



GC-MS Evolutionary Technology & the Art of POPs Analysis

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Persistent Organic Pollutants

Organochlorine pesticides



Aldrin, Chlordane, DDT, Dieldrin, Endrin, Heptachlor, Hexachlorobenzene, Mirex and Toxaphene

Polychlorinated biphenyls

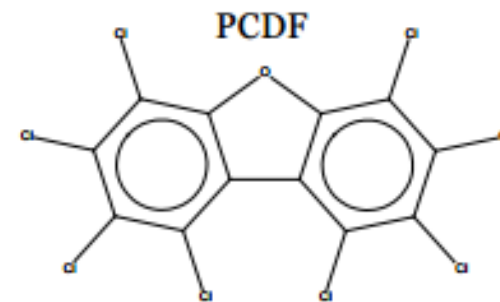
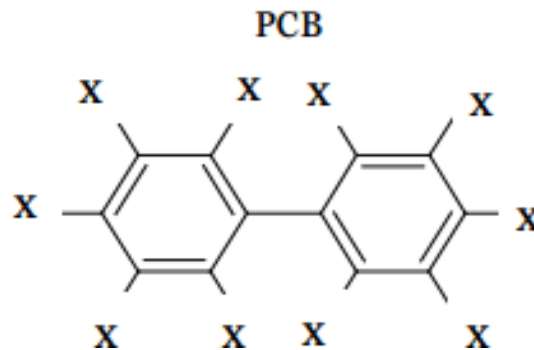
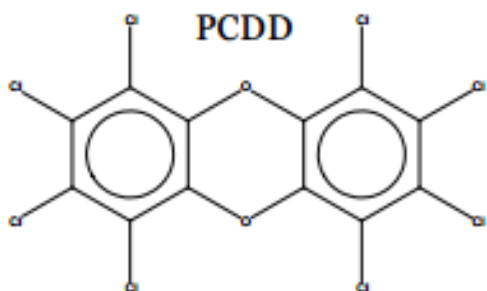


They are used as dielectric oil in transformers and accumulators and other uses

Dioxins



Anthropogenic substances coming from different sources, despite a minor natural contribution



Dioxin Incidents” – Huge Economical Impact

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Wednesday, June 2, 1999 Published at 12:10 GMT 13:10 UK

World: Europe

Belgian chickens banned in Europe



Belgium exports about half its egg and chicken produce

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Italy recalls tainted mozzarella

The Italian government has recalled from sale the mozzarella cheese linked to dioxin contamination.



Italy's health ministry said the affected cheese came from 25 producers in the Campania region near Naples, where buffalo mozzarella is made.

France has now lifted a ban on sales

Mozzarella production is big business in Italy

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Irish pork products recalled after dioxin is found

Published: Sunday, December 7, 2008

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From The Times
December 8, 2008

Contaminated pork from Ireland: your questions answered

“Dioxin Incidents” – Huge Economical Impact

- **Dioxin and PCB incidents**
- **... are expensive.**
 - Egg production enterprise with 5 million hens would cost in excess of \$30 million.
 - Broiler enterprise producing 3 million broilers per week would exceed \$85 million.
 - Times Beach, Missouri, road dust covered by oil spread. The cleanup cost with a total of \$110 million.
- **Sources to the Environment**
 - Combustion processes
 - Landfill sites
 - Improper waste management
 - Sewedge sludge
- **Accumulation in Food Chain**
 - Regular control required
 - In environment, feed, and food

Repeated “Dioxin” cases from only recent years:

- 1998 Milk from citrus pulp pellet feed
- 1999 Belgian PCB/dioxin in eggs, poultry
- 1999 Clay and zeolithes for feed
- 2000 Choline chloride
- 2002 “Carbosan Copper”
- 2004 Potato pulp
- 2005 Hydrochloric acid
- 2007 Indian Guar Gum thickener
- 2008 Irish Pork Meat
- 2008 Italian Mozzarella Cheese ...
- ... still on today around Naples.



Different dynamic and abatement strategies:

- **Global Scale – Stockholm Convention on POPs**
- **EU Regulation:**
 - Setting limits in well-known sources
 - Setting limits in feed and food for dioxins and PCBs
 - Defining not only the limits but also developing methods for sampling and analysis for the official control of levels of dioxins and PCBs in food and feed:
Confirmatory and Screening methods





HOME | THE CONVENTION | PROCEDURES | IMPLEMENTATION | COUNTRIES | PARTNERS

You are here: Stockholm Convention Clearing House > The Convention > The POPs > The 12 Initial POPs | Login

The POPs

- What are POPs
- The 12 Initial POPs**
- The New POPs
- Listing of POPs in the Stockholm Convention
- Chemicals proposed for listing under the Convention







The 12 initial POPs under the Stockholm Convention




Initially, twelve POPs have been recognized as causing adverse effects on humans and the ecosystem and these can be placed in 3 categories:

- **Pesticides:** aldrin, chlordane, DDT, dieldrin, endrin, heptachlor, hexachlorobenzene, mirex, toxaphene;
- **Industrial chemicals:** hexachlorobenzene, polychlorinated biphenyls (PCBs); and
- **By-products:** hexachlorobenzene; polychlorinated dibenzo-*p*-dioxins and polychlorinated dibenzofurans (PCDD/PCDF), and PCBs.

Annex C (Unintentional production)

Parties must take measures to reduce the **unintentional releases** of chemicals listed under Annex C with the goal of continuing minimization and, where feasible, ultimate elimination.

Hexachlorobenzene (HCB) 	Pentachlorobenzene 	Polychlorinated biphenyls (PCB) 
Polychlorinated dibenzo-<i>p</i>-dioxins (PCDD) 	Polychlorinated dibenzofurans (PCDF) 	Polychlorinated naphthalenes 

 Pesticide  Industrial chemical  Unintentional production

EU Commission Regulation – Past/Present

Past

Commission Regulation (EU) No 252/2012

- Specifies use of GC-HRMS for confirmatory dioxin analysis in food and feed.
- **GC-MS/MS** allowed as screening technique.



Present

Commission Regulation (EU) No 589/2014

- Specifies use of **GC-HRMS** or **GC-MS/MS** for confirmatory dioxin analysis in food and feed.

Commission Regulation (EU) No 589/2014 of 2 June 2014

- Specifies use of **GC-HRMS** or **GC-MS/MS** for confirmatory dioxin analysis.
- **GC-MS/MS** is “an appropriate confirmatory method for checking compliance with the maximum”, only.
- **GC-HRMS** remains the recommended technique for “determination of low background levels in food monitoring, following of time trends, exposure assessment of the population”.

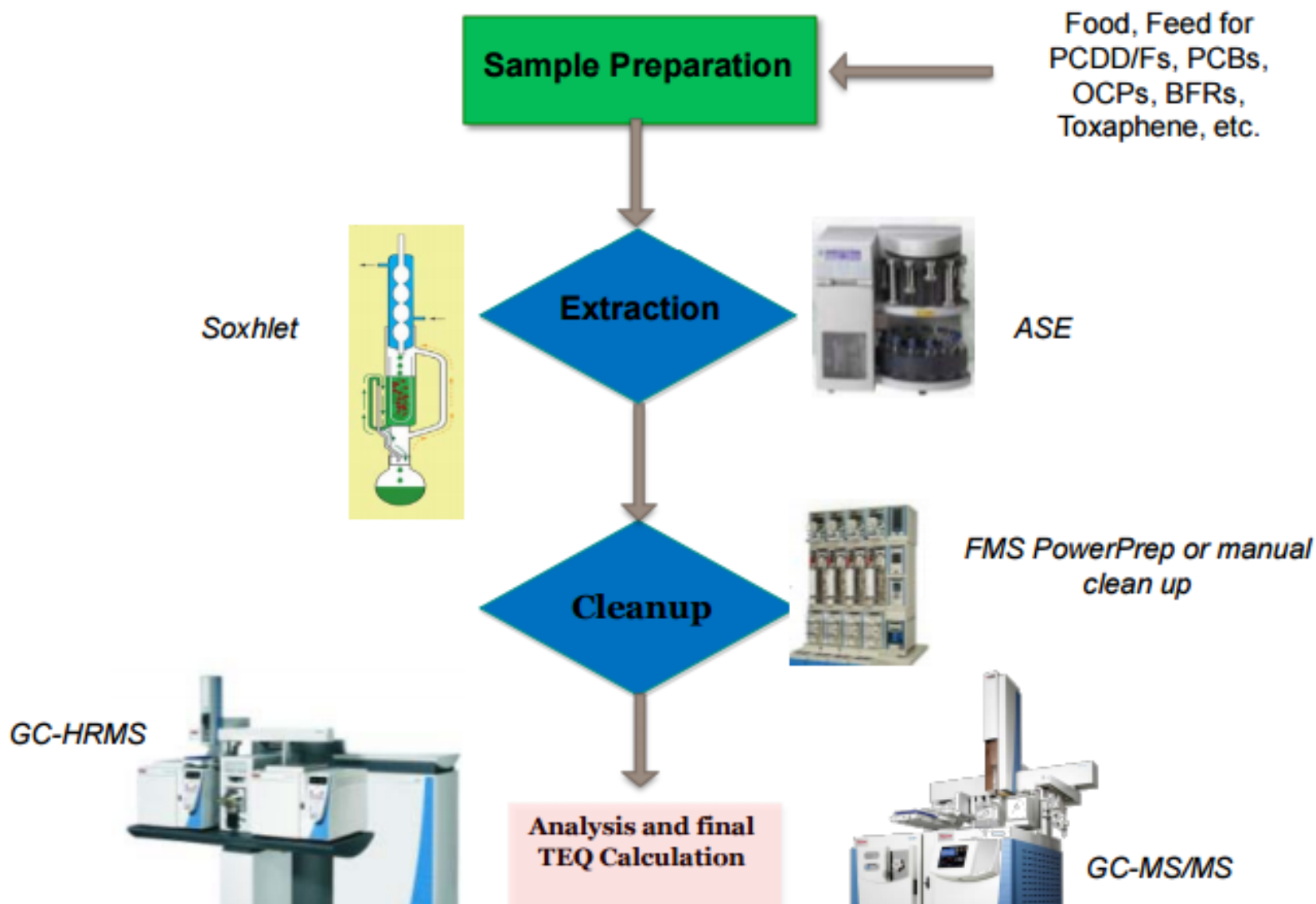
HRMS is recognized to deliver superior sensitivity, as required for **low level background studies**

HRMS fulfills all requirements for all types of dioxin applications, is considered the **Reference** standard for dioxin analysis.

SPECIFIC REQUIREMENTS FOR **GC-MS/MS METHODS** TO BE COMPLIED WITH FOR **CONFIRMATORY PURPOSES** (in food and feed samples):

- Monitoring of at least **2 specific precursor ions, each with one specific corresponding transition product ion**, for all labelled and unlabelled analytes...
- Maximum permitted tolerance of **relative ion intensities of $\pm 15\%$ for selected transition product ions** in comparison to calculated or measured values (average from calibration standards), applying identical MS/MS conditions, in particular collision energy and gas pressure,...
- **Resolution for each quadrupole** to be set **equal to or better than unit mass resolution** (unit mass resolution: sufficient resolution to separate two peaks one mass unit apart)...
- **LOQ**: The method must demonstrate that it is able to distinguish between the blank and cut-off value. In reporting a value, a notification level should be established to decide what to do with the samples with a response below this level.
 - For PCDD/PCDF, LOD should be in the range of higher femtograms (10^{-15} g). For most congeners of PCBs, is sufficient LOQ in the nanogram (10^{-9} g). However, to measure congeners similar to dioxin-like PCBs (in particular non-ortho substituted congeners) the lower limit of the working range must reach the lowest picogram (10^{-12} g).

POPs Analysis Workflow



Magnetic Sector GC-HRMS

Ultimate sensitivity & Maximum robustness

The **Thermo Scientific™ DFS™ Magnetic Sector GC-HRMS**, the Gold Standard for Dioxins & POPs, is the only GC-MS specifically designed for Dioxin and POPs analysis. The DFS GC-HRMS offers worldwide full compliance with any official Dioxin, PCB or PBDE method (for example, EPA 1613, 1668, 1614). Exploit the benefits of highest available Dioxin sensitivity and robustness, delivered by our large-volume ion source.



Thermo Scientific DFS Magnetic Sector GC-HRMS

- World-Wide Method compliance (e.g. EPA 1613)
- Best Dioxin installation spec
- Highest available sensitivity
- Robust-by-size with large volume ion source

GC-MS Triple Quadrupole

High sensitivity & Ease-of-use

The **Thermo Scientific™ TSQ™ 8000 Evo GC-MS/MS Analyzer** brings a compelling productivity opportunity to your laboratory. It offers you fully compliant analysis of Dioxins in food and feed in respect to the latest EU regulations, meeting stringent European Union performance criteria. High productivity within more efficient lab workflows accelerates sample turnaround times.



Thermo Scientific TSQ 8000 Evo GC-MS/MS Analyzer

- Ease-of-use
- Compliant with EU regulations for Dioxin food & feed analysis
- Great price/performance ratio

Orbitrap GC-MS

The ultimate tool for unknown identification

The **Thermo Scientific™ Q Exactive™ GC Orbitrap™ GC-MS/MS** represents the first-ever combination of high-resolution gas chromatography and high-resolution, accurate-mass (HRAM) Orbitrap mass spectrometry. It is an easy-to-use, dedicated benchtop GC-MS system that provides highest confidence for emerging POPs research with unmatched performance in compound discovery, identification and quantitation for a comprehensive understanding of your samples.



Thermo Scientific Q Exactive GC Orbitrap GC-MS/MS

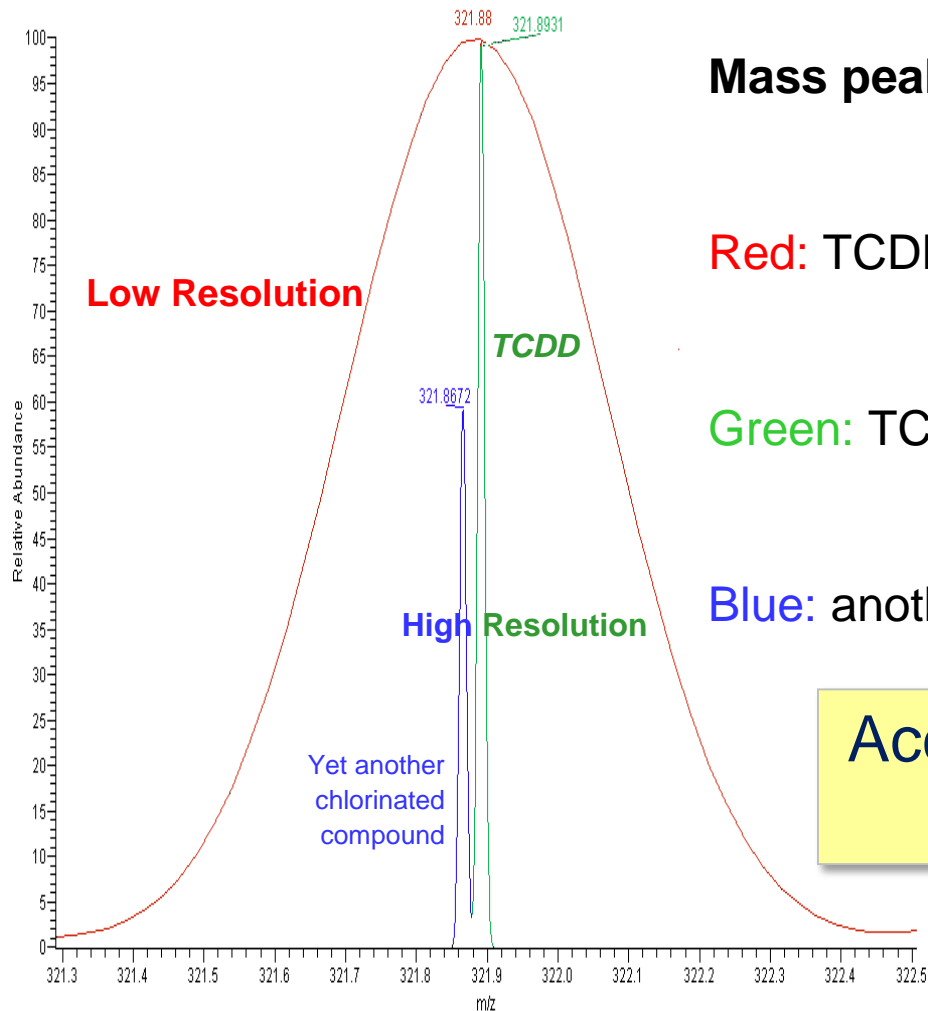
- The ultimate research tool
- Highest resolution and mass accuracy
- Unique unknown identification capabilities

Thermo Scientific DFS Method 1631 for GC-HRMS

- **Worldwide method compliance**
 - EPA 1613,1668,1614 & others
 - Use PFK/PFTBA , lock mass trace, etc.
- **Analytical Performance**
 - best Dioxin spec available
 - large ion source: ultimate sensitivity *combined with* highest **robustness**
 - Unique Dual GC configuration
- **References (10 years DFS!)**
 - Worldwide use in Dioxin expert labs

Confirmation Analysis with *High Resolution*

C12H4O2Cl4*1.00 + C12H3Cl5*1.00: p(gss, s/p.40) Chrg 10...



Mass peaks at Different Resolution Settings:

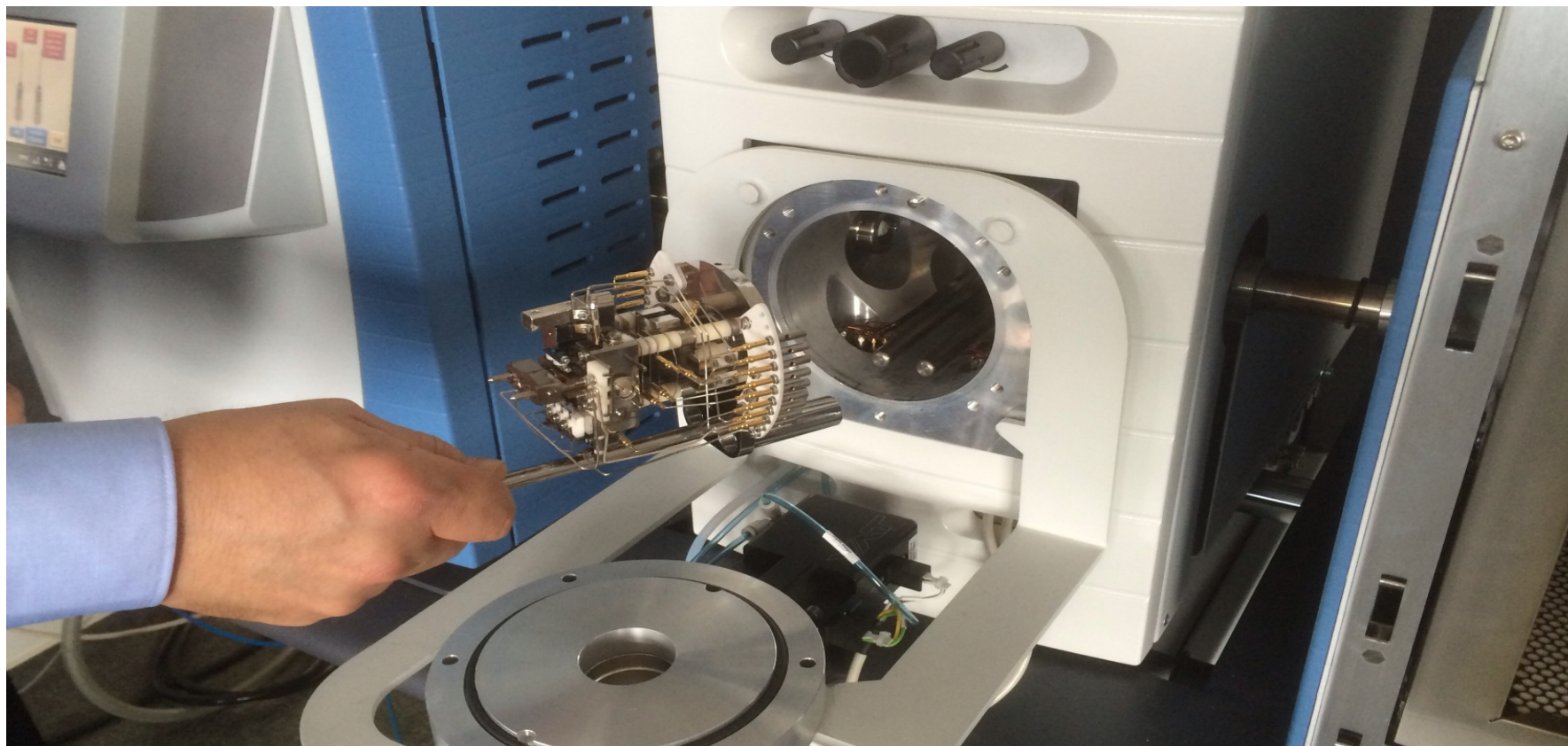
Red: TCDD mass trace at R 1.000 (GC-MS/MS)

Green: TCDD mass trace at R 10.000 (DFS)

Blue: another chlorinated compound mass trace

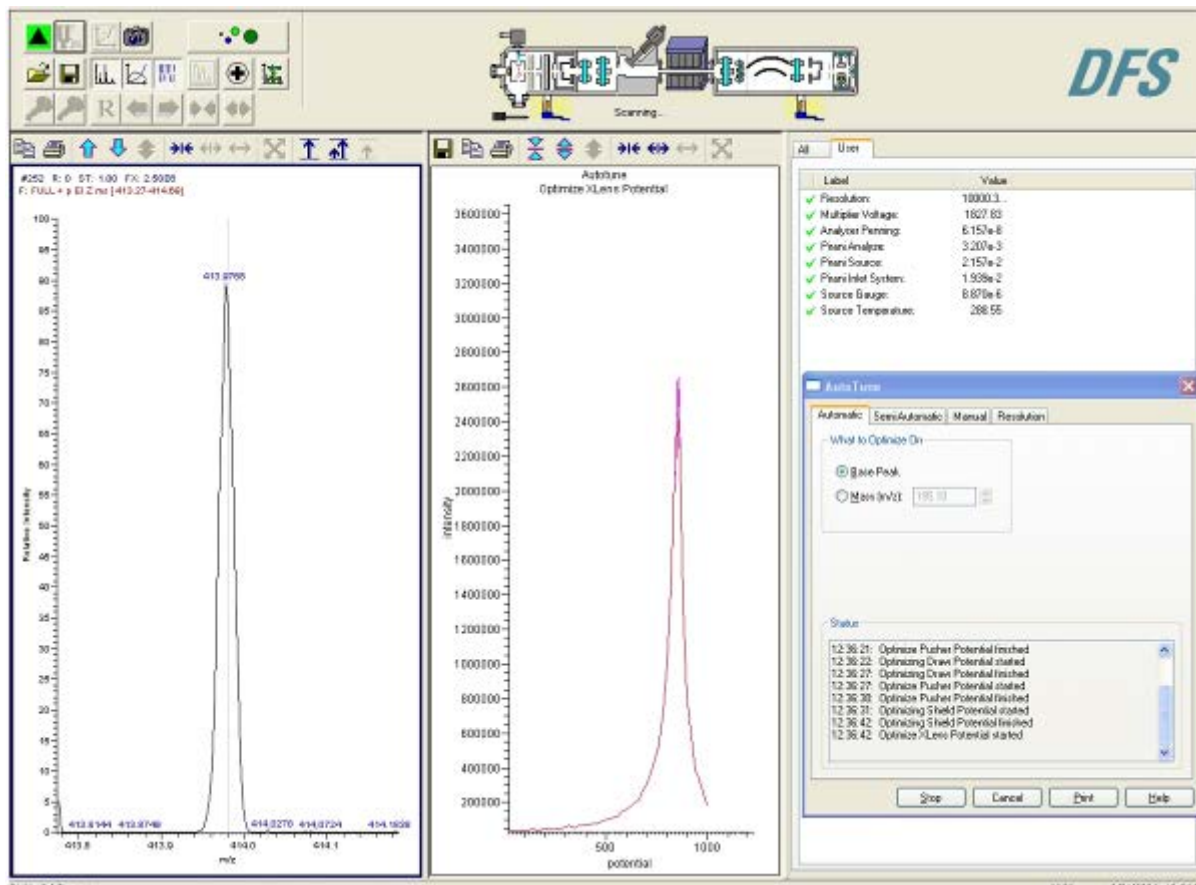
Accurate mass SIM is used for
Confirmation Analysis

DFS Ion Source – Robustness by Size



DFS ion source - designed for combining ultimate sensitivity with robustness, because sensitivity alone is not enough

Makes the DFS as easy to operate as a Benchtop MS



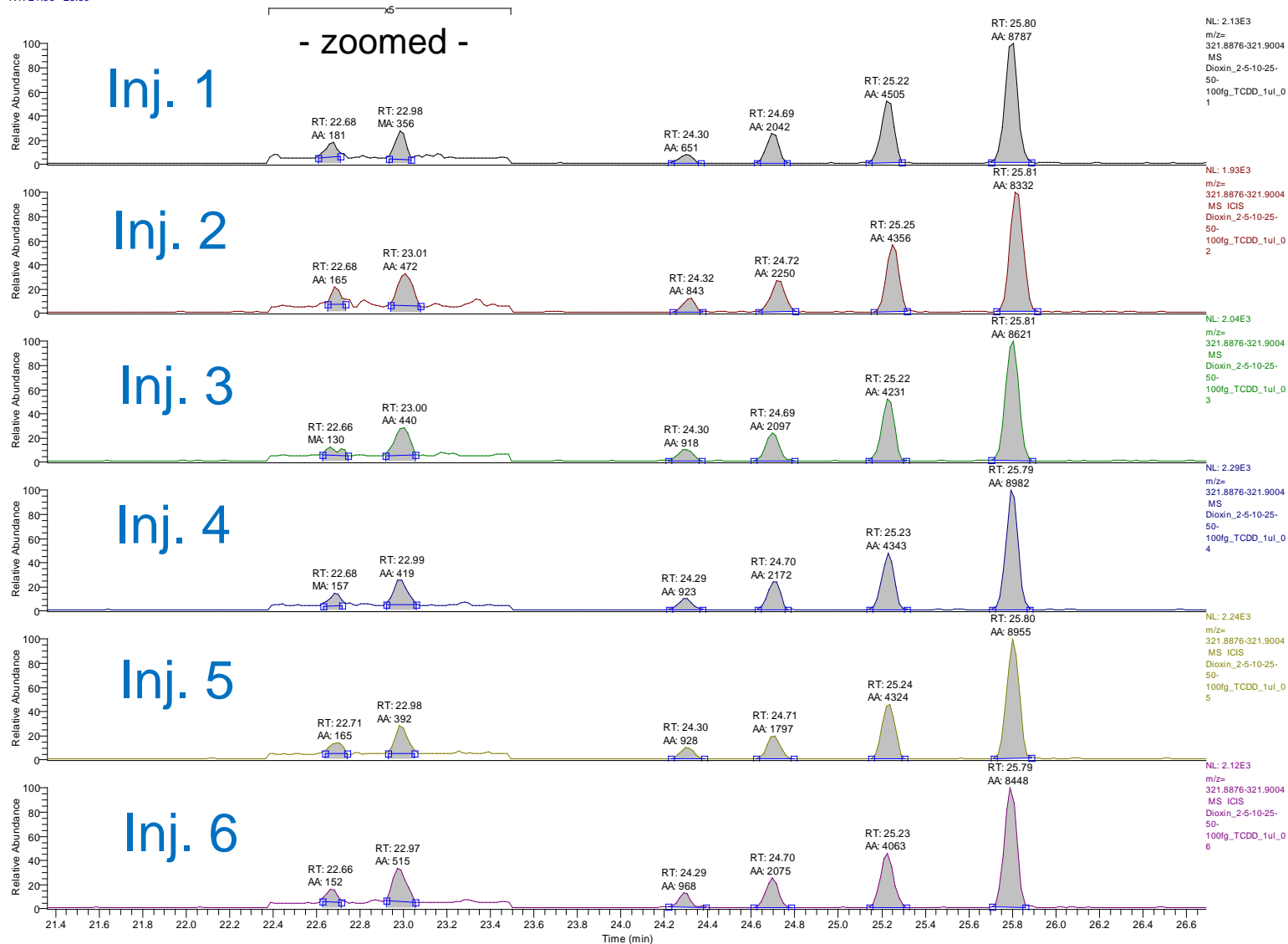
- Common User Interface known from DSQ, Polaris Q or LCQ/LTQ.

- **AutoTune**

- **Resolution setting**

2fg -100 fg TCDD on 60 m column

RT: 21.36 - 26.69

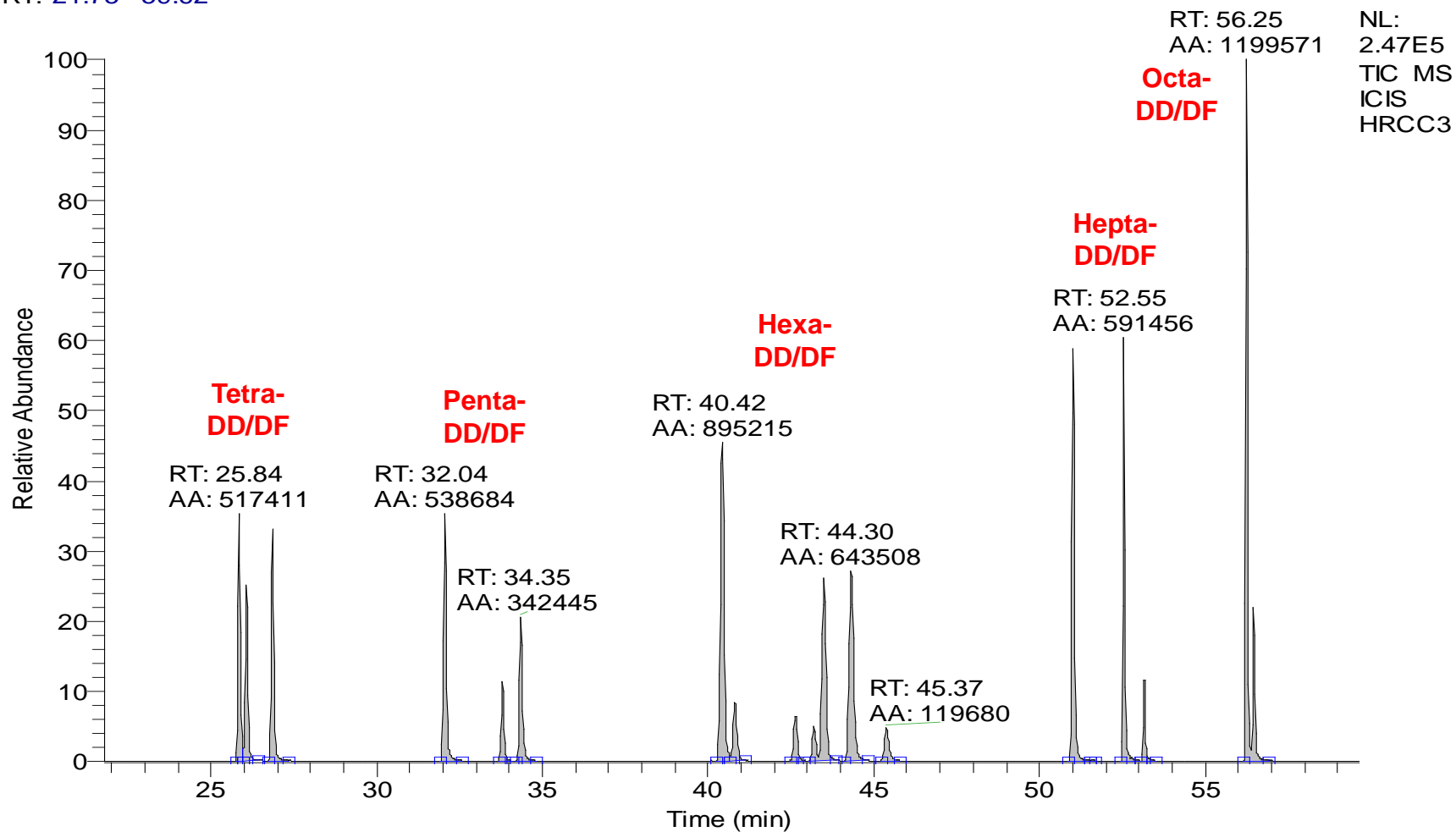


Dioxin & Furan Analysis

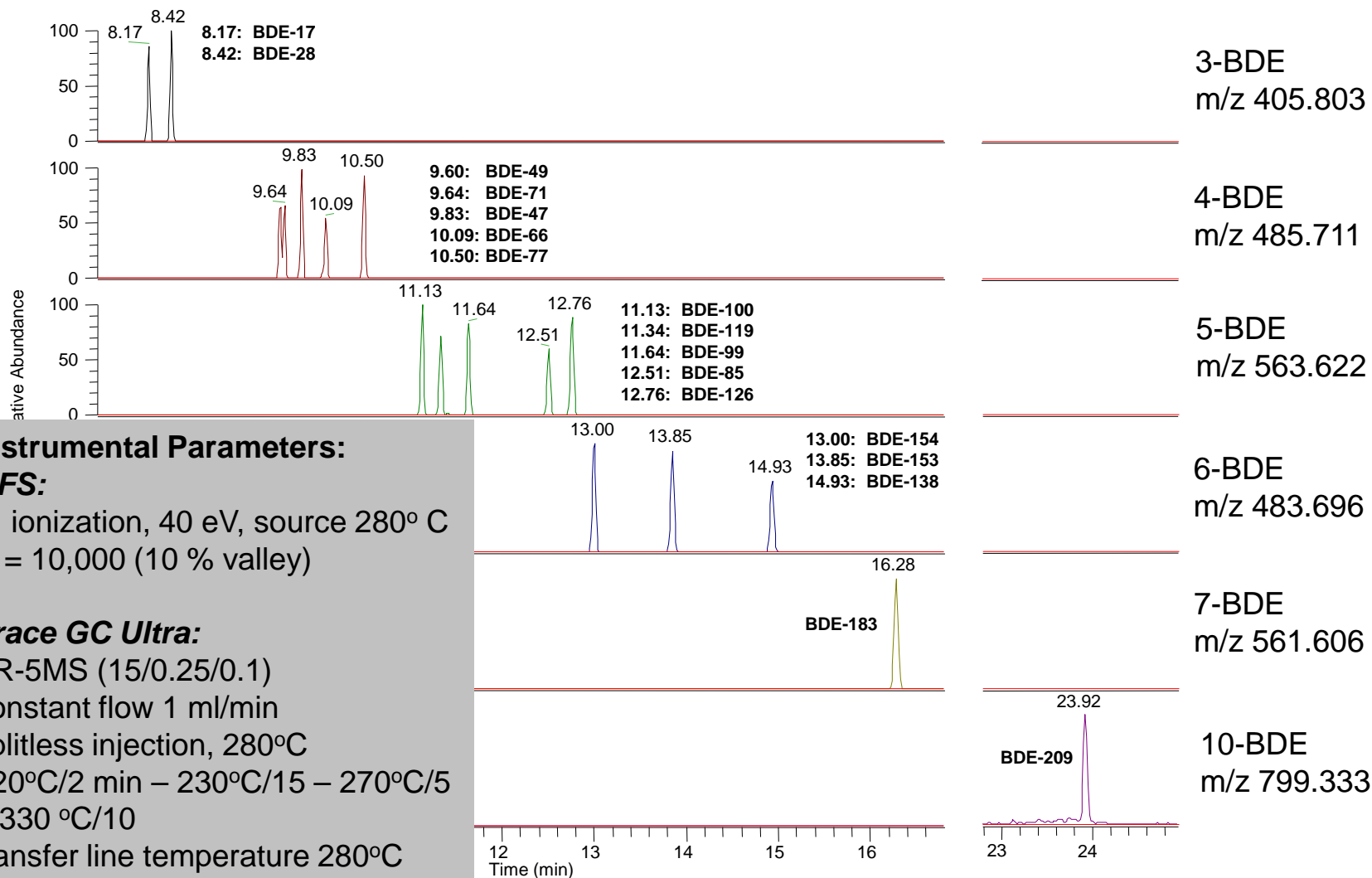
D:\Demo\...\calib8290\100514PMHRCC3

5/14/2010 7:25:30 PM

RT: 21.75 - 59.62



PBDE Brominated Flame Retardants (BFRs)



TSQ 8000 EVO – State of Art GC-MS/MS

- Thermo Scientific™ TSQ™ 8000 Evo triple quadrupole GC-MS
- State-of-the-art triple quadrupole GC-MS system introduced at ASMS 2014
- Fast collision cell instrument with EvoCell technology for high SRM transition speeds, precision, and sensitivity



- Thermo Scientific™ ExtractaBrite™ ion source is heated throughout for high matrix tolerance
- Ion source is fully removable, hot, under vacuum when cleaning is necessary or swapping with a spare

EU Regulation 589/2014 Requirements

6.5. Specific criteria for confirmatory methods

— For GC-HRMS:

In HRMS, the resolution shall typically be greater than or equal to 10 000 for the entire mass range at 10 % valley.

Fulfilment of further identification and confirmation criteria as described in internationally recognised standards, for example, in standard EN 16215:2012 (Animal feed — Determination of dioxins and dioxin-like PCBs by GC/HRMS and of indicator PCBs by GC/HRMS) and/or in EPA methods 1613 and 1668 as revised.

— For GC-MS/MS:



Monitoring of at least 2 specific precursor ions, each with one specific corresponding transition product ion, for all labelled and unlabelled analytes in the scope of analysis.



Maximum permitted tolerance of relative ion intensities of $\pm 15\%$ for selected transition product ions in comparison to calculated or measured values (average from calibration standards), applying identical MS/MS conditions, in particular collision energy and collision gas pressure, for each transition of an analyte.



Resolution for each quadrupole to be set equal to or better than unit mass resolution (unit mass resolution: sufficient resolution to separate two peaks one mass unit apart) in order to minimise possible interferences on the analytes of interest.



Fulfilment of the further criteria as described in internationally recognised standards, for example, in standard EN 16215:2012 (Animal feed — Determination of dioxins and dioxin-like PCBs by GC/HRMS and of indicator PCBs by GC/HRMS) and/or in EPA methods 1613 and 1668 as revised, except the obligation to use GC-HRMS.

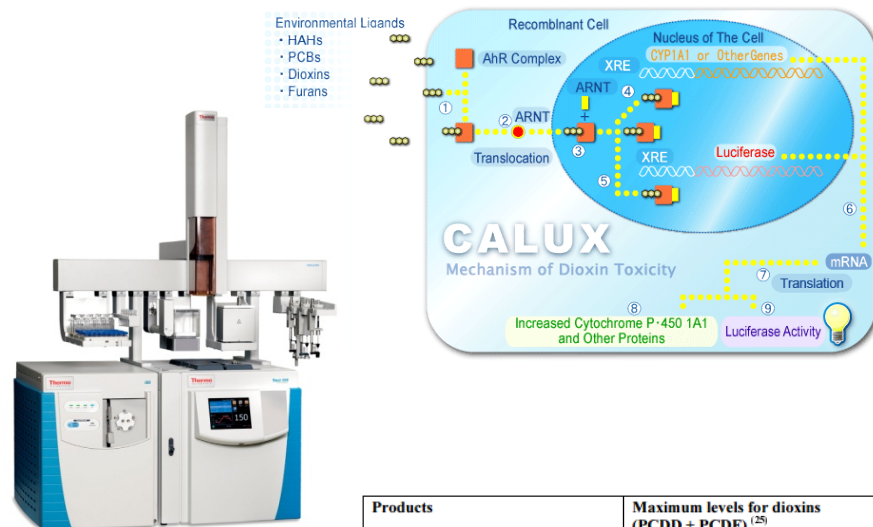
Methods for Dioxin/Furans & PCBs Measurement

SCREENING:

- GC-MS
- GC-MS/MS
- Magnetic Sector
- Bioassay (e.g. CALUX)

CONFIRMATION:

- Magnetic Sector – MLs and background
- GC-MS/MS – MLs as outlined in EU regulations 709/2014 and 589/2014



Products	Maximum levels for dioxins (PCDD + PCDF) ⁽²⁶⁾ (pg WHO-PCDD/F-TEQ/g fat or product)
5.1.1 Meat and meat products ⁽²⁸⁾ originating from - Ruminants (bovine animals, sheep) - Poultry and farmed game - Pigs	3 pg WHO-PCDD/F-TEQ /g fat ^(26,27) 2 pg WHO-PCDD/F-TEQ /g fat ^(26,27) 1 pg WHO-PCDD/F-TEQ /g fat ^(26,27)
5.1.2 Liver and derived products	6 pg WHO-PCDD/F-TEQ /g fat ^(26,27)
5.2. Muscle meat of fish and fishery products ⁽²⁹⁾ and products thereof	4 pg WHO-PCDD/F-TEQ /g fresh weight ⁽²⁶⁾
5.3. Milk ⁽³⁰⁾ and milk products, including butter fat	3 pg WHO-PCDD/F-TEQ /g fat ^(26,27)
5.4 Hen eggs and egg products ^(31,32)	3 pg WHO-PCDD/F-TEQ /g fat ^(26,27)
5.5. Oils and fats - Animal fat - from ruminants - from poultry and farmed game - from pigs - mixed animal fat - Vegetable oil - fish oil intended for human consumption	3 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾ 2 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾ 1 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾ 2 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾ 0.75 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾ 2 pg WHO-PCDD/F-TEQ /g fat ⁽²⁶⁾

Experimental conditions (GC & MS)

TRACE 1310 GC Parameters

Injection Volume (μL):	2.0
Liner	SSL Single tapered P/N: 453A2342
Inlet (°C):	280
Inlet Module and Mode:	SSL splitless
Carrier Gas, (mL/min):	He, 1.2

Oven Temperature Program:

Temperature 1 (°C):	120
Hold Time (min):	2
Temperature 2 (°C):	250
Rate (°C/min)	25
Hold Time (min):	0
Temperature 3 (°C):	285
Rate (°C/min)	2.5
Hold Time (min):	0
Temperature 4 (°C):	320
Rate (°C/min)	10
Hold Time (min):	15

Column

Type	Length (m)	I.D.(mm)	Film (μm)	P/N
TG-5SilMS	60	0.25	0.25	26066-1540

TSQ 8000 Evo Mass Spectrometer Parameters

Transfer line (°C):	280
Ionization type:	EI
Ion source(°C):	300
Electron energy (eV):	40
Acquisition Mode:	SRM (Timed Acquisition)
Collision gas:	Argon (60 psig)
Resolution Q1:	0.7 u (FWHM)
Resolution Q3:	0.7 u (FWHM)



All 17 PCDD/F and 12 dl-PCBs native and labeled congeners can be acquired in one method

Sample Preparation

- PCDD/Fs and dl-PCB standards containing the native and the ^{13}C -labelled compounds were obtained from **Wellington Laboratories** Inc. The following food and feed extracted samples were provided by the Institute of Environmental Assessment and Water Research, CSIC Barcelona, Spain:

dl-PCBs

- Adipose tissue
- Fish liver
- Fish oil
- Dry fishspiked feed sample

PCDD/Fs

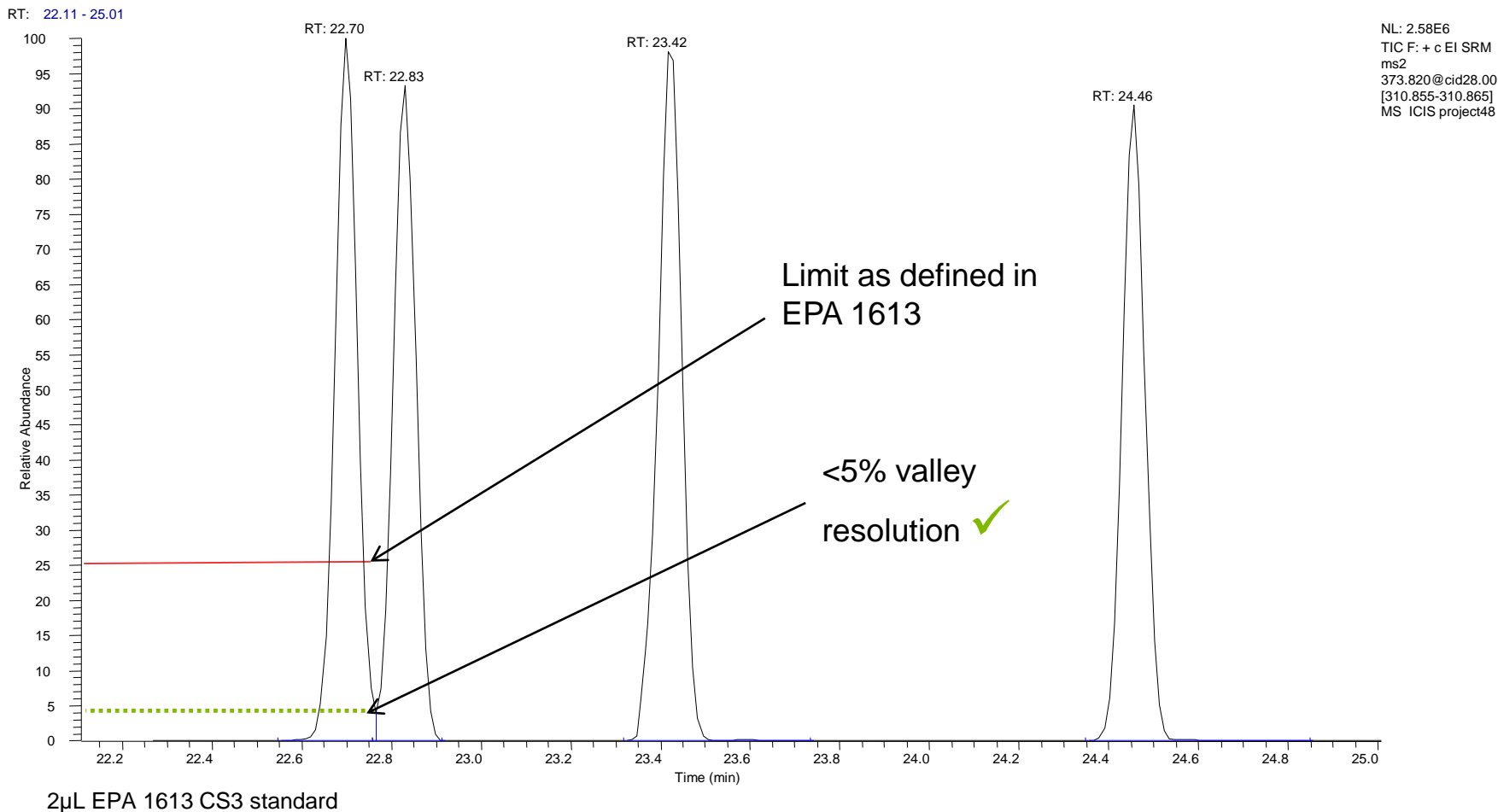
- milk powder sample (certified reference material).
- 3x dry fish samples (previously used in inter-laboratory studies)
- one feed sample (internal reference material).
- one milk powder sample (certified reference material).



- Extraction and clean-up of the matrix samples was performed either by PowerPrep™ SPE system (feed sample) or using a manual clean-up with multilayer silica, followed by basic alumina and a final carbon column (milk, tissue and fish samples).

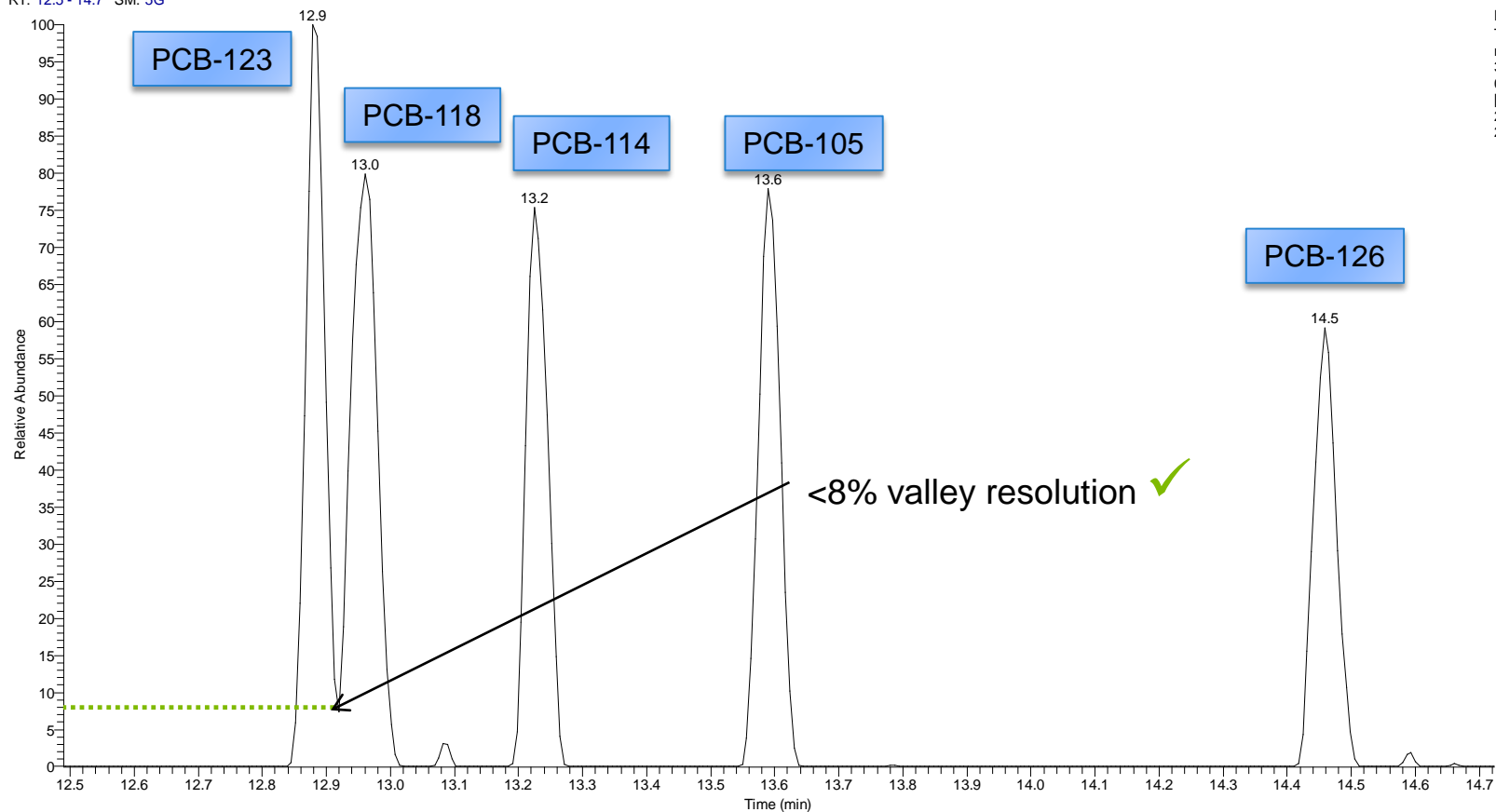
Chromatography - PCDD/Fs

Excellent resolution of the isomeric pair 1,2,3,4,7,8 and 1,2,3,6,7,8 HxCDF



Chromatography - dl-PCBs

RT: 12.5 - 14.7 SM: 5G



NL: 1.85E3
TIC F: + c EI SRM
ms2
323.900@cid22.0
0
[253.895-
253.905] MS
280115_025

2µL 68C-CVS CS1 standard

Resolution of 'Penta' dl-PCBs – well below 25% valley

The level of interest...

Unlike GC-HRMS, often in SRM data there is no noise and a signal to noise calculation based on LOD/LOQ can be meaningless. Therefore a statistical method was used.

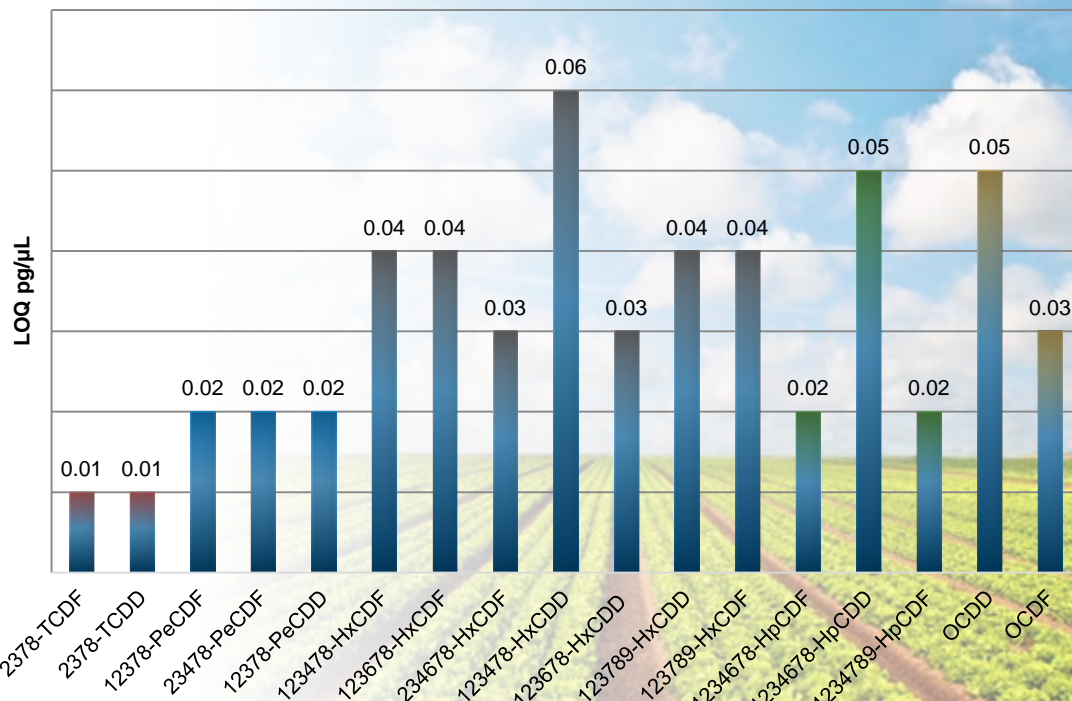
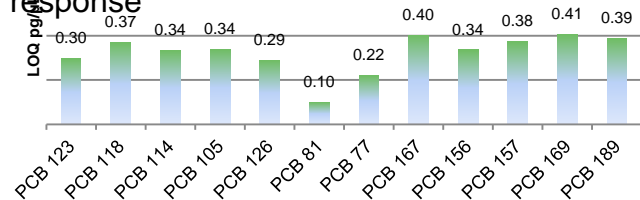
$$\text{LOQ} = t * \text{Amount} * \% \text{RSD}$$

Where,

t = student t -value for **one-tailed** distribution:
for $n = 10$ injections; $t = 2.821$

Amount = amount of analyte (on-column)

%RSD = relative standard deviation of the response



All ion ratios and response factor values met the required criteria at these limits

Linearity – Response factors (RF)

Linearity / Calibration (5 point)

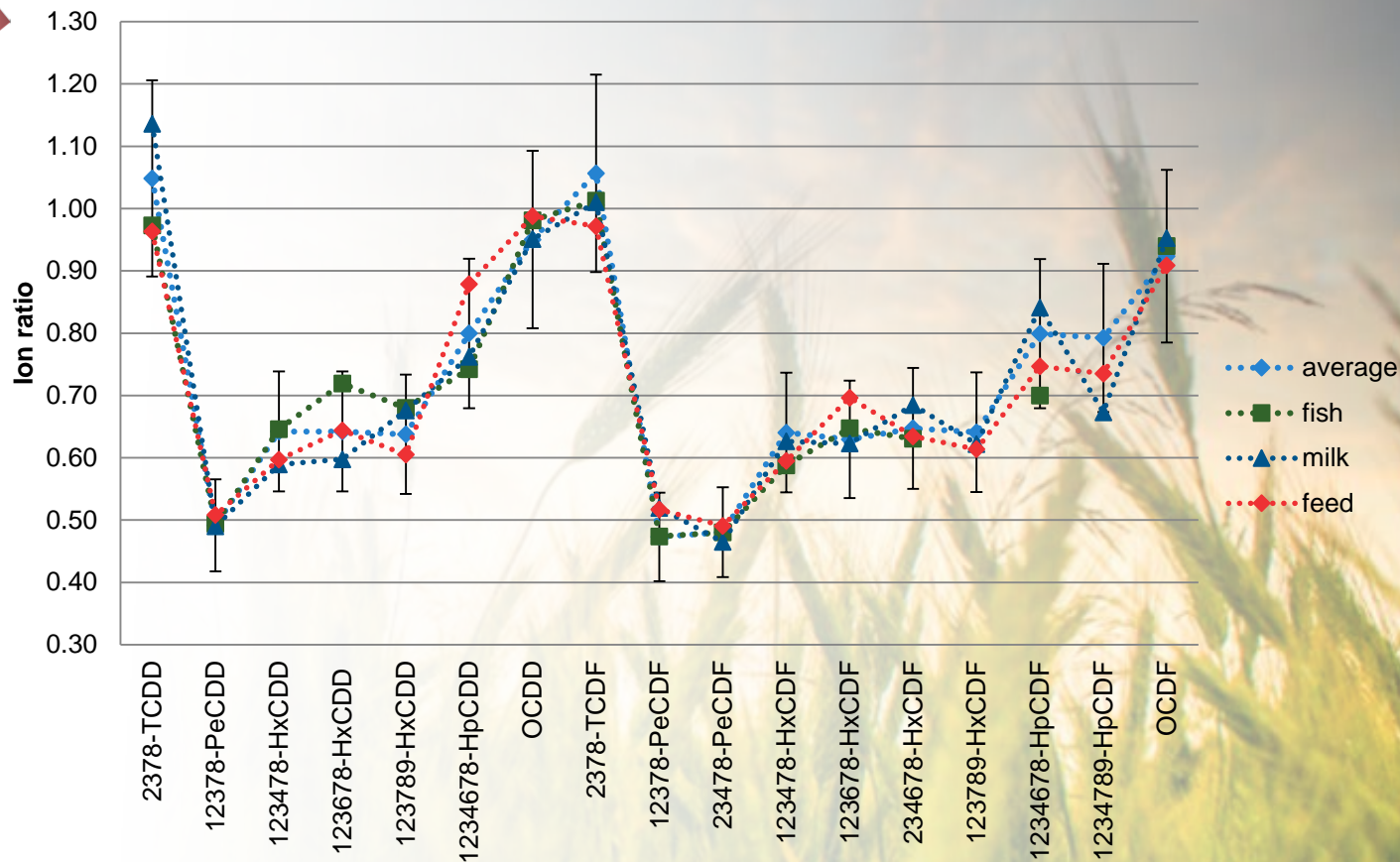
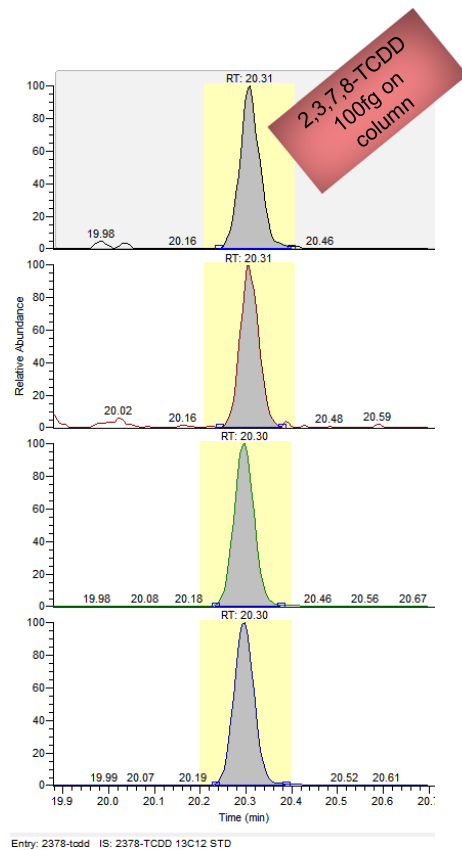
Compound	Concentration range (pg/μL)	Average RF	stdev	RF % RSD
PCB 77	1 - 2000	1.11	0.01	1.0
PCB 81	1 - 2000	1.10	0.01	1.0
PCB 105	1 - 2000	0.98	0.03	2.6
PCB 114	1 - 2000	1.08	0.04	3.7
PCB 118	1 - 2000	1.04	0.01	0.8
PCB 123	1 - 2000	1.01	0.02	1.7
PCB 126	1 - 2000	1.07	0.03	2.7
PCB 156	1 - 2000	1.20	0.02	1.4
PCB 157	1 - 2000	1.15	0.03	2.8
PCB 167	1 - 2000	1.22	0.07	5.6
PCB 169	1 - 2000	1.00	0.12	2.5
PCB 189	1 - 2000	1.01	0.04	4.3

Linearity / Calibration (6 point)

Compound	Concentration range (pg/μL)	Average RF	stdev	RF % RSD
2378-TCDF	0.1 - 40	1.04	0.02	1.9
2378-TCDD	0.1 - 40	1.12	0.02	2.2
12378-PeCDF	0.5 - 200	1.01	0.02	1.5
23478-PeCDF	0.5 - 200	1.03	0.02	1.6
12378-PeCDD	0.5 - 200	1.08	0.01	1.4
123478-HxCDF	0.5 - 200	1.03	0.01	1.2
123678-HxCDF	0.5 - 200	1.02	0.01	1.4
234678-HxCDF	0.5 - 200	1.06	0.03	3.2
123478-HxCDD	0.5 - 200	0.96	0.01	1.6
123678-HxCDD	0.5 - 200	1.21	0.04	3.6
123789-HxCDD	0.5 - 200	1.21	0.04	3.6
123789-HxCDF	0.5 - 200	1.36	0.11	0.8
1234678-HpCDF	0.5 - 200	1.06	0.02	1.7
1234678-HpCDD	0.5 - 200	1.07	0.02	2.1
1234789-HpCDF	0.5 - 200	1.12	0.02	2.2
OCDD	1.0 - 400	1.54	0.04	2.4
OCDF	1.0 - 400	1.09	0.03	3.2

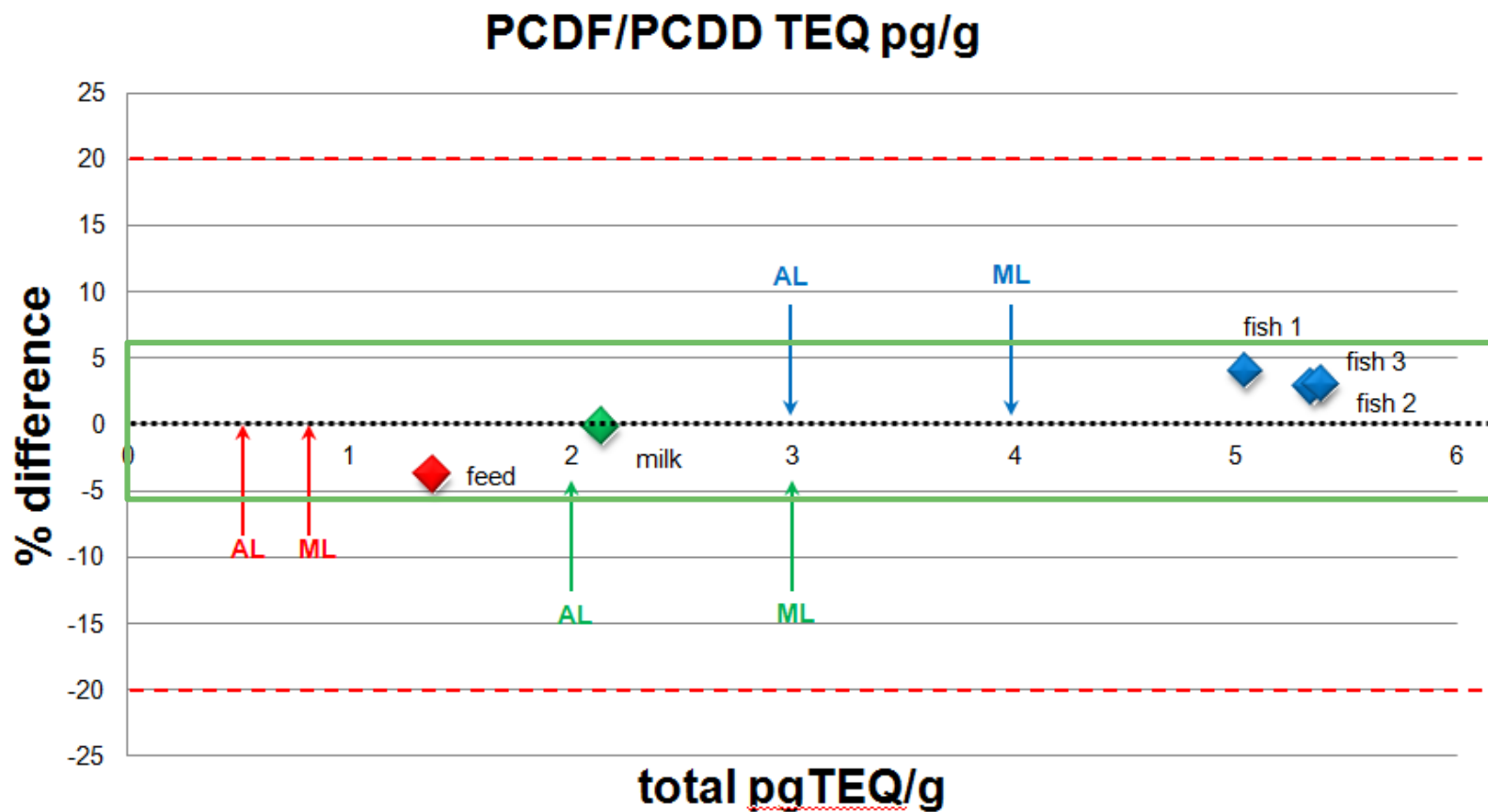
All RFs are well with the limits defined by US EPA 1613 ($\pm 15\%$)

Ion Ratio Abundance – PCDD/Fs



All ion ratios were within $\pm 15\%$ tolerance – meeting EU criteria

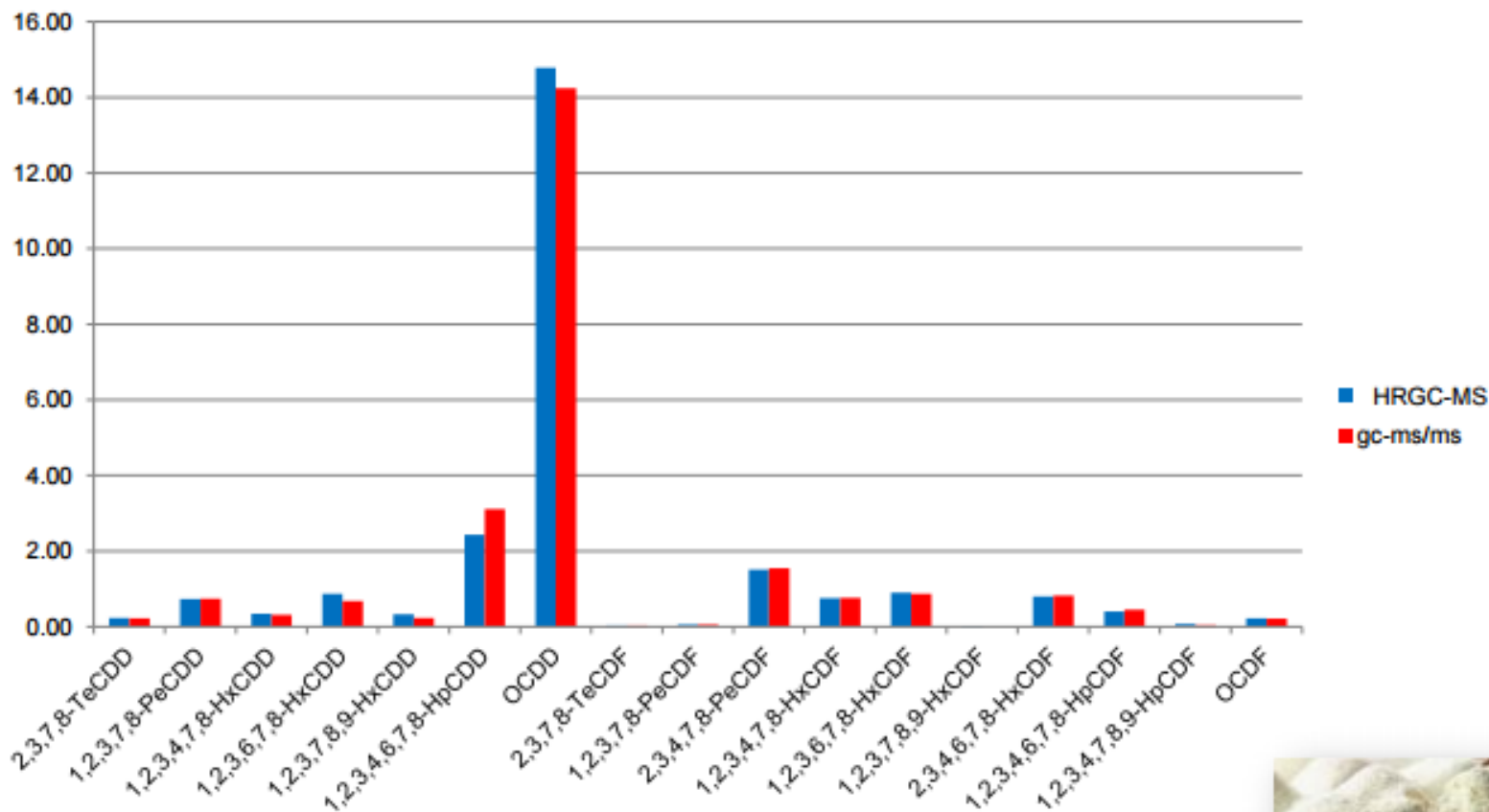
Agreement with GC-HRMS (PCDD/Fs)



<5% difference between total dioxin concentration (WHO-PCDD/F-TEQ pg/g)

Determination of DioxinFurans in Milk Powder

Comparison GC-MS/MS with GC-HRMS in Milk powder Sample

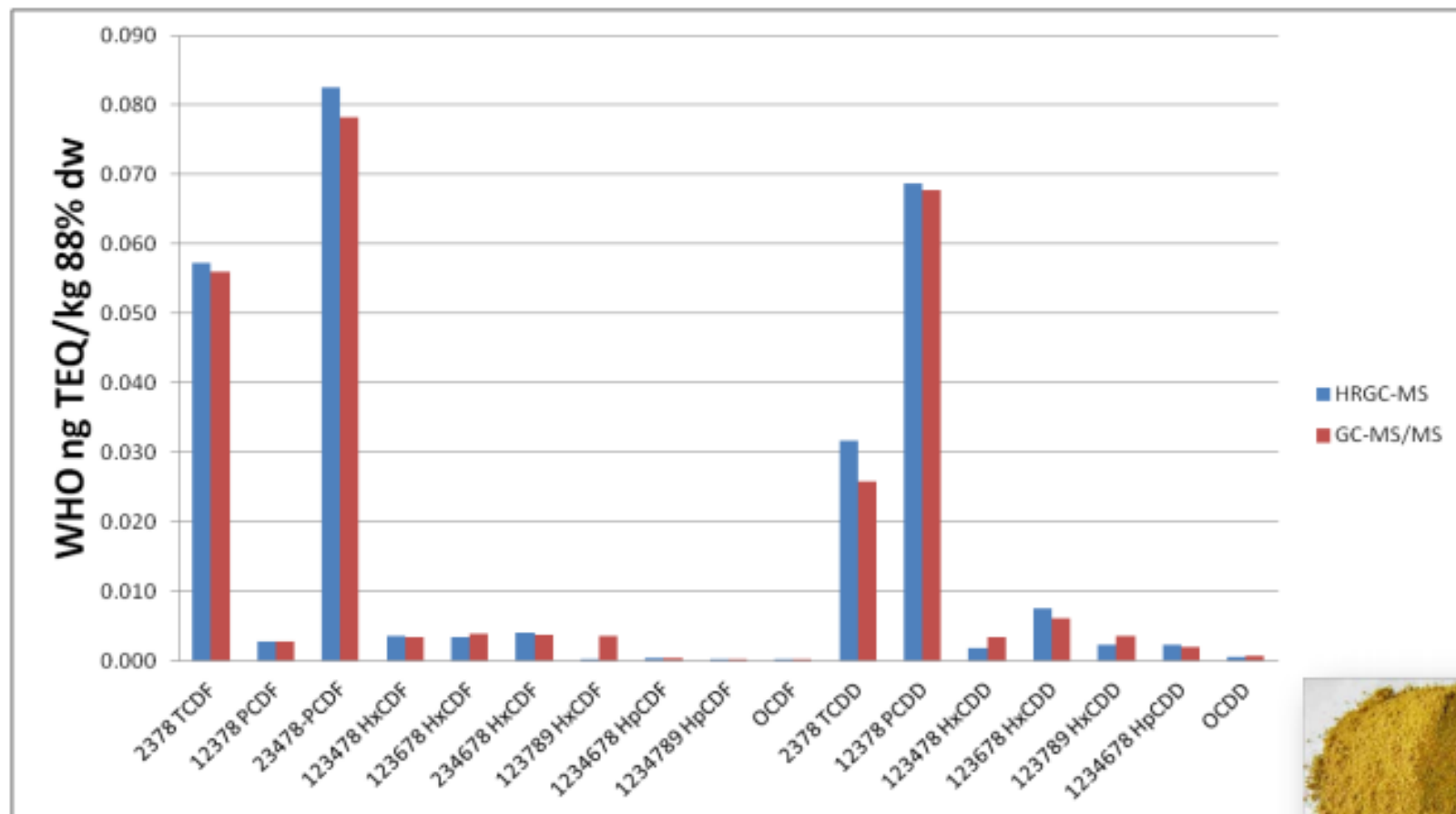


Good correlation between GC-QqQ and GC-HRMS



Determination of Dioxin/Furans in Animal Feed

Comparison GC-MS/MS with GC-HRMS in Feed Sample



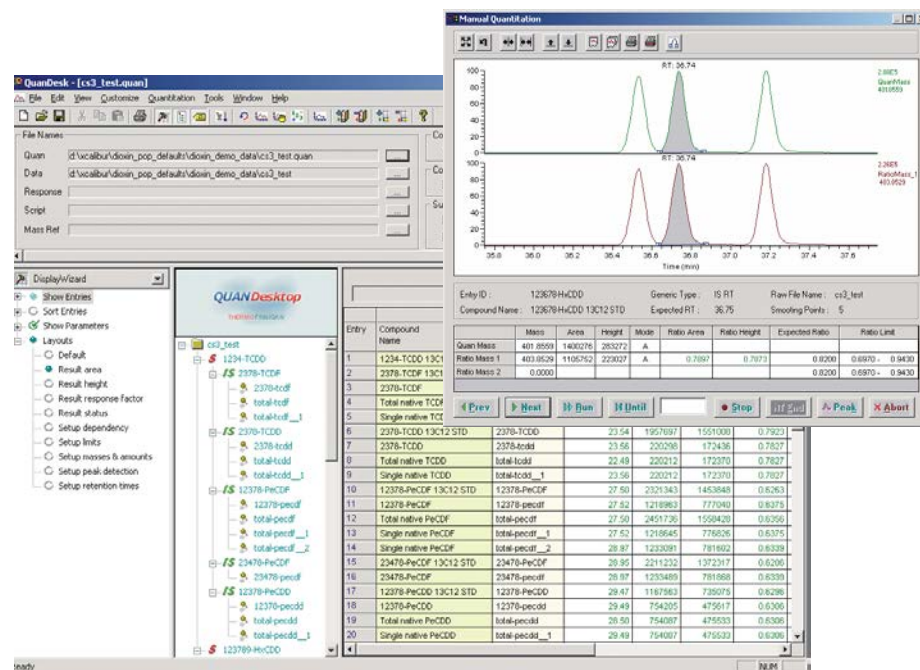
Good correlation between GC-QqQ and GC-HRMS



TargetQuan – POPs Quantitation for TSQ and DFS

Features for dioxin applications:

- Toxicity equivalents (TEQ)
 - according to WHO definition
 - including lower, medium and upper boundary calculation
- User definable summation
 - Of calculated amounts or TEQs for reporting of sum TEQ values
- Ion ratio confirmation
 - One quantitation mass and up to two masses based on abundance
- EU and EPA 1613, 1668 compliant
 - Allows quantification based on average response of selected compound Retention time correction



TargetQuan – POPs Quantitation for TSQ and DFS

The screenshot displays the TargetQuan software interface with the following components:

- Chromatogram (Top Left):** Shows a peak at RT: 23.56. Parameters: NL: 5.56E4, m/z= 321.9, 321.9 MS, cs3_test.
- Chromatogram (Middle Left):** Shows a peak at RT: 23.56. Parameters: NL: 4.55E4, m/z= 319.9, 319.9 MS, cs3_test.
- Chromatogram (Bottom Left):** Shows peaks at RT: 22.96 and RT: 23.54. Parameters: NL: 4.86E5, m/z= 333.9, 334.0 MS, cs3_test.
- Sequence List (Top Right):** Lists data files for various compounds and standards.
- Data Table (Bottom Right):** Contains the following data:

Entry	Compound Name	Entry Identifier	Quan. Mass	GM1
1	1234-TCDD 13C12 STD	1234-TCDD	333.9339 +/- 50 ppm	
2	2378-TCDF 13C12 STD	2378-TCDF	317.9389 +/- 50 ppm	
3	2378-TCDF	2378-tcdf	305.8987 +/- 50 ppm	
4	2378-TCDD 13C12 STD	2378-TCDD	333.9339 +/- 50 ppm	
5	2378-TCDD	2378-tcdd	321.8936 +/- 50 ppm	
6	12378-PeCDF 13C12 STD	12378-PeCDF	351.9000 +/- 50 ppm	
7	12378-PeCDF	12378-pecdf	339.8597 +/- 50 ppm	
8	23478-PeCDF 13C12 STD	23478-PeCDF	351.9000 +/- 50 ppm	
9	23478-PeCDF	23478-pecdf	339.8597 +/- 50 ppm	
10	12378-PeCDD 13C12 STD	12378-PeCDD	351.9000 +/- 50 ppm	
11	12378-PeCDD	12378-pe		
12	123789-HxCDD 13C12 STD	123789-HxCDD		
13	123478-HxCDF 13C12 STD	123478-HxCDF		
14	123478-HxCDF	123478-h		
15	123678-HxCDF 13C12 STD	123678-HxCDF		
16	123678-HxCDF	123678-h		
17	234678-HxCDF 13C12 STD	234678-HxCDF		
18	234678-HxCDF	234678-h		
19	123789-HxCDF 13C12 STD	123789-HxCDF		
20	123789-HxCDF	123789-h		
21	123478-HxCDD 13C12 STD	123478-HxCDD		
22	123478-HxCDD	123478-h		
23	123678-HxCDD 13C12 STD	123678-HxCDD	401.8559 +/- 50 ppm	
24	123678-HxCDD	123678-hxcdd	389.8157 +/- 50 ppm	
25	123789-HxCDD	123789-hxcdd	389.8157 +/- 50 ppm	
26	1234678-HpCDF 13C12 STD	1234678-HpCDF	419.8220 +/- 50 ppm	
27	1234678-HpCDF	1234678-hpcdf	407.7818 +/- 50 ppm	
28	1234789-HpCDF 13C12 STD	1234789-HpCDF	419.8220 +/- 50 ppm	
29	1234789-HpCDF	1234789-hpcdf	407.7818 +/- 50 ppm	
30	1234678-HpCDD 13C12 STD	1234678-HpCDD	435.8169 +/- 50 ppm	
31	1234678-HpCDD	1234678-hpcdd	423.7766 +/- 50 ppm	
32	OCDD 13C12 STD	OCDD	471.7750 +/- 2000.0 mmu	
33	OCDF	ocdf	443.7399 +/- 50 ppm	
34	OCDD	ocdd	459.7348 +/- 50 ppm	
*		<New>		

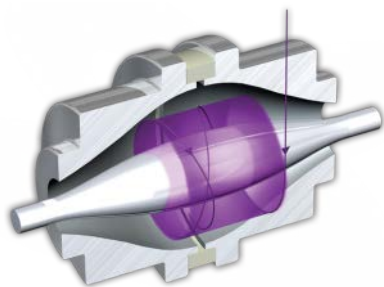
By Compound One Analysis Sequentially Selected Entry

By Analysis One Compound Sequentially Selected Files

Table Data:

Area	Height	Integra...	Ratio Area	Ratio Height	Ratio Limit	Expect
220297.590	55538.157	A				
Ratio Mass 1 172435.926	45419.885	A	0.783	0.818	0.6545 - 0.8855	0.77
Ratio Mass 2 0.000	0.000	---	0.000	0.000	0.6545 - 0.8855	0.77

Integration of 3 highly successful technologies



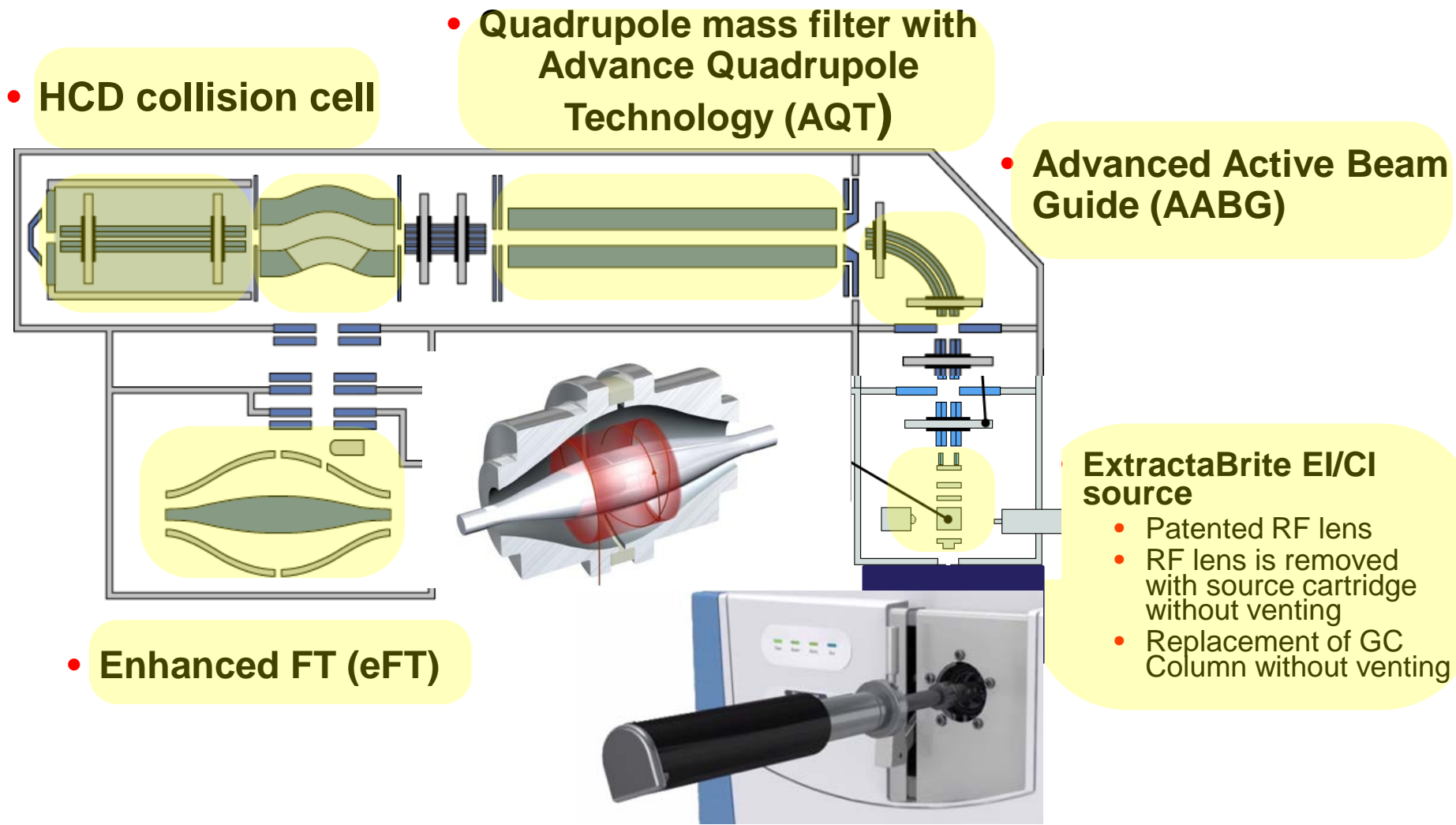
Orbitrap mass analyzer technology

Incredible HR/AM performance

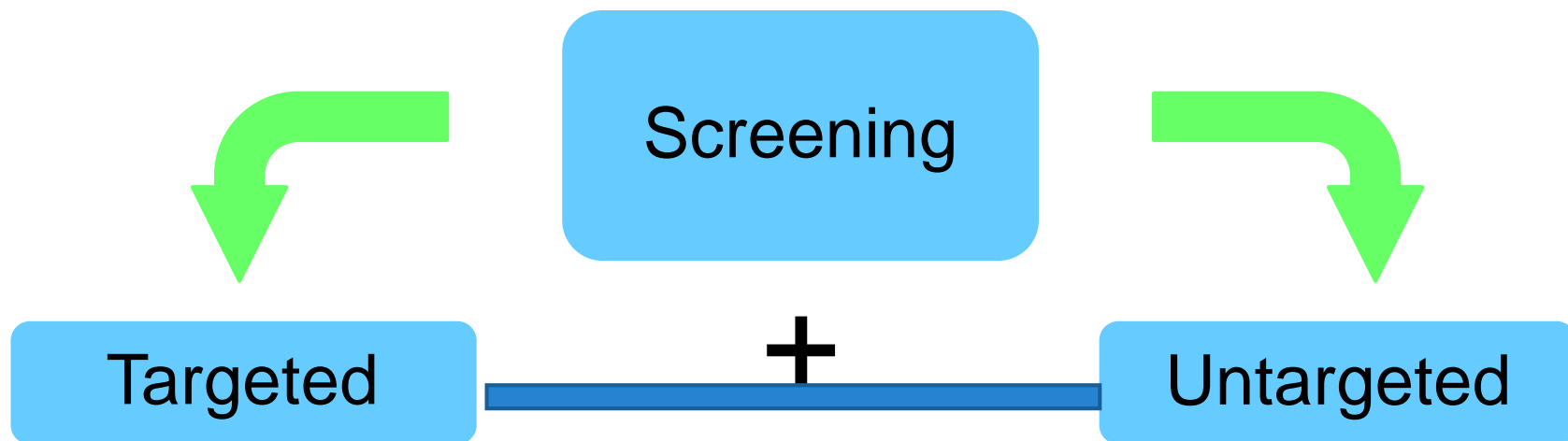
Highly regarded QExactive platform



Q Exactive GC Mass Spectrometer: Hardware Innovations



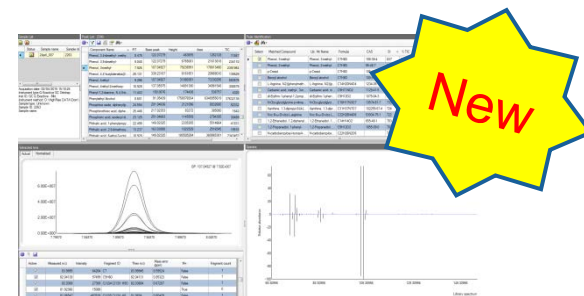
QExactive GC - Application Workflow Capability & Flexibility



**Multiple High Resolution
Databases**

**High Resolution Mass
Deconvolution**

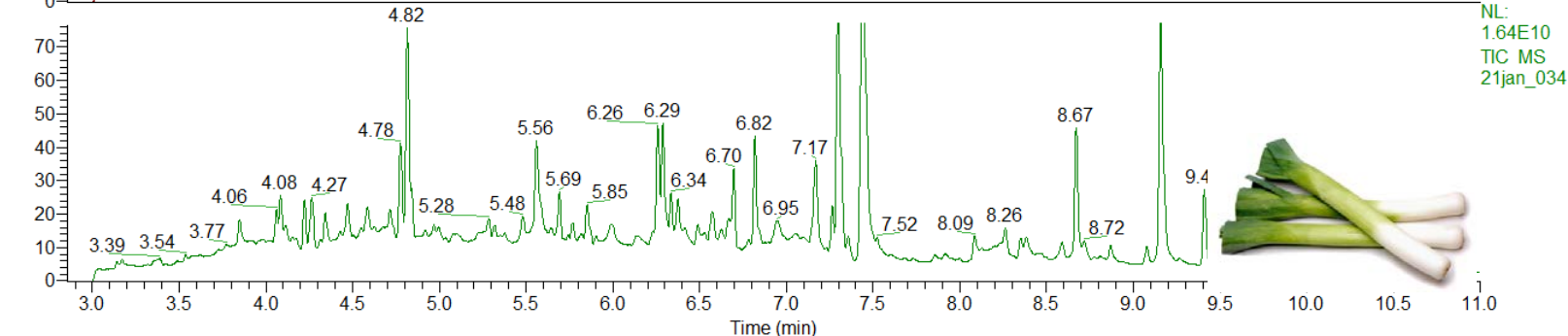
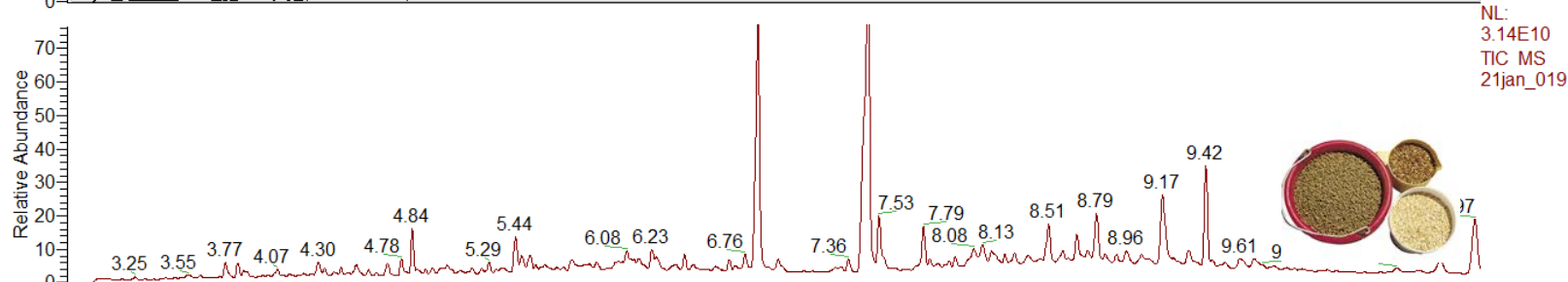
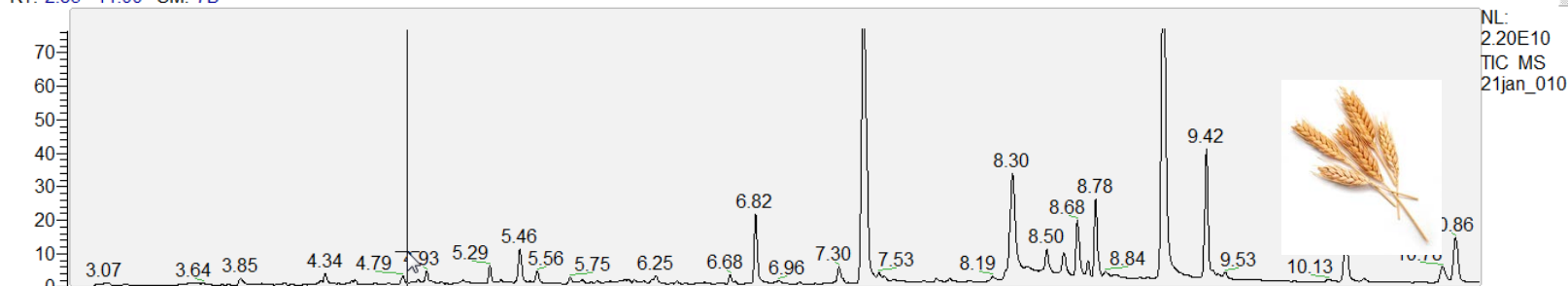
Multiple Libraries
(NIST has over 276,248 spectra
Wiley 719,000 spectra
Wiley MS Designer Drugs, 19,000 spectra
Wiley MS Of Drugs, Poisons, Pesticides
Wiley MS Steroids ...)



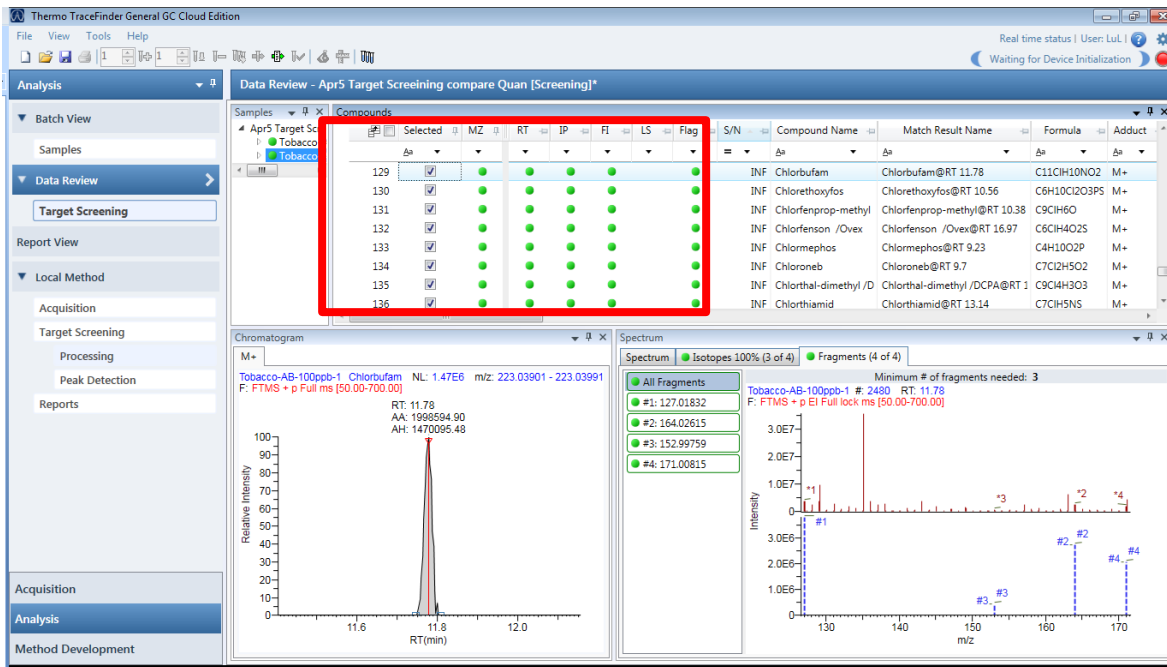
QExactive GC – Full Scan 60,000 RP Chromatography

- Full scan TIC for 55 pesticides spiked into wheat, horse feed and leek extracts.

RT: 2.85 - 11.00 SM: 7B

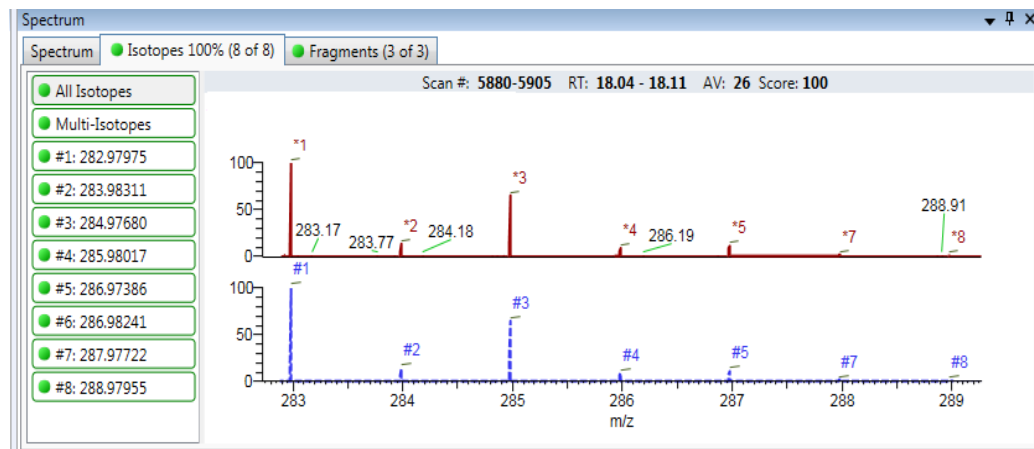


Targeted Screening Review – Pesticides Identification



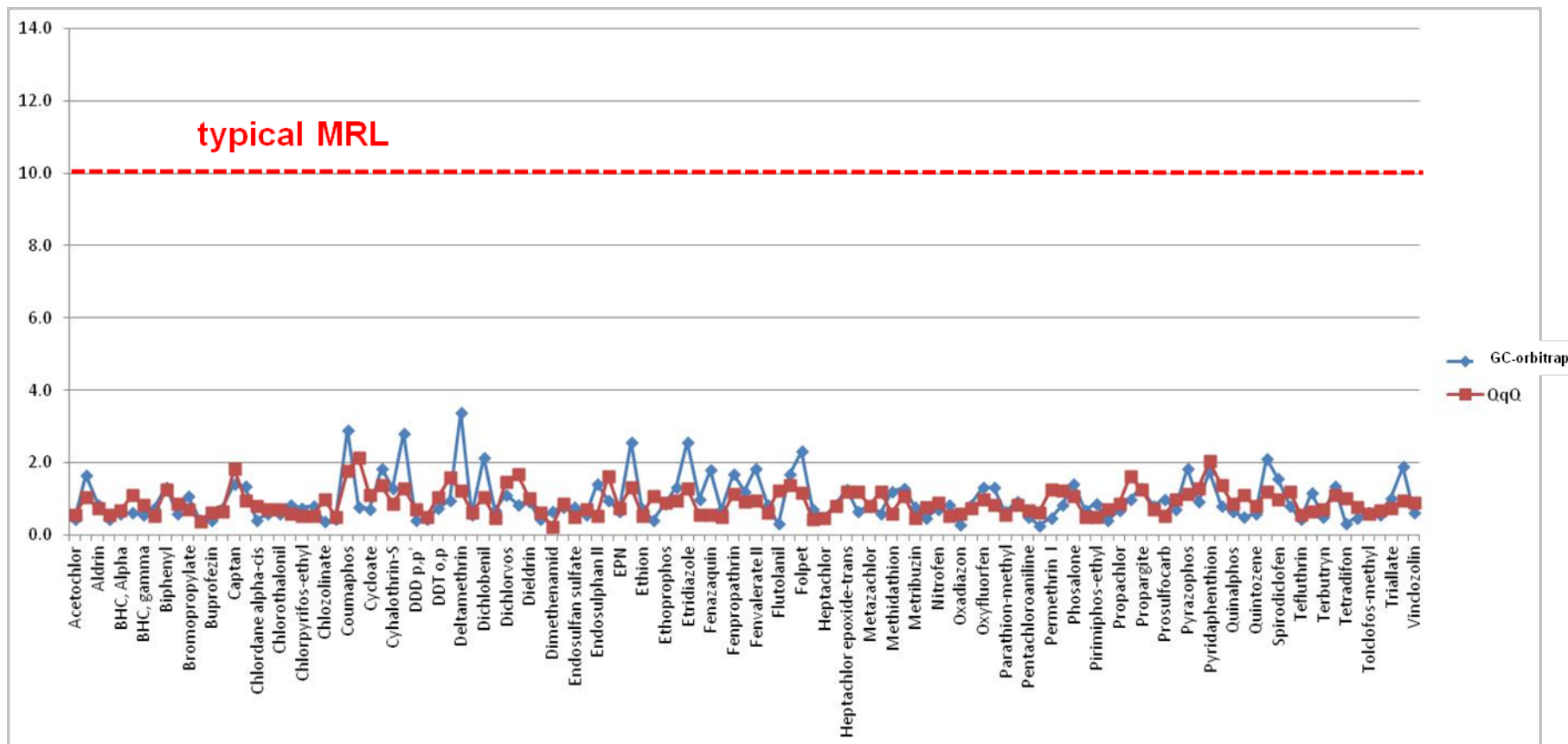
Confirmation Criteria

- 2-5 EI Fragment Ions
- Accurate Tolerance < 2ppm
- Retention Time
- Isotopic pattern

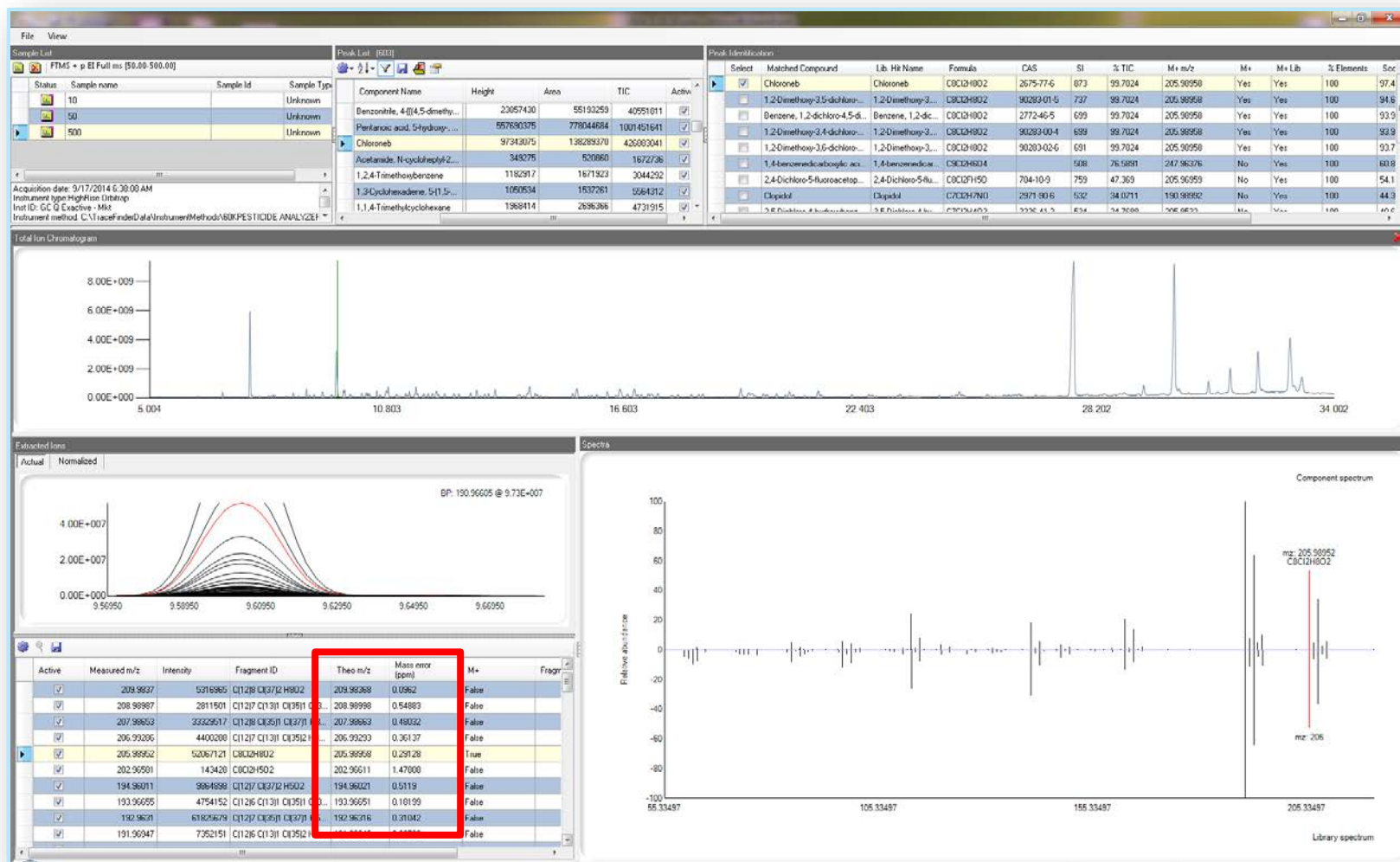


Achieving Pesticides MRL with Full Scan @ 60,000 RP

QE-GC: Full Scan at Resolution of 60,000 Resolving Power QQQ: SRM



EI Spectra Deconvolution & Library Search



Library search compatible with existing unit resolution libraries

Summary – Take Away Message

- **GC-MS/MS** – Cost Efficient Alternative
 - EvoCell technology – High Speed SRM with Precision & Sensitivity
 - Removable ExtractaBrite™ Ion Source - easy maintenance
 - Viable and cost efficient alternative for food & feed analysis
 - Fast - with high throughput capability, automated runs
- **GC-HRMS** – The GOLD Standard Confirmation Solution
 - DFS Magnetic Sector - Compliant with international regulations
 - QExactive GC - New benchmark for powerful unknown compound identification for GC-MS, including POPs analysis
 - Highest sensitivity and precision for low level analyses
- **GC-MS/MS + GC-HRMS** – Work Seamlessly Together
 - High precision data using labelled internal standards
 - Common TargetQuan software platform + LIMS sample manager