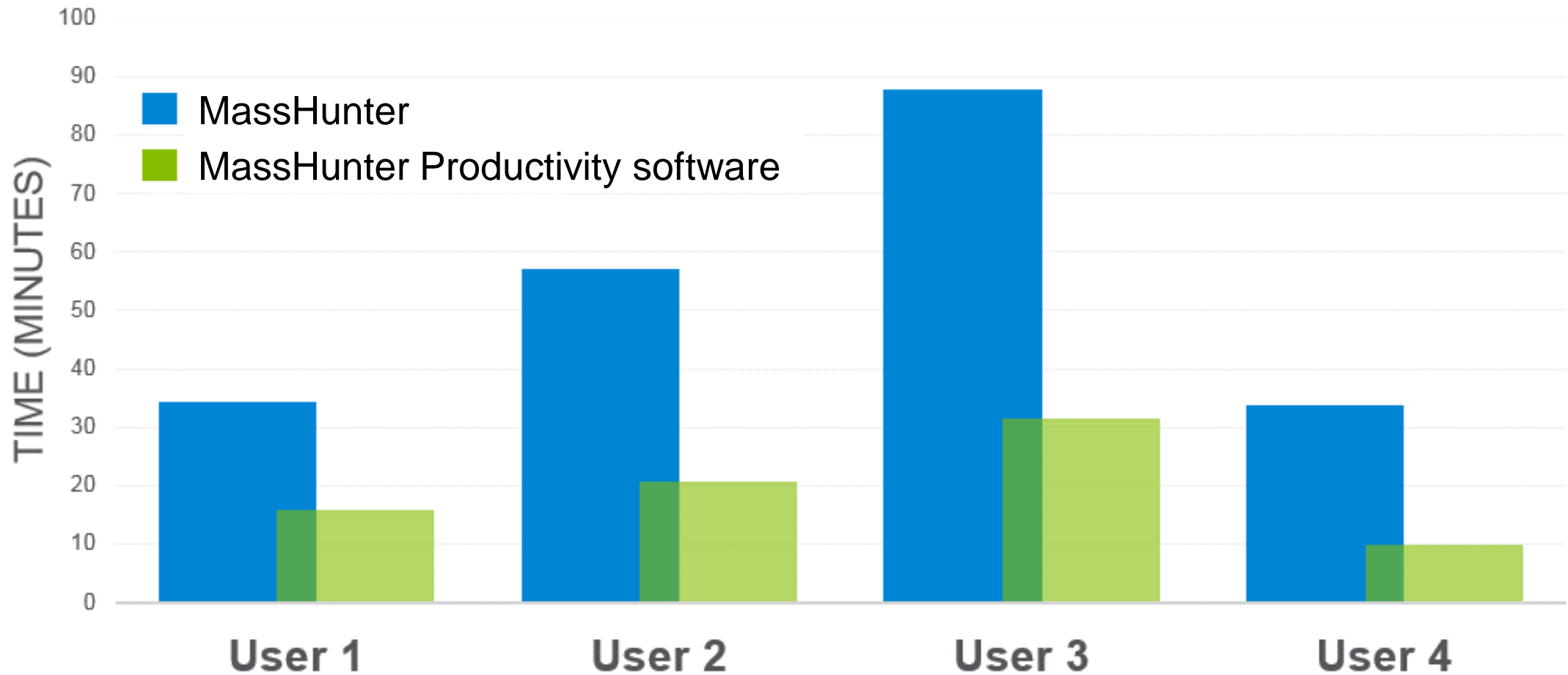
Options Time Range(s)

Workflow Target/Suspect Screening

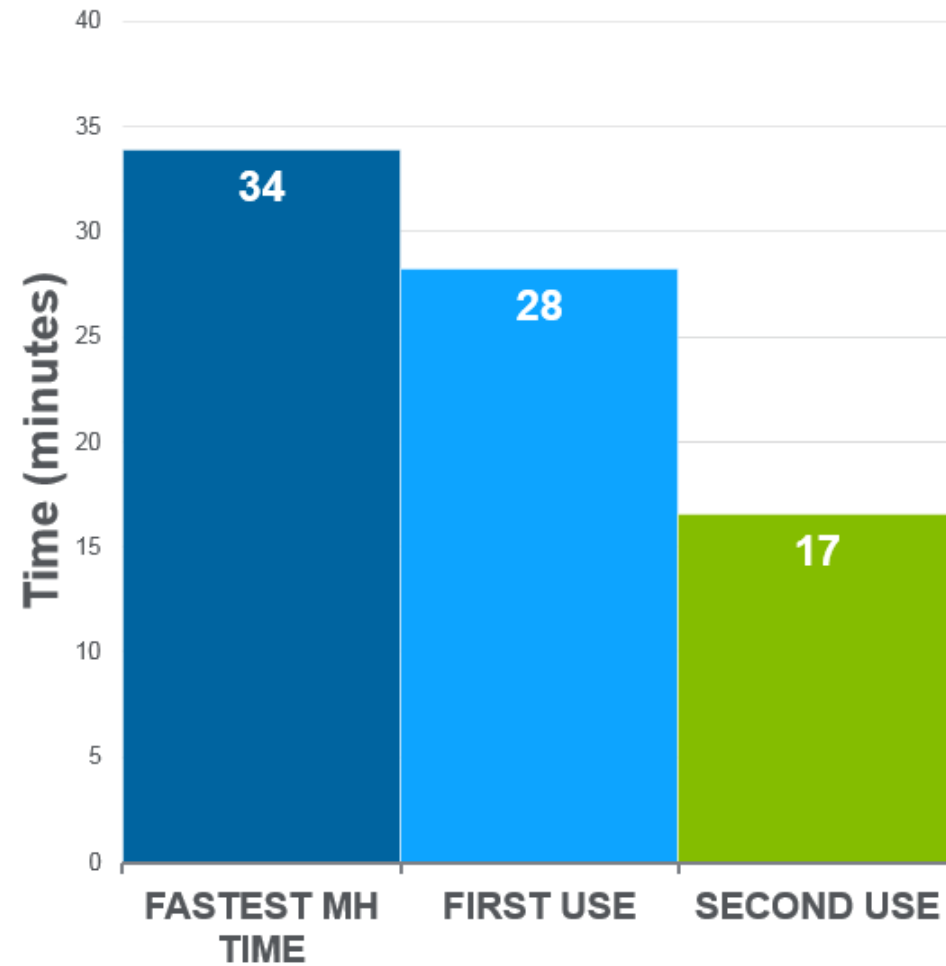
Compound mining Find by MRM



What about production laboratories?



Time to execute standard targeted quant workflow



Increased productivity gains after only two uses

Build Sequence

Select Compounds

Sequence



Runtime: 16:05:59

| Name | Position | Type | Group | Volume | Inj/Sample | Comments | Compound Groups |
|-----------------------------|----------|--------|-------|--------|------------|----------|-----------------|
| 1 X Sample 1 blank | P1-A9 | Sample | | 20 | 1 | | + |
| 2 X Sample 2 cpd 1 | P1-B1 | Sample | | 2 | 1 | | + |
| 3 X Sample 3 cpd 2 | P1-B2 | Sample | | 2 | 1 | | + |
| 4 X Sample 4 blank | P1-B3 | Sample | | 20 | 1 | | + |
| 5 X Sample 5 blank | P1-B4 | Sample | | 20 | 1 | | + |
| 6 X Sample 6 cpd 3 | P1-B5 | Sample | | 2 | 1 | | + |
| 7 X Sample 7 blank | P1-B6 | Sample | | 20 | 1 | | + |
| 8 X Sample 8 cpd 1_2 | P1-B7 | Sample | | 2 | 1 | | + |
| 9 X Sample 9 cpd 4 | P1-B8 | Sample | | 2 | 1 | | + |
| 10 X Sample 10 cpd 5 | P1-B9 | Sample | | 2 | 1 | | + |
| 11 X Sample 11 cpd 4_5 | P1-C2 | Sample | | 2 | 1 | | + |
| 12 X Sample 12 cpd 1_5 | P1-C3 | Sample | | 2 | 1 | | + |
| 13 X Sample 13 blank | P1-C4 | Sample | | 20 | 1 | | + |
| 14 X Sample 14 blank | P1-C5 | Sample | | 20 | 1 | | + |
| 15 X Sample 15 cpd 2_3 | P1-C6 | Sample | | 2 | 1 | | + |
| 16 X Sample 16 cpd 1 | P1-C7 | Sample | | 2 | 1 | | + |
| 17 X Sample 17 cpd 1_4 | P1-C8 | Sample | | 2 | 1 | | + |
| 18 X Sample 18 blank | P1-C9 | Sample | | 20 | 1 | | + |
| 19 X Sample 19 cpd 4 200ppb | P1-D1 | Sample | | 2 | 1 | | + |
| 20 X Sample 20 blank | P1-D2 | Sample | | 20 | 1 | | + |
| 21 X Sample 21 blank | P1-D4 | Sample | | 20 | 1 | | + |
| 22 X Sample 22 cpd 2 | P1-D5 | Sample | | 2 | 1 | | + |
| 23 X Sample 23 cpd 3 | P1-D6 | Sample | | 2 | 1 | | + |
| 24 X Sample 24 cpd 1_2_3 | P1-D7 | Sample | | 2 | 1 | | + |
| 25 X Sample 25 blank | P1-D8 | Sample | | 20 | 1 | | + |
| 26 X Sample 26 cpd 2_4_5 | P1-D9 | Sample | | 2 | 1 | | + |
| 27 X Sample 27 blank | P1-E1 | Sample | | 20 | 1 | | + |

Apply Sequence Template

Apply Sequence Template

Template: ...

Use sample list: ...

Create samples:

Use current sequence

Shutdown instrument after run completes

File Home

Standby Start Stop Setup Monitor Review RT Tune Log In

Instrument Run Control Views Actions Admin

Select Compounds



Runtime: 16:05:59

| Name | Position | Type | Group | Volume | Inj/Sample | Comments | Compound Groups |
|-----------------------------|----------|--------|-------|--------|------------|----------|-----------------|
| 1 X Sample 1 blank | P1-A9 | Sample | | 20 | 1 | | + |
| 2 X Sample 2 cpd 1 | P1-B1 | Sample | | 2 | 1 | | + |
| 3 X Sample 3 cpd 2 | P1-B2 | Sample | | 2 | 1 | | + |
| 4 X Sample 4 blank | P1-B3 | Sample | | 20 | 1 | | + |
| 5 X Sample 5 blank | P1-B4 | Sample | | 20 | 1 | | + |
| 6 X Sample 6 cpd 3 | P1-B5 | Sample | | 2 | 1 | | |
| 7 X Sample 7 blank | P1-B6 | Sample | | 20 | 1 | | |
| 8 X Sample 8 cpd 1_2 | P1-B7 | Sample | | 2 | 1 | | |
| 9 X Sample 9 cpd 4 | P1-B8 | Sample | | 2 | 1 | | |
| 10 X Sample 10 cpd 5 | P1-B9 | Sample | | 2 | 1 | | |
| 11 X Sample 11 cpd 4_5 | P1-C2 | Sample | | 2 | 1 | | |
| 12 X Sample 12 cpd 1_5 | P1-C3 | Sample | | 2 | 1 | | |
| 13 X Sample 13 blank | P1-C4 | Sample | | 20 | 1 | | |
| 14 X Sample 14 blank | P1-C5 | Sample | | 20 | 1 | | |
| 15 X Sample 15 cpd 2_3 | P1-C6 | Sample | | 2 | 1 | | |
| 16 X Sample 16 cpd 1 | P1-C7 | Sample | | 2 | 1 | | |
| 17 X Sample 17 cpd 1_4 | P1-C8 | Sample | | 2 | 1 | | + |
| 18 X Sample 18 blank | P1-C9 | Sample | | 20 | 1 | | + |
| 19 X Sample 19 cpd 4 200ppb | P1-D1 | Sample | | 2 | 1 | | + |
| 20 X Sample 20 blank | P1-D2 | Sample | | 20 | 1 | | + |
| 21 X Sample 21 blank | P1-D4 | Sample | | 20 | 1 | | + |
| 22 X Sample 22 cpd 2 | P1-D5 | Sample | | 2 | 1 | | + |
| 23 X Sample 23 cpd 3 | P1-D6 | Sample | | 2 | 1 | | + |
| 24 X Sample 24 cpd 1_2_3 | P1-D7 | Sample | | 2 | 1 | | + |
| 25 X Sample 25 blank | P1-D8 | Sample | | 20 | 1 | | + |
| 26 X Sample 26 cpd 2_4_5 | P1-D9 | Sample | | 2 | 1 | | + |
| 27 X Sample 27 blank | P1-E1 | Sample | | 20 | 1 | | + |

Apply Sequence Template

Apply Sequence Template

Template: Pesticide Analysis Sequence Template

Use sample list: []

Create samples: 10

Use current sequence

Apply Cancel

Shutdown instrument after run completes

HiP Sampler | Binary Pump | Column Comp. | QQQ

Idle

Current user: admin

Build Sequence

Select Compounds

Sequence



Runtime: 16:05:59

| Name | Position | Type | Group | Volume | Inj/Sample | Comments | Compound Groups |
|-----------------------------|----------|--------|-------|--------|------------|----------|-----------------|
| 1 X Sample 1 blank | P1-A9 | Sample | | 20 | 1 | | + |
| 2 X Sample 2 cpd 1 | P1-B1 | Sample | | 2 | 1 | | |
| 3 X Sample 3 cpd 2 | P1-B2 | Sample | | 2 | 1 | | |
| 4 X Sample 4 blank | P1-B3 | Sample | | 20 | 1 | | |
| 5 X Sample 5 blank | P1-B4 | Sample | | 20 | 1 | | |
| 6 X Sample 6 cpd 3 | P1-B5 | Sample | | 2 | 1 | | |
| 7 X Sample 7 blank | P1-B6 | Sample | | 20 | 1 | | |
| 8 X Sample 8 cpd 1_2 | P1-B7 | Sample | | 2 | 1 | | |
| 9 X Sample 9 cpd 4 | P1-B8 | Sample | | 2 | 1 | | |
| 10 X Sample 10 cpd 5 | P1-B9 | Sample | | 2 | 1 | | |
| 11 X Sample 11 cpd 4_5 | P1-C2 | Sample | | 2 | 1 | | |
| 12 X Sample 12 cpd 1_5 | P1-C3 | Sample | | 2 | 1 | | |
| 13 X Sample 13 blank | P1-C4 | Sample | | 20 | 1 | | |
| 14 X Sample 14 blank | P1-C5 | Sample | | 20 | 1 | | |
| 15 X Sample 15 cpd 2_3 | P1-C6 | Sample | | 2 | 1 | | |
| 16 X Sample 16 cpd 1 | P1-C7 | Sample | | 2 | 1 | | |
| 17 X Sample 17 cpd 1_4 | P1-C8 | Sample | | 2 | 1 | | |
| 18 X Sample 18 blank | P1-C9 | Sample | | 20 | 1 | | |
| 19 X Sample 19 cpd 4 200ppb | P1-D1 | Sample | | 2 | 1 | | |
| 20 X Sample 20 blank | P1-D2 | Sample | | 20 | 1 | | + |
| 21 X Sample 21 blank | P1-D4 | Sample | | 20 | 1 | | + |
| 22 X Sample 22 cpd 2 | P1-D5 | Sample | | 2 | 1 | | + |
| 23 X Sample 23 cpd 3 | P1-D6 | Sample | | 2 | 1 | | + |
| 24 X Sample 24 cpd 1_2_3 | P1-D7 | Sample | | 2 | 1 | | + |
| 25 X Sample 25 blank | P1-D8 | Sample | | 20 | 1 | | + |
| 26 X Sample 26 cpd 2_4_5 | P1-D9 | Sample | | 2 | 1 | | + |
| 27 X Sample 27 blank | P1-E1 | Sample | | 20 | 1 | | + |

Apply Sequence Template

Template:

Use sample list:

Create samples:

Use current sequence

Shutdown instrument after run completes

Review Analysis

Review Control

Design Report

Results - 06 May 2018 - 11_02_16

Select results for report

Filters

- Select filter set...
 Select filter set...
 Filter Flag-1
 Filter Flag-2
 Filter Flag-3

Flags

- No flags (18)
- Accuracy out of range (55)
- Above calibration range (2)
- R2 below threshold (180)
- Qualifier peak not found (6)
- Qualifier ratio out of range (33)
- QC out of limits (2)
- Retention time out of range (81)
- CC response ratio out of range (18)
- Matrix spike percent recovery out of range (18)

Sample Type

- Sample (54)
- Blank (18)
- Double Blank (18)
- Calibration (54)
- QC (18)
- CC (18)
- Matrix Blank (18)
- Matrix Spike (18)

Report Status

- Selected (216)

Samples

- Search...
- Blank 0.d (18)
Blank, PLA1, Acquired Sunday, May 06, 2018 11:03 AM, 5.00uL
 - Calibration 1.d (18)
Calibration, PLA2, Acquired Sunday, May 06, 2018 11:25 AM, 0.30uL
 - Calibration 2.d (18)
Calibration, PLA3, Acquired Sunday, May 06, 2018 11:47 AM, 0.30uL
 - Calibration 3.d (18)
Calibration, PLA4, Acquired Sunday, May 06, 2018 12:09 PM, 0.30uL
 - Sample 1.d (18)
Sample, PLA5, Acquired Sunday, May 06, 2018 12:32 PM, 0.30uL
 - Sample 2.d (18)
Sample, PLA5, Acquired Sunday, May 06, 2018 12:54 PM, 0.30uL
 - Sample 3.d (18)
Sample, PLA5, Acquired Sunday, May 06, 2018 1:16 PM, 0.30uL
 - QC 1.d (18)
QC, PLA6, Acquired Sunday, May 06, 2018 1:39 PM, 0.30uL
 - CC 1.d (18)
CC, PLA7, Acquired Sunday, May 06, 2018 2:01 PM, 0.30uL
 - Matrix Blank 10 (18)
Matrix Blank, PLA8, Acquired Sunday, May 06, 2018 2:23 PM, 0.30uL
 - Matrix Spike 11 (18)

Compounds

Search...

Beflubutamid (Sample 2.d) Below Threshold (2.96 ng/ml) ⚠

R2 = 0.83, 9.00 - 47.00 ng/ml

356.0 -> 91.0 356.0 -> 65.1 356.0 -> 162.1

9.72

356.0 -> 65.1
356.0 -> 162.1

Qualifier Ratios

Tags: R2 below threshold, Retention time out of range

Benalaxyl (Sample 2.d) Below Threshold (2.93 ng/ml) ⚠

R2 = 0.87, 9.00 - 47.00 ng/ml

326.1 -> 148.1 326.1 -> 91.2 326.1 -> 208.2

10.27

326.1 -> 91.2
326.1 -> 208.2

Qualifier Ratios

Tags: R2 below threshold, Retention time out of range

Bifenazate (Sample 2.d) Below Threshold (3.01 ng/ml) ⚠

R2 = 0.97, 9.00 - 47.00 ng/ml

301.1 -> 198.2 301.1 -> 170.1 301.1 -> 153.1

8.42

301.1 -> 170.1
301.1 -> 153.1

Qualifier Ratios

Tags: R2 below threshold, Retention time out of range

Filters

Select filter set...

- Select filter set...
- Filter Flags-1
- Filter Flags-2
- Filter Flag-3

Flags

- No flags (18)
- Accuracy out of range (55)
- Above calibration range (2)
- R2 below threshold (180)
- Qualifier peak not found (6)
- Qualifier ratio out of range (33)
- QC out of limits (2)
- Retention time out of range (81)
- CC response ratio out of range (18)
- Matrix spike percent recovery out of range (18)

Sample Type

- Sample (54)
- Blank (18)
- Double Blank (18)
- Calibration (54)
- QC (18)
- CC (18)

Review Control
Design Report

Select results for report

Compounds

Search...

Beflubutamid (Sample 2.d) Below Threshold (2.96 ng/ml) 🔍 📄 📄

R2 = 0.83, 9.00 - 47.00 ng/ml

356.0 -> 91.0 356.0 -> 65.1 356.0 -> 162.1

356.0 -> 65.1
356.0 -> 162.1

Qualifier Ratios

Tags: R2 below threshold Retention time out of range

Benalaxyl (Sample 2.d) Below Threshold (2.93 ng/ml) 🔍 📄 📄

R2 = 0.87, 9.00 - 47.00 ng/ml

326.1 -> 148.1 326.1 -> 91.2 326.1 -> 208.2

326.1 -> 91.2
326.1 -> 208.2

Qualifier Ratios

Tags: R2 below threshold Retention time out of range

Bifenazate (Sample 2.d) Below Threshold (3.01 ng/ml) 🔍 📄 📄

R2 = 0.97, 9.00 - 47.00 ng/ml

301.1 -> 198.2 301.1 -> 170.1 301.1 -> 153.1

301.1 -> 170.1
301.1 -> 153.1

Qualifier Ratios

Tags: R2 below threshold Retention time out of range

Results - 06 May 2018 - 11_02_16

Select results for report

Samples

- Search...
- Blank 0.d** (18)
Blank, PLA1, Acquired Sunday, May06, 2018 11:03 AM, 5.00uL
 - Calibration 1.d** (18)
Calibration, PLA2, Acquired Sunday, May06, 2018 11:25 AM, 0.30uL
 - Calibration 2.d** (18)
Calibration, PLA3, Acquired Sunday, May06, 2018 11:47 AM, 0.30uL
 - Calibration 3.d** (18)
Calibration, PLA4, Acquired Sunday, May06, 2018 12:09 PM, 0.30uL
 - Sample 1.d** (18)

Compounds

Search...

- Beflubutamid (Sample 2.d)** Below Threshold (2.96 ng/ml) ⚠

R2 = 0.83, 9.00 - 47.00 ng/ml

356.0 -> 91.0 356.0 -> 65.1 356.0 -> 162.1

356.0 -> 65.1
356.0 -> 162.1

Tags: R2 below threshold Retention time out of range
Add comment... Add comment...

- Calibration (54)
 - QC (18)
 - CC (18)
 - Matrix Blank (18)
 - Matrix Spike (18)
 - Report Status**
 - Selected (216)
- QC 1.d** (18)
QC, PLA6, Acquired Sunday, May06, 2018 1:39 PM, 0.30uL
 - CC 1.d** (18)
CC, PLA7, Acquired Sunday, May06, 2018 2:01 PM, 0.30uL
 - Matrix Blank 10** (18)
Matrix Blank, PLA8, Acquired Sunday, May06, 2018 2:23 PM, 0.30uL
 - Matrix Spike 11** (18)

- Bifenazate (Sample 2.d)** Below Threshold (3.01 ng/ml) ⚠

R2 = 0.97, 9.00 - 47.00 ng/ml

301.1 -> 198.2 301.1 -> 170.1 301.1 -> 153.1

301.1 -> 170.1
301.1 -> 153.1

Tags: R2 below threshold Retention time out of range
Add comment... Add comment...

Review Analysis

Review Control

Design Report

Report Options

Template

Default Report

Options

Generate chromatogram images

Report Preview

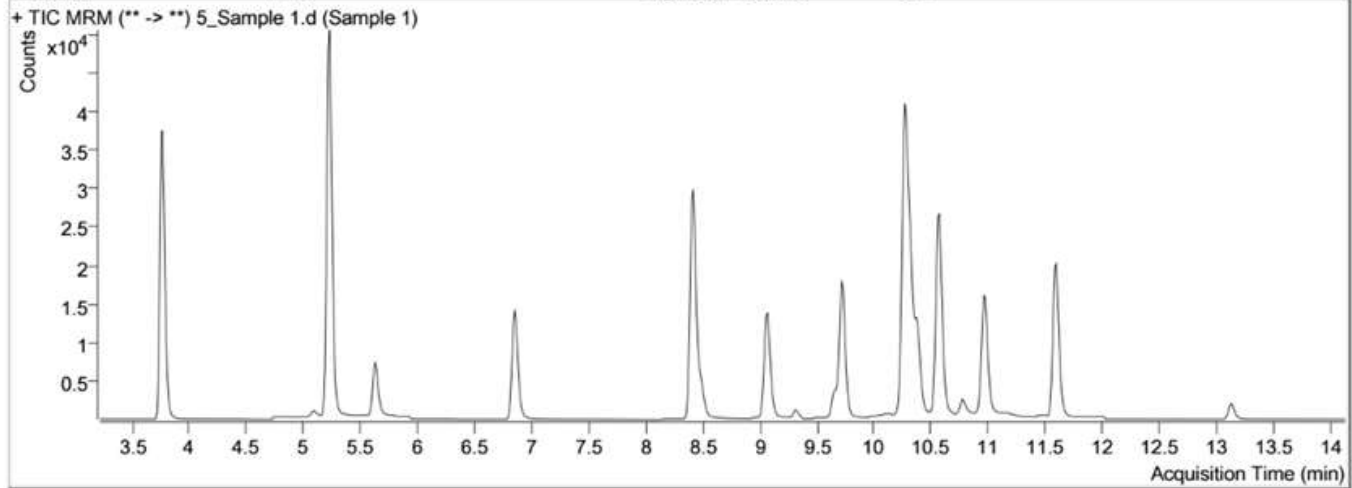
9 / 24 98% Find:

Bookmarks

- Blank 0 (1_Blank 0.d)
- Calibration 1 (2_Calibration 1.d)
- Calibration 2 (3_Calibration 2.d)
- Calibration 3 (4_Calibration 3.d)
- Sample 1 (5_Sample 1.d)
- Sample 2 (6_Sample 2.d)
- Sample 3 (7_Sample 3.d)
- QC1 (8_QC 1.d)
- QC1 (9_QC 1.d)
- Matrix Blank 10 (10_Matrix Blank 10.d)
- Matrix Spike 11 (11_Matrix Spike 11.d)
- Double Blank 9 (12_Double Blank 9.d)

Analysis Info

| | | | |
|--------------------|-------------------------|-------------------------|-------------------|
| Instrument | Instrument 1 | Operator | |
| Data File | 5_Sample 1.d | Sample Name | Sample 1 |
| Sample Type | Sample | Dilution | 1 |
| Acq. Method | AcquisitionTempMethod.m | Acq. Date | 5/6/2018 12:32 PM |
| Position | P1A5 | Injection Volume | 0.3 |



| Compound | RT | Ref RT | Transition(T) | Transition(Q) | T-Resp | Q-Resp | QRatio | Ref QRatio | Final Conc. | Units |
|--------------------|-------|--------|----------------|----------------|--------|--------|--------|------------|-------------|-------|
| Dimethoate | 3.756 | 3.820 | 230.0 -> 125.0 | 230.0 -> 199.0 | 41325 | 36083 | 87.3 | 87.8 | 3.1821 | ng/ml |
| R2 below threshold | | | | | | | | | | |
| Dichlorvos | 5.095 | 5.200 | 221.0 -> 109.0 | 221.0 -> 79.0 | 1888 | 242 | 12.8 | 12.2 | 3.5974 | ng/ml |
| Carbofuran | 5.232 | 5.340 | 222.1 -> 123.1 | 222.1 -> 165.1 | 79339 | 71692 | 90.4 | 90.2 | 2.9662 | ng/ml |
| R2 below threshold | | | | | | | | | | |
| Imazalil | 5.629 | 5.700 | 297.0 -> 159.0 | 297.0 -> 255.1 | 9571 | 4873 | 50.9 | 52.6 | 3.1928 | ng/ml |
| R2 below threshold | | | | | | | | | | |

MassHunter Productivity Software Application

Targeted Quant with Pesticides MRM database

Use your methods and MRM database or use ours!

- MassHunter Productivity Software
- Pesticides dMRM DB for LC/TQ
 - >600 commonly used pesticides with RT
 - 2-4 optimized MRM transitions per compound
 - 3-year free updates
- Workflow Guide
- Acquisition and Quant Methods
- Suite of Customizable Report Templates
- Optional Consumables

Demonstrations available
this week!



Analytical Determination of Small Molecules Drugs in Serum and Amino Acids in Animal Feed Matrices using Ultivo LC/TQ

Yanan Yang, PhD
LCMS Application Scientist
Agilent Technologies, Inc.
Santa Clara, CA



Sample Preparation and LCMS Method



Sample Preparation

- 0.1 g homogenized animal feed sample was weighed in a 250 mL Erlenmeyer flask and hydrolyzed using 25 mL 6N HCl with phenol.
- Incubation under nitrogen flow for 23 hours.
- 0.1N HCl was filled to 50 mL and mixed. Aliquots were transferred to sample vials for introduction into the LC/TQ system.

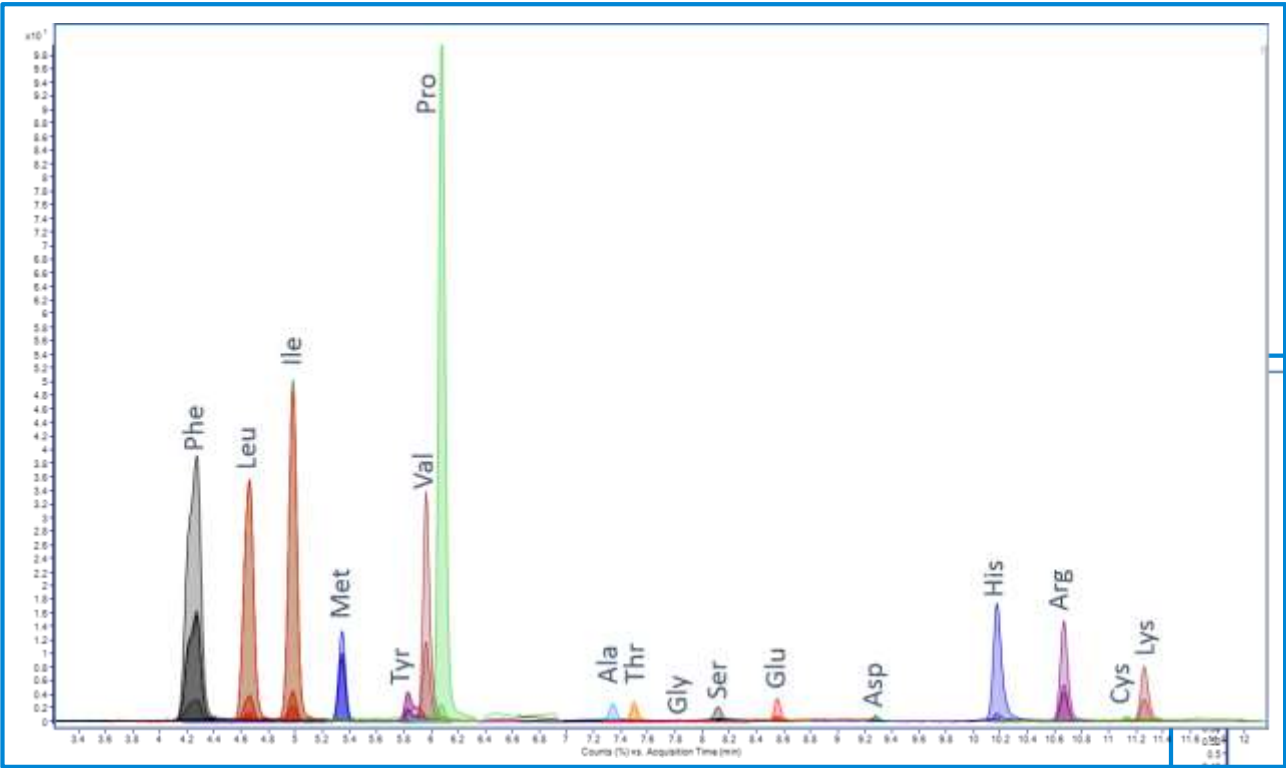
| Column | Poroshell 120 HILIC-Z 2.1 x 150 mm, 2.7µm (p/n 6x3775-924) | | | | | | | | | | | | | |
|---------------------------|--|------|----|---|-----|------|----|------|-----|------|----------------------|------|----------------------|--|
| Column temp | 25°C | | | | | | | | | | | | | |
| Injection volume | 2 µL | | | | | | | | | | | | | |
| Mobile phase | A: 20 mM ammonium acetate + 0.1% formic acid in water, pH 3 B: 20 mM ammonium acetate in 90% acetonitrile, pH 3 | | | | | | | | | | | | | |
| Flow rate | 0.50 mL/min | | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time</th> <th>B%</th> </tr> </thead> <tbody> <tr> <td>0</td> <td>100</td> </tr> <tr> <td>11.5</td> <td>70</td> </tr> <tr> <td>12.0</td> <td>100</td> </tr> <tr> <td>15.0</td> <td>100 <i>stop-time</i></td> </tr> <tr> <td>19.0</td> <td>100 <i>post time</i></td> </tr> </tbody> </table> | Time | B% | 0 | 100 | 11.5 | 70 | 12.0 | 100 | 15.0 | 100 <i>stop-time</i> | 19.0 | 100 <i>post time</i> | |
| Time | B% | | | | | | | | | | | | | |
| 0 | 100 | | | | | | | | | | | | | |
| 11.5 | 70 | | | | | | | | | | | | | |
| 12.0 | 100 | | | | | | | | | | | | | |
| 15.0 | 100 <i>stop-time</i> | | | | | | | | | | | | | |
| 19.0 | 100 <i>post time</i> | | | | | | | | | | | | | |
| Drying gas temp | 150 °C | | | | | | | | | | | | | |
| Drying gas flow | 15 L/min | | | | | | | | | | | | | |
| Sheath gas temp | 400 °C | | | | | | | | | | | | | |
| Sheath gas flow | 12 L/min | | | | | | | | | | | | | |
| Nebulizer pressure | 20 psi | | | | | | | | | | | | | |
| Capillary voltage | 2000 V(+) | | | | | | | | | | | | | |
| Nozzle voltage | 0 V(+) | | | | | | | | | | | | | |
| Cycle Time | 500 ms | | | | | | | | | | | | | |

MRM Transition List

| Compound Name | Precursor (m/z) | Product (m/z) | RT (min) | RT Window (min) | Fragmentor (V) | CE (V) |
|---------------|-----------------|---------------|----------|-----------------|----------------|--------|
| Alanine | 90.1 | 62.1 | 6.83 | 2 | 75 | 0 |
| Alanine | 90.1 | 44.1 | 6.83 | 2 | 75 | 0 |
| Arginine | 175.1 | 116.1 | 10.43 | 2 | 105 | 2 |
| Arginine | 175.1 | 70.1 | 10.43 | 2 | 105 | 8 |
| Arginine | 175.1 | 60.1 | 10.43 | 2 | 105 | 4 |
| Aspartic Acid | 134.1 | 88.0 | 9.03 | 2 | 75 | 0 |
| Aspartic Acid | 134.1 | 74.0 | 9.03 | 2 | 75 | 4 |
| Aspartic Acid | 134.1 | 70.0 | 9.03 | 2 | 75 | 6 |
| Cystine | 241.0 | 241.0 | 11.16 | 2 | 105 | 0 |
| Cystine | 241.0 | 152.0 | 11.16 | 2 | 105 | 0 |
| Cystine | 241.0 | 120.0 | 11.16 | 2 | 105 | 0 |
| Cystine | 241.0 | 74.1 | 11.16 | 2 | 105 | 25 |
| Glutamic Acid | 148.1 | 148.1 | 8.27 | 2 | 85 | 0 |
| Glutamic Acid | 148.1 | 84.0 | 8.27 | 2 | 85 | 6 |
| Glutamic Acid | 148.1 | 56.1 | 8.27 | 2 | 85 | 22 |
| Glutamic Acid | 148.1 | 41.0 | 8.27 | 2 | 85 | 18 |
| Glycine | 76.0 | 48.0 | 7.36 | 2 | 65 | 0 |
| Glycine | 76.0 | 30.0 | 7.36 | 2 | 65 | 0 |
| Histidine | 156.1 | 110.1 | 9.81 | 2 | 95 | 4 |
| Histidine | 156.1 | 95.1 | 9.81 | 2 | 95 | 6 |
| Isoleucine | 132.1 | 86.1 | 4.90 | 2 | 85 | 0 |
| Isoleucine | 132.1 | 44.1 | 4.90 | 2 | 85 | 16 |
| Isoleucine | 132.1 | 41.0 | 4.90 | 2 | 85 | 18 |
| Isoleucine | 132.1 | 30.0 | 4.90 | 2 | 85 | 6 |
| Leucine | 132.1 | 86.1 | 4.62 | 2 | 85 | 0 |

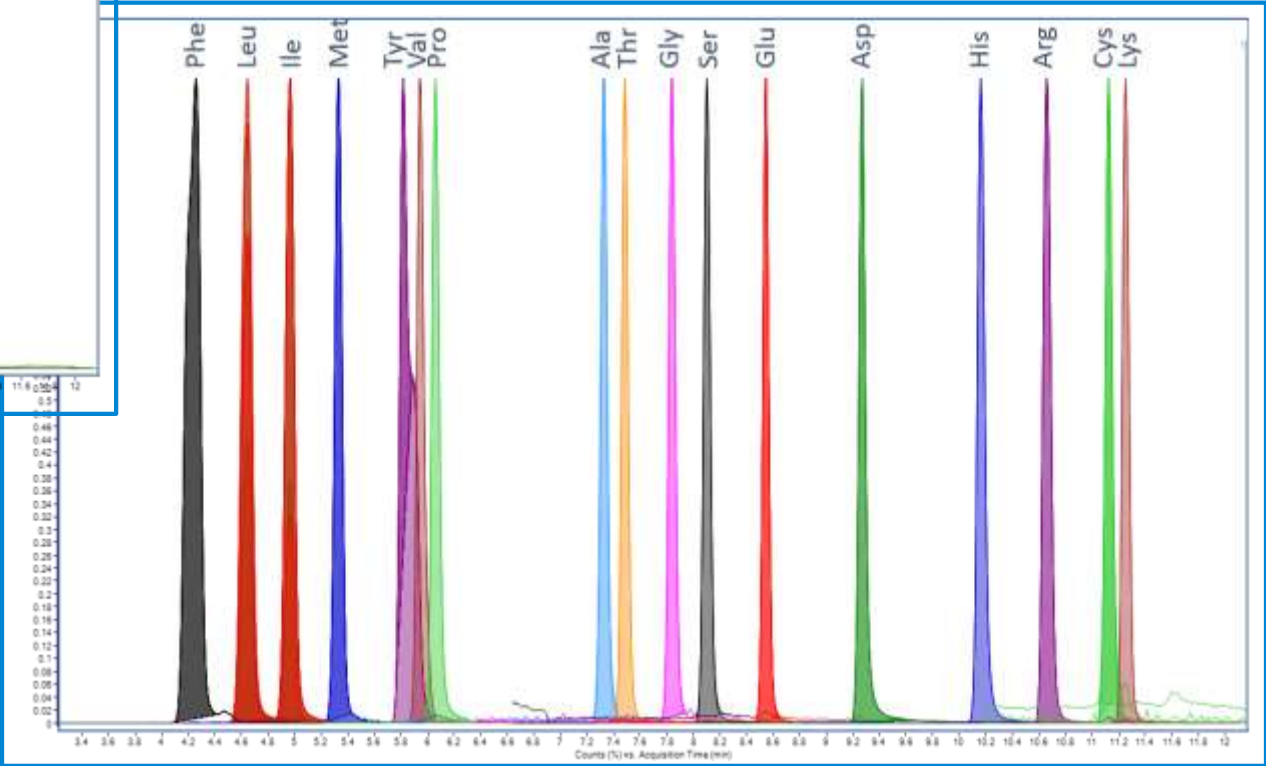
| Compound Name | Precursor (m/z) | Product (m/z) | RT (min) | RT Window (min) | Fragmentor (V) | CE (V) |
|---------------|-----------------|---------------|----------|-----------------|----------------|--------|
| Leucine | 132.1 | 44.1 | 4.62 | 2 | 85 | 14 |
| Leucine | 132.1 | 41.0 | 4.62 | 2 | 85 | 25 |
| Leucine | 132.1 | 30.0 | 4.62 | 2 | 85 | 4 |
| Lysine | 147.1 | 130.1 | 11.12 | 2 | 85 | 0 |
| Lysine | 147.1 | 84.1 | 11.12 | 2 | 85 | 6 |
| Methionine | 150.1 | 104.1 | 5.16 | 2 | 75 | 0 |
| Methionine | 150.1 | 61.0 | 5.16 | 2 | 75 | 14 |
| Methionine | 150.1 | 56.1 | 5.16 | 2 | 75 | 6 |
| Methionine | 150.1 | 28.0 | 5.16 | 2 | 75 | 26 |
| Phenylalanine | 166.1 | 120.1 | 4.23 | 2 | 85 | 4 |
| Phenylalanine | 166.1 | 103.1 | 4.23 | 2 | 85 | 22 |
| Phenylalanine | 166.1 | 91.1 | 4.23 | 2 | 85 | 32 |
| Phenylalanine | 166.1 | 77.0 | 4.23 | 2 | 85 | 36 |
| Proline | 116.1 | 70.1 | 6.01 | 2 | 85 | 6 |
| Proline | 116.1 | 43.1 | 6.01 | 2 | 85 | 25 |
| Serine | 103.1 | 88.1 | 7.63 | 2 | 65 | 0 |
| Serine | 103.1 | 60.1 | 7.63 | 2 | 65 | 0 |
| Threonine | 120.0 | 74.1 | 6.98 | 2 | 75 | 0 |
| Threonine | 120.0 | 56.1 | 6.98 | 2 | 75 | 6 |
| Tyrosine | 182.1 | 136.1 | 5.53 | 2 | 95 | 0 |
| Tyrosine | 182.1 | 119.1 | 5.53 | 2 | 95 | 10 |
| Tyrosine | 182.1 | 91.1 | 5.53 | 2 | 95 | 22 |
| Tyrosine | 182.1 | 77.0 | 5.53 | 2 | 95 | 34 |
| Valine | 118.1 | 72.1 | 5.84 | 2 | 75 | 0 |
| Valine | 118.1 | 55.1 | 5.84 | 2 | 75 | 14 |

Composite dMRM Chromatogram of Amino Acids

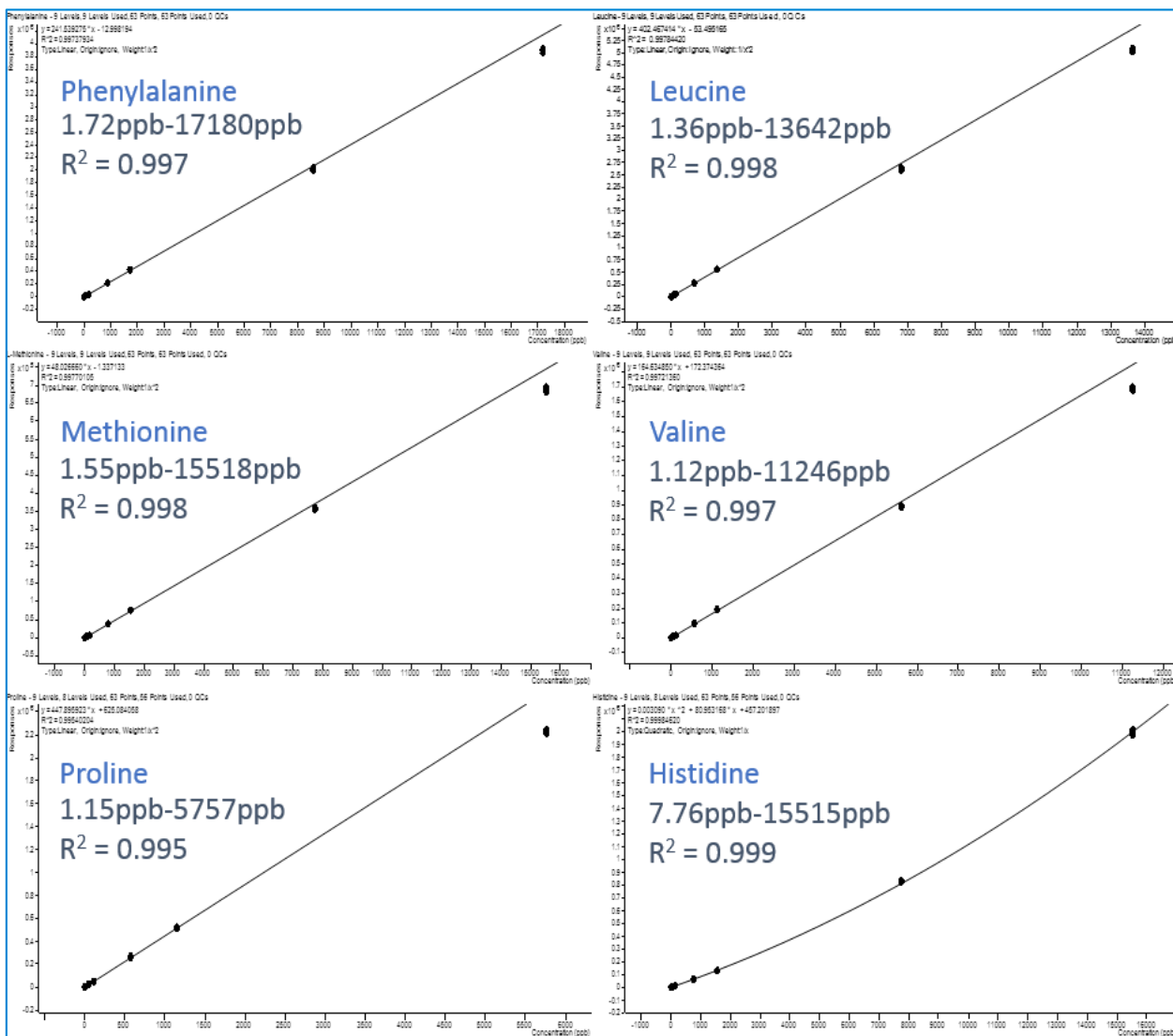


Overlaid dMRM chromatograms of amino acids

Peak height normalized

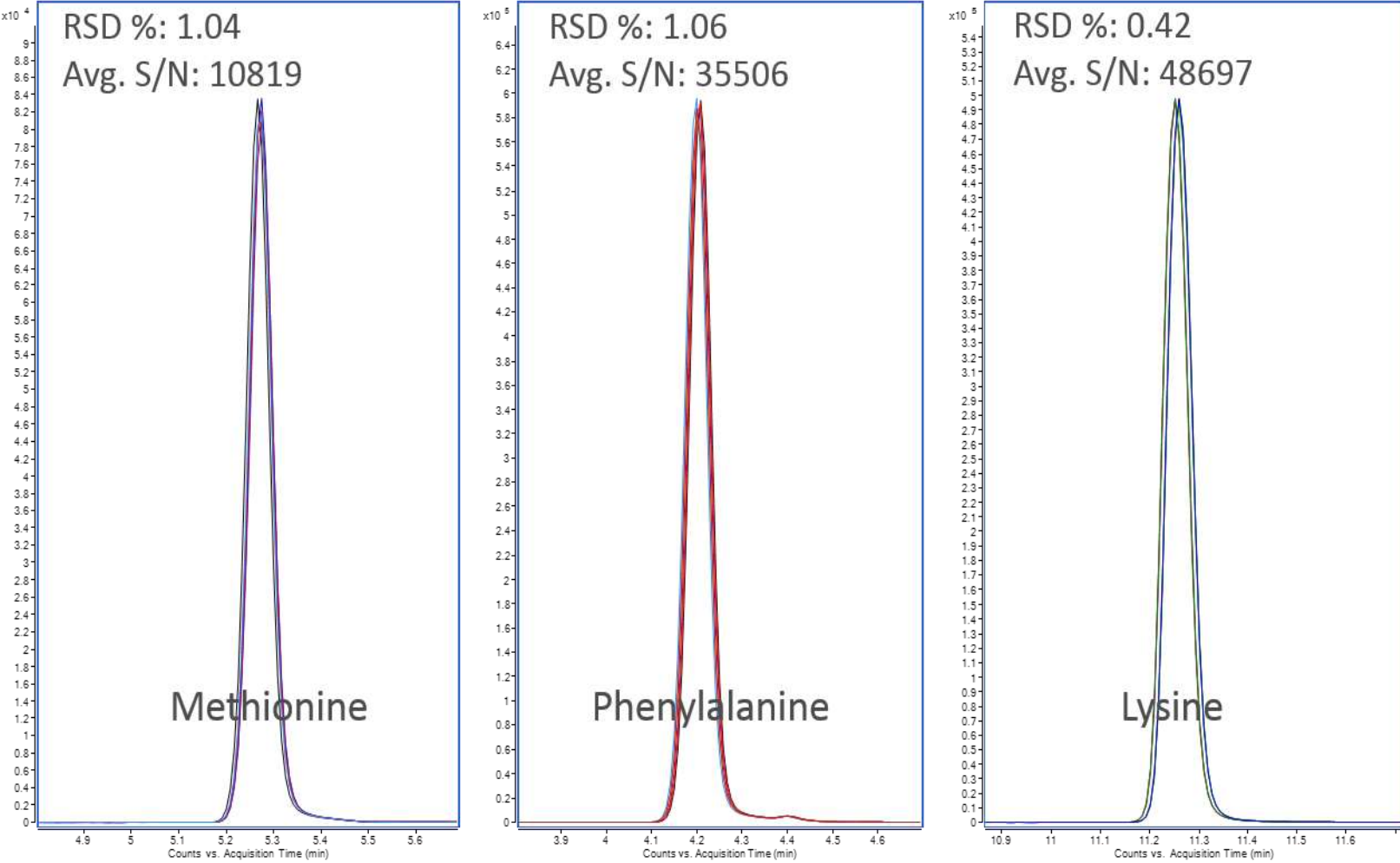


Calibration Curves and LOQs



| Compound Name | Curve Fit | R ² | LOQ (ppb) | S/N at LOQ |
|---------------|-----------|----------------|-----------|------------|
| Alanine | Linear | 0.9997 | 46.33 | 4.35 |
| Arginine | Quadratic | 0.9920 | 8.71 | 10.75 |
| Aspartic Acid | Linear | 0.9971 | 13.31 | 5.76 |
| Cystine | Linear | 0.9969 | 61.04 | 13.51 |
| Glutamic Acid | Quadratic | 0.9998 | 7.65 | 5.37 |
| Glycine | Linear | 0.9986 | 75.07 | 3.51 |
| Histidine | Quadratic | 0.9998 | 7.76 | 15.92 |
| Isoleucine | Linear | 0.9961 | 1.36 | 5.88 |
| Leucine | Linear | 0.9978 | 1.36 | 4.03 |
| Lysine | Quadratic | 0.9934 | 7.60 | 18.94 |
| Methionine | Linear | 0.9977 | 1.55 | 14.57 |
| Phenylalanine | Linear | 0.9974 | 1.72 | 4.82 |
| Proline | Linear | 0.9954 | 1.15 | 6.03 |
| Serine | Quadratic | 0.9987 | 10.51 | 5.43 |
| Threonine | Linear | 0.9995 | 11.91 | 4.35 |
| Tyrosine | Linear | 0.9943 | 1.88 | 3.86 |
| Valine | Linear | 0.9972 | 1.12 | 6.19 |

Reproducibility: Six Replicates in Sample Matrix



New Ultivo Acquisition Feature: Mixed Mode

Agilent MassHunter Workstation Data Acquisition

File View Sample Worklist Method Tools Help

Layout

Instrument

Multi

0.00

Chromatogram

Abundance

35000
30000
25000
20000
15000
10000
5000

Method

Properties

Sample Run

Method

Acquisition Source

Chromatograms

Timetable

Tune

Autotune

Ion source: AJS ESI

Stop time: As pump/No limit Limit (min) 1

Time filter window (min) 0.02

Time Segments

Start time (min) Scan type

0 Mixed

Acquisition Parameters

| Scan type | Compound/Segment name | Precursor (m/z) | MS1 res | Product (m/z) | MS2 start mass (m/z) | MS2 end mass (m/z) | MS2 res | Scan Dwell time (ms) | Frag (V) | ISTD? | CAV (V) | CE (V) | Thre |
|-------------|-----------------------|-----------------|---------|---------------|----------------------|--------------------|---------|----------------------|----------|--------------------------|---------|--------|------|
| Product Ion | | 132.1 | Unit | | 20 | 150 | | 250 | 85 | <input type="checkbox"/> | 9 | 0 | 0 |
| Product Ion | | 132.1 | Unit | | 20 | 150 | | 250 | 85 | <input type="checkbox"/> | 9 | 25 | 0 |
| MRM | Compound1 | 132.1 | Unit | 86 | | | Unit | 20 | 85 | <input type="checkbox"/> | 9 | 10 | |
| MRM | Compound1 | 132.1 | Unit | 68 | | | Unit | 20 | 85 | <input type="checkbox"/> | 9 | 15 | |

Estimated cycle time (ms/cycle) 543

Estimated max scan speed (Da/s) 600

Data storage Profile

able:
nsitions

Agilent MassHunter Workstation Data Acquisition

File View Sample Worklist Method Tools Help

Layout

Instrument

Multi

0.00

Chromatogram

Abundance

35000
30000
25000
20000
15000
10000
5000

Method

Properties

Sample Run

Method

Acquisition Source

Chromatograms

Timetable

Tune

Autotune

Ion source: AJS ESI

Stop time: As pump/No limit Limit (min) 1

Time filter window (min) 0.02

Time Segments

Start time (min) Scan type

0 Mixed

Acquisition Parameters

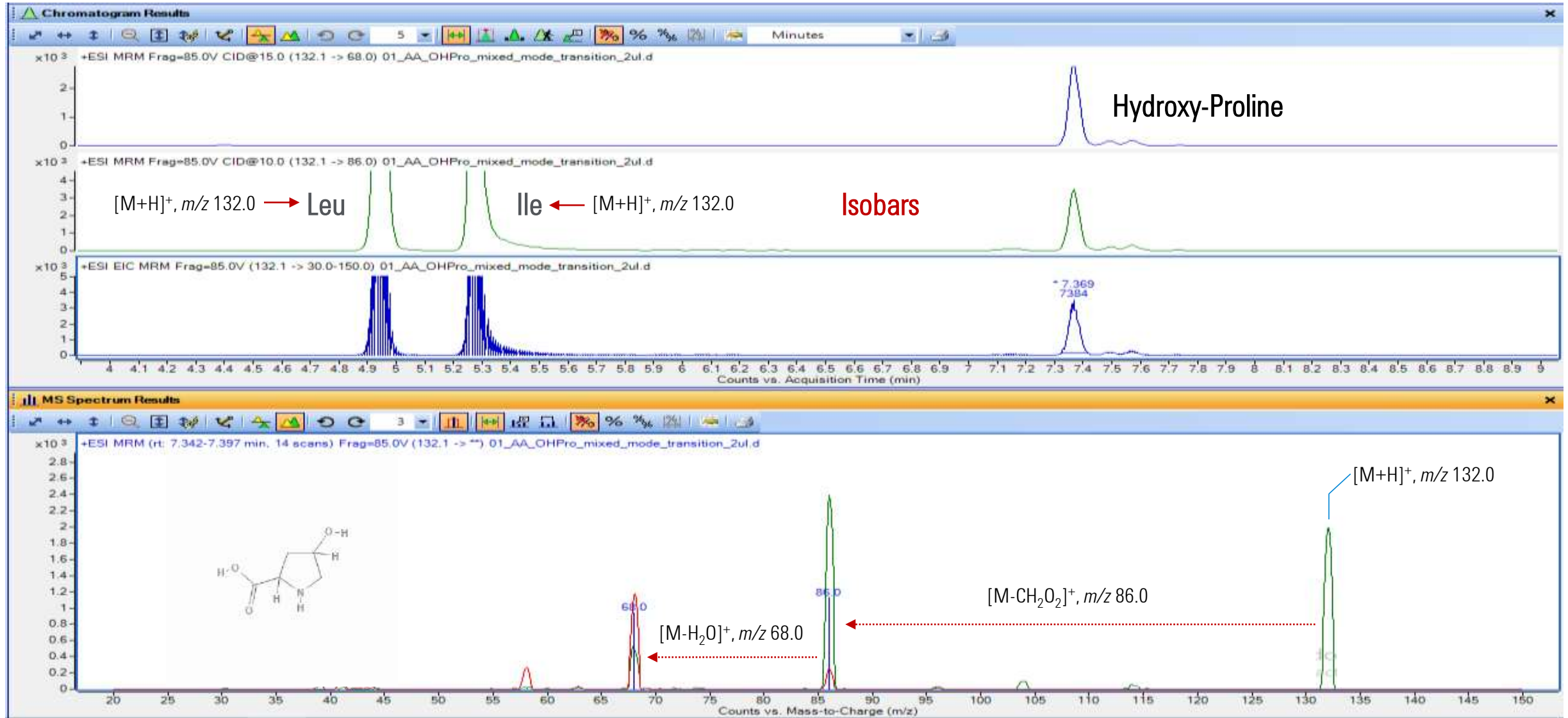
| Scan type | Compound/Segment name | Precursor (m/z) | MS1 res | Product (m/z) | MS2 start mass (m/z) | MS2 end mass (m/z) | MS2 res | Scan Dwell time (ms) | Frag (V) | ISTD? | CAV (V) | CE (V) | Threshold | Polarity |
|-------------|-----------------------|-----------------|---------|---------------|----------------------|--------------------|---------|----------------------|----------|--------------------------|---------|--------|-----------|----------|
| Product Ion | | 132.1 | Unit | | 20 | 150 | | 250 | 85 | <input type="checkbox"/> | 9 | 0 | | Positive |
| Product Ion | | 132.1 | Unit | | 20 | 150 | | 250 | 85 | <input type="checkbox"/> | 9 | 25 | | Positive |
| MRM | Compound1 | 132.1 | Unit | 86 | | | Unit | 20 | 85 | <input type="checkbox"/> | 9 | 10 | | Positive |
| MRM | Compound1 | 132.1 | Unit | 68 | | | Unit | 20 | 85 | <input type="checkbox"/> | 9 | 15 | | Positive |

Estimated cycle time (ms/cycle) 543

Estimated max scan speed (Da/s) 600

Data storage Profile

Mixed-Mode Acquisition: Hydroxyproline MRM and Product Ion Scan



Eight Small Molecule Drugs in Serum

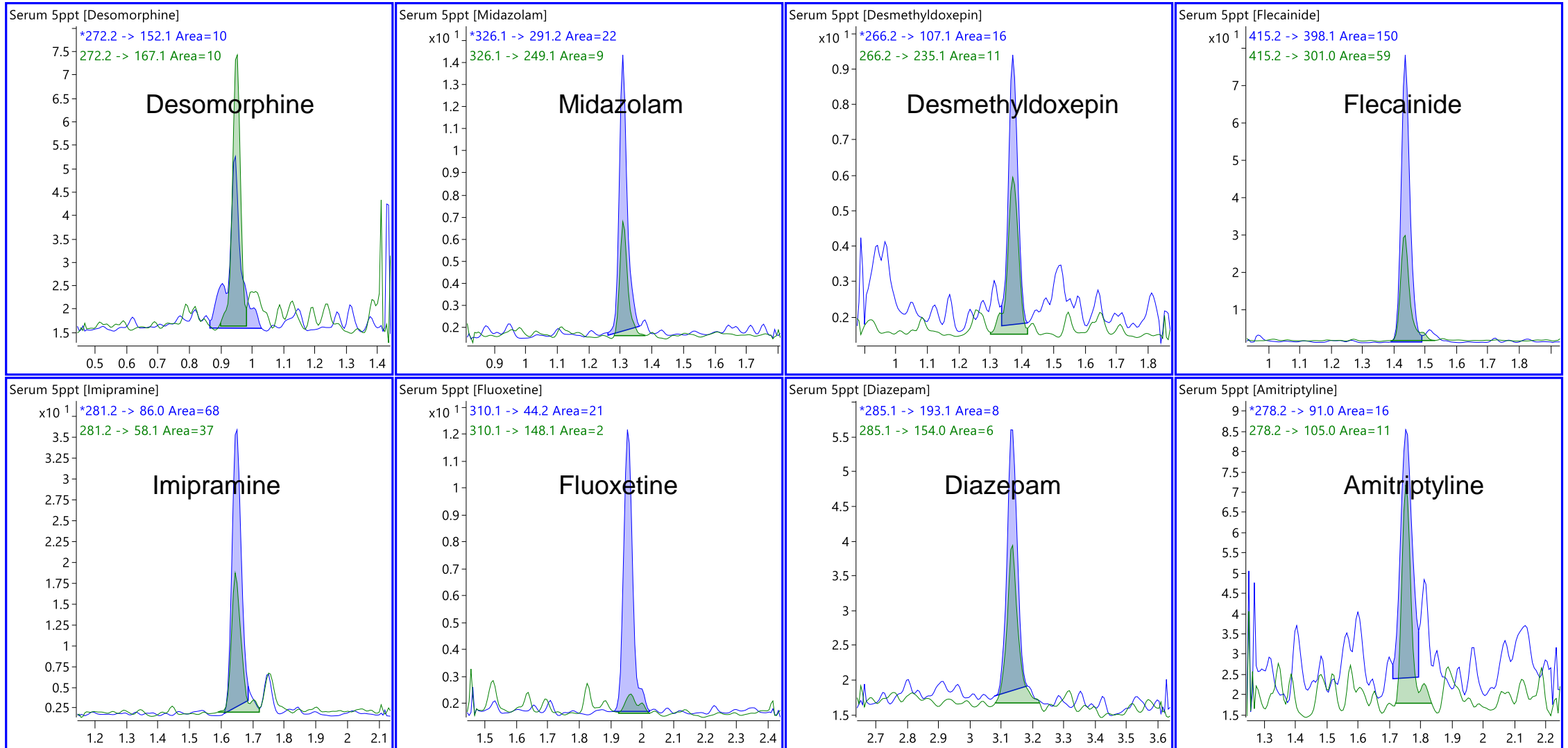


Sample Preparation

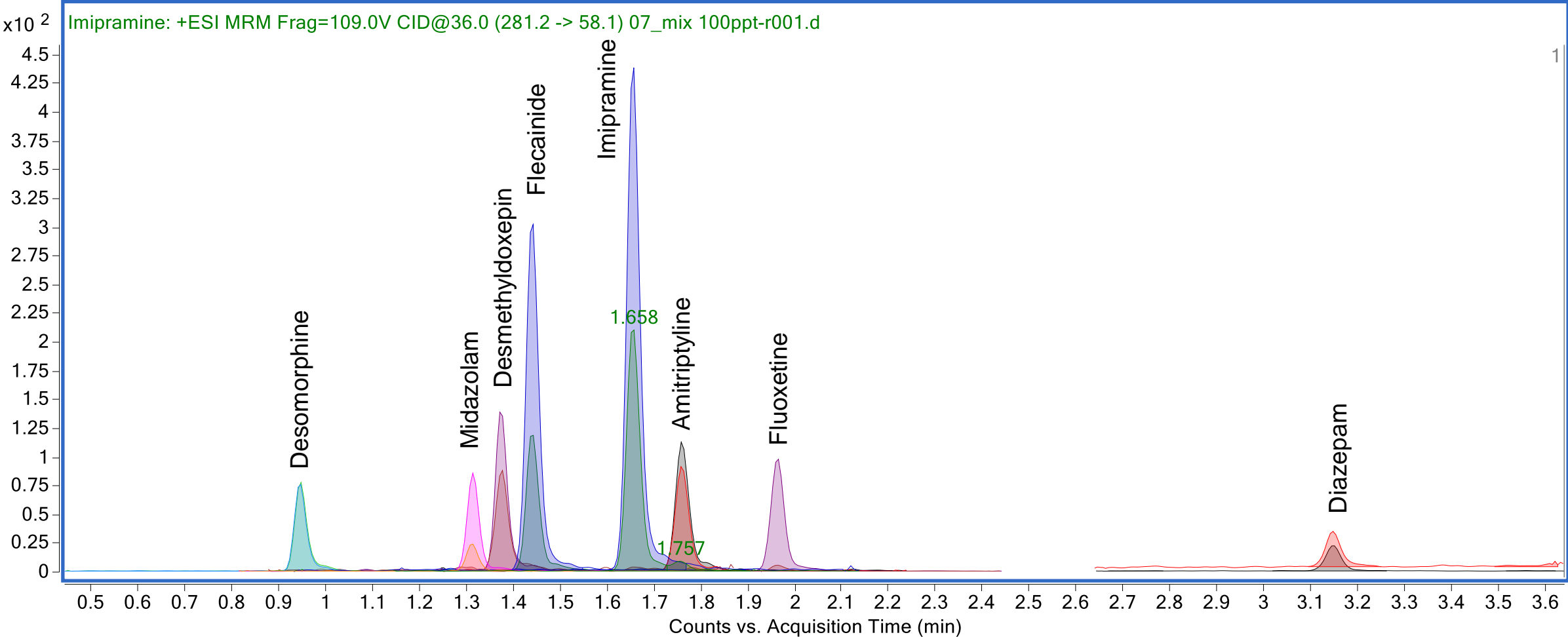
- 250 uL human serum was crashed with 500 uL acetonitrile, vortexed for 1 min and centrifuged for 4 min at 10,000 rpm.
- 500 uL supernatant was transferred and diluted with 500 uL of water.
- The 11-point calibration ranges of all drugs was from 0.001 to 100 ng/mL with 25 ng/mL sufficient internal standard across all sample types injected.

| | | |
|---------------------------|---|-----------|
| Column | Zorbax Eclipse Plus C8, 2.1 x 100 mm, 1.8 µm (p/n 959758-906) | |
| Column temp | 40°C | |
| Injection volume | 2 µL | |
| Mobile phase | A: Water B: Acetonitrile | |
| Flow rate | 0.6 mL/min | |
| Gradient | <u>Time</u> | <u>B%</u> |
| | 0 | 5 |
| | 0.5 | 35 |
| | 2 | 40 |
| | 3 | 50 |
| | 3.5 | 75 |
| | 3.51 | 95 |
| 4 | 95 (stop time) | |
| 5 | (post time) | |
| Drying gas temp | 200 °C | |
| Drying gas flow | 11 L/min | |
| Sheath gas temp | 400 °C | |
| Sheath gas flow | 12 L/min | |
| Nebulizer pressure | 35 psi | |
| Capillary voltage | 2750 V(+) | |
| Nozzle voltage | 0 V(+) | |
| Cycle Time | 350 ms | |

LODs: 5 pg/mL in dilute and shoot serum



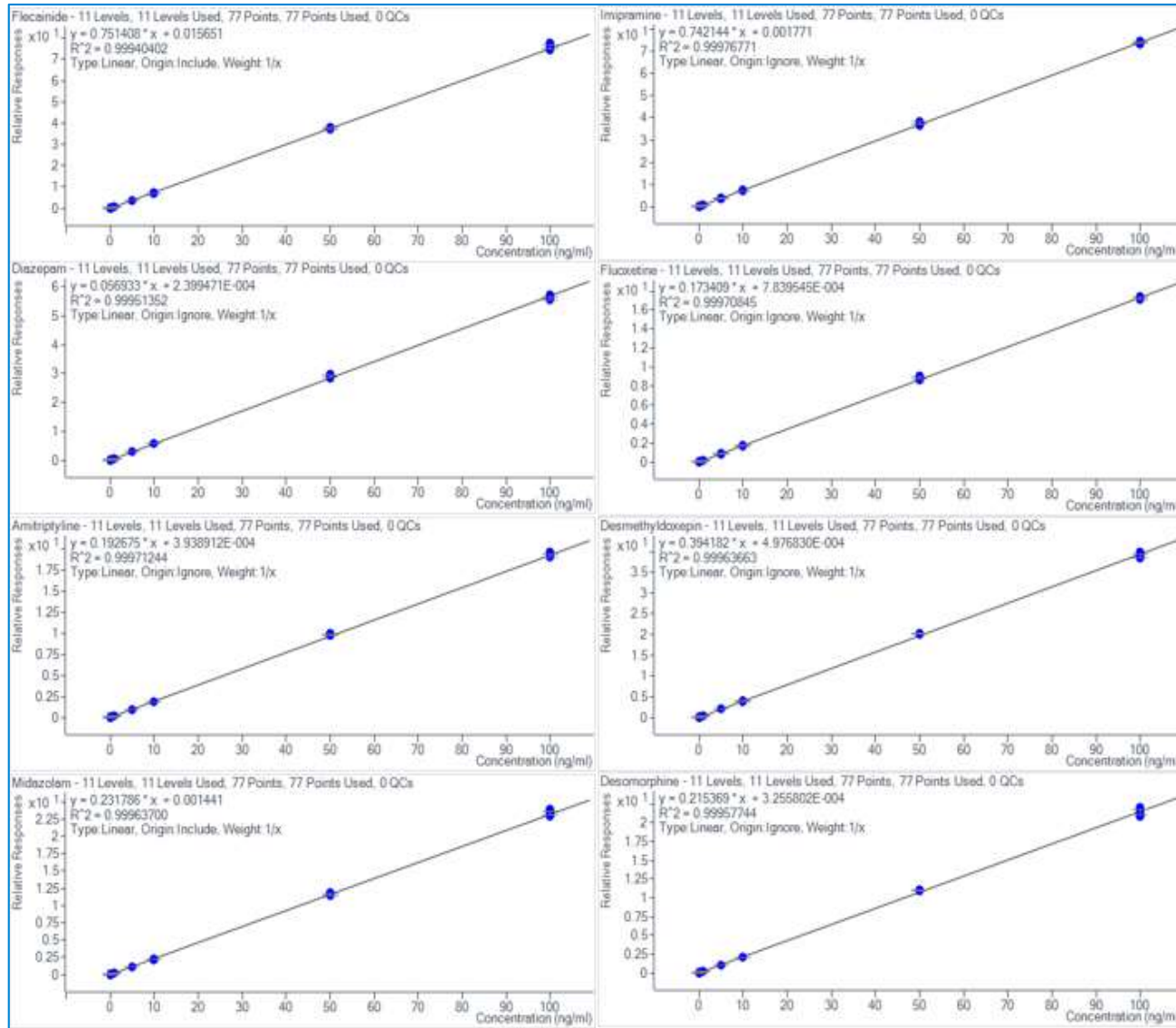
Analytical Analysis of Drugs in Human Serum: 0.1 ng/mL



Accuracy and Reproducibility (n=7)

| Level ng/mL | Desomorphine | | Midazolam | | Desmethyldoxepin | | Flecainide | | Imipramine | | Amitriptyline | | Fluoxetine | | Diazepam | |
|----------------|--------------|------|-----------|-------|------------------|-------|------------|-------|------------|------|---------------|------|------------|------|----------|------|
| | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD | Accuracy | %RSD |
| 0.005 | 101.0 | 9.31 | 102.9 | 44.66 | 87.6 | 14.97 | 85.6 | 6.33 | 83.0 | 3.55 | 78.3 | 7.65 | 100.1 | 8.39 | 129.1 | 9.91 |
| 0.01 | 105.1 | 6.96 | 156.1 | 10.78 | 103.9 | 8.44 | 96.1 | 11.29 | 99.0 | 5.90 | 97.6 | 4.53 | 110.4 | 8.46 | 147.3 | 6.19 |
| 0.05 | 94.4 | 3.66 | 91.6 | 2.69 | 99.5 | 6.12 | 88.0 | 4.17 | 96.7 | 2.52 | 101.9 | 3.03 | 101.1 | 2.21 | 101.5 | 4.59 |
| 0.1 | 96.7 | 3.26 | 93.1 | 3.55 | 99.9 | 3.70 | 93.4 | 3.63 | 98.5 | 1.11 | 98.1 | 4.41 | 99.2 | 2.43 | 103.3 | 2.92 |
| 0.5 | 97.1 | 1.90 | 99.0 | 1.49 | 102.7 | 2.06 | 96.2 | 2.08 | 101.0 | 1.34 | 108.2 | 1.47 | 103.1 | 1.98 | 106.6 | 2.79 |
| 1 | 96.8 | 1.49 | 97.5 | 2.28 | 99.5 | 2.25 | 93.8 | 1.99 | 96.6 | 0.95 | 96.0 | 1.69 | 100.4 | 1.47 | 103.0 | 1.23 |
| 5 | 97.2 | 0.79 | 97.3 | 1.85 | 101.6 | 1.86 | 94.7 | 0.86 | 98.6 | 1.33 | 98.3 | 1.07 | 100.0 | 1.54 | 102.8 | 1.40 |
| 10 | 96.6 | 1.18 | 95.3 | 0.82 | 99.3 | 0.65 | 93.4 | 1.03 | 96.7 | 1.20 | 96.7 | 0.95 | 97.6 | 0.85 | 100.1 | 0.68 |
| 50 | 102.2 | 0.83 | 100.2 | 1.22 | 102.2 | 0.44 | 99.8 | 0.94 | 101.3 | 1.63 | 101.6 | 1.03 | 102.0 | 1.59 | 102.5 | 1.25 |
| 100 | 99.5 | 1.57 | 100.5 | 1.28 | 98.9 | 1.41 | 101.1 | 1.27 | 99.8 | 0.58 | 99.6 | 1.01 | 99.2 | 0.57 | 98.5 | 0.92 |

Calibration Curves and LOQs



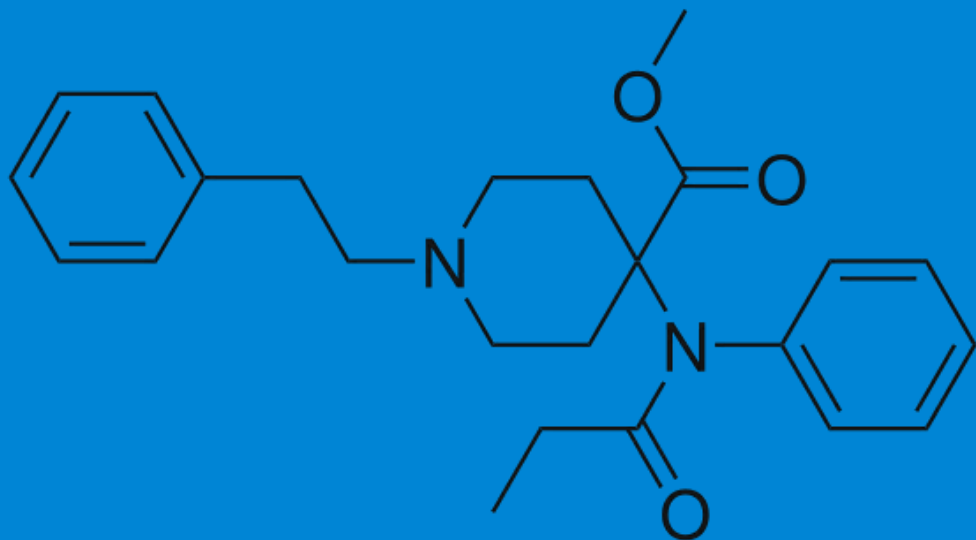
| Compound name | Limits of Quantification | |
|------------------|--------------------------|----------------|
| | Ultivo (fg on column) | Ultivo (pg/mL) |
| Desomorphine | 17.3 | 1.73 |
| Desmethyldoxepin | 27.8 | 2.78 |
| Flecaïnide | 11.7 | 1.17 |
| Midazolam | 40.0 | 4.0 |
| Imipramine | 6.57 | 0.66 |
| Amitriptyline | 14.2 | 1.42 |
| Fluoxetine | 15.6 | 1.56 |
| Diazepam | 18.4 | 1.84 |

Summary

- Agilent Ultivo LC/TQ system demonstrated the accurate and sensitive detection of commonly small molecule monitored amino acids in a pet food matrix, while the use of the Poroshell 120 HILIC-Z column enabled underivatized analytes to be monitored, resulting in a simplified workflow.
- A fast, sensitive, accurate method was presented for the analytical determination of eight drugs in human serum on the new Ultivo TQ LC/MS.
- Ultivo is an exceptionally innovative new mass spectrometer, which can minimize laboratory workspace needs, as well as reduce maintenance challenges, creating a productive work environment for high-throughput laboratories.



Analytical Determination of New Synthetic Fentanyls and Pesticides in Cannabis Using Ultivo LC/TQ



Jennifer Hitchcock, PhD
LCMS Application Scientist
Agilent Technologies, Inc.
Santa Clara, CA



OR the Pete Stone Showcase

Presented by Jennifer Hitchcock

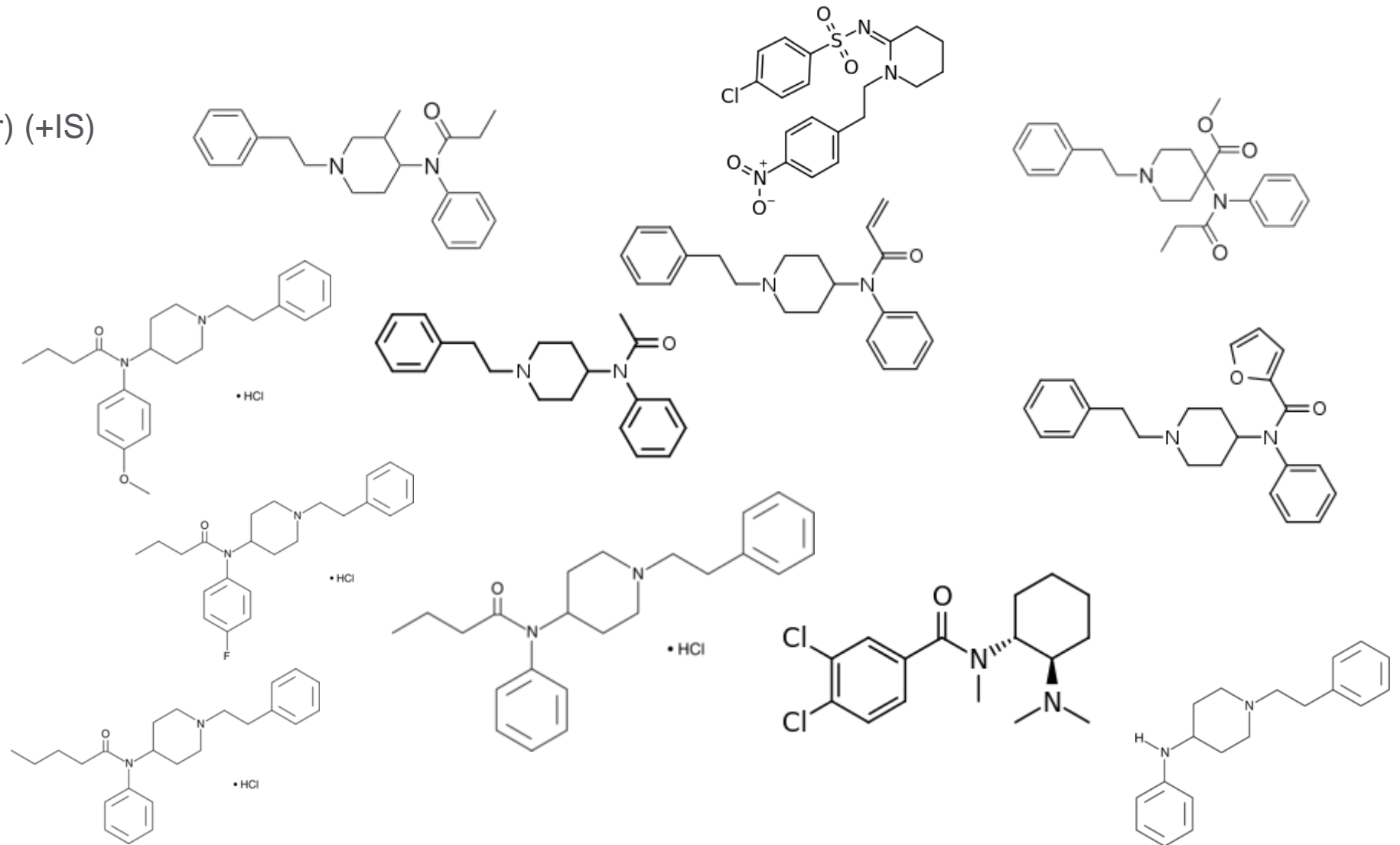


Introduction

- Opioid misuse is severe, with approximately 100 reported overdose fatalities per day across the US
- It's thought that historically abused opiates like heroin are being “cut” with many new synthetic, more potent opioids like fentanyl
- Keeping up with new synthetic illicit drugs and analyzing for them is difficult
- Research method tested on Ultivo for a list of 12 new synthetic fentanyl opioids and 4-ANNP, the DEA-scheduled precursor

Synthetic fentanyls used in research study

- 4-ANNP (synthetic precursor) (+IS)
- 3-methylfentanyl
- Acetylfentanyl (+IS)
- Acetylnorfentanyl (+IS)
- Acrylfentanyl (+IS)
- Butyrylfentanyl (+IS)
- Carfentanyl (+IS)
- Furanylfentanyl
- N-desmethyl U-47700
- Norcarfentanyl
- Para-fluorobutyrylfentanyl
- Valerylfentanyl (+IS)
- U-47700 (+IS)
- **W-18 RM**



Sample preparation

Standards and ISTDs from Cerilliant and Cayman Chemicals
Serum and negative urine from Golden West Biologicals

Urine Matrix:

1. Negative urine diluted 10x using DI water
2. Diluted negative urine centrifuged at 10,000 rpm through nanosep 3K filters, then supernatant combined as stock solution
3. ISTDs spiked into diluted negative urine stock at concentration of 25 ng/mL
4. Serial dilutions made for calibrators from 500 ng/mL to 1 pg/mL using ISTD-spiked filtered stock (equivalence dilution 1/10)

Serum Matrix:

1. Gold serum (10 mL) mixed with 20 mL cold acetonitrile
2. Serum/ACN mix vortexed for 2 min, then centrifuged at 5000 rpm for 15 min
3. Supernatant (20 mL) spiked with ISTD at concentration of 50 ng/mL
4. Supernatant and ISTDs diluted with 20 mL DI water (ISTD conc. 25 ng/mL)
5. Serial dilutions made for calibrators from 500 ng/mL to 1 pg/mL (equivalence dilution 1/6)

Synthetic fentanyl in serum and urine



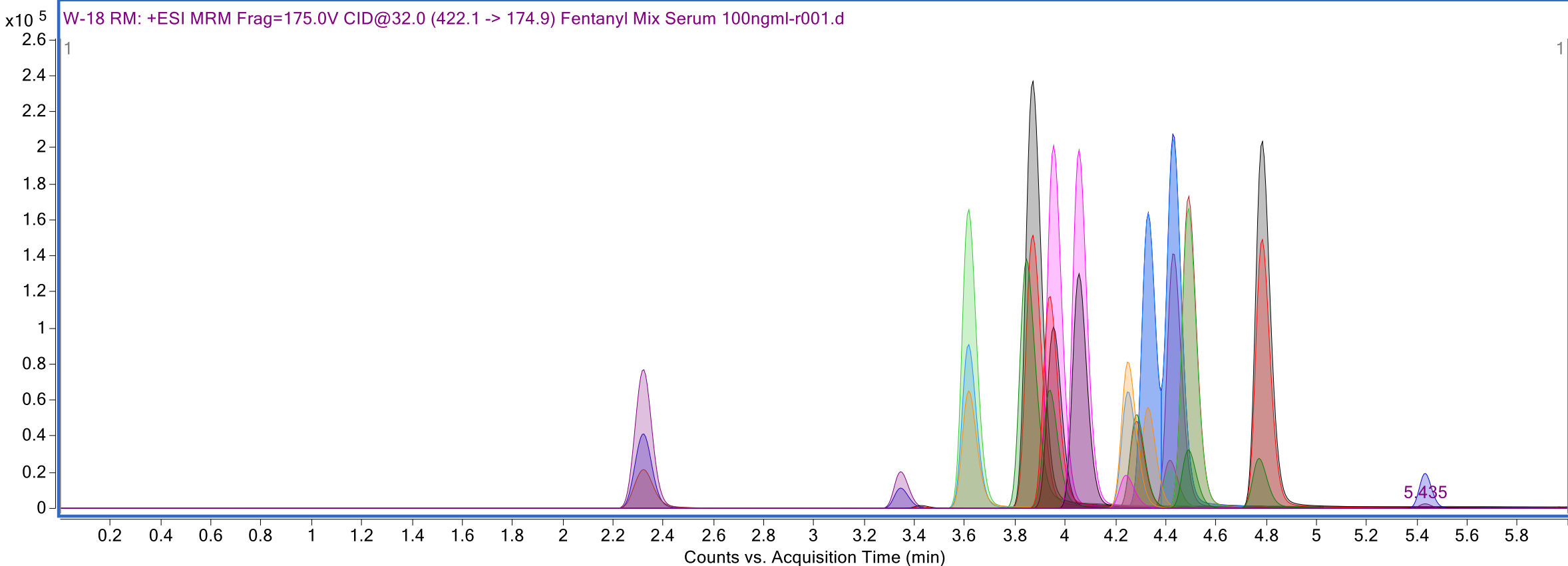
| Column | Poroshell 120 EC-C18, 2.7 μ m, 2.1x50 mm (pn 699775-902) | | | | | | | | | | | | | | | |
|---------------------------|--|-------------|-----------|------|----|------|----|------|----|------|----|------|----------------|------|----------------|--|
| Column temp | 55°C | | | | | | | | | | | | | | | |
| Injection volume | 5 μ L | | | | | | | | | | | | | | | |
| Mobile phase | <u>A</u> : 5mM ammonium formate + 0.01% formic acid in water <u>B</u> : 0.01% formic acid in methanol | | | | | | | | | | | | | | | |
| Flow rate | 0.5 mL/min | | | | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th><u>Time</u></th> <th><u>B%</u></th> </tr> </thead> <tbody> <tr> <td>0.00</td> <td>10</td> </tr> <tr> <td>0.50</td> <td>15</td> </tr> <tr> <td>3.50</td> <td>50</td> </tr> <tr> <td>5.50</td> <td>95</td> </tr> <tr> <td>6.00</td> <td>95 (stop time)</td> </tr> <tr> <td>7.00</td> <td>10 (post time)</td> </tr> </tbody> </table> | <u>Time</u> | <u>B%</u> | 0.00 | 10 | 0.50 | 15 | 3.50 | 50 | 5.50 | 95 | 6.00 | 95 (stop time) | 7.00 | 10 (post time) | |
| <u>Time</u> | <u>B%</u> | | | | | | | | | | | | | | | |
| 0.00 | 10 | | | | | | | | | | | | | | | |
| 0.50 | 15 | | | | | | | | | | | | | | | |
| 3.50 | 50 | | | | | | | | | | | | | | | |
| 5.50 | 95 | | | | | | | | | | | | | | | |
| 6.00 | 95 (stop time) | | | | | | | | | | | | | | | |
| 7.00 | 10 (post time) | | | | | | | | | | | | | | | |
| Drying gas temp | 325 °C | | | | | | | | | | | | | | | |
| Drying gas flow | 9 L/min | | | | | | | | | | | | | | | |
| Sheath gas temp | 350 °C | | | | | | | | | | | | | | | |
| Sheath gas flow | 11 L/min | | | | | | | | | | | | | | | |
| Nebulizer pressure | 35 psi | | | | | | | | | | | | | | | |
| Capillary voltage | 3500 V(+) | | | | | | | | | | | | | | | |
| Nozzle voltage | 0 V(+) | | | | | | | | | | | | | | | |
| Cycle Time | 400 ms | | | | | | | | | | | | | | | |

MRM transition list

| Compound Name | Precursor (m/z) | Product (m/z) | Dwell Time (ms) | Frag (V) | CE (V) |
|-------------------------|-----------------|---------------|-----------------|----------|--------|
| W-18 RM | 422.1 | 174.9 | 10 | 175 | 32 |
| W-18 RM | 422.1 | 110.9 | 10 | 175 | 56 |
| Furanyl fentanyl | 375.2 | 188.1 | 10 | 170 | 24 |
| Furanyl fentanyl | 375.2 | 105 | 10 | 170 | 48 |
| Valeryl fentanyl-D5 | 370.3 | 105 | 10 | 180 | 48 |
| p-fluorobutyrylfentanyl | 369.2 | 188.1 | 10 | 180 | 24 |
| p-fluorobutyrylfentanyl | 369.2 | 105 | 10 | 180 | 52 |
| Valeryl fentanyl | 365.3 | 188.1 | 10 | 180 | 24 |
| Valeryl fentanyl | 365.3 | 105 | 10 | 180 | 48 |
| Butyrylfentanyl-D5 | 356.3 | 188.1 | 10 | 180 | 24 |
| Butyrylfentanyl-D5 | 356.3 | 105.1 | 10 | 180 | 48 |
| Cis-3-methyl fentanyl | 351.2 | 202.1 | 10 | 180 | 24 |
| Butyrylfentanyl | 351.2 | 188.1 | 10 | 180 | 24 |
| Butyrylfentanyl | 351.2 | 105 | 10 | 180 | 48 |
| Cis-3-methyl fentanyl | 351.2 | 105 | 10 | 180 | 48 |
| Acrylfentanyl-D5 | 340.2 | 188.1 | 10 | 165 | 24 |
| Acrylfentanyl-D5 | 340.2 | 105 | 10 | 165 | 44 |
| U-47700-D6 | 335.2 | 284 | 10 | 120 | 16 |
| Acrylfentanyl | 335.2 | 188.1 | 10 | 165 | 24 |
| Acrylfentanyl | 335.2 | 105 | 10 | 165 | 44 |

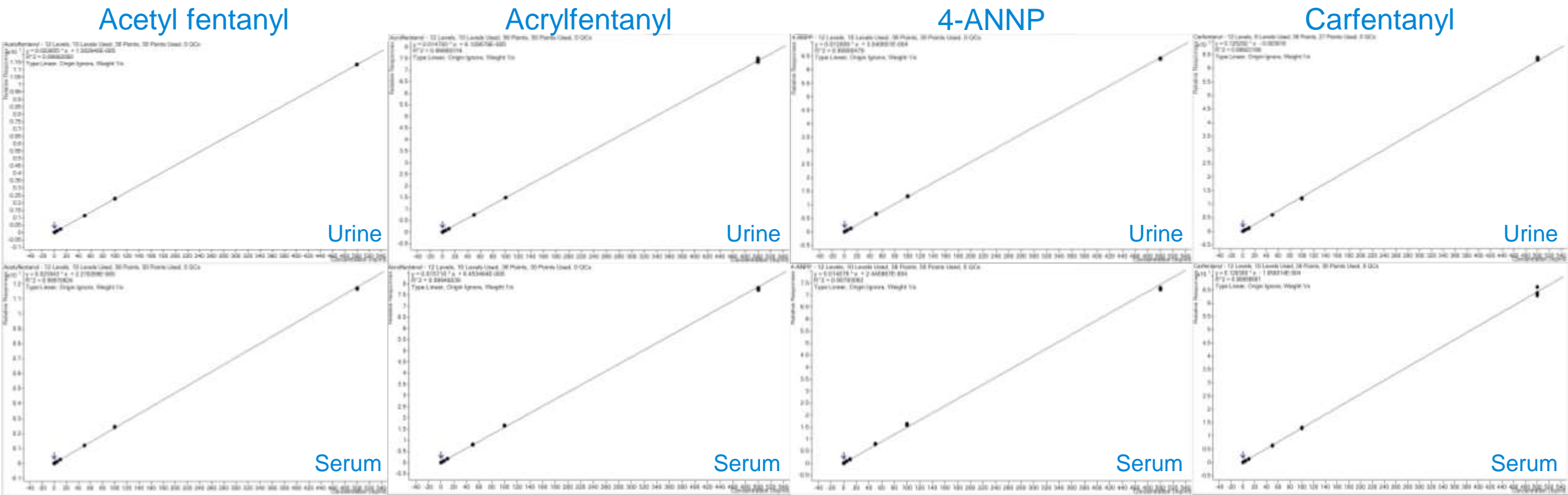
| Compound Name | Precursor (m/z) | Product (m/z) | Dwell Time (ms) | Frag (V) | CE (V) |
|-------------------------|-----------------|---------------|-----------------|----------|--------|
| Acetylfentanyl-13C6 | 329.2 | 105 | 10 | 170 | 44 |
| U-47700 | 329.1 | 284 | 10 | 120 | 16 |
| U-47700 | 329.1 | 172.9 | 10 | 120 | 36 |
| Acetyl fentanyl | 323.2 | 188.1 | 10 | 170 | 24 |
| Acetyl fentanyl | 323.2 | 105.1 | 10 | 170 | 44 |
| N-desmethyl U-47700 | 315.1 | 284 | 10 | 120 | 16 |
| N-desmethyl U-47700 | 315.1 | 172.9 | 10 | 120 | 36 |
| Norcarfentanyl | 291.2 | 231.1 | 10 | 95 | 12 |
| Norcarfentanyl | 291.2 | 142 | 10 | 95 | 16 |
| 4-ANNP-D5 | 286.2 | 105.1 | 10 | 140 | 36 |
| 4-ANNP | 281.2 | 188.1 | 10 | 140 | 16 |
| 4-ANNP | 281.2 | 105 | 10 | 140 | 36 |
| Acetyl norfentanyl-13C6 | 225.2 | 84.1 | 10 | 120 | 20 |
| Acetyl norfentanyl | 219.1 | 84.1 | 10 | 120 | 20 |
| Acetyl norfentanyl | 219.1 | 55.2 | 10 | 120 | 44 |
| Carfentanyl-D5 | 400.3 | 113 | 10 | 130 | 36 |
| Carfentanyl | 395.2 | 335.2 | 10 | 135 | 16 |
| Carfentanyl | 395.2 | 113 | 10 | 135 | 36 |

Composite EIC of each fentanyl opioid transition (Serum mix calibrant 100 ng/mL)



Selected calibration curves

1pg/mL → 500ng/mL



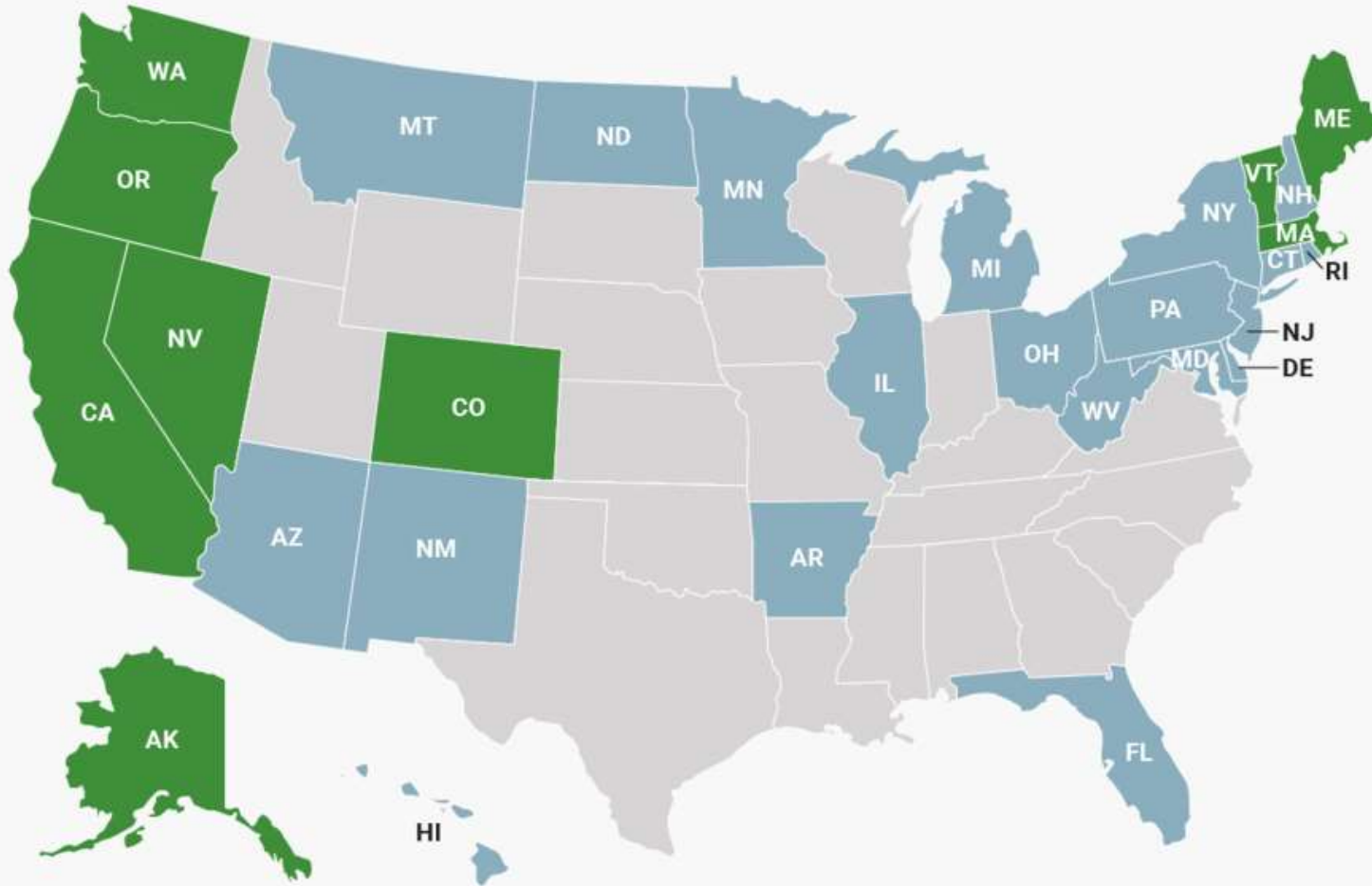
LLOQ results table for synthetic fentanyls

| Analyte | Actual LLOQ (Urine) pg/ml | Actual LLOQ (Serum) pg/ml | Equivalence LLOQ (Urine) pg/ml (1/10 dilution) | Equivalence LLOQ (Serum) pg/ml (1/6 dilution) |
|----------------------------|---------------------------|---------------------------|--|---|
| 4-ANPP | 100 | 10 | 1000 | 60 |
| 3-methylfentanyl | 100 | 50 | 1000 | 300 |
| Acetylfentanyl | 50 | 10 | 500 | 60 |
| Acetylnorfentanyl | 100 | 50 | 1000 | 300 |
| Acrylfentanyl | 50 | 10 | 500 | 60 |
| Butyrylfentanyl | 100 | 50 | 1000 | 300 |
| Carfentanyl | 50 | 10 | 500 | 60 |
| Furanylfentanyl | 50 | 50 | 500 | 300 |
| Para-fluorobutyrylfentanyl | 50 | 50 | 500 | 300 |
| Norcarfentanyl | 100 | 50 | 1000 | 300 |
| N-desmethyl U-47700 | 100 | 50 | 1000 | 300 |
| Valerylfentanyl | 100 | 50 | 1000 | 300 |
| U-47700 | 100 | 50 | 1000 | 300 |
| W-18 RM | 50 | 50 | 500 | 300 |

*Equivalence refers to the actual concentration of the urine or serum prior to dilution

States where marijuana is legal

■ Legalized recreational and medical marijuana ■ Legalized medical marijuana



BUSINESS INSIDER

Cannabis pesticide testing

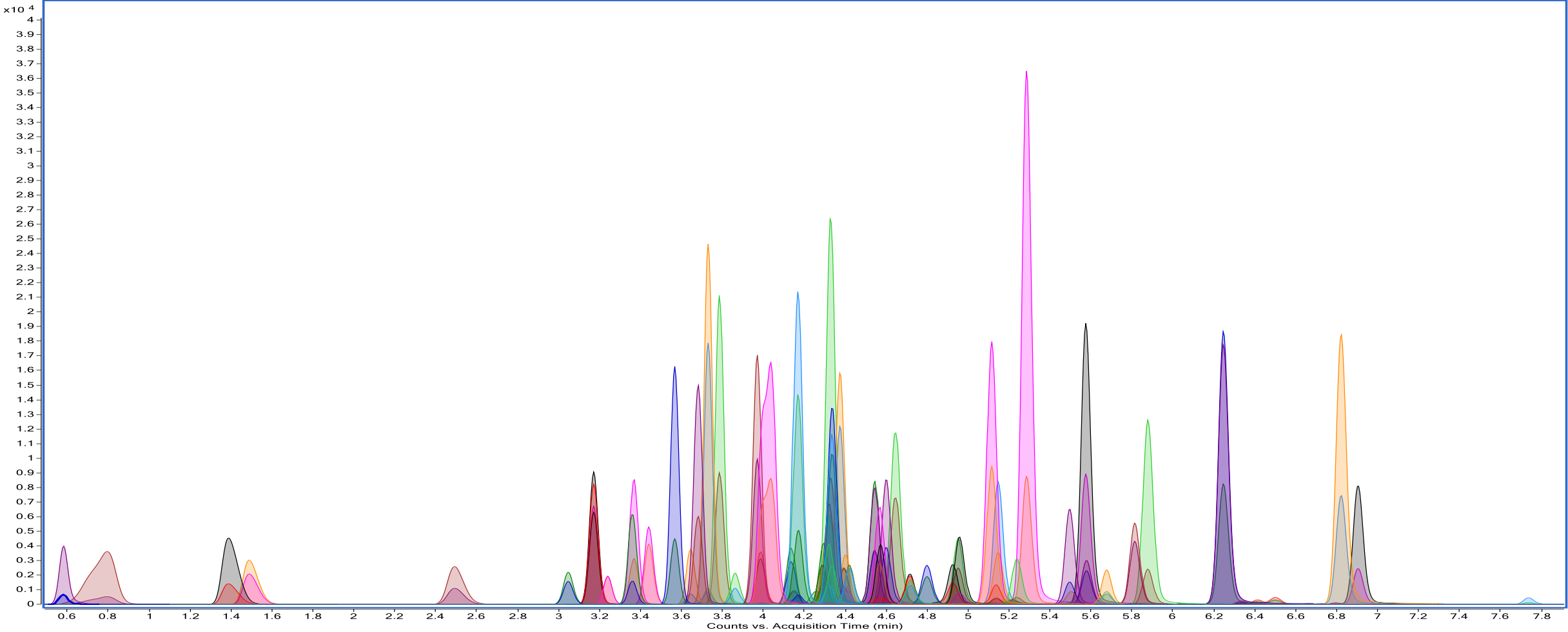
- Cannabis is a complex botanical with many endogenous compounds that can make analytical detection of trace amounts of pesticides difficult
- Each state determines which pesticides to analyze and what appropriate action limits are
- Primary focus has been on California list as most comprehensive in US for cannabis pesticide testing
 - 66 total pesticides
 - Category I (detected or not) has 21
 - Category II (specific action levels for different matrices) has 45
- Canada has proposed regulations containing an additional 33 compounds when compared to the California panel
- Levels are best guesses at this point and may change as more information is acquired

Pesticides in cannabis

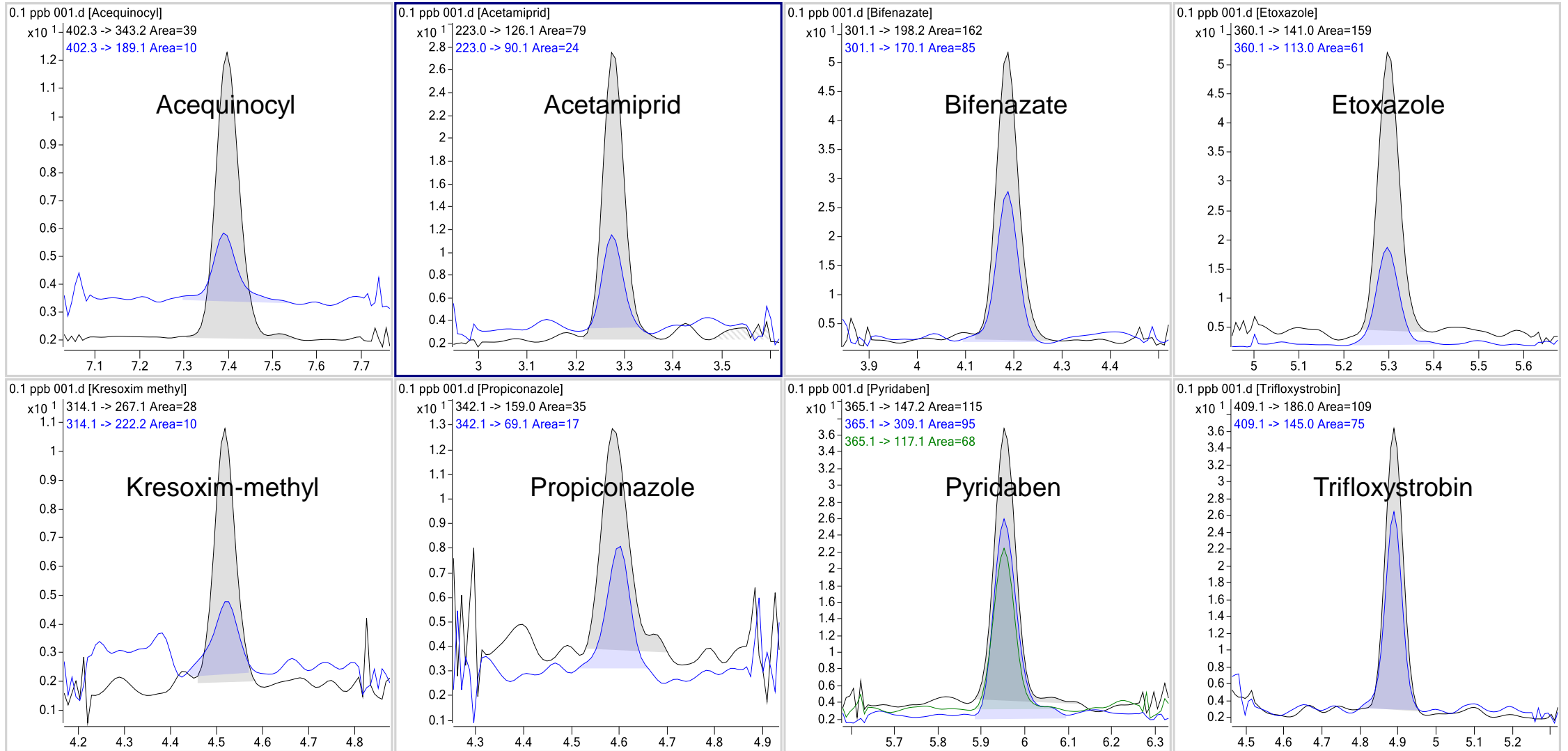


| Column | Poroshell 120 Phenyl-Hexyl 2.7 μ m, 3.0x100 mm (pn 695975-312) | | | | | | | | | | | | | | | | | | | |
|---------------------------|---|------|----|------|----|------|----|------|----|------|----|------|-----|------|-----|------|----|-------|--------------|--|
| Column temp | 55°C | | | | | | | | | | | | | | | | | | | |
| Injection volume | 2 μ L | | | | | | | | | | | | | | | | | | | |
| Mobile phase | A: 5 mM ammonium formate + 0.1% formic acid in water B: 0.1% formic acid in methanol | | | | | | | | | | | | | | | | | | | |
| Flow rate | 0.50 mL/min | | | | | | | | | | | | | | | | | | | |
| Gradient | <table border="1"> <thead> <tr> <th>Time</th> <th>B%</th> </tr> </thead> <tbody> <tr><td>0.00</td><td>20</td></tr> <tr><td>1.00</td><td>20</td></tr> <tr><td>2.00</td><td>75</td></tr> <tr><td>8.00</td><td>96</td></tr> <tr><td>9.00</td><td>100</td></tr> <tr><td>9.50</td><td>100</td></tr> <tr><td>9.51</td><td>20</td></tr> <tr><td>11.00</td><td>20 stop time</td></tr> </tbody> </table> | Time | B% | 0.00 | 20 | 1.00 | 20 | 2.00 | 75 | 8.00 | 96 | 9.00 | 100 | 9.50 | 100 | 9.51 | 20 | 11.00 | 20 stop time | |
| Time | B% | | | | | | | | | | | | | | | | | | | |
| 0.00 | 20 | | | | | | | | | | | | | | | | | | | |
| 1.00 | 20 | | | | | | | | | | | | | | | | | | | |
| 2.00 | 75 | | | | | | | | | | | | | | | | | | | |
| 8.00 | 96 | | | | | | | | | | | | | | | | | | | |
| 9.00 | 100 | | | | | | | | | | | | | | | | | | | |
| 9.50 | 100 | | | | | | | | | | | | | | | | | | | |
| 9.51 | 20 | | | | | | | | | | | | | | | | | | | |
| 11.00 | 20 stop time | | | | | | | | | | | | | | | | | | | |
| Drying gas temp | 200 °C | | | | | | | | | | | | | | | | | | | |
| Drying gas flow | 13 L/min | | | | | | | | | | | | | | | | | | | |
| Sheath gas temp | 200 °C | | | | | | | | | | | | | | | | | | | |
| Sheath gas flow | 10 L/min | | | | | | | | | | | | | | | | | | | |
| Nebulizer pressure | 55 psi | | | | | | | | | | | | | | | | | | | |
| Capillary voltage | 5500 V (+) / 3500 V (-) | | | | | | | | | | | | | | | | | | | |
| Nozzle voltage | 2000 V (+) / 500 V (-) | | | | | | | | | | | | | | | | | | | |
| Cycle Time | 500 ms | | | | | | | | | | | | | | | | | | | |

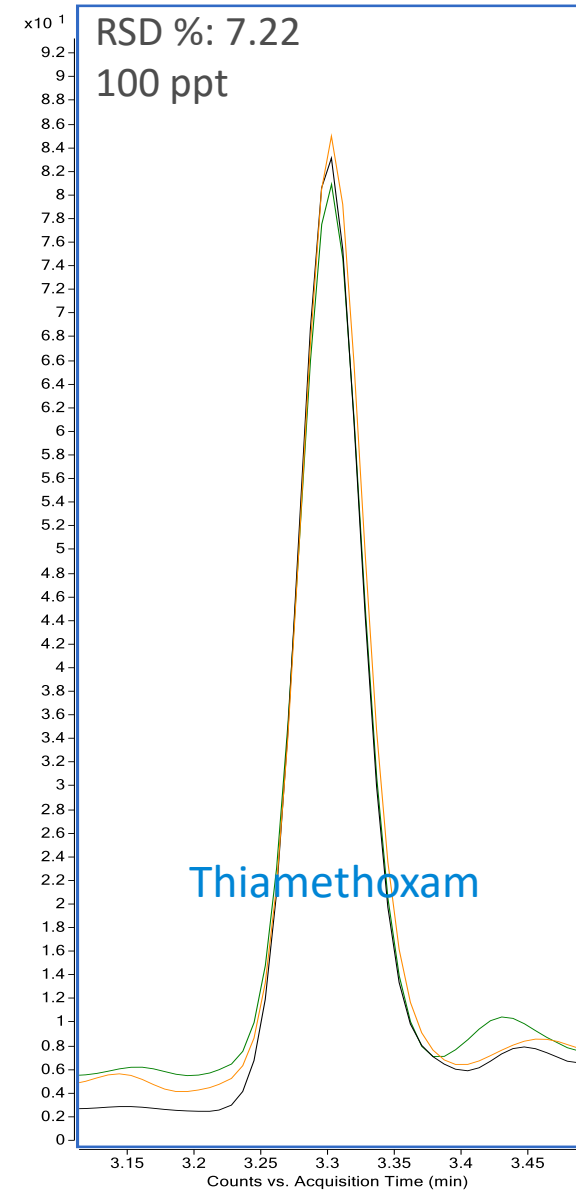
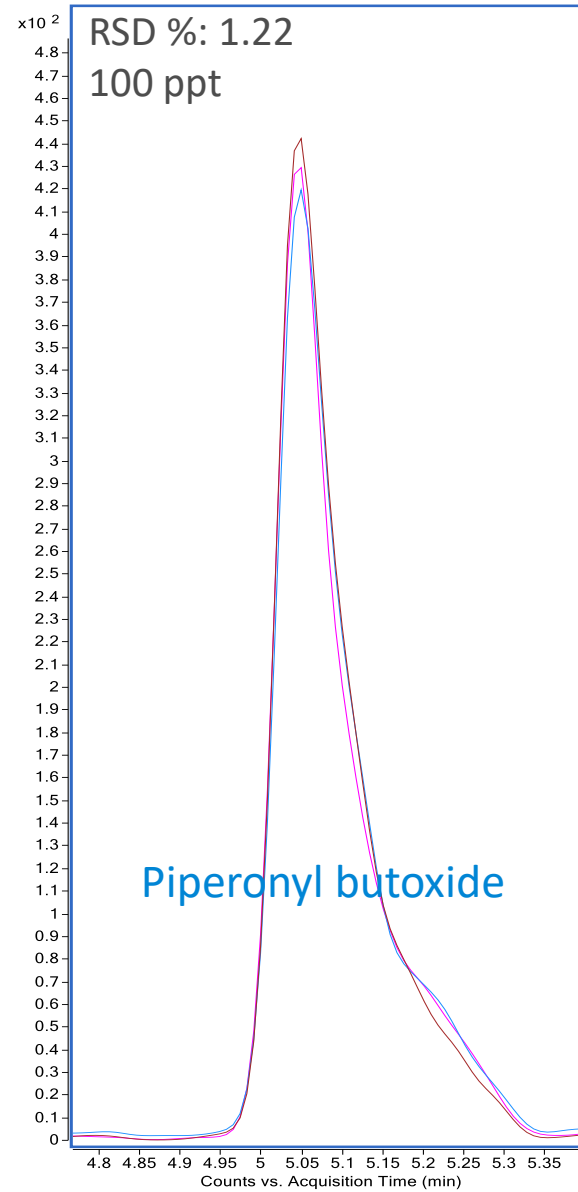
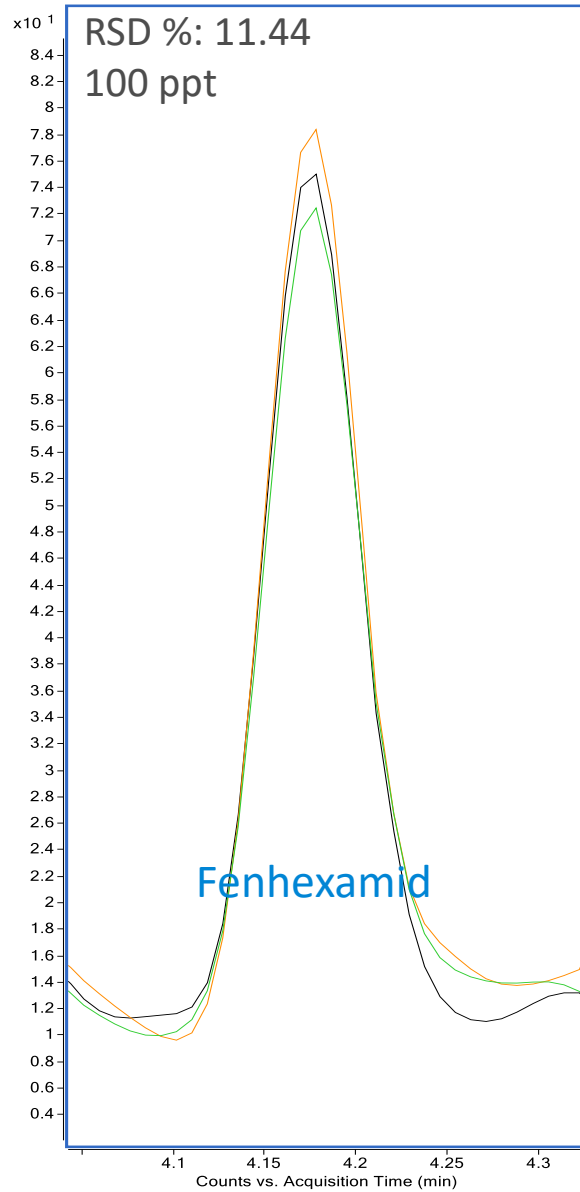
Composite EIC of pesticides and aflatoxins: 10 ppb



Example chromatograms at 0.1 ppb

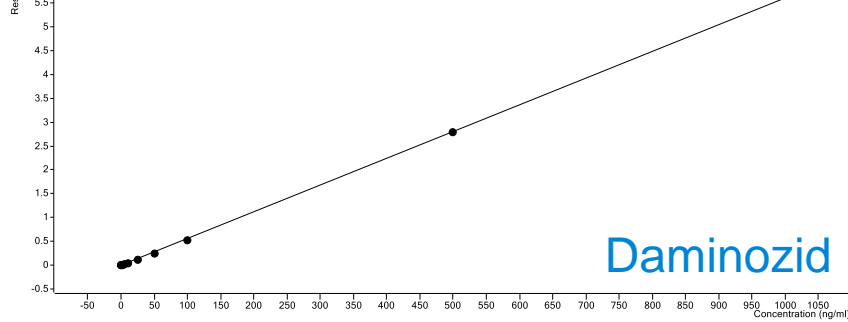


Reproducibility in matrix (n=3)

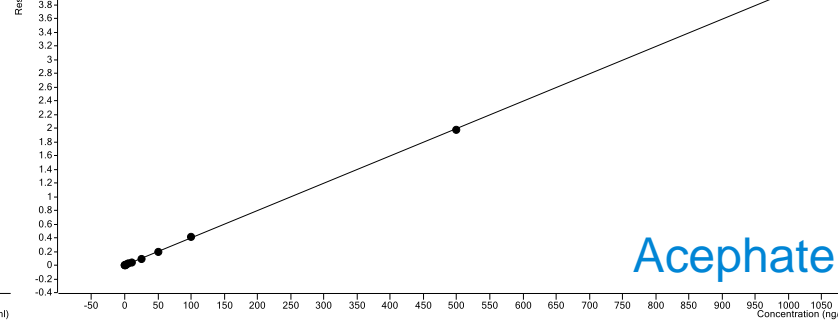


Representative calibration curves

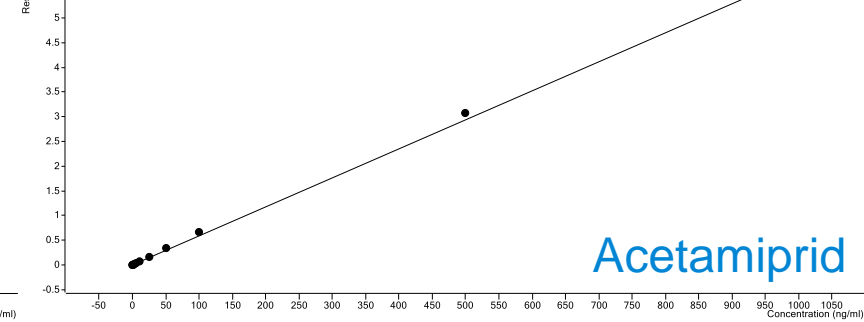
Daminozid - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 56.032506 \cdot x + 6.920863$
 $R^2 = 0.99678174$
 Type:Linear, Origin:Ignore, Weight:1/x



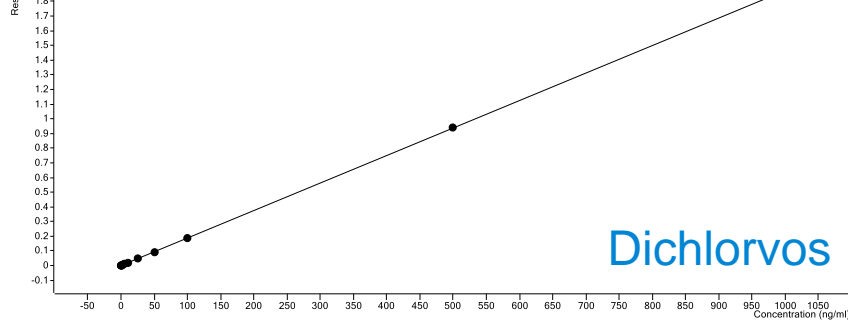
Acephate - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 399.321334 \cdot x + 1.407795$
 $R^2 = 0.99991243$
 Type:Linear, Origin:Ignore, Weight:1/x



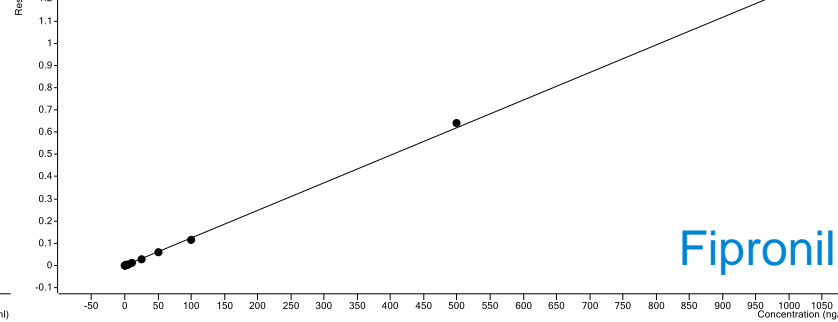
Acetamidrid - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 587.411345 \cdot x + 76.613032$
 $R^2 = 0.99530712$
 Type:Linear, Origin:Ignore, Weight:1/x



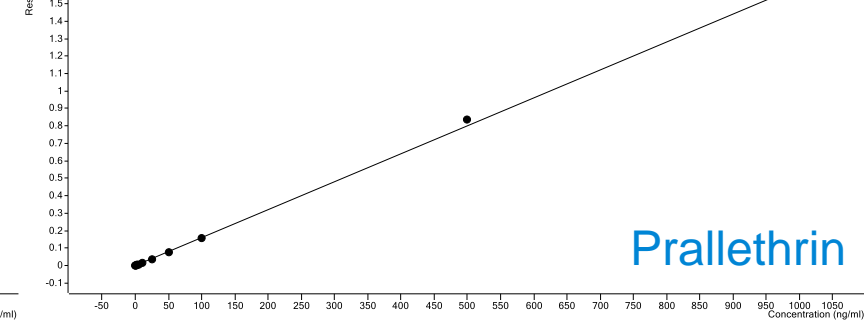
Dichlorvos - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 187.392745 \cdot x + 2.980355$
 $R^2 = 0.99694725$
 Type:Linear, Origin:Ignore, Weight:1/x



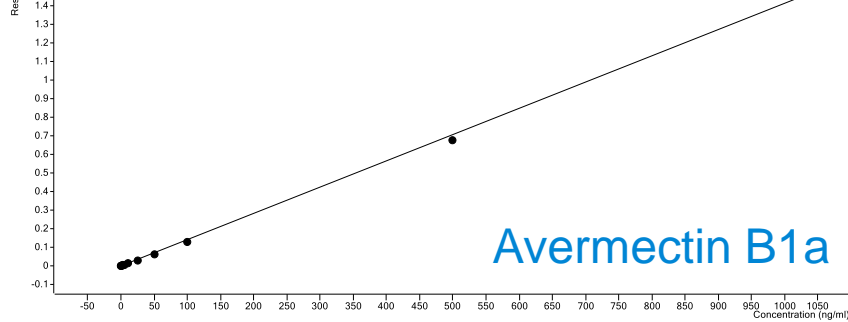
Fipronil - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 124.025333 \cdot x - 11.345027$
 $R^2 = 0.99918240$
 Type:Linear, Origin:Ignore, Weight:1/x



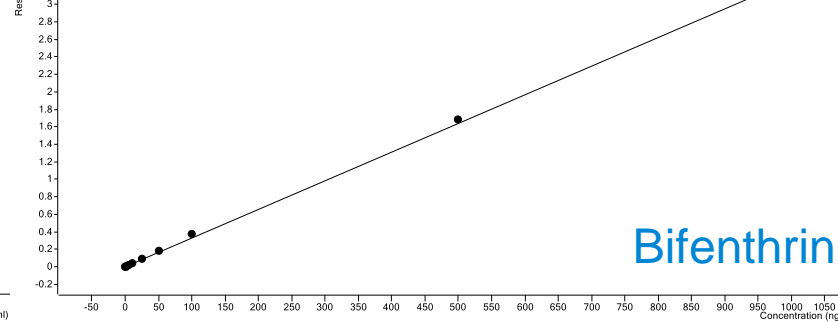
Prallethrin - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 100.100666 \cdot x + 0.631968$
 $R^2 = 0.99913856$
 Type:Linear, Origin:Ignore, Weight:1/x



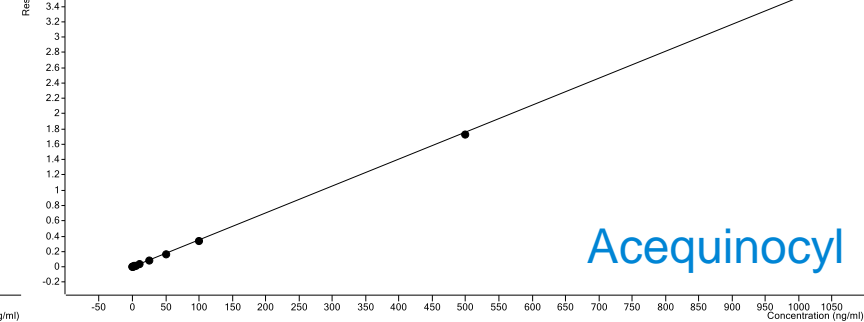
Avermectin B1a - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 14.138533 \cdot x - 0.322910$
 $R^2 = 0.99698622$
 Type:Linear, Origin:Ignore, Weight:1/x



Bifenthrin - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 327.684493 \cdot x + 39.111115$
 $R^2 = 0.99653197$
 Type:Linear, Origin:Ignore, Weight:1/x



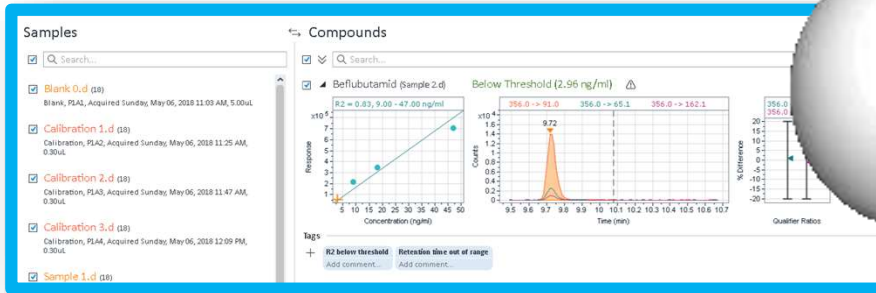
Acequinocyl - 11 Levels, 11 Levels Used, 11 Points, 11 Points Used, 0 QCs
 $y = 351.870571 \cdot x - 9.201364$
 $R^2 = 0.99909879$
 Type:Linear, Origin:Ignore, Weight:1/x



Summary

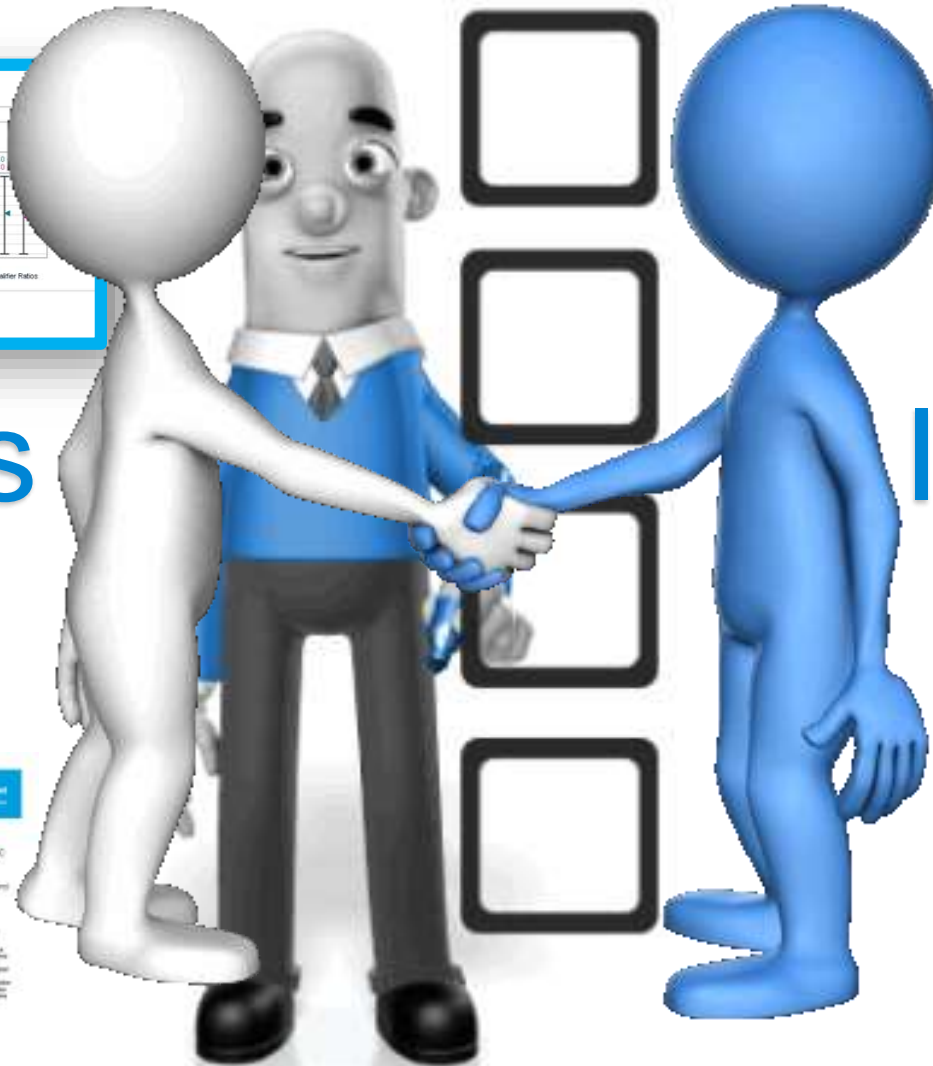
- Excellent sensitivity has been demonstrated in a variety of matrices and applications, including synthetic fentanyl in biological fluids and pesticides in cannabis
- In these applications, results on Ultivo are similar to those determined on the 6470
- Ultivo innovates the mass spec lab by minimizing bench space requirements and reducing maintenance challenges

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A collage of four Agilent application notes. The first is titled 'Robustness of the Agilent Ultivo Triple Quadrupole I Analysis in Food SA'. The second is 'Multiclass Residue Analysis of Veterinary Drugs in Pork and Hen Eggs Using an Agilent Ultivo Triple Quadrupole LC/MS/MS'. The third is 'Sensitive Detection of Trace Organic Contaminants in Water Using Agilent 1090 Infinity Plus/1260 Infinity HPLC and Agilent Ultivo Triple Quadrupole LC/MS/MS'. Each note includes a title, introduction, and application information.



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