

New Possibilities for Targeted and Untargeted Contaminant Analysis in Environmental Samples

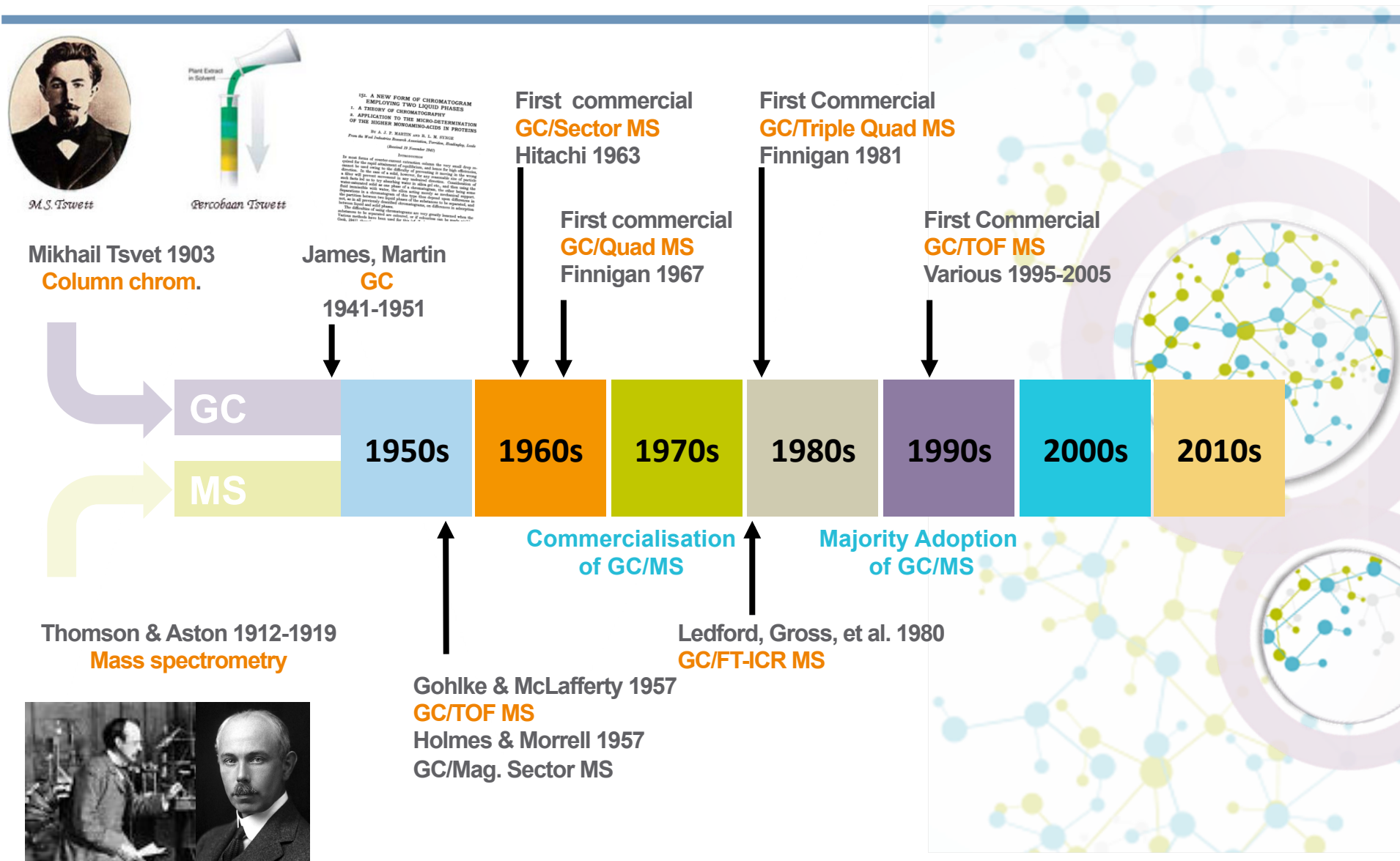
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 The world leader in serving science

GC-MS Until Now



GC-MS Until Now



Challenge

Limited GC-MS analysis

- No comprehensive quantitative & qualitative analysis
- For quantitation: targeted approaches required
- HR/AM Qualitative approaches compromised



Compromised Options

Multi-instrument approach

- Can be inefficient, convoluted and complicated
- GC single and triple quadrupole MS for target quantitation
- GC- Time-of-flight (Tof) MS with limited performance



Breakthrough in GC-MS capability

The Power of Q Exactive GC

Resolution

Up to
120,000 at
m/z 200

- Highest available
- Maximum selectivity
- Fast enough for GC!

Mass Accuracy

< 1ppm

- Every scan
- All concentrations
- In complex matrix
- Across the mass range
- Everyday!

Sensitivity

ppt

- In full scan
- High selectivity
- High spectral fidelity

Dynamic Range

>6 orders

- Excellent coverage in sample profiling
- "Triple quad grade" quantitation in full scan

Breakthrough in GC-MS capability



Challenge

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- HR/AM Qualitative approaches compromised
- For quantitation: targeted approaches required



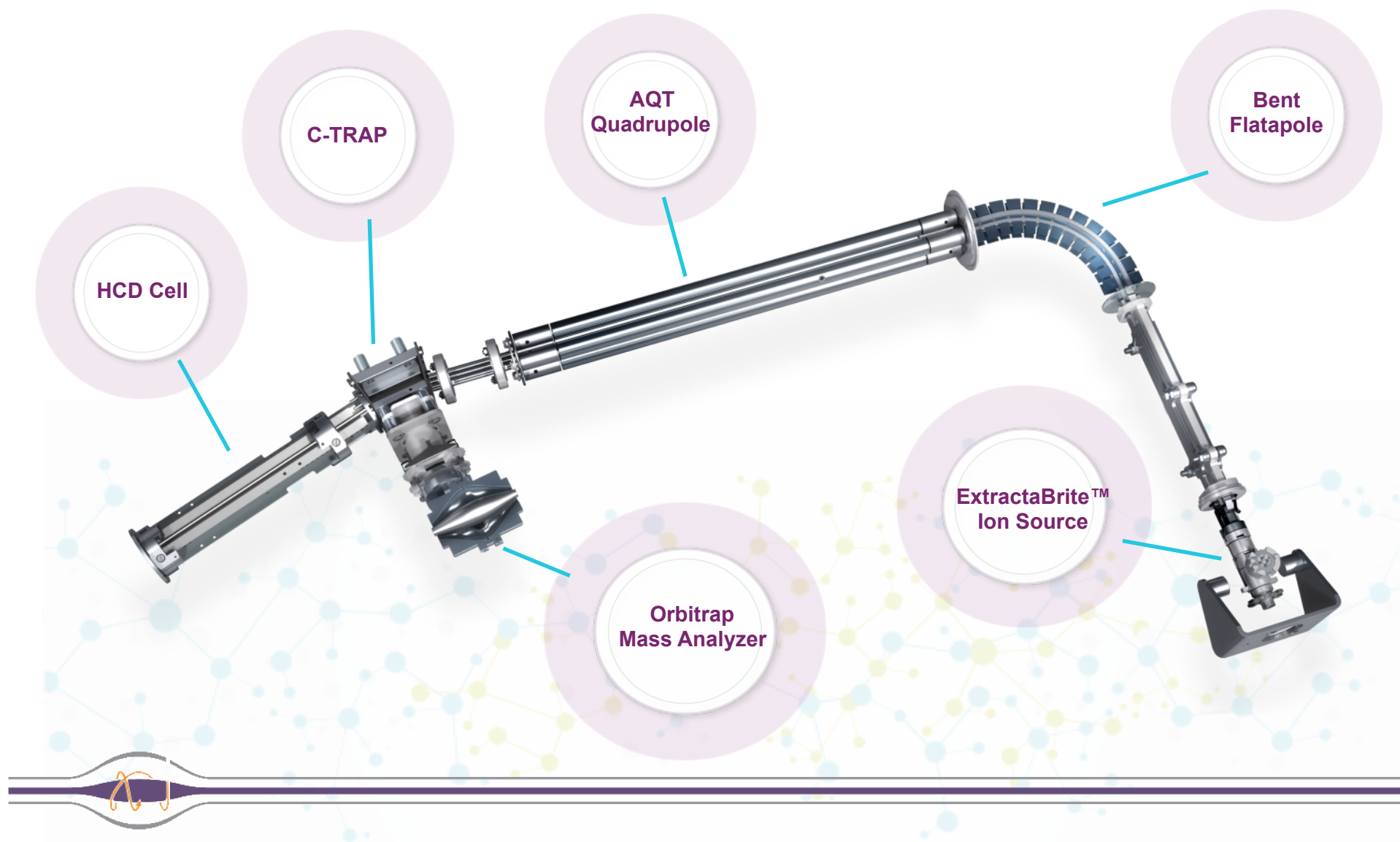
Q Exactive GC

Powerful GC-MS analysis

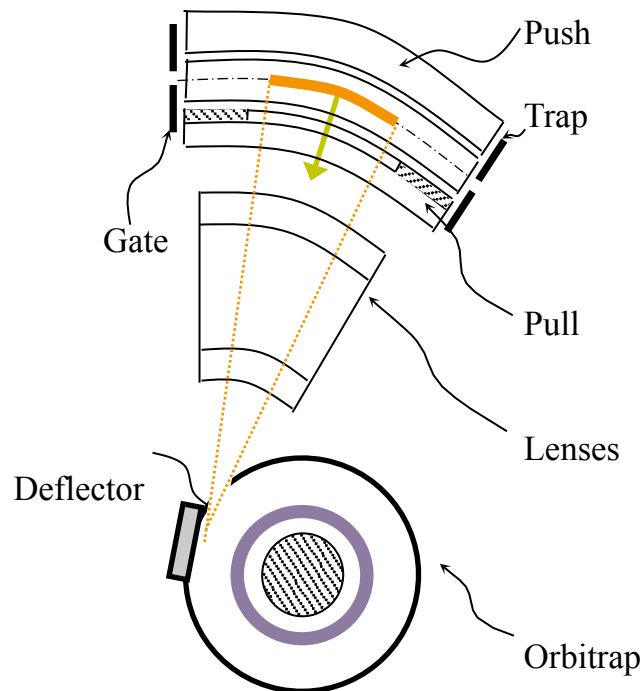
- Easy, full quantitative & qualitative analysis
- Superior qualitative analysis and uncompromised HR/AM performance
- Quantitation as good as best performing GC Triples



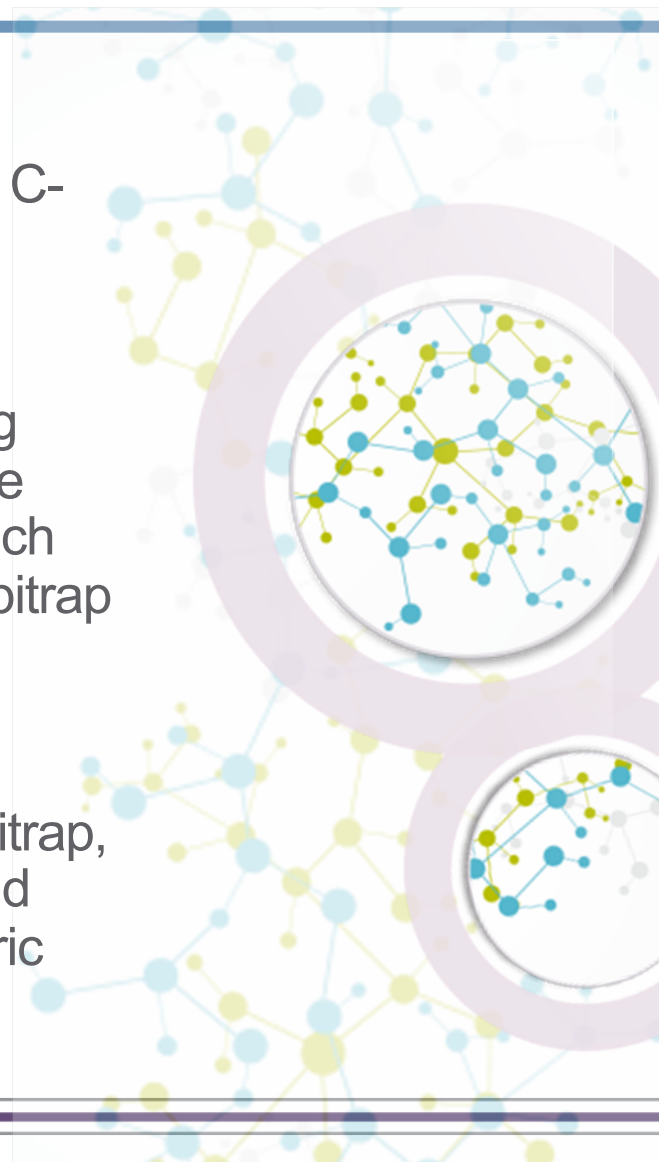
Bringing GC and Orbitrap Technology Together



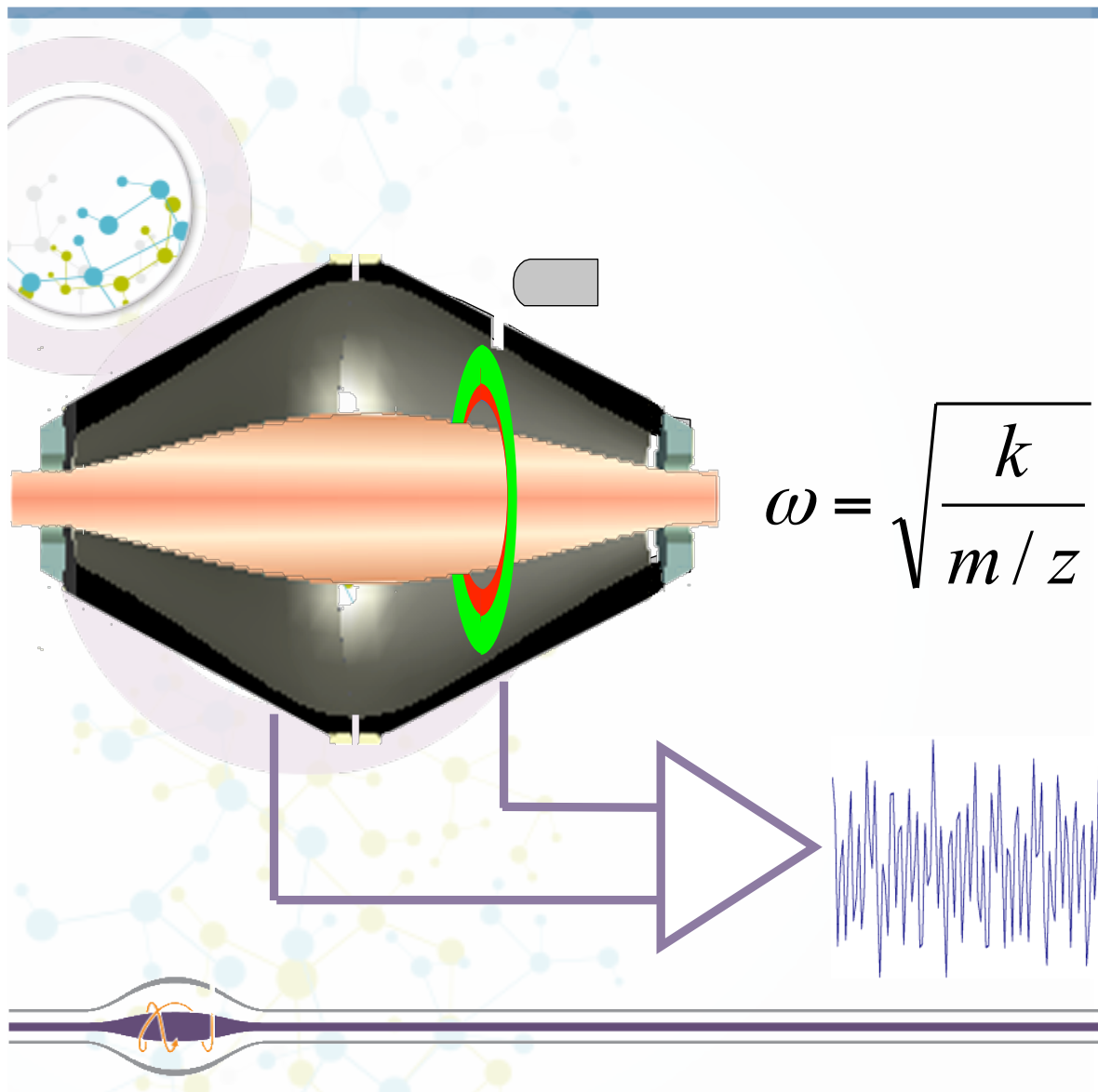
Pulsing Ions into the Orbitrap: Curved Linear Trap (C-trap)



- Ions are stored and cooled in the RF-only C-trap
- Ions are ejected along lines converging to the pole of curvature (which coincides with the Orbitrap entrance).
- As ions enter the Orbitrap, they are picked up and squeezed by its electric field



Retaining the Ions in the Orbitrap



- Frequency of axial oscillations are independent of initial conditions of ions entering trap
- Therefore these oscillations used for mass determination
- Image current measured by outer split electrodes (no electron multiplier to replace!)
- Ion frequencies determine by complex superposition of measured ring oscillations through Fourier transform

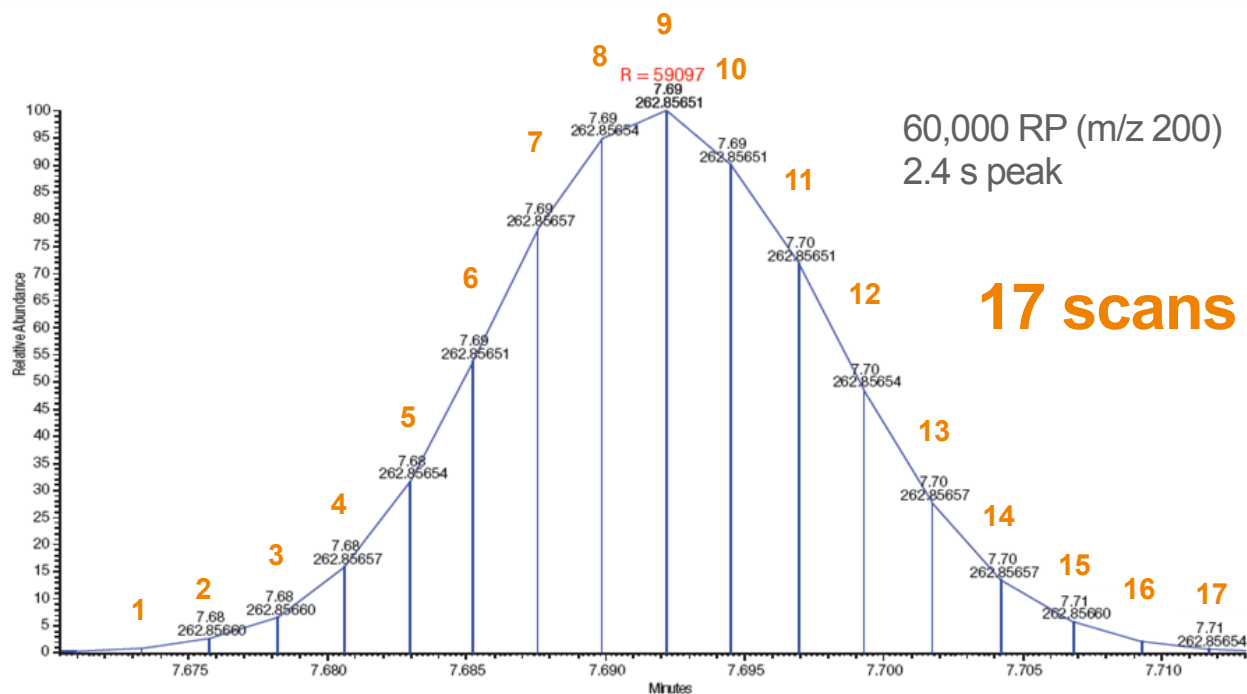
Breakthrough in GC-MS Performance

Highest selectivity and confidence with high resolving power

Fast acquisitions

- Important for accurate profiling of narrow GC peaks
- Full scan with resolving power of 60 (FWHM @ m/z 200) generates **17 scans**
- Fast enough for GC!

XIC of dieldrin in baby food (m/z 262.85642) 100ppb



Breakthrough in GC-MS Performance

Consistently excellent mass accuracy

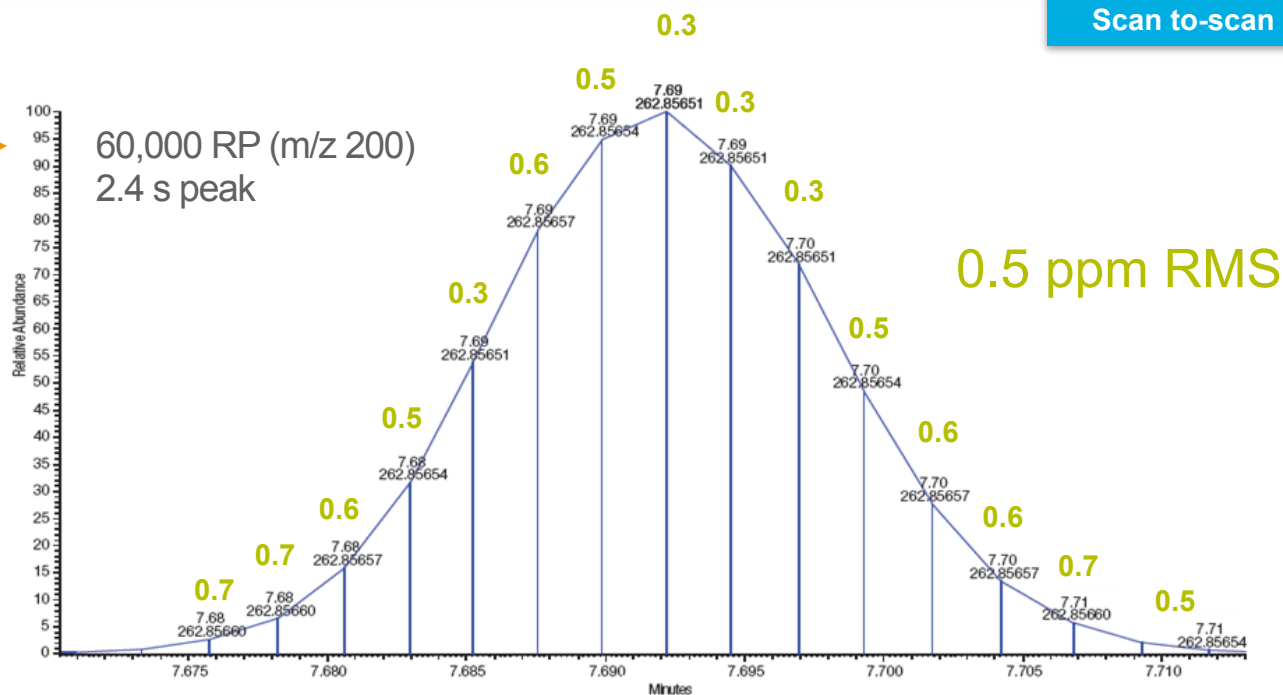
< 1ppm (internal)

< 3ppm (external)

- Scan-to-scan
- Low level in matrix
- Over full mass range
- Over full concentration range
- No need for continuously calibrating in sequence

XIC of dieldrin in baby food (m/z 262.85642) 100ppb

Scan to-scan



Untargeted screening of environmental contaminants: water iodinated disinfection by-products (DBPs)

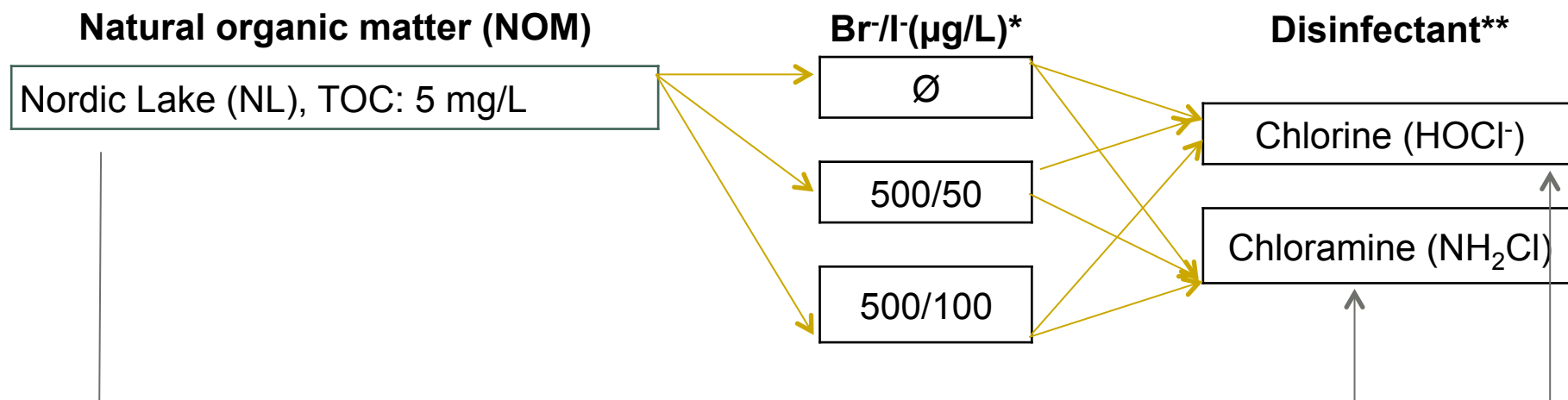
Cristina Postigo Rebollo
Damia Barcelo



Screening of environmental contaminants: water DBPs

- Water DBPs are formed when a chemical disinfectant reacts with the natural organic matter present in water.
- To date more than 600 DBPs have been already identified, and some of them pose a health risk as indicated by epidemiological studies.
- However, about 50% of the total halogenated material formed during disinfection treatments is still unknown.
- Iodo-DBPs formed in chloramination processes are particularly cytotoxic and poorly characterized.
- These compounds are of concern for the environment and human health due to their potential toxicity.

GENERATION OF DBP MIXTURES (I)



*Bromine and iodine were added as KBr and KI, respectively.

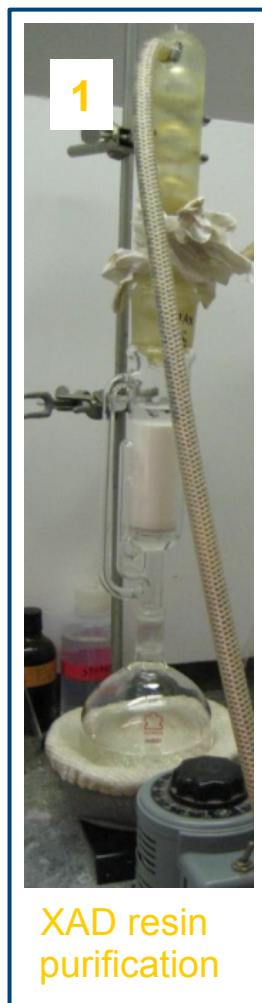
**[Cl₂]=5 mg/L for SR, 4 mg/L for NL, 7.5 mg/L for LLOB, after chlorine demand test.

Reaction conditions:

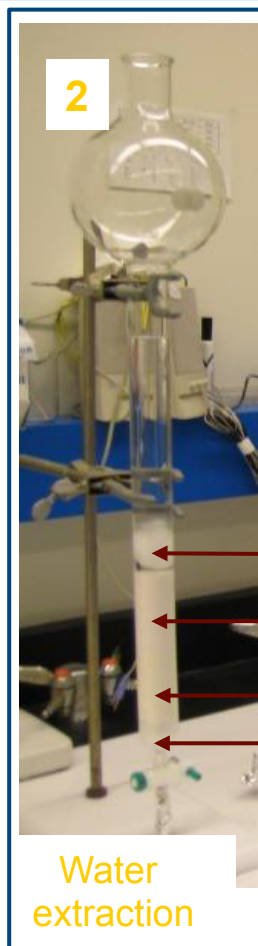
- **room T^a**
- **pH = 7.5** (phosphate buffer)
- **stirring for 72 h**
- **in the dark**
- **volumen = 17 L**

GENERATION OF DBP MIXTURES (II)

Sample extraction – XAD resin extraction



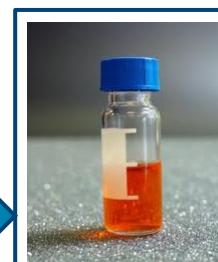
Chlorinated/
chloraminated
water



Glass wool
XAD8 (24 mL)
XAD2 (24 mL)
Glass wool

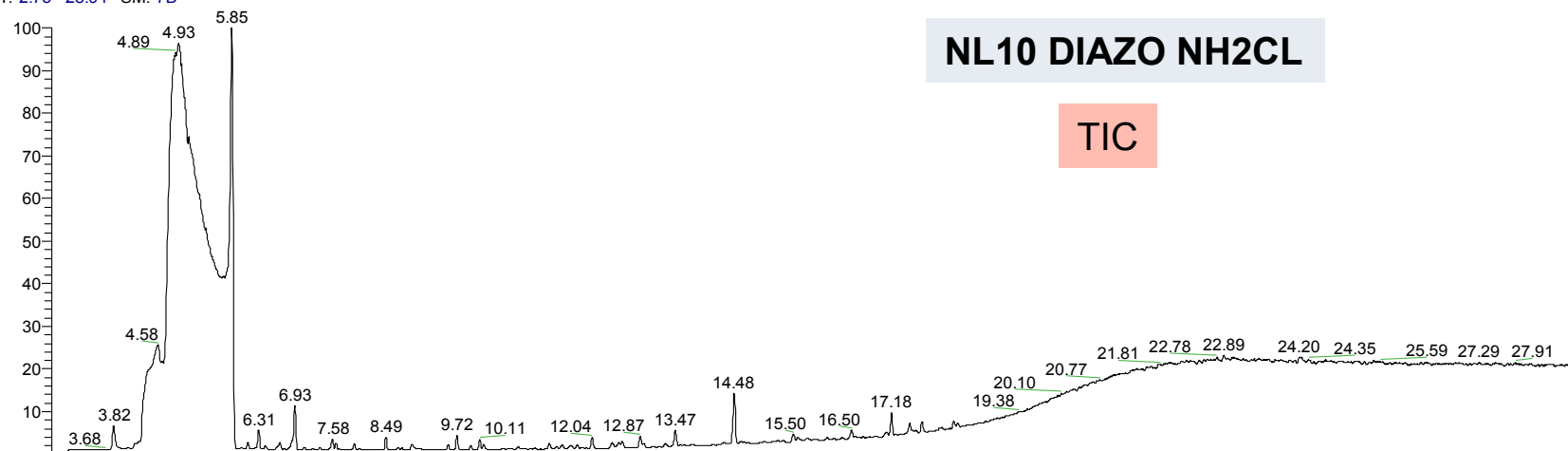
XAD resin extraction

1. Soxhlet extraction for purification of XAD resins (24 h with MeOH, 24 h with ethyl acetate, 24 h with MeOH).
2. Water extraction (16L) onto purified XAD resins.
3. Elution of analytes with ethyl acetate (224 mL).
4. Rotary evaporation of ethyl acetate extract (140 rpm, 35 °C).
5. Evaporation under N₂ until 0.8 mL.
6. Diazomethane derivatization of half of the extract.

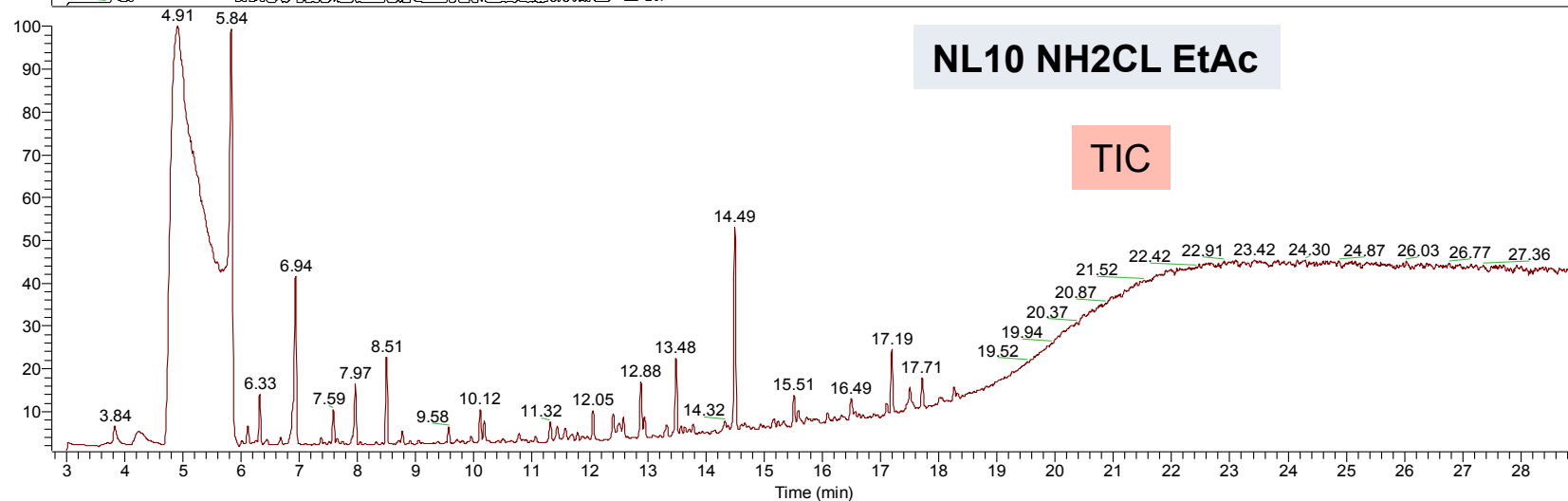


Samples were screened for iodine-DBPs

RT: 2.75 - 28.94 SM: 7B



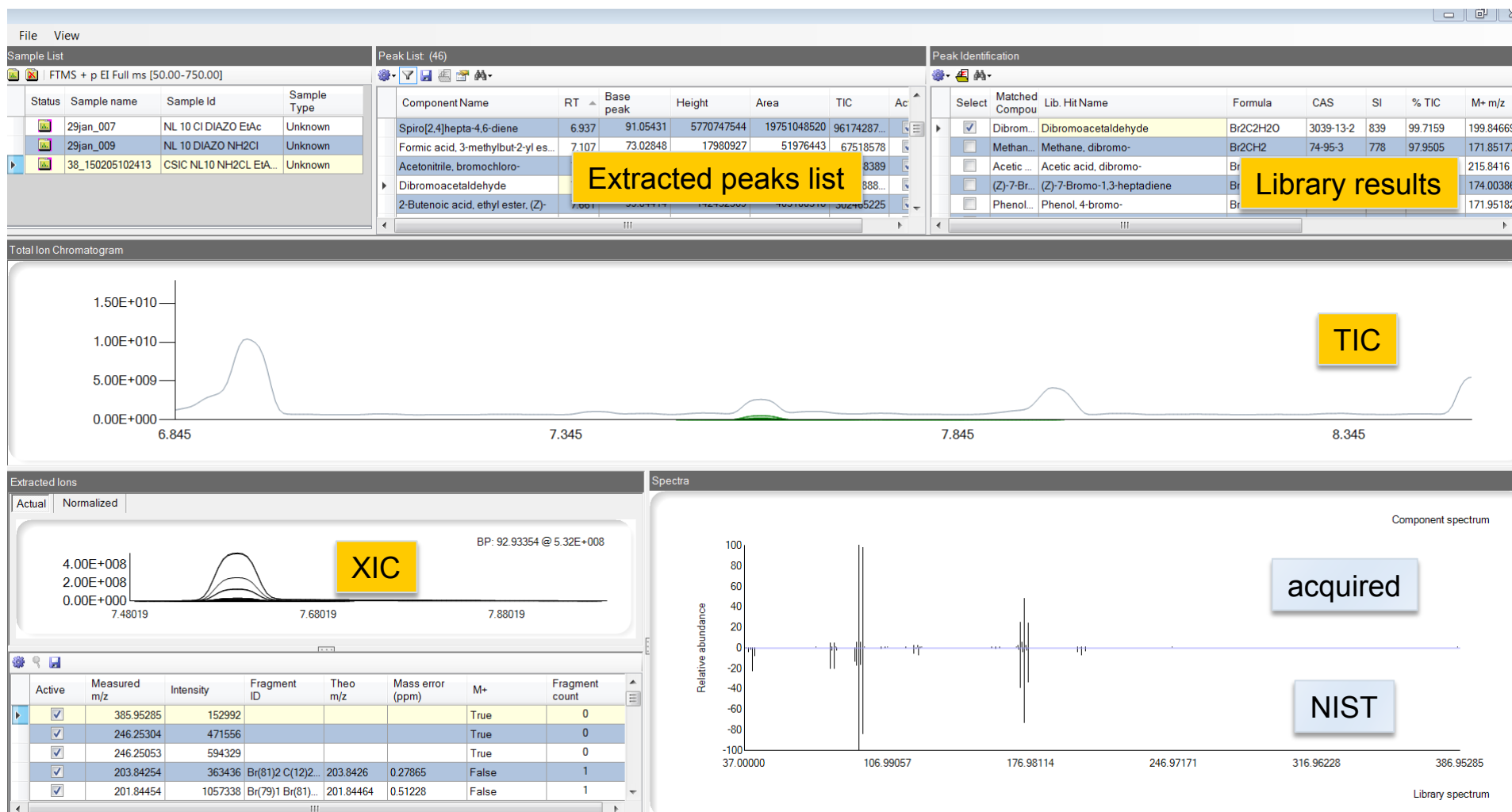
NL:
3.17E10
TIC MS
39_150205
114306



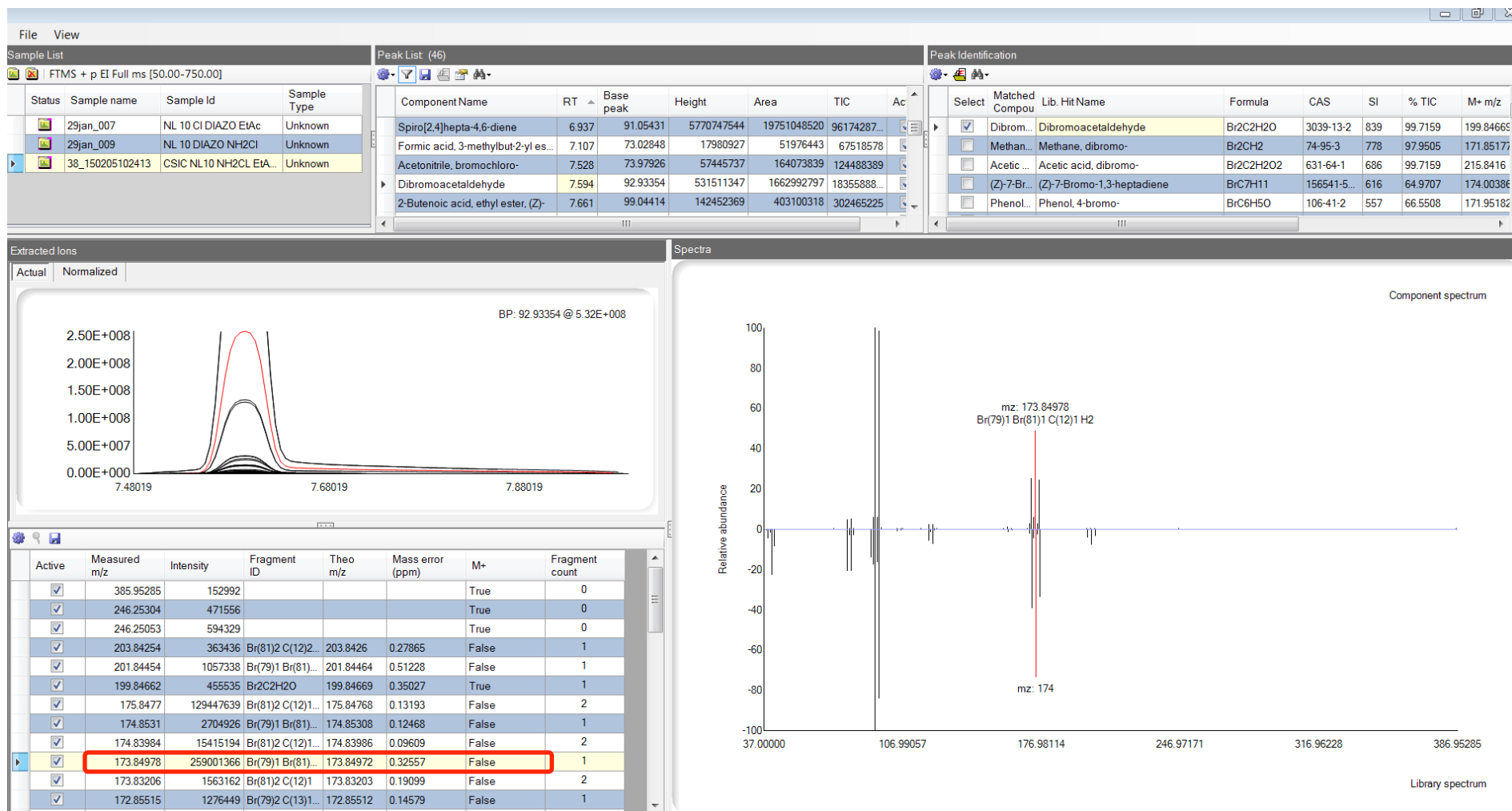
NL:
2.97E10
TIC MS
38_150205
102413

Deconvolution software

Results Browser



Dibromoacetaldehyde: intelligent candidate verification



Library search compatible with existing unit res libraries

Example of compounds found in sample NL10 NH2CL EtAc (NIST ID)

25 unique compounds identified in the treated sample compared to blank

| Component Name | RT | m/z | Area |
|--|-------|-----------|------------|
| Methane, bromo- | 3.84 | 93.94136 | 2287289010 |
| Ethyl bromide | 4.19 | 109.95492 | 70878448 |
| Dichloroacetaldehyde | 4.21 | 83.95286 | 239462033 |
| Methane, iodo- | 4.26 | 141.92743 | 6835098508 |
| Methane, bromo- | 4.32 | 93.94136 | 79510322 |
| Methane, bromodichloro- | 6.45 | 82.94501 | 231940325 |
| Acetonitrile, dichloro- | 6.47 | 81.9372 | 27542835 |
| Acetonitrile, bromochloro- | 7.53 | 73.97926 | 164073839 |
| Methane, dibromo- | 7.59 | 92.93354 | 1662992797 |
| Acetonitrile, dibromo- | 8.78 | 119.92674 | 784206464 |
| Ethyl iodoacetate | 9.58 | 185.91718 | 1060392837 |
| 1-Bromo-3,3,3-trifluoroacetone | 10.25 | 122.92638 | 17442292 |
| 3,4-Dibromo-1-pentene | 11.31 | 67.05429 | 428945278 |
| Benzene, 1-bromo-2,3-dimethyl- | 11.44 | 105.06999 | 495067921 |
| Threo-1,4-dibromo-3-pentanol | 11.51 | 136.95973 | 103169909 |
| Benzoic acid | 11.58 | 105.0336 | 1259211272 |
| Methane, bromodiiodo- | 11.70 | 218.83012 | 288558451 |
| Acetic acid, bromochloro-, methyl ester | 12.45 | 128.89227 | 510614169 |
| Benzene, 1-bromo-2,4,5-trimethyl- | 12.87 | 119.08565 | 2366794432 |
| Benzene, 2-bromo-4-methyl-1-(1-methylethyl)- | 13.57 | 196.99606 | 331797666 |
| Phenol, 2,4-dibromo- | 13.78 | 251.86038 | 582559994 |
| Phenol, 3-(2-bromoethyl)- | 14.39 | 199.98312 | 32854422 |
| Benzene, 2,4-dibromo-1,3,5-trimethyl- | 15.51 | 196.99606 | 854581173 |
| Phenol, 2,4,6-tribromo- | 16.49 | 329.77087 | 637209104 |
| Phthalic acid, 6-ethyl-3-octyl butyl ester | 18.26 | 149.02339 | 1402969802 |

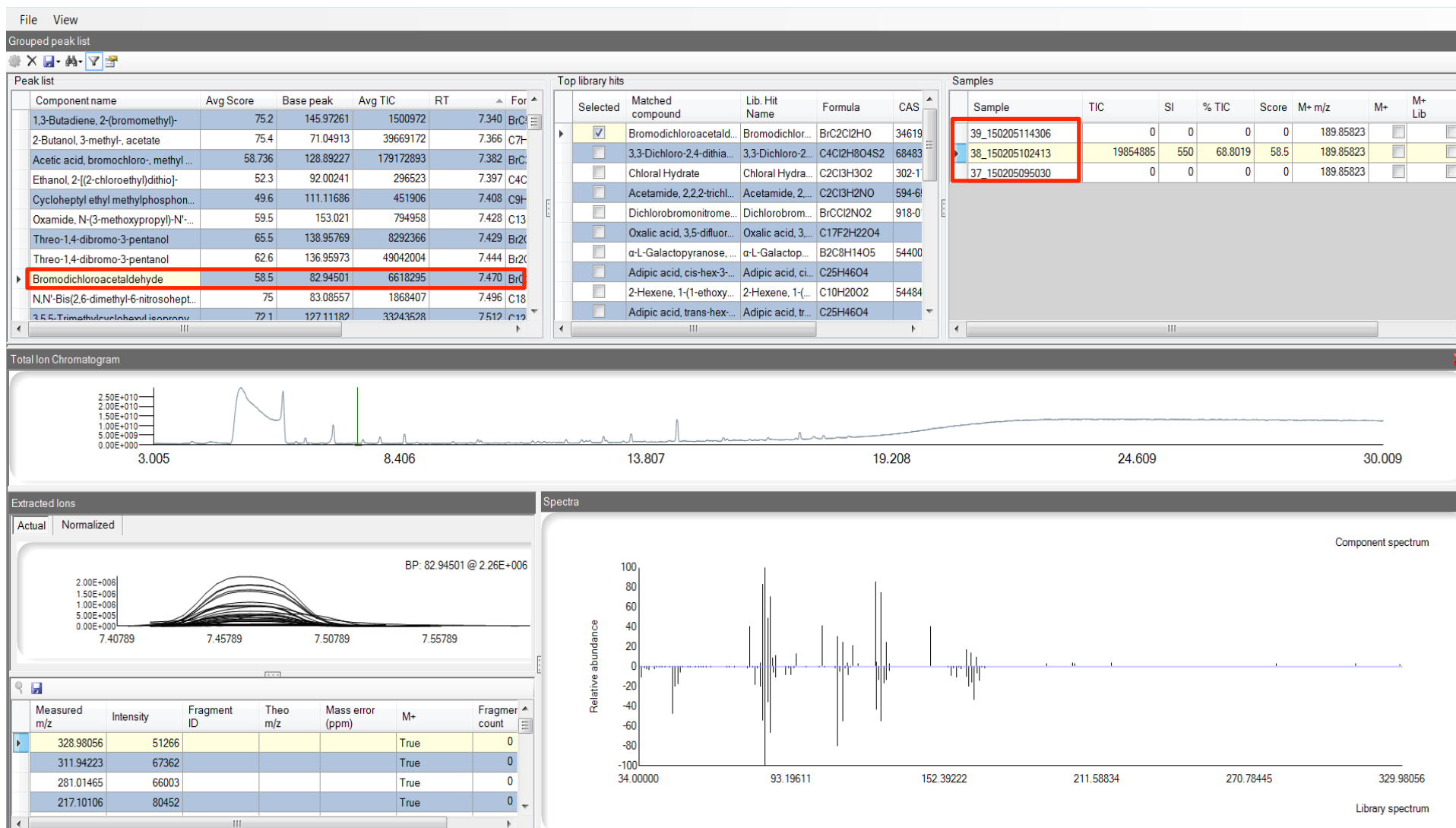
Example of compounds found in sample NL10 DIAZO NH2CL (NIST ID)

23 unique compounds identified in the treated sample compared to blank

| Component Name | RT | m/z | Area |
|--|----------|----------|----------|
| Methane, bromo- | 3.81893 | 93.94133 | 2.59E+09 |
| Methane, iodo- | 4.18849 | 141.9274 | 1.01E+09 |
| Methane, bromo- | 6.12199 | 93.94133 | 3.83E+08 |
| Propanoic acid, ethyl ester | 6.30544 | 74.03629 | 1.3E+09 |
| Methane, dibromo- | 6.40918 | 92.93354 | 24929829 |
| 2-Pentyne, 4,4-dimethyl- | 6.41953 | 81.06994 | 2.54E+08 |
| Chloriodomethane | 6.61803 | 175.8884 | 1.22E+08 |
| 2-Pentyne, 4,4-dimethyl- | 6.81009 | 81.06994 | 13107733 |
| Propanoic acid, 2-methyl- | 7.09094 | 88.05194 | 68333595 |
| Methane, dibromo- | 7.57956 | 92.93354 | 5.45E+08 |
| Acetonitrile, dibromo- | 8.77223 | 117.9288 | 1.31E+08 |
| Ethyl iodoacetate | 9.56759 | 185.9173 | 3.22E+08 |
| Methyl sorbate | 10.90216 | 95.0492 | 4954951 |
| Benzene, 1-bromo-2,3-dimethyl- | 11.43655 | 105.07 | 75604516 |
| Benzoic acid | 11.52464 | 105.0336 | 4.08E+08 |
| Methane, bromodiiodo- | 11.69547 | 218.8301 | 1.06E+08 |
| Benzene, 1-(1,1-dimethylethyl)-4-ethoxy- | 11.76807 | 151.1118 | 12353169 |
| Benzene, 2-bromo-1,3,5-trimethyl- | 12.57058 | 119.0857 | 51473818 |
| 2,6-Diisopropylanisole | 13.06213 | 119.0857 | 14988011 |
| Benzene, 2-bromo-4-methyl-1-(1-methylethyl)- | 13.62064 | 196.9961 | 72300603 |
| Phenol, 2,4-dibromo- | 13.7705 | 251.8604 | 1.61E+08 |
| Benzene, 3-bromo-1,2,4,5-tetramethyl- | 14.65367 | 133.1012 | 98075372 |
| 2,4-Di-tert-butylphenol | 14.70524 | 191.143 | 80705238 |

Direct data comparison

Presence/absence of a chromatographic peak in all the sample analysed with abundance information



Quantification of environmental contaminants: NDMA



Experimental

- Data was acquired at 60,000 resolution (FWHM @ m/z 200).
- GC column: TG-200 ms, 30m x 0.25 mm x 0.25 μ m
- TraceFinder software for automated data acquisition and processing.

TRACE 1310 GC Parameters

| | |
|--------------------------------|-----------------------------------|
| Injection Volume (μ L): | 1.0 |
| Liner | Single gooseneck carbofrit packed |
| Inlet ($^{\circ}$ C): | 220 |
| Inlet Module and Mode: | Splitless |
| Oven Temperature | |
| Program: | |
| Temperature 1 ($^{\circ}$ C): | 35 |
| Hold Time (min): | 1 |
| Temperature 2 ($^{\circ}$ C): | 130 |
| Rate ($^{\circ}$ C/min) | 25 |
| Temperature 3 ($^{\circ}$ C): | 230 |
| Rate ($^{\circ}$ C/min) | 125 |
| Hold Time (min): | 1 |

Mass Spectrometer Parameters

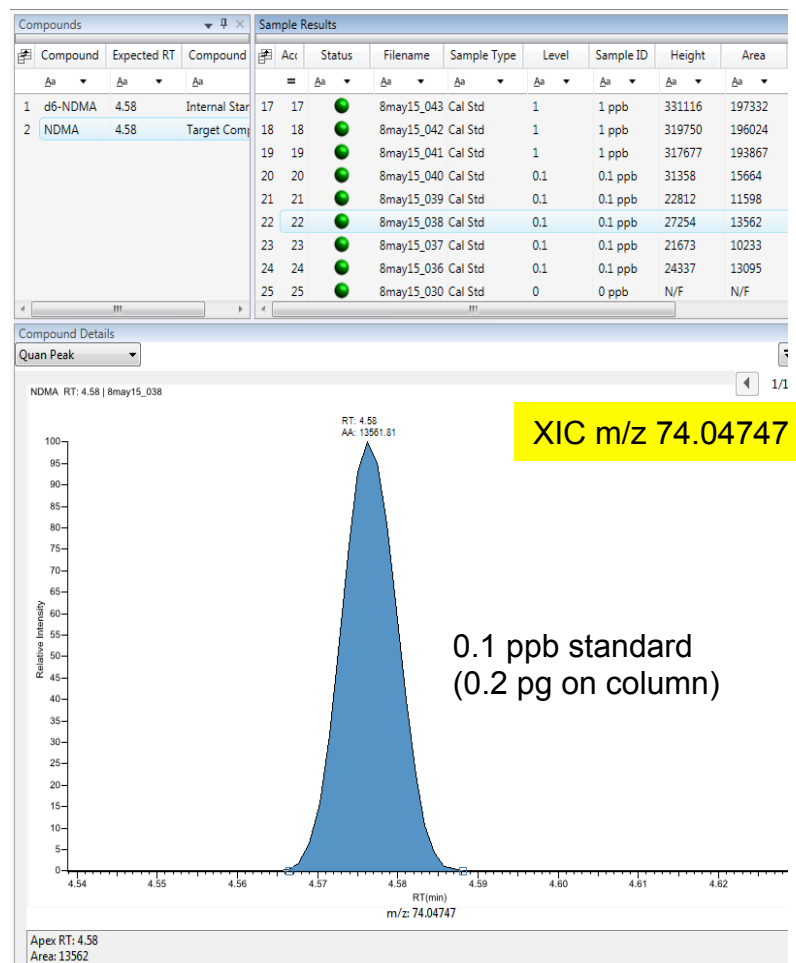
| | |
|--------------------------------|-----------------|
| Transfer line ($^{\circ}$ C): | 260 |
| Ionization type: | El |
| Ion source($^{\circ}$ C): | 230 |
| Electron energy (eV): | 70 |
| Acquisition Mode: | Full Scan & SIM |
| Mass range (m/z): | 50-750 |
| Lockmass (m/z): | 207.03235 |

Samples

- Solvent standards (in dichloromethane) each spiked with 20 pg/ μ L (ppb) d6-NDMA:
 - 0 ppb
 - 0.01 ppb
 - 0.1 ppb
 - 1 ppb
 - 10 ppb
 - 100 ppb

Chromatography @ 0.1 ppb level

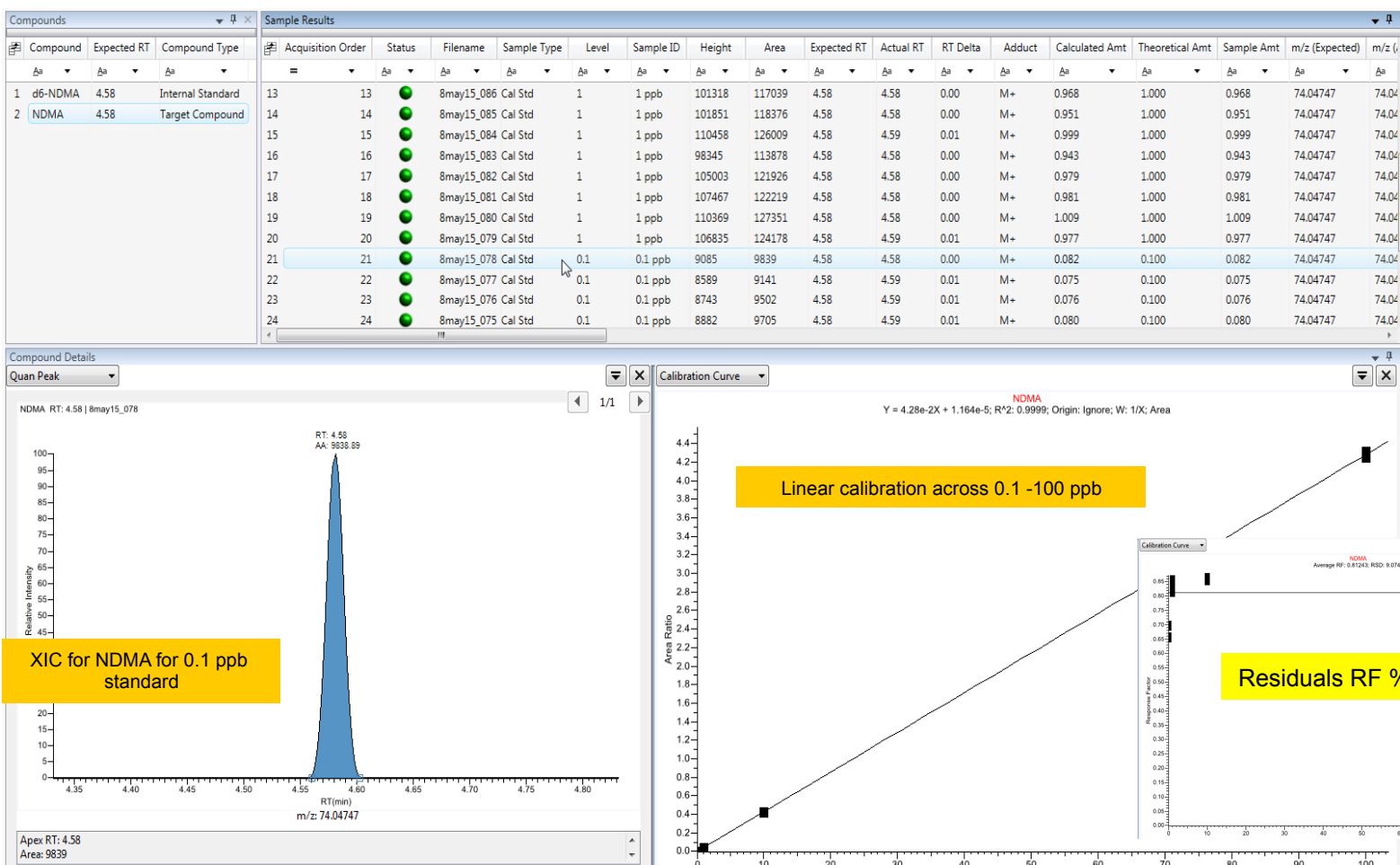
Full scan



TraceFinder browser

Linearity: 0.1 – 100 ppb

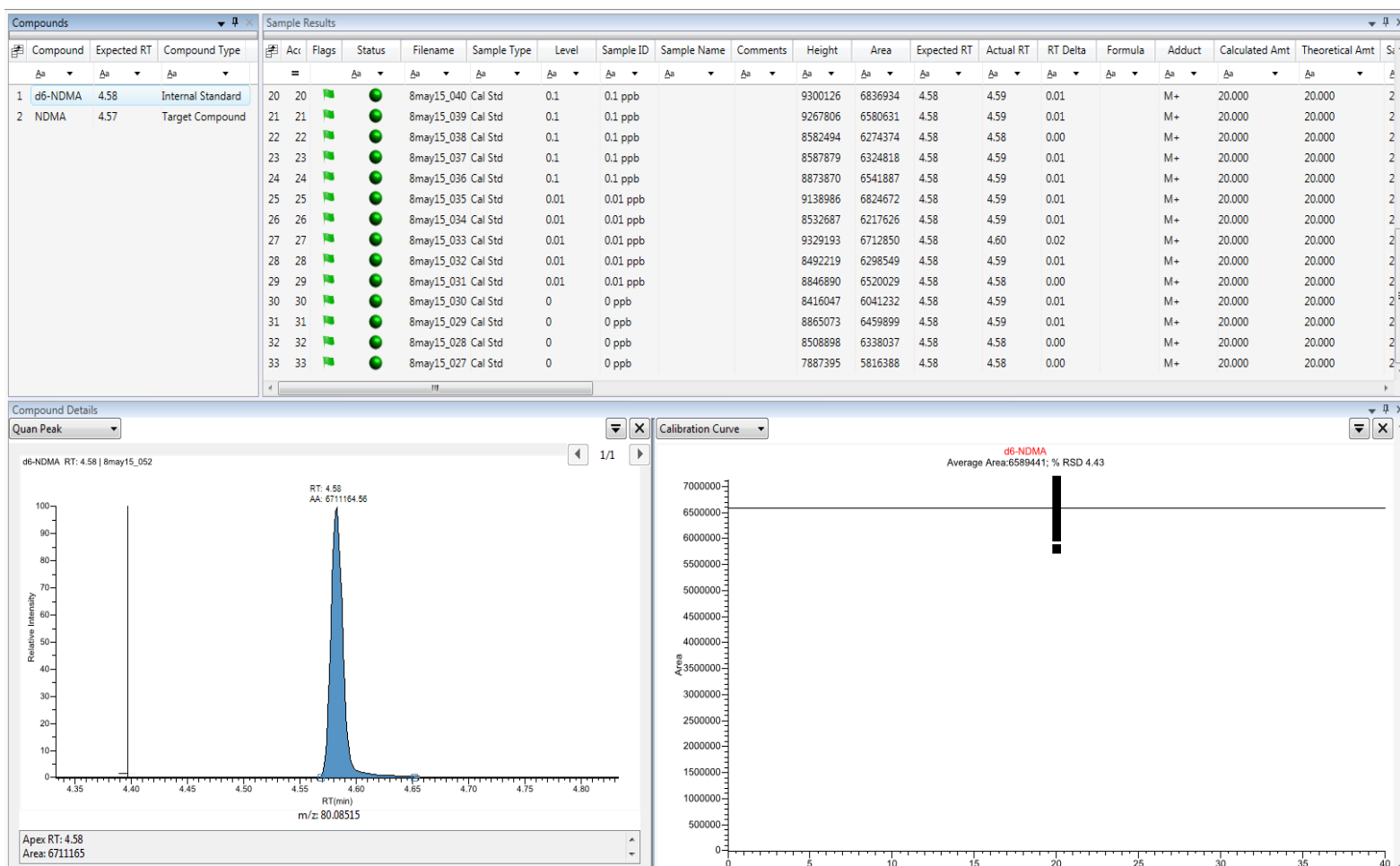
- Linearity was assessed across 0.1 – 100 ppb. d6-NDMA used as internal standard.
- Residuals RF %RSD is also shown (bottom right)



Internal standard (d6-NDMA)

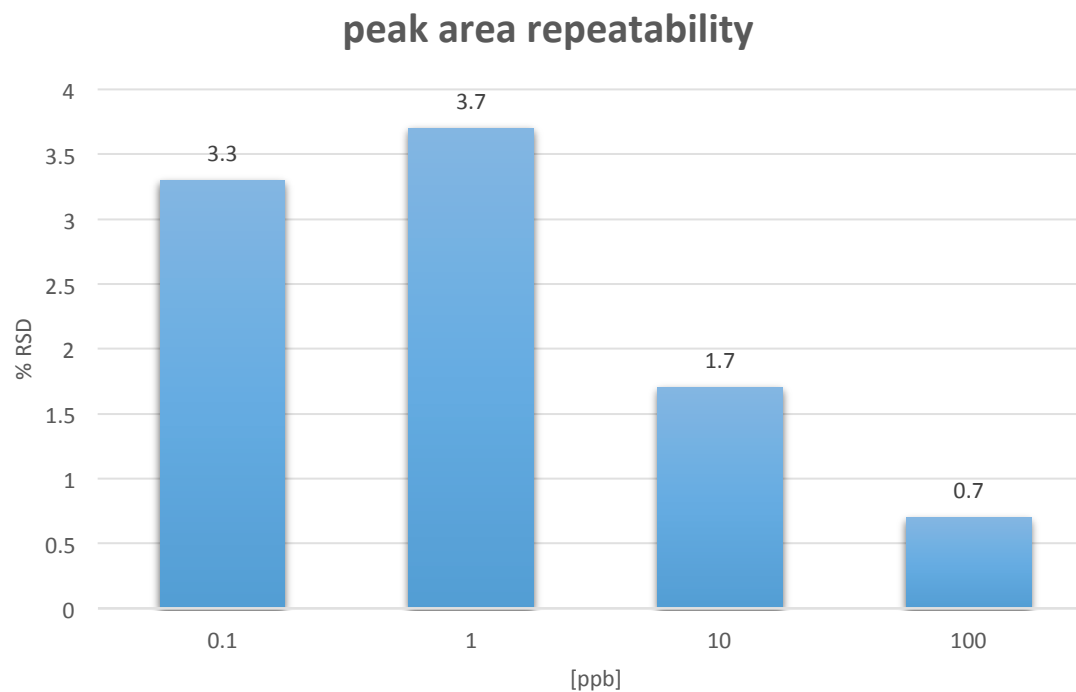
Repeatability of peak area response

- Calculated d6-NDMA peak area %RSD for n = 33 injections: **4.4%**



Peak area repeatability

- No manual adjustment of peak integration was used.

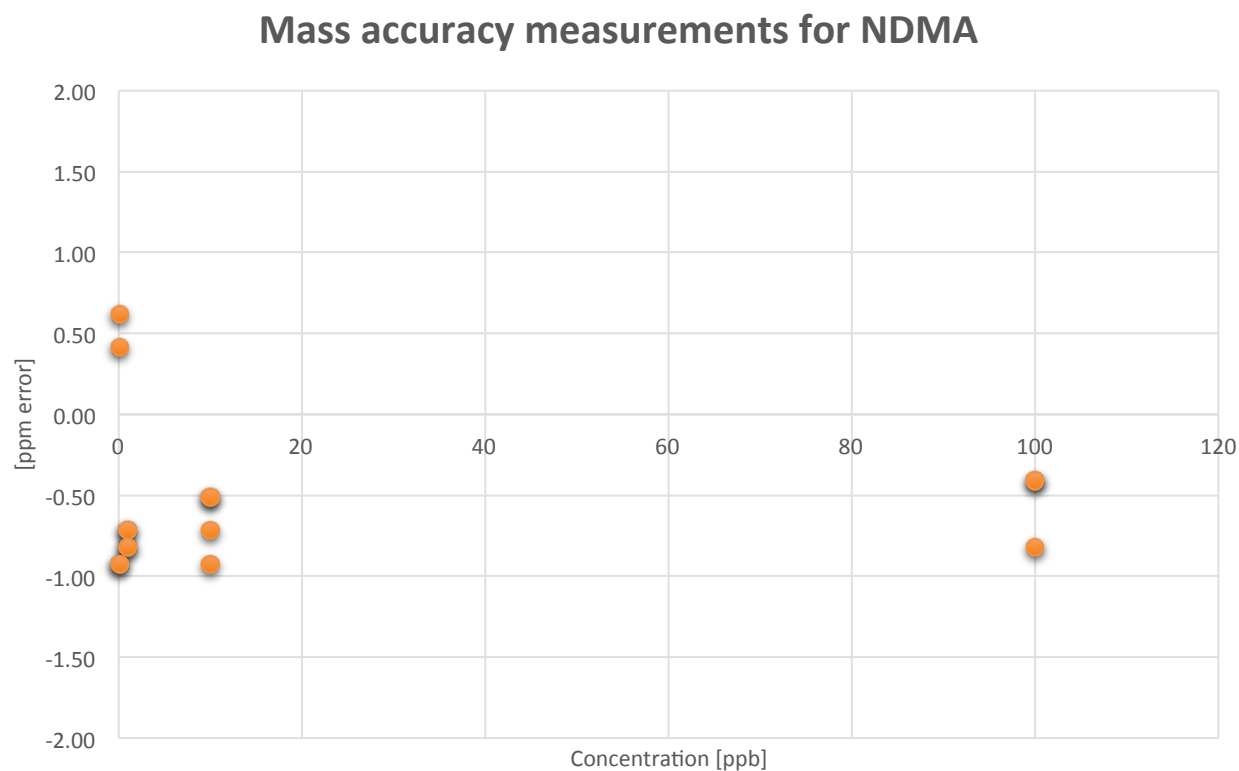


* *n=5 injections per calibration standard were used except 1.0 ppb level; where n=9 inj. were used*

SIM data

Mass accuracy measurements for m/z 74.04747 (NDMA ion)

- Mass error [ppm] across concentration range. Each dot represent a single injection.
- NDMA exact mass: m/z 74.04747.



Thermo Scientific Q Exactive GC Hybrid Quadrupole-Orbitrap GC-MS/MS System

www.thermoscientific.com/QExactiveGC



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