

The Discovery of Potential Cancer Biomarkers in Human Plasma Using GC- and GCxGC-TOFMS

David E. Alonso¹, Habtom Resson², Cristina Di Poto², and Joseph E. Binkley¹; ¹LECO Corporation, St. Joseph, MI; ²Georgetown University Medical School, Washington D.C.

Introduction

- Hepatocellular carcinoma (HCC) is the leading cause of cancer-related deaths worldwide (9th in the US)
- Risk Factors:** Hepatitis (B & C), alcoholism, diabetes, obesity, and nonalcoholic fatty liver disease
- Critical Need:** Early stage intervention and effective medical treatment for HCC

Objectives

- To complement a much larger study investigating metabolite levels in HCC versus liver cirrhosis (CIRR) patients
- To identify potential HCC biomarkers using a non-targeted, multiplatform approach

Samples

Plasma
Metabolite Extraction (ACN/i-PrOH/H₂O)
64 Patient Samples HCC & CIRR)

Derivatization

Methoximation: 30 μL of methoxyamine (20 mg/mL in Pyridine), heat/agitate at 60 °C for 30 minutes
Silylation: 70 μL of MSTFA, heat/agitate at 60 °C for 30 minutes

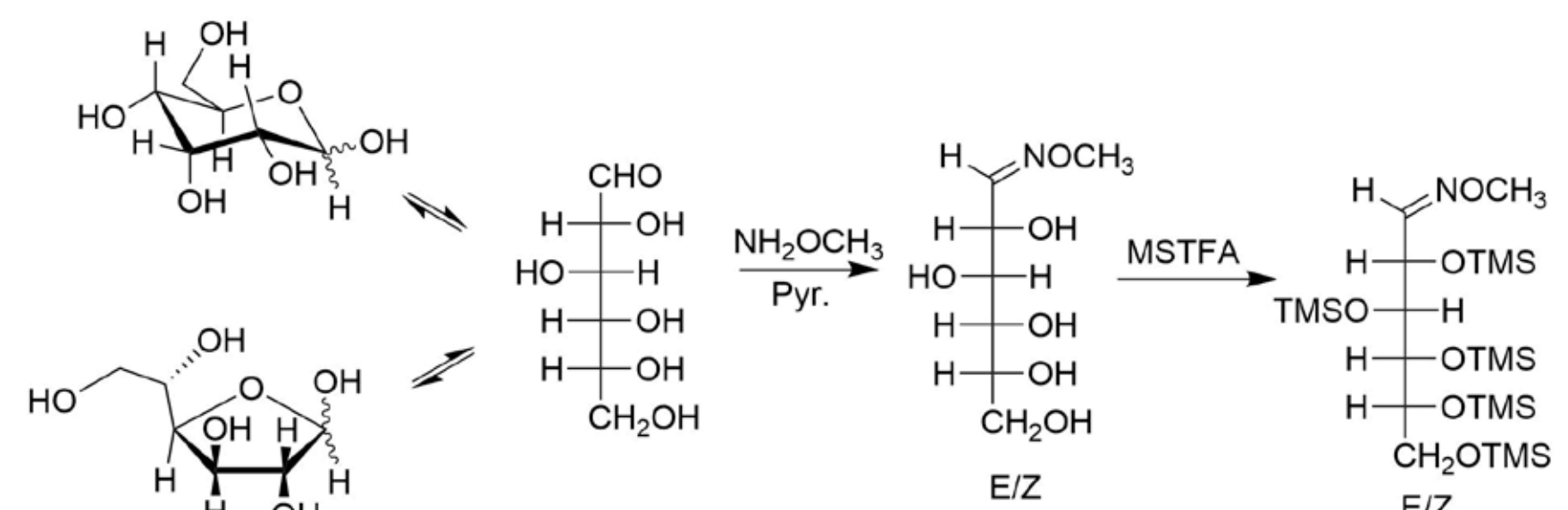


Figure 1: Two-step derivatization of glucose

Instrument Parameters

Gas Chromatograph	LECO GCxGC (Dual Stage Quad Jet Thermal Modulator)
Autosampler	LECO L-PAL 3
Injection	1 μL, Split 20:1, 250 °C
Carrier Gas	He @ 1.4 mL/min, Constant Flow
Column Set	Rxi-5ms, 30 m x 0.25 mm i.d. x 0.25 μm (Restek) Rxi-17sil ms, 0.6 m x 0.25 mm i.d. x 0.25 μm (Restek)
Temperature Program	50 °C (1 min), ramped 10 °C/min to 300 °C (12 min) Secondary oven offset was 5 °C relative to primary oven
Modulation	3s modulation; temperature offset 15 °C relative to secondary oven
Mass Spectrometer	LECO Pegasus® BT
Ion Source Temperature	250 °C
Ionization Mode	EI
Mass Range (m/z)	45-750
Acquisition Rate	10 spectra/s (200 spectra/s GCxGC-TOFMS)

Pegasus BT 4D



GC-TOFMS Results

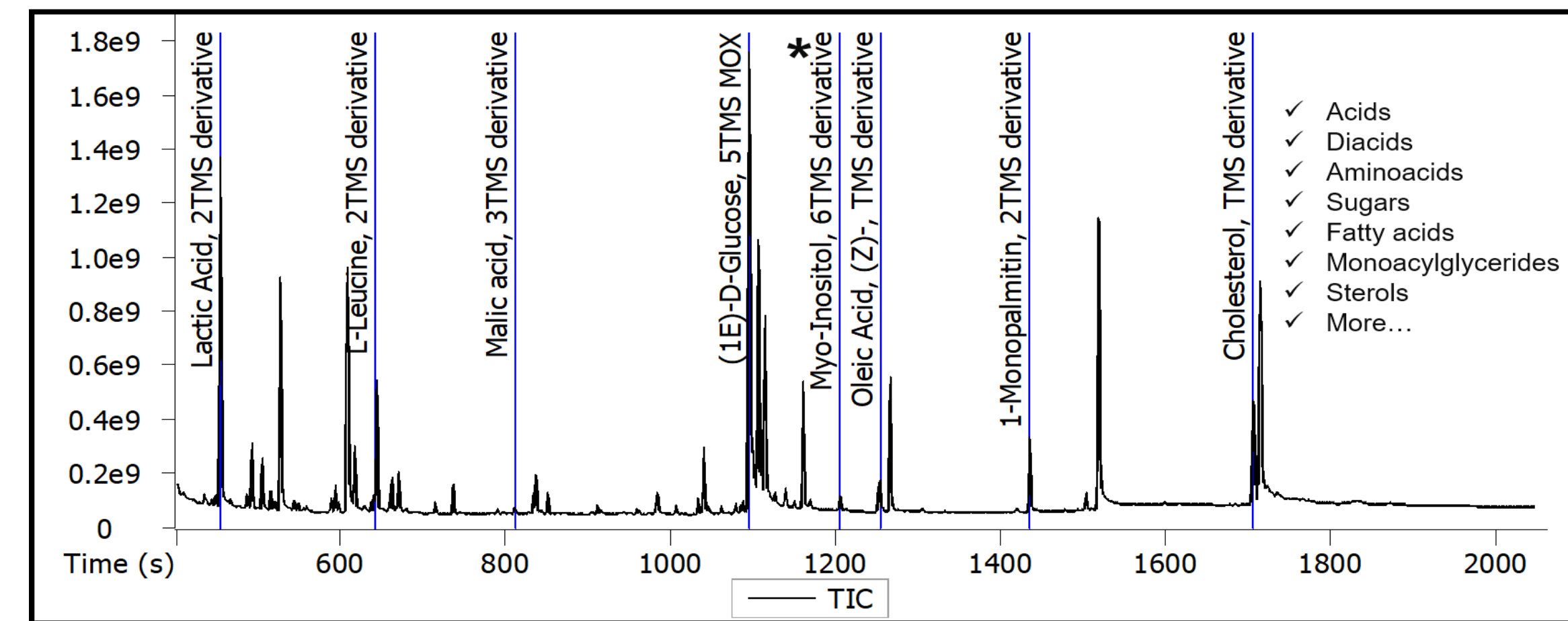


Figure 2: TIC showing the chemical diversity of plasma sample metabolites identified by GC-TOFMS.

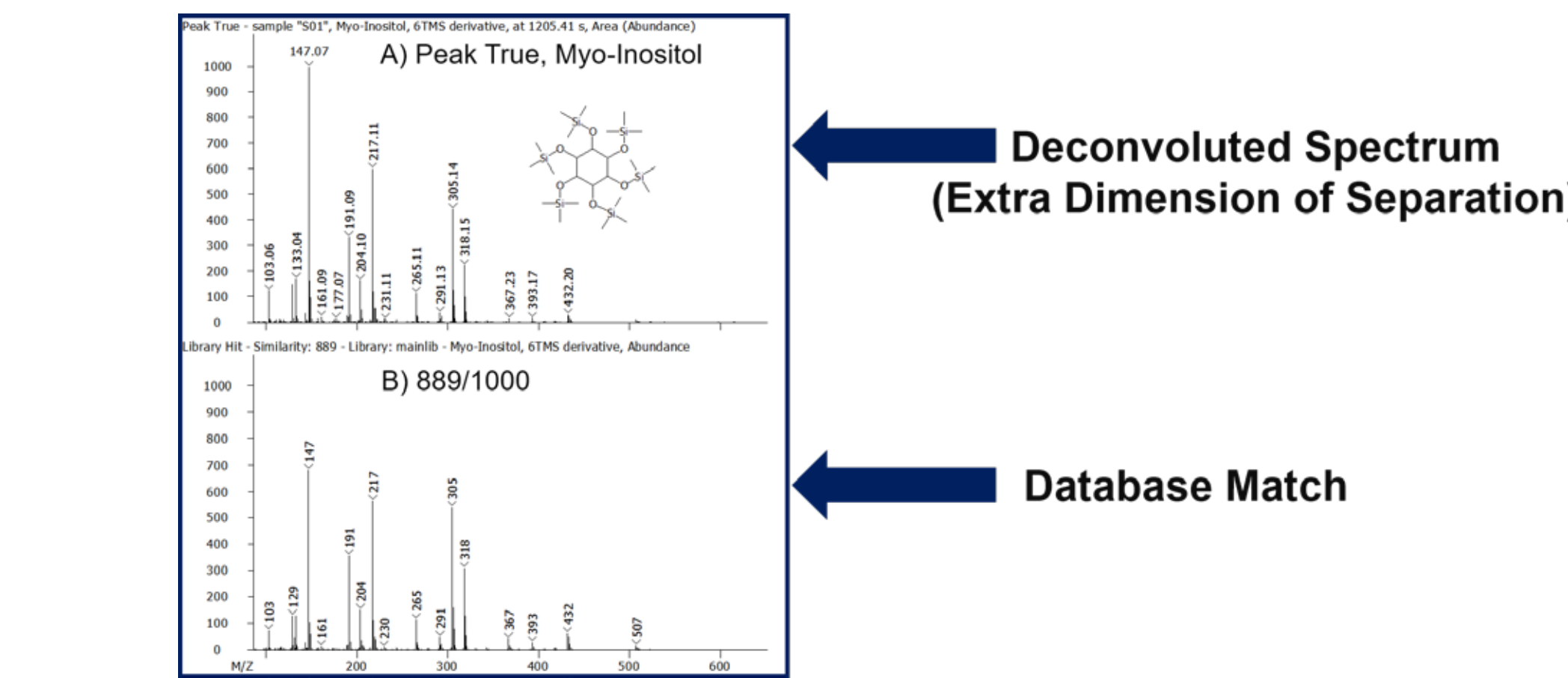


Figure 3: A) Peak True (Deconvoluted), and B) library spectra for myo-inositol.

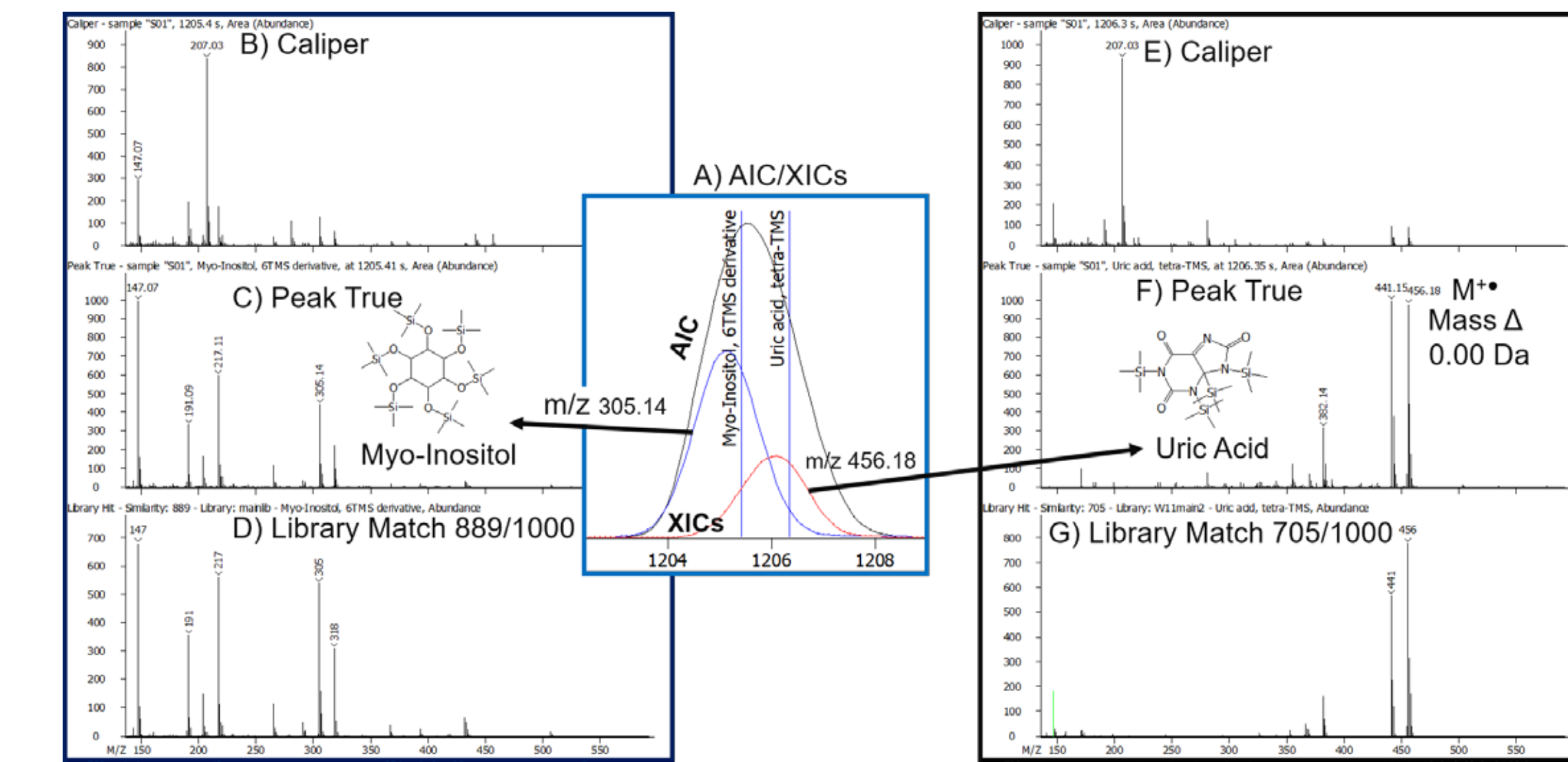


Figure 4: A) Analytical Ion Chromatogram (AIC) showing extracted ions (XICs) for coeluting myo-inositol & uric acid metabolites. Caliper, Peak True and Library spectra for myo-inositol (B-D) and uric acid (E-G). Identification of uric acid is supported by mass Δ calculations (ΔM = m/z_{obs} - m/z_{calc}).

Table 1: Representative Compounds in plasma (Untargeted Peak Find Processing)

Name	Formula	R.T. (s)	Similarity	Name	Formula	R.T. (s)	Similarity	Name	Formula	R.T. (s)	Similarity
Lactic Acid, 2TMS	C ₉ H ₁₄ O ₅ Si ₂	454	941	Erythritol, 4TMS	C ₁₂ H ₂₀ O ₅ Si ₄	831	879	(1E)-D-Galactose, 5TMS MOX	C ₁₂ H ₂₀ O ₅ Si ₅	1119	817
Glyceric acid, 2TMS	C ₇ H ₁₀ O ₅ Si ₂	467	882	L-Proline, 5-oxo-, 5TMS	C ₁₁ H ₁₇ NO ₅ Si ₅	834	927	Palmitic acid, TMS	C ₁₇ H ₃₂ O ₂ Si	1150	921
α-Alanine, 2TMS	C ₅ H ₉ NO ₃ Si ₂	492	927	Hydroxyproline, 3TMS	C ₇ H ₁₁ NO ₃ Si ₃	837	931	Myo-Inositol, 6TMS	C ₁₈ H ₃₂ O ₅ Si ₆	1161	950
2-Hydroxypropanoic acid, 2TMS	C ₅ H ₉ NO ₃ Si ₂	515	909	Uric acid, 4TMS	C ₅ H ₄ N ₂ O ₄ Si ₄	891	821	Uric acid, 4TMS	C ₅ H ₄ N ₂ O ₄ Si ₄	1205	889
Cholic acid, 2TMS	C ₂₇ H ₄₅ O ₂ Si ₂	518	897	2-Ketoglutaric acid, TMS MOX	C ₈ H ₁₂ NO ₅ Si ₅	874	705	Margaric acid, TMS	C ₁₇ H ₃₂ O ₂ Si	1214	837
2-Hydroxyvaleric acid, 2TMS	C ₆ H ₁₀ O ₅ Si ₂	549	952	L-Ornithine, 3TMS	C ₆ H ₁₁ N ₂ O ₃ Si ₃	901	775	Linoleic acid, TMS	C ₁₈ H ₃₂ O ₂ Si	1251	929
2-Aminobutanoic acid, 2TMS	C ₅ H ₉ NO ₃ Si ₂	555	883	Glutamic acid, 3TMS	C ₇ H ₁₁ NO ₄ Si ₃	905	717	Oleic acid, (Z)-, TMS	C ₁₈ H ₃₄ O ₂ Si	1254	941
3-Hydroxyvaleric acid, 2TMS	C ₆ H ₁₀ O ₅ Si ₂	587	787	Phenylalanine, 2TMS	C ₉ H ₉ NO ₃ Si ₂	911	922	Stearic acid, TMS	C ₁₈ H ₃₆ O ₂ Si	1266	946
2-Keto-N-caproic acid, 2TMS MOX	C ₁₀ H ₁₅ NO ₅ Si ₅	591	867	L-Glutamine, 3TMS	C ₇ H ₁₁ N ₂ O ₃ Si ₃	1007	877	Arachidonic acid, TMS	C ₂₀ H ₃₆ O ₂ Si	1363	835
L-Valine, 2TMS	C ₆ H ₁₁ NO ₃ Si ₂	594	901	Lauric acid, TMS	C ₁₄ H ₂₆ O ₂ Si	920	699	l-Glycerol arachidate, 2TMS	C ₃₄ H ₆₄ O ₅ Si ₂	1387	745
Urea, N,N'-bis(trimethylsilyl)-	C ₃ H ₁₀ N ₂ O ₅ Si ₃	608	787	Asparagine, 3TMS	C ₈ H ₁₃ N ₂ O ₃ Si ₃	941	855	L-Cystine, 4TMS	C ₉ H ₁₄ N ₂ O ₃ Si ₄	1305	825
L-Leucine, 2TMS	C ₁₁ H ₂₁ NO ₃ Si ₂	641	919	5-Hydroxytryptophan, 4TMS	C ₁₀ H ₁₄ N ₂ O ₅ Si ₄	943	722	Pseudo uridine, 3TMS	C ₁₀ H ₁₆ N ₂ O ₅ Si ₃	1329	756
Glycerol, 3TMS	C ₉ H ₁₈ O ₃ Si ₃	644	931	Ribitol, 3TMS	C ₆ H ₁₀ O ₅ Si ₃	973	882	Arachidonic acid, TMS	C ₂₀ H ₃₆ O ₂ Si	1332	928
Isoleucine, 2TMS	C ₁₁ H ₂₁ NO ₃ Si ₂	659	876	Glutamine, 4TMS	C ₈ H ₁₃ N ₂ O ₃ Si ₄	984	833	1-Monomyristin, 2TMS	C ₁₄ H ₂₆ O ₅ Si ₂	1346	728
L-Proline, 2TMS	C ₅ H ₉ NO ₃ Si ₂	662	874	L-Glutamine, 3TMS	C ₇ H ₁₁ N ₂ O ₃ Si ₃	1007	877	Arachidonic acid, TMS	C ₂₀ H ₃₆ O ₂ Si	1504	864
Glycine, 3TMS	C ₂ H ₃ NO ₂ Si ₃	670	915	l-Ornithine, 4TMS	C ₆ H ₁₁ N ₂ O ₃ Si ₄	1037	894	Embramine	C ₁₀ H ₁₇ N ₃ O	1374	916
Succinic acid, 2TMS	C ₄ H ₆ O ₄ Si ₂	672	753	Citric acid, 4TMS	C ₆ H ₈ O ₇ Si ₄	1040	871	Uridine, 3TMS	C ₉ H ₁₃ N ₂ O ₅ Si ₃	1380	781
Glyceric acid, 3TMS	C ₇ H ₁₀ O ₅ Si ₃	691	869	Myristic acid, TMS	C ₁₄ H ₂₆ O ₂ Si	1046	861	2-Palmitoylglycerol, 2TMS	C ₃₀ H ₅₈ O ₅ Si ₂	1420	849
Fumaric acid, 2TMS	C ₄ H ₄ O ₄ Si ₂	698	829	Caffeine	C ₈ H ₁₀ N ₄ O ₂	1056	919	1-Monopalmitin, 2TMS	C ₂₂ H ₄₀ O ₅ Si ₂	1435	908
2,3-Dihydroxybutanoic acid, 3TMS	C ₆ H ₁₀ O ₅ Si ₃	703	784	L,β-Alanine, 3TMS	C ₃ H ₅ NO ₂ Si ₃	1062	871	2-Monosteirin, 2TMS	C ₁₈ H ₃₂ O ₅ Si ₂	1504	864
Norbornic acid, TMS	C ₇ H ₁₀ O ₅ Si	707	880	D-Fructose, 5TMS MOX	C ₆ H ₁₀ NO ₅ Si ₅	1079	922	l-Glycerol arachidate, 2TMS	C ₃₄ H ₆₄ O ₅ Si ₂	1509	804
Serine, 3TMS	C ₃ H ₅ NO ₃ Si ₃	715	896	(1E)-D-Galactose, 5TMS MOX	C ₁₂ H ₂₀ NO ₅ Si ₅	1088	924	γ-Tocopherol, TMS	C ₂₉ H ₅₀ O ₂ Si	1616	806
L-Threonine, 3TMS	C ₄ H ₉ NO ₃ Si ₃	737	946	(1E)-D-Glucose, 5TMS MOX	C ₆ H ₁₀ NO ₅ Si ₅	1095	821	l-Tocopherol, TMS	C ₂₉ H ₅₀ O ₂ Si	1632	839
S-Methionine, 3TMS	C ₄ H ₉ NO ₃ Si ₃	762	833	L-Lysine, 4TMS	C ₆ H ₁₁ N ₂ O ₃ Si ₄	1099	820	Cholesterol, TMS	C ₂₇ H ₄₆ O ₂ Si	1707	923
β-Alanine, 3TMS	C ₃ H ₅ NO ₂ Si ₃	765	774	L-Histidine, 4TMS	C ₆ H ₉ N ₃ O ₃ Si ₄	1100	752	Campesterol, 3TMS	C ₂₈ H ₄₈ O ₂ Si ₃	1774	723
Aminomalonic acid, 3TMS	C ₄ H ₇ NO ₅ Si ₃	800	832	L-Tyrosine, 3TMS	C ₉ H ₉ NO ₃ Si ₃	1120	855				
Malic acid, 3TMS	C ₄ H ₆ O ₅ Si ₃	812	849	Mannitol, 6TMS	C ₆ H ₁₂ O ₆ Si ₆	1114	947				

Ave. Similarity = 855/1000

GCxGC-TOFMS Results (Untargeted)

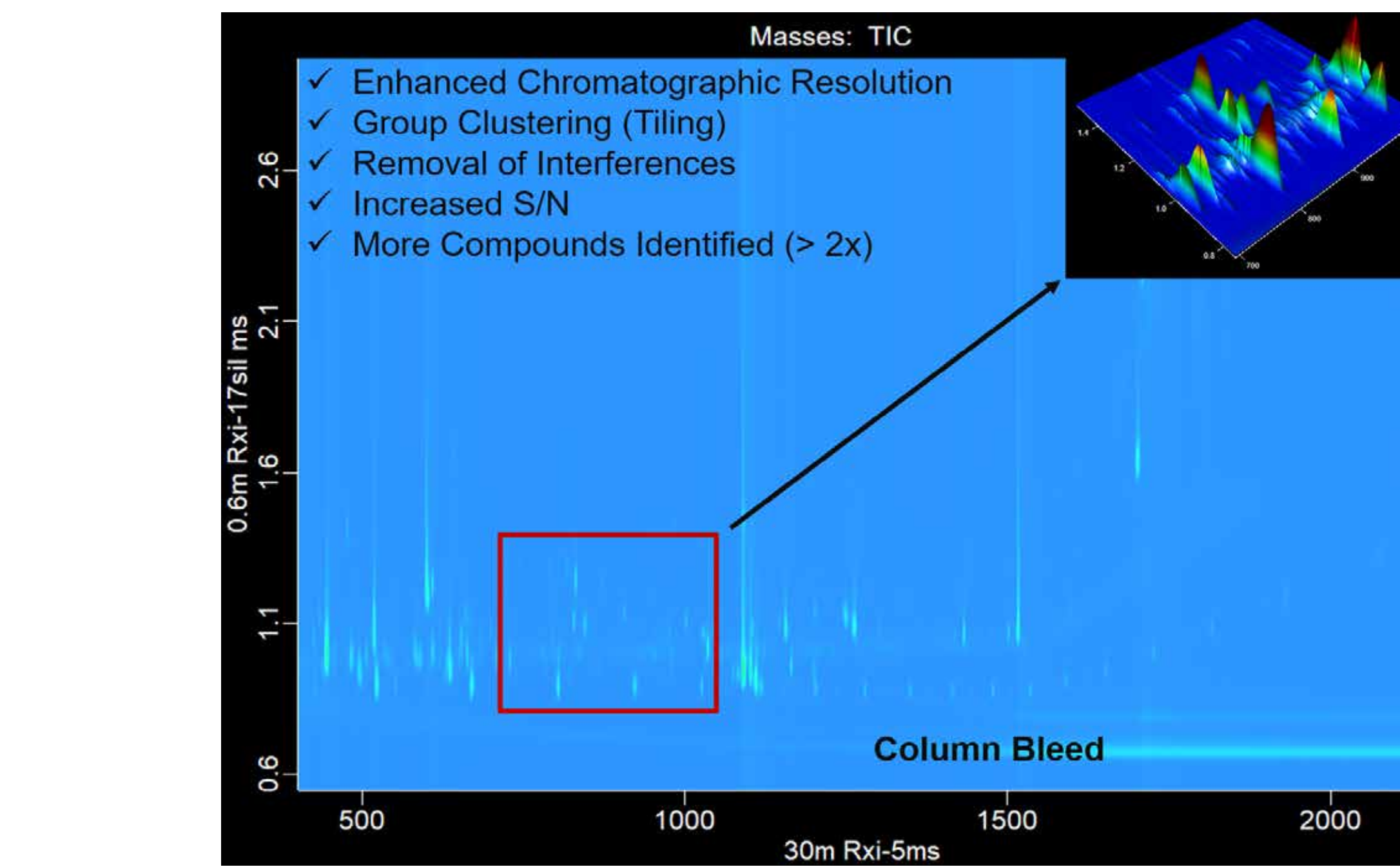


Figure 5: GCxGC-TOFMS contour plot and surface plot expansion (inset).

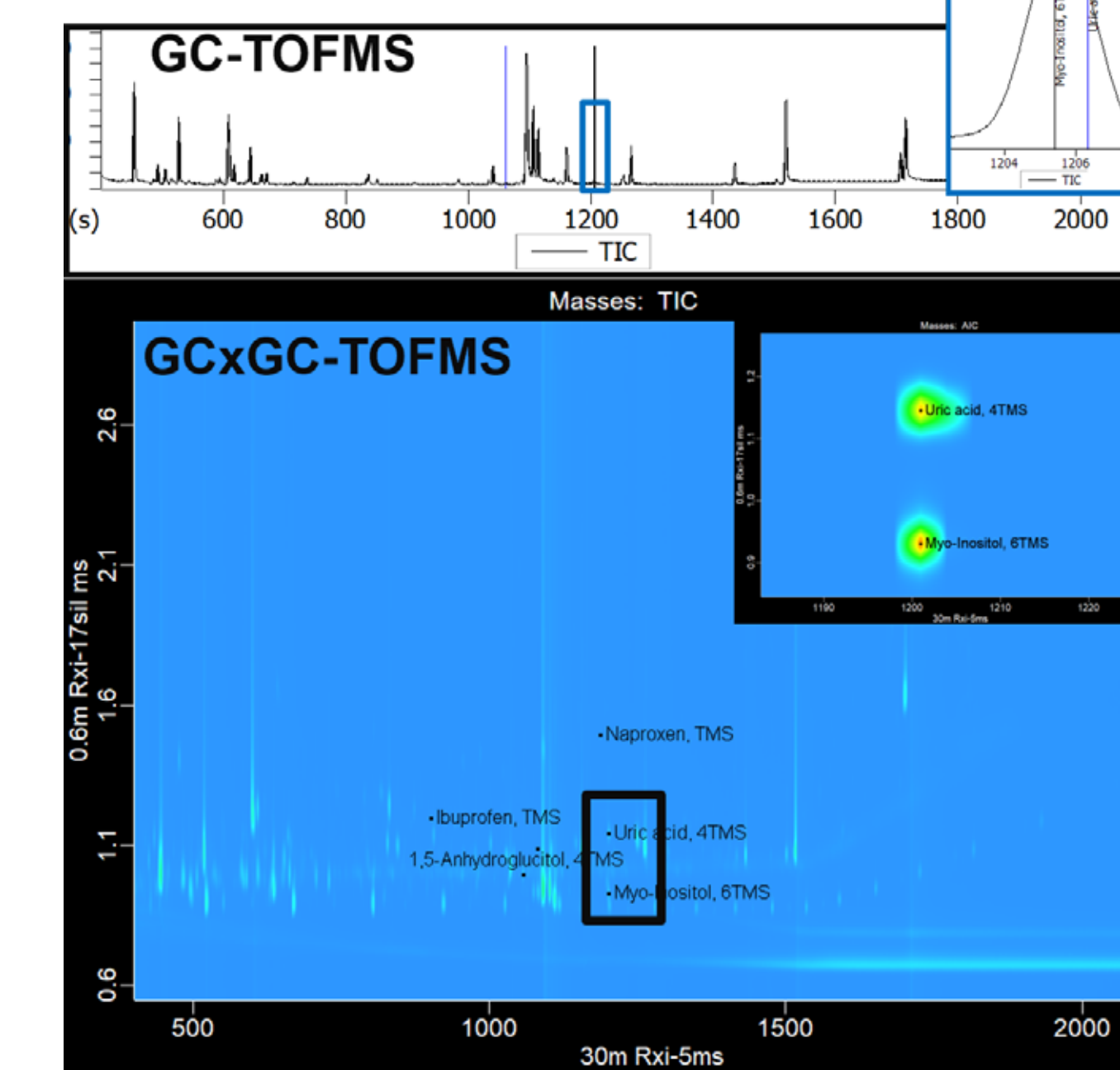


Figure 6: Comparison of GC- (top) and GCxGC-TOFMS (bottom) results. The enhanced chromatographic resolution in transitioning from 1D to 2D results in an improvement in S/N and similarity scores (upper/lower data tables).

Name	R.T. (s)	Peak S/N	Similarity
Ibuprofen, TMS	Not Detected		
1,5-Anhydroglucitol, 4TMS	1061.64	2070	871
4-Hydroxyphenylacetic acid, 3TMS	Not Detected		
Naproxen, TMS	Not Detected		
Myo-Inositol, 6TMS	1205.41	2436	889
Uric acid, 4TMS	1206.35	375	705

Name	R.T. (s)	Peak S/N	Similarity
Ibuprofen, TMS	901 s, 1.199 s	137	828
1,5-Anhydroglucitol, 4TMS	1057 s, 1.000 s	1830	926
4-Hydroxyphenylacetic acid, 3TMS	1081 s, 1.090 s	873	874
Naproxen, TMS	1186 s, 1.495 s	97	828
Myo-Inositol, 6TMS	1201 s, 0.931 s	4642	951
Uric acid, 4TMS	1201 s, 1.147 s	4864	863

Unknowns → Knowns
More Compounds Identified

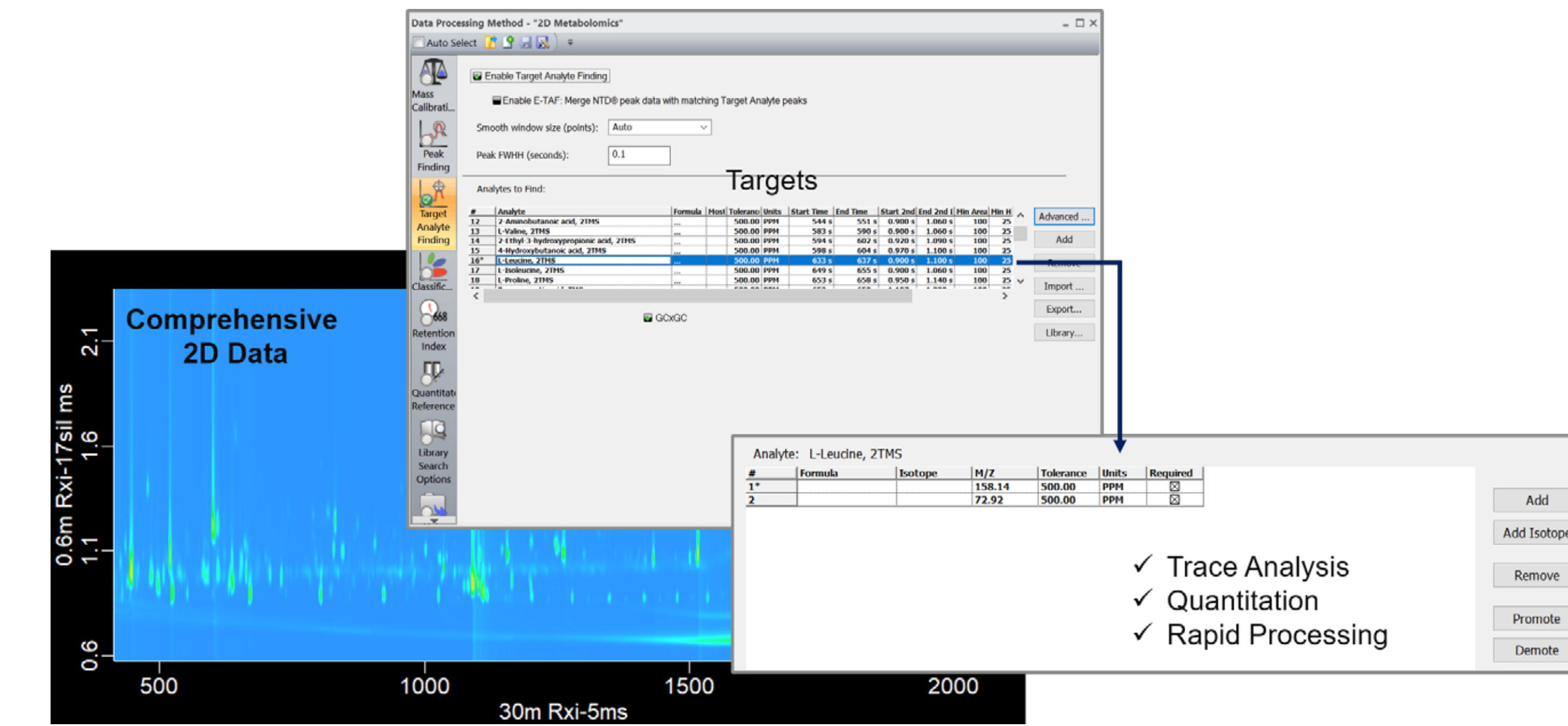


Figure 7: Target Analyte Finding (TAF) data processing method with metabolite list and target ions with tolerances.

GCxGC-TOFMS Target Analyte Processing

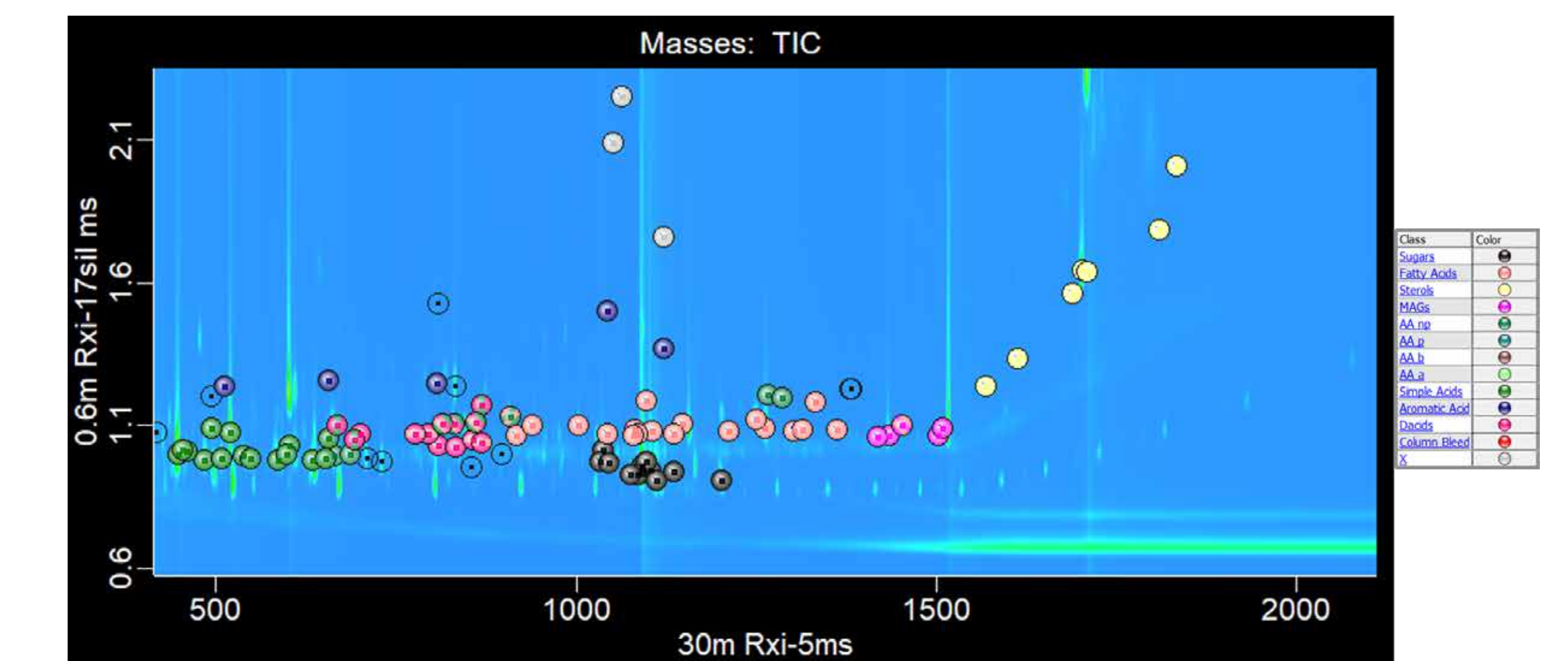


Figure 8: Contour plot displaying the results of Target Analyte Finding (TAF) of comprehensive GCxGC-TOFMS data. The color coded bubbles represent different classes of compounds identified.

Table 2: TAF results table listing targeted metabolites, 2D retention times and areas.

Name	R.T. (s)	Area	Name	R.T. (s)	Area
Lactic Acid, 2TMS	454 s, 1.001 s	115781630	L-Methionine, 3TMS	879 s, 1.101 s	116166501
Hexanoic acid, TMS	454 s, 1.031 s	13162022	L-Aspartic acid, 3TMS	1111 s, 0.911 s	3156604
Glycolic acid, 2TMS	465 s, 1.015 s	68842184	l-Cysteine, 3TMS	852 s, 1.245 s	488124860
Alanine, 2TMS	464 s, 0.980 s	46296600	l-Cysteine, 4TMS	853 s, 0.960 s	12004485
Butanoic acid, 2-(methylamino) 3-methyl-, TMS	494 s, 1.095 s	13622013	L-Cystine, 3TMS	856 s, 0.955 s	12124911
D-Hydroxyvaleric acid, 2TMS	505 s, 0.990 s	18207963	Creatinine, 4TMS	858 s, 1.118 s	93800248
Levulinic acid, TMS	511 s, 1.040 s	438778	Pentanedioic acid, 2-hydroxy-, TMS	868 s, 1.045 s	940099
Quinic acid, TMS	520 s, 1.080 s	78433938	Pentanedioic acid, 2-oxo-, 2TMS MOX	868 s, 1.175 s	8610200
5-Hydroxyvaleric acid, 2TMS	538 s, 1.001 s	68805035	l-Ornithine, 3TMS	895 s, 1.125 s	116103076
2-Aminobutanoic acid, 2TMS	547 s, 0.990 s	79609072	Phenylalanine, 2TMS	907 s, 1.131 s	106023775
L-Valine, 2TMS	565 s, 1.001 s	12949700	Biotin, 3TMS	918 s, 1.079 s	110688502
1,5-Anhydroglucitol, 4TMS	598 s, 1.005 s	30007	Asparagine, 3TMS	937 s, 1.105 s	12024351
4-Hydroxyphenylacetic acid, 3TMS	601 s, 0.939 s	810291	L-Glutamine, 3TMS	1023 s, 1.025 s	101939192
Leucine, 2TMS	634 s, 0.985 s	23506845	l-Ornithine, 4TMS	1023 s, 1.000 s	1111335
L-Isoleucine, 2TMS	652 s, 0.990 s	83617835	Citric acid, 4TMS	1026 s, 1.020 s	121423396
L-Proline, 2TMS	655 s, 1.001 s	30976200	Arachidonic acid, TMS	1045 s, 0.975 s	23015114
Benzammonic acid, TMS	659 s, 1.065 s	537811	Malic acid, TMS	1042 s, 1.065 s	882709
Glycine, 3TMS	664 s, 1.001 s	26859565	D-Fructose, 1, 5TMS MOX	1047 s, 1.075 s	622792
Succinic acid, 2TMS	667 s, 1.05 s	31859277	D-Fructose, 2, 5TMS MOX	1075 s, 0.955 s	116178769
Fumaric acid, 2TMS	686 s, 1.005 s	42811519	Phenylalanine, 2TMS	1078 s, 1.070 s	379261
Formic acid, 3TMS	691 s, 1.001 s	889112	l-Hydroxyphenylacetic acid, 3TMS	1078 s, 1.070 s	2396120
Benzoic acid, TMS	700 s, 1.075 s	1962605	D-Galactose (E), 5TMS MOX	1084 s, 0.935 s	39729792
Serine, 3TMS	709 s, 0.990 s	23669936	Benzoic acid, TMS	1084 s, 1.075 s	738440
L-Threonine, 3TMS	730 s, 0.980 s	18897565	D-Galactose (Z), 5TMS MOX	1090 s, 0.955 s	154365312
Decanoic acid, TMS	775 s, 1.075 s	3605185	Sulfhydrylglycerol, 5TMS MOX	1090 s, 0.965 s	20774227
2-Aminomalonic acid,					