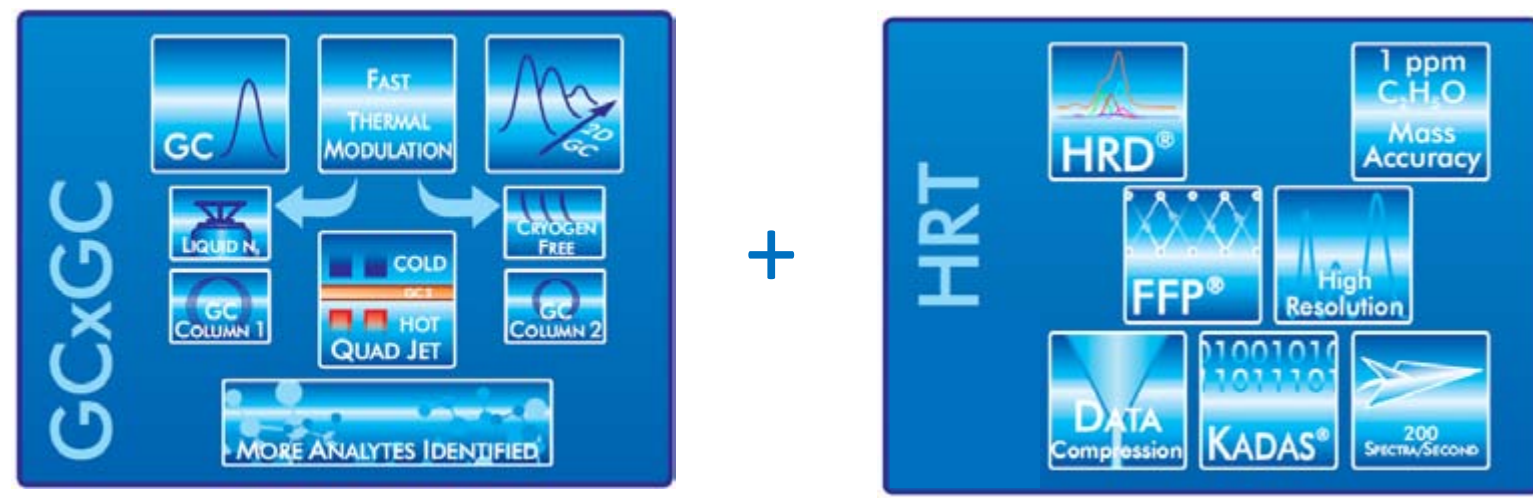


The Use of Accurate Mass Peak Filters to Detect and Identify New Disinfection Byproducts in Swimming Pools and Spas

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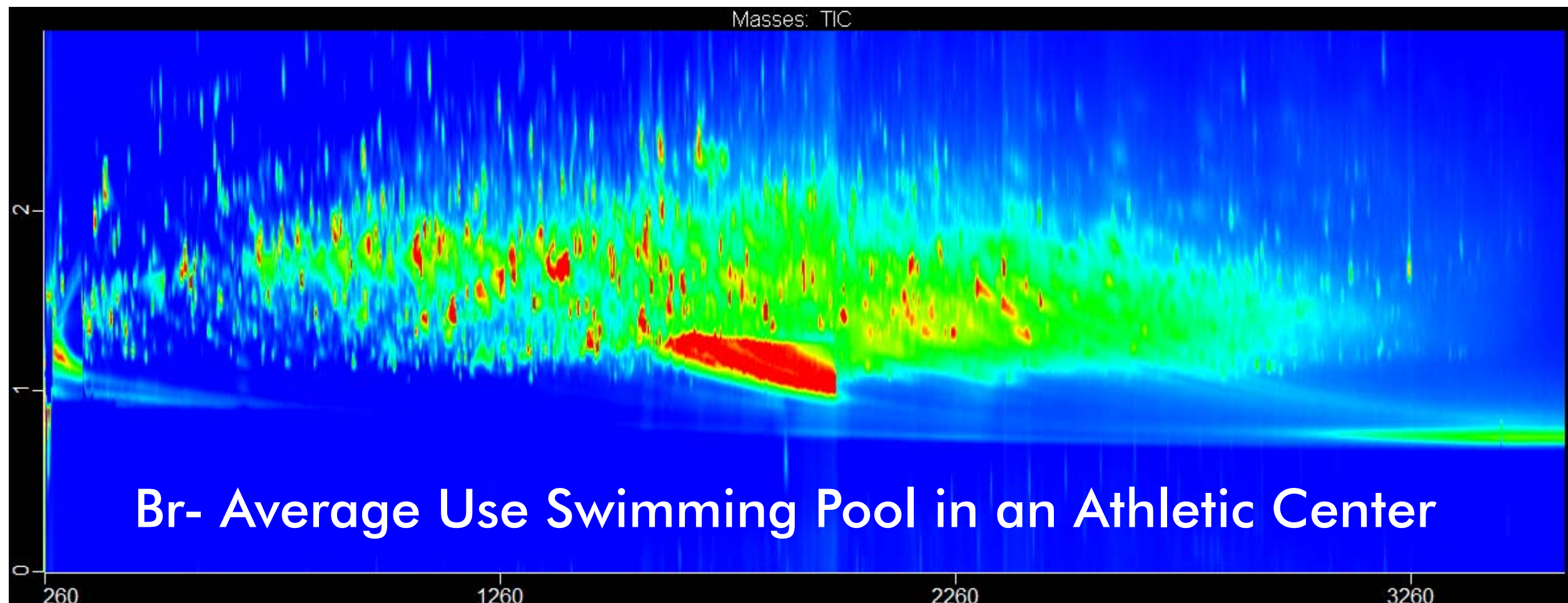
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Sample Workflow



Data Acquisition

Collect electron ionization (EI) or chemical ionization (CI) data in high resolution (Resolving Power = 25K FWHH) or ultra high resolution (RP = 50K FWHH) mode at an acquisition rate of up to 200 spectra/second. Combines LECO's market-leading GCxGC technology with high resolution time-of-flight mass spectrometry, all in one integrated platform and software package.



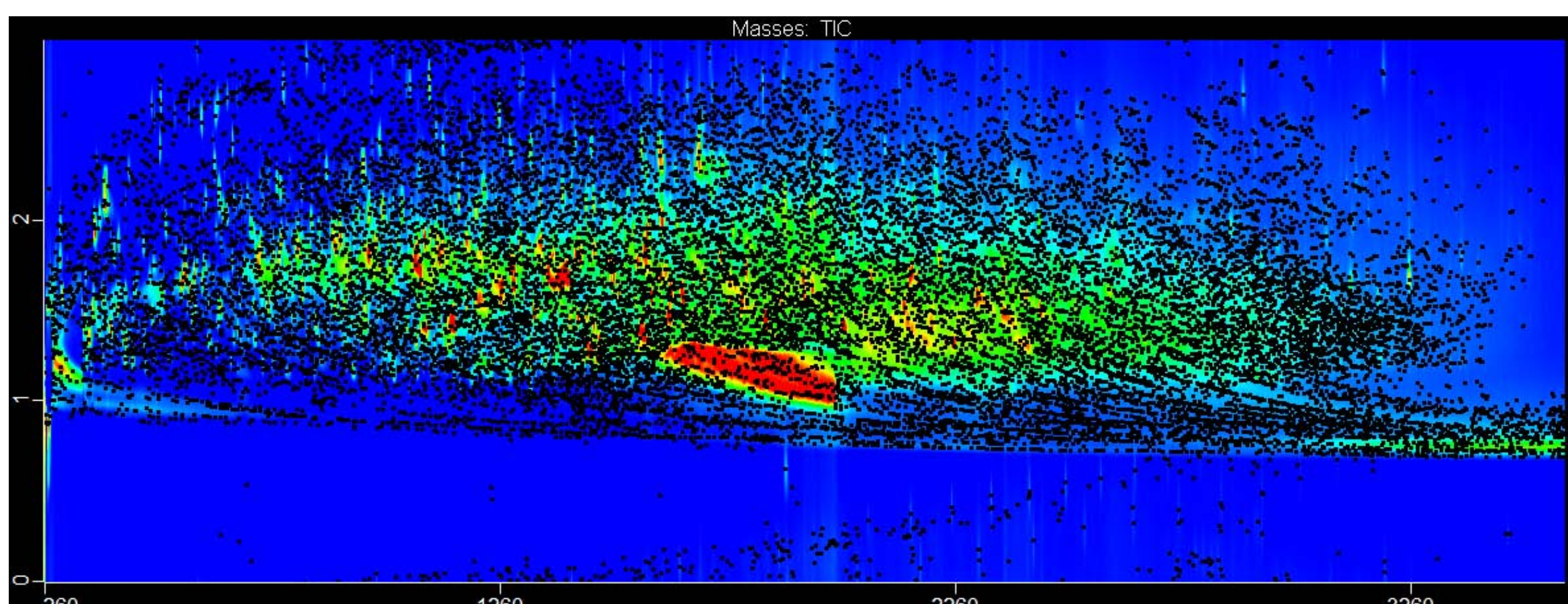
Mass Calibration

A reference compound such as perfluorotributylamine may be introduced throughout the chromatographic run to enhance the mass accuracy with post-acquisition mass calibration. If the TOF drift correction is stable throughout the run, then a region without analyte interference, such as during the hold time at the end of the run, may be used for post-acquisition mass calibration. The mass calibration table demonstrates a mass accuracy RMS = 0.47 ppm.

Name	Formula	Species	Charge	Mass Window	Start Time (s)	End Time (s)	Priority	Expected M/Z	Observed M/Z	Mass Delta (mDa)	Mass Accuracy (ppm)	Resolution
1*	Carbon dioxide	CO ₂	1	0.5000	3396	3510	1	43.9893	43.9893	0.05	1.03	18055
2	PFTBA069	CF ₃	1	1.0000	3396	3510	1	68.9947	68.9946	-0.02	-0.32	23900
3	PFTBA131	CF ₃	1	1.0000	3396	3510	1	130.9915	130.9914	-0.04	-0.32	24619
4	PFTBA219	CF ₃	1	1.0000	3396	3510	1	218.9851	218.9851	0.02	0.09	25964
5	PFTBA264	C ₂ F ₅ N	1	2.0000	3396	3510	1	263.9866	263.9867	0.12	0.47	24492
6	PFTBA414	C ₂ F ₅ N	1	2.0000	3396	3510	1	413.9770	413.9769	-0.06	-0.14	21318
7	PFTBA502	C ₂ F ₅ N	1	2.0000	3396	3510	1	501.9706	501.9705	-0.11	-0.23	21311

Automatic Peak Finding with Mass Spectral HRD[®]

Industry leading High Resolution Deconvolution[®] (HRD) identifies chromatographic peaks automatically. In this swimming pool treated with bromine as the disinfectant, more than 10,000 individual chromatographic peaks were indicated; more than 10x the number of peaks detected in the 1D chromatogram. About 3% of the peaks had a library similarity score greater than 850 and a mass accuracy ±2 ppm, indicating that the large majority of the peaks in the sample were unknowns.



Accurate Mass Peak Filters

A compound class(es) may be identified by leveraging unique features in its mass spectrum; e.g. accurate mass, isotope pattern, and/or fragmentation pattern. Spectral peak filters were designed for Br₂, Cl₂, Br₂Cl₂, and I containing compounds; where x = 1 to 5 and y = 1 to 7. The I⁺ ion was targeted using m/z 126.9039 ± 5 ppm.

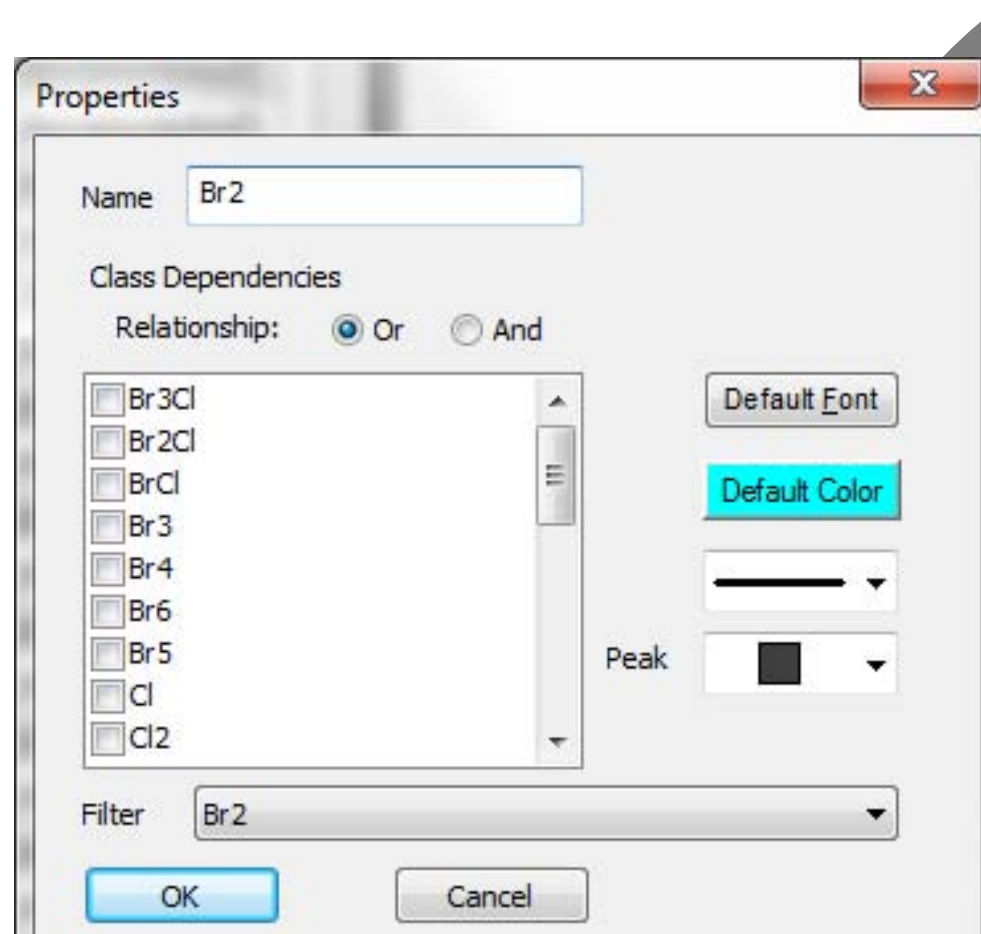
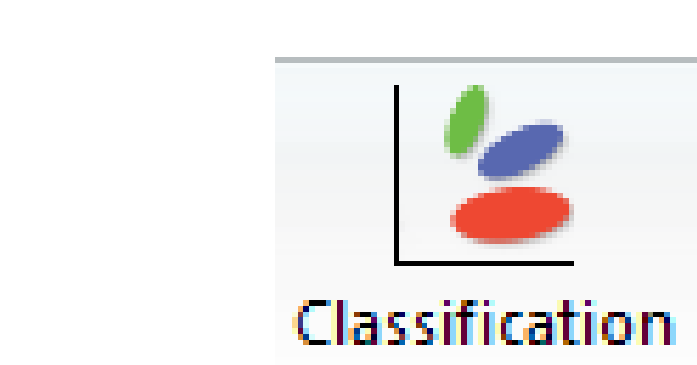
Spectral Peak Filter for Br₂ compound:

- Intensity of Any Monoisotope is > 100
- AND
- Abundance of Any Monoisotope is > 44% of abundance of Any Monoisotope ± m/z 1.9979±0.0015Da
- AND
- Abundance of Any Monoisotope is < 58% of abundance of Any Monoisotope ± m/z 1.9979±0.0015Da
- AND
- Abundance of Any Monoisotope ± m/z 3.9985±0.0015Da is > 42% of abundance of Any Monoisotope ± m/z 1.9979±0.0015Da
- AND
- Abundance of Any Monoisotope ± m/z 3.9985±0.0015Da is < 56% of abundance of Any Monoisotope ± m/z 1.9979±0.0015Da
- AND
- Abundance of Any Monoisotope ± m/z 1.0028±0.0015Da is < abundance of Any Monoisotope
- AND
- Abundance of Any Monoisotope ± m/z 5.9937±0.0015Da is < 2% of abundance of Any Monoisotope ± m/z 1.9979±0.0015Da
- AND
- Intensity of Any Monoisotope + Br±0.0015Da is < 20
- AND
- Intensity of Any Monoisotope + Br₂±0.0015Da is < 20
- AND
- Intensity of Any Monoisotope + Cl±0.0015Da is < 20
- AND
- Intensity of Any Monoisotope + Cl₂±0.0015Da is < 20

Classification Table

Create a classification table for the compound class(es) that the spectral filters were designed to identify, inserting the appropriate filter into the properties option for the class. One or more summary columns may be added for each class. In this case, the summary will be based on the total area of all of the ions in the deconvoluted (peak true) spectra for all of the chromatographic peaks of each class.

#	Show	Class	Component	Region	Color	dTIC
1	<input checked="" type="checkbox"/>	Br2C1			Yellow	tic(deconvoluted)
2	<input checked="" type="checkbox"/>	Br2C2			Red	tic(deconvoluted)
3	<input checked="" type="checkbox"/>	Br2C3			Green	tic(deconvoluted)
4	<input checked="" type="checkbox"/>	Br2C4			Blue	tic(deconvoluted)
5	<input checked="" type="checkbox"/>	Br2C5			Purple	tic(deconvoluted)
6	<input checked="" type="checkbox"/>	Br2C6			Orange	tic(deconvoluted)
7	<input checked="" type="checkbox"/>	Br2C7			Pink	tic(deconvoluted)
8	<input checked="" type="checkbox"/>	Br2C8			Light Blue	tic(deconvoluted)
9	<input checked="" type="checkbox"/>	Cl			Light Green	tic(deconvoluted)
10	<input checked="" type="checkbox"/>	Cl2			Light Purple	tic(deconvoluted)
11	<input checked="" type="checkbox"/>	Cl3			Light Orange	tic(deconvoluted)
12	<input checked="" type="checkbox"/>	Cl4			Light Yellow	tic(deconvoluted)
13	<input checked="" type="checkbox"/>	Cl5			Light Blue	tic(deconvoluted)
14	<input checked="" type="checkbox"/>	Cl6			Light Green	tic(deconvoluted)
15	<input checked="" type="checkbox"/>	Cl7			Light Purple	tic(deconvoluted)
16	<input checked="" type="checkbox"/>	I			Light Orange	tic(deconvoluted)
17	<input checked="" type="checkbox"/>	Br			Light Green	tic(deconvoluted)
18	<input checked="" type="checkbox"/>	Br2Cl2			Light Purple	tic(deconvoluted)
19	<input checked="" type="checkbox"/>	Br2Cl3			Light Orange	tic(deconvoluted)
20	<input checked="" type="checkbox"/>	Br2Cl4			Light Yellow	tic(deconvoluted)
21	<input checked="" type="checkbox"/>	Br2Cl5			Light Blue	tic(deconvoluted)
22	<input checked="" type="checkbox"/>	Br2Cl6			Light Green	tic(deconvoluted)
23	<input checked="" type="checkbox"/>	Br2Cl7			Light Purple	tic(deconvoluted)
24	<input checked="" type="checkbox"/>	Br2Cl8			Light Orange	tic(deconvoluted)
25	<input checked="" type="checkbox"/>	Br2Cl9			Light Green	tic(deconvoluted)
26*	<input checked="" type="checkbox"/>	BrCH			Light Purple	tic(deconvoluted)
27	<input checked="" type="checkbox"/>	Unclassified			Light Orange	tic(deconvoluted)
28	<input checked="" type="checkbox"/>	Unclassified			Light Green	tic(deconvoluted)



Unknown Identification



A putative identification can be made for an unknown compound based on the mass spectral interpretation of accurate mass data. After the elemental composition of the molecular ion was determined below, ChemSpider was launched directly from ChromTOF-HRT[®] brand software to search for compounds matching the chemical formula assigned. Of the 16 results in the database, only two compounds had an appropriate structure that corresponded with the observed mass spectrum. Note that they are positional isomers.

ChemSpider

Search term: C₈H₄Br₂N₂O₂ (Found by molecular formula)

ID	Structure	Molecular Formula	Molecular Weight	# of Data Sources	# of References	# of PubMed	# of RSC
13789183		C ₈ H ₄ Br ₂ N ₂ O ₂	319.9376	5	5	0	0
13789325		C ₈ H ₄ Br ₂ N ₂ O ₂	319.9376	1	1	0	0

Elemental Composition Determined from Monoisotopic Mass

Elements from which to build the formula:

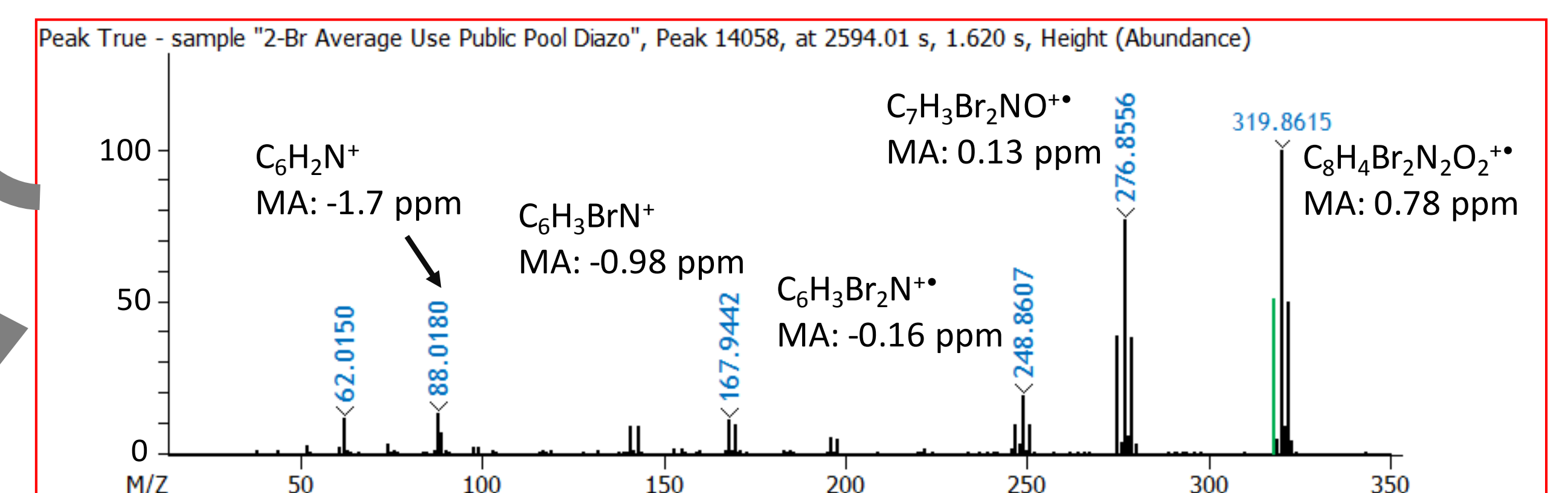
Element	Mass	Minimum	Maximum	Valences
C*	12.0000	1	50	4
H	1.0078	0	100	1
Br	79.9183	2	2	1
Cl	34.9689	0	10	1
I	126.9045	0	10	1
N	14.0031	0	10	3
O	15.9949	0	10	2
S	31.9721	0	10	2

Search Mass: 317.863516

Neutral Mass m/z

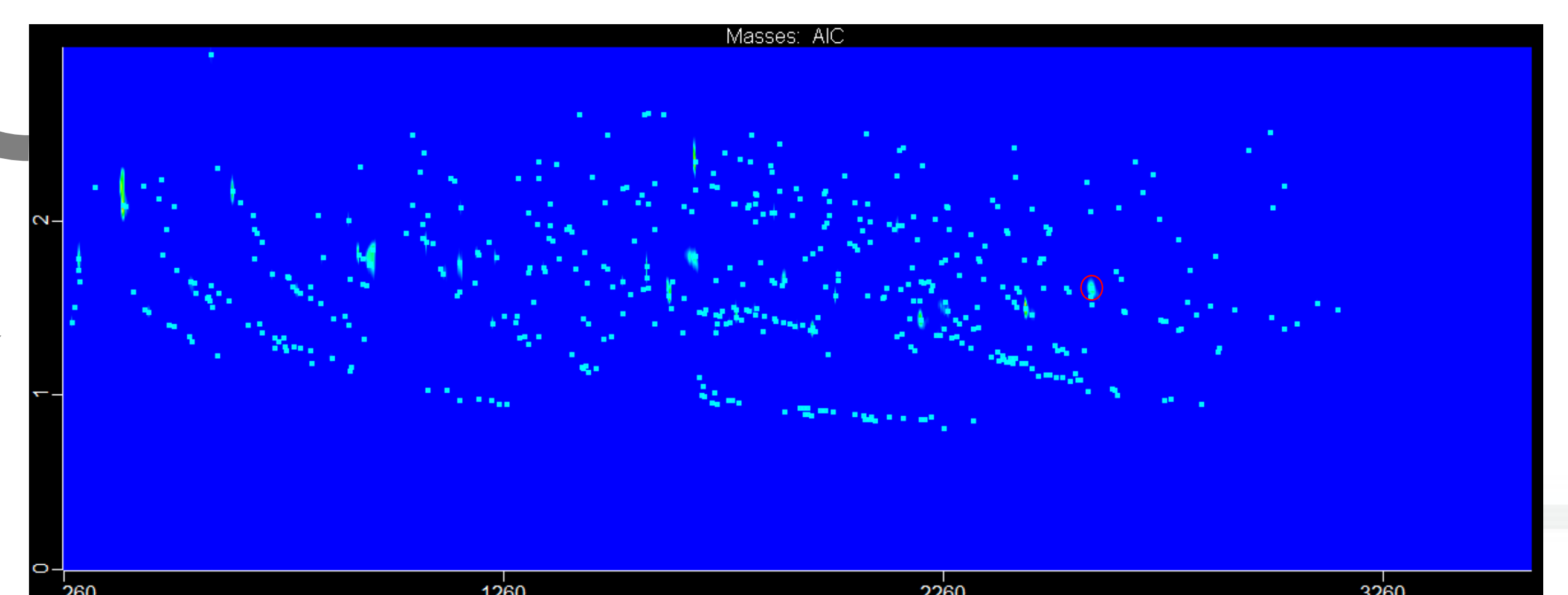
Primary Selection Criteria: mDa Isotope Pattern Mass Difference

Hit	Formula	Expected Ion m/z	Observed Ion m/z	Mass Accuracy (ppm)	Similarity	RDBE
>1	C ₈ H ₄ Br ₂ N ₂ O ₂	317.8634	317.8637	0.78	993	7
	2-C ₈ H ₄ Br ₂ N ₂ O ₂	317.8641	317.8637	1.90	951	3
	3-C ₈ H ₄ Br ₂ N ₂ O ₂	317.8639	317.8637	0.75	824	2.5



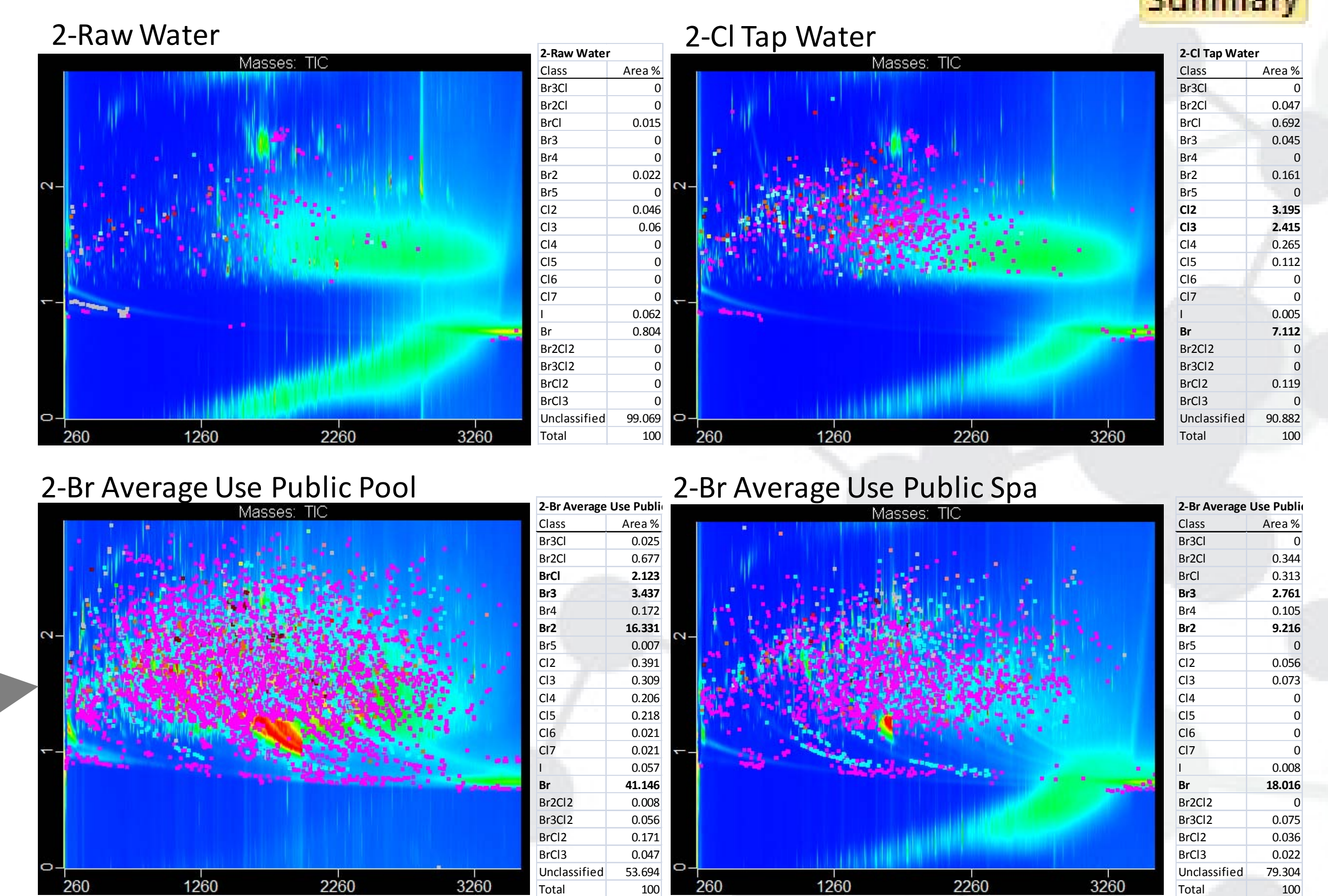
Classification with Peak Filters: Br₂ Class Compounds

Analytical ion chromatogram (AIC) showing only the peaks that were classified as containing 2-Br₂, defined by the spectral filters (almost 500 peaks). Notice how even though the TIC is extremely complex, the AIC shows high quality chromatographic peaks determined automatically by the software. If we select a peak, in this case one of the most intense peaks by area, we can view the deconvoluted mass spectrum and library search results. Notice the high quality of the mass spectrum from a very complex region of the chromatogram.



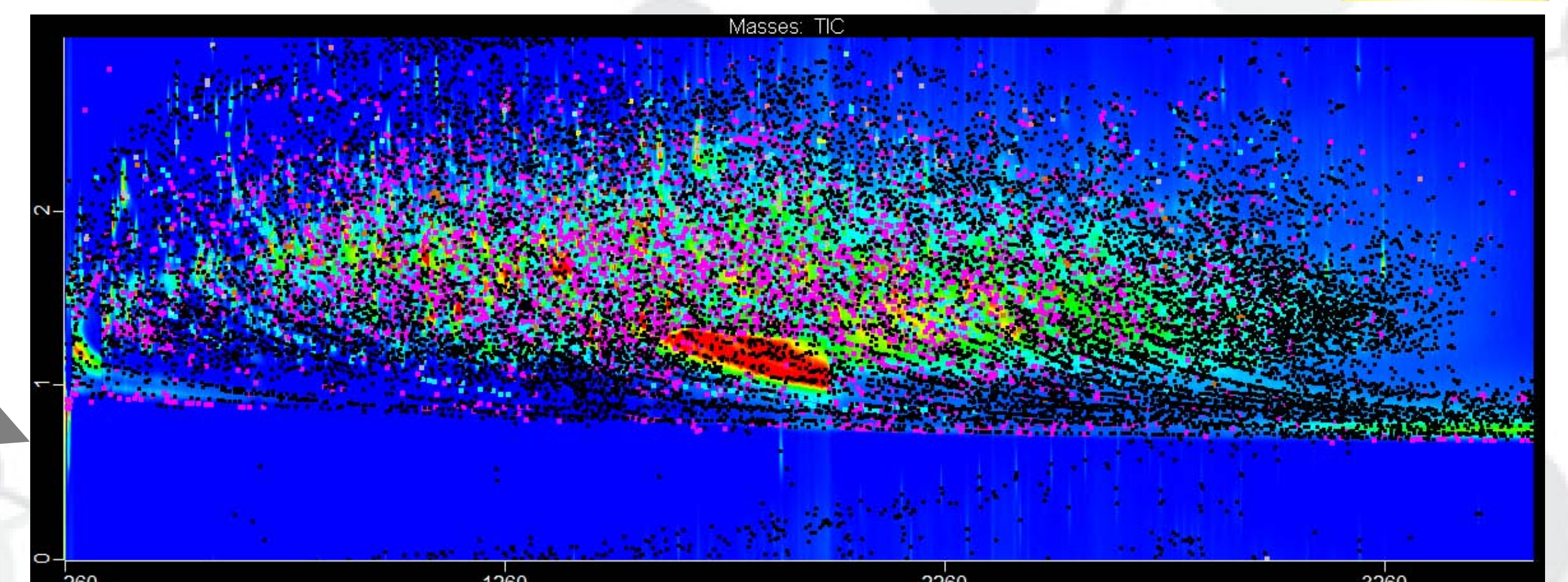
Classification Summary Tables

Area % as measured by the deconvoluted TIC for each class was compared for four sample types from Raw Water to Tap Water to Swimming Pool Water and Spa Water. The data show that the abundance of brominated compounds increased significantly in the swimming pool and spa treated with bromine as the disinfectant compared to the tap water used to fill the pools. Also, the tap water treated with chlorine had more chlorinated compounds than the untreated raw water [1].



Classification with Accurate Mass Peak Filters

Classify sample peaks to tabulate data for the classification table. The color of the peak markers correspond to the assignment based on the properties for the class(es) in the classification table.



Reference

1. Daiber, E.J. et al. 2016. Environ Sci Technol. Progressive Increase in Disinfection Byproducts and Mutagenicity from Source to Tap to Swimming Pool and Spa Water: Impact of Human Inputs. <http://pubs.acs.org/doi/pdf/10.1021/acs.est.6b00808>.

