

Poster Reprint

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An Innovative Approach to Q-TOF High Resolution Accurate Mass Analyte Screening Using an Improved Software Algorithm and Screener Tool

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Introduction

High resolution, accurate mass screening (HRAM) for high numbers of targets using a Q-TOF Mass Spectrometer can be difficult to reproducibly achieve in a single analysis due to the variability of sample matrices, the concentration ranges and the nature of the data mining algorithms.

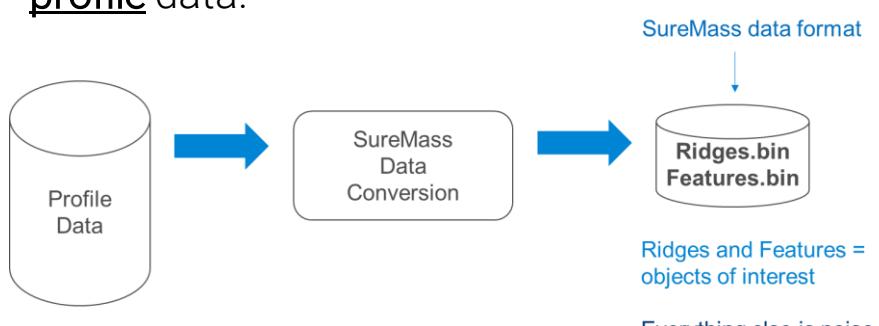
This poster illustrates a proof of principle ability to detect a panel of over 200+ targets and suspects reproducibly in a single 8-minute analysis at high and low concentrations using a newly improved software mining algorithm SureMass.

The results presented herein this poster demonstrate near 100% identification of a comprehensive panel reproducibly to below 5ppm mass accuracy over a wide range of typical working concentrations.

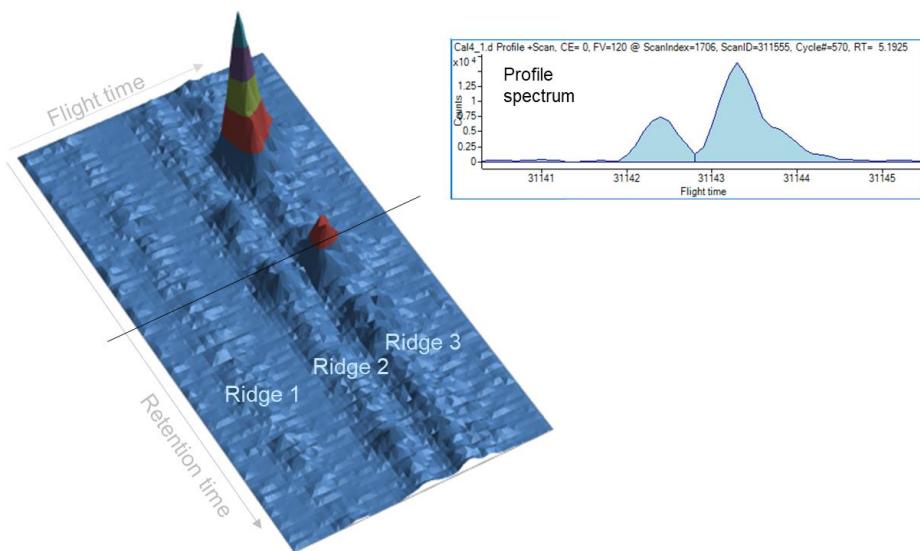
SureMass Data Mining

What is SureMass?

- A low-level data processing algorithm designed to extract maximum information from accurate mass profile data.



Ridges and Features



Attributes of a Feature:

- Scan space (Collision Energy, Fragmentor Voltage)
- Assigned flight time and Retention Time
- Abundance profile and Flight time profile
- Assigned m/z

SureMass Data Mining (cont'd.)

Why SureMass?

Both Profile and Centroid data formats have individual issues.

1. **Speed** - Profile data analysis is painfully slow
2. **Mass Accuracy** - Centroid data analysis is fast, but gives poor mass accuracy at high abundance levels
3. **Saturation** – Centroid can't handle Saturation (Mass shifts, Split peaks, False negatives)

SureMass handles saturation gracefully (No high-abundance negatives)

Experimental

Outline.

Repeat injections ($N=10$) containing 200+ positive polarity targets were injected onto an 8-minute reverse phase chromatography gradient for separation from isobars and interferences that may be present. Column phase was **Poroshell 120 EC C18 100 x 2.1mm**. (Aqueous MP= 0.01% FA, 5mM Amm F).

Pure solvent standard mixes at high and low concentrations relative to minimal and high working concentrations for the respective target classes included in this study were analyzed using an Agilent 6546 Q-TOF LC/MS operating in MS-only or All Ions positive polarity mode to attain the identification results.

Data mining software SureMass was utilized to identify and report low false negative and positive percentage rates via a novel screener software tool.

HPLC Gradient Conditions.

0.00 min 10% Methanol; 1.00 min 15% Methanol;
4.00 min 50% Methanol; 6.00 min 95% Methanol;
8.00 min 95% Methanol. 8 min total run time.

MS Conditions (AJS-ESI Positive Polarity).

Data collection Rate:

- 3x data points/s (MS-only)
- 9x data points/s (All Ions, CE = 0, 20, 30 eV)

Source Settings at HPLC Flow Rate of 0.5 mL/min:

- | | |
|-------------------------|-----------------|
| • Nozzle Voltage: | 500V |
| • Sheath Gas Temp/Flow: | 360°C; 11L/min |
| • Dry Gas Temp/Flow: | 300°C; 10 L/min |
| • Nebulizer Pressure: | 55 psi |
| • Capillary Voltage: | 3750 V |

Discussion

Screener Tool.

The Screener Tool Graphical User Interface is an integral element of MassHunter Quantitative Analysis Software. It displays each sample in the batch interactively and simplifies the analyst review process by displaying the targets and suspects using a color-coded 'traffic light' system of green, amber and red. Green=positive, amber=review required, Red= negative. Filters are provided to simplify the review process further.

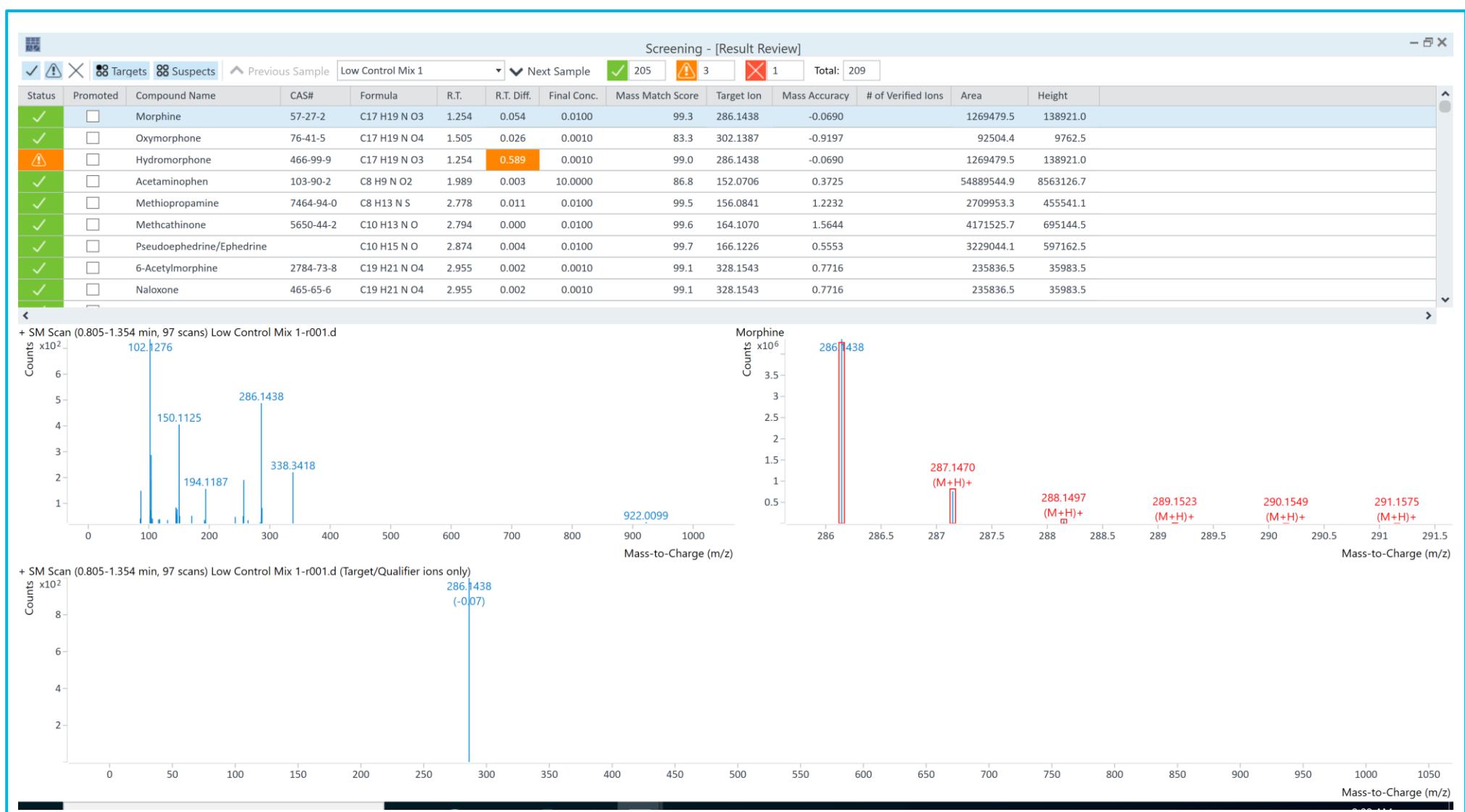


Figure 1. Screener Tool Graphical User Interface.

Mass Match Score %.

The Mass Match Score in Table 1 is derived from the contributions of three scoring elements:

- Monoisotopic Mass Accuracy Score (50%)
- Isotope Spacing Score (25%)
- Isotope Abundance Score compared with Theoretical (25%)
 - Retention time Delta window also used for Identification.
 - Coelution Score available when using All Ions mode of acquisition.

Conclusions

- It has been shown that Comprehensive and large panels of Targets and Suspects can be accurately and reliably screened using SureMass Data Mining combined with the Agilent 6546 Q-TOF LC/MS.
- Mass match Scores are consistently in the high 90% ranges for high and low concentration replicate injections. .
- Mass Accuracy measurements for very high and very low concentration ranges within target classes are consistently below 5 ppm.
- Fast data processing speed for SureMass converted data files of large target and Suspect lists.
- Easy and intuitive Screener Tool for fast data review.
- MS-only and All Ions data formats compatible and reliable.

References

1/ Karen E. Yannell,, and Manuel Gomez, Agilent Technologies, Inc., Pub 5994-1744EN

Results

Mass Accuracy Results for 200+ Target Mixes.

- Of the 200+ Targets and Suspects analyzed as part of this proof of principle study, the vast majority exhibited a Mass Match Score of High 90% values for Low and High Concentration Controls (Averaged over 10x different batches.)
- Mass Accuracy measurements were less than 5ppm for All Targets at low and high concentrations.
- MS-only data results are illustrated due to space constraints, however All Ions data yielded comparable results.

Compound	Low Controls N=10				High Controls N=10				Compound	Low Controls N=10				High Controls N=10					
	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	Name	Mass Accuracy Average (ppm)	Mass Match Score Average	
258-NBOMe	Transition	380.0856	0.7303	99.4506	0.5289	99.0208	Isobutrylfentanyl/beta-Methylfentanyl	351.2431	0.0865	99.1605	0.7147	99.4432	99.4432	Transition	351.2431	0.0865	99.1605	0.7147	99.4432
25C-NBOMe		336.1361	0.9524	98.8240	1.2707	97.6275	Ketamine	238.0993	1.3008	99.2295	1.3093	98.9875	98.9875	Lacosamide	251.1390	1.4773	99.1081	1.4790	99.2036
25H-NBOMe		302.1751	0.9877	99.5339	0.7183	99.4547	Lamotrigine	256.0151	0.5244	98.1497	1.2585	98.1517	98.1517	Lidocaine	235.1805	0.9896	99.2873	0.8662	95.3565
25i-NBF		416.0517	0.9026	99.5882	0.8628	99.5358	Loperamide	477.2303	1.4817	95.4532	0.6926	96.0532	96.0532	Loratadine	383.1521	1.5419	96.2688	1.0261	97.7322
25i-NBMD		442.0510	0.8085	99.5329	0.3036	99.6687	Loxapine	328.1211	1.1375	99.1810	0.7724	98.6391	98.6391	LSD	324.2070	-0.1899	91.0280	1.2810	83.8689
25i-NBOH		414.0561	0.9066	99.4942	0.5042	99.5230	Maprotiline	278.1903	1.4441	98.9089	0.8646	99.3495	99.3495	MDA	180.1019	1.4183	99.5409	1.4508	99.5028
25i-NBOMe		428.0717	0.9317	99.5551	0.4652	99.4947	MDEA	208.1332	1.5558	99.3208	1.5388	98.9703	98.9703	MDMA/Methedrone	194.1176	1.4041	98.8112	1.3169	96.5051
2-MAPB		190.1226	1.0261	99.6256	1.5285	99.3207	MDPV	276.1594	1.1180	99.6175	0.8843	99.5324	99.5324	Meperidine	248.1645	1.4982	99.4144	1.0344	99.2055
3-Chloroethcathinone		212.0836	1.9641	99.4559	1.1749	99.4396	Mephedrone/6-APDB	178.1226	1.1267	99.6459	1.0252	97.9841	97.9841	Meowfentanyl	247.1805	1.4236	99.3452	0.9966	99.5207
3-Chloromethcathinone		198.0680	1.5290	99.5521	1.4755	98.7582	Mepivacaine	310.2165	1.1909	99.6076	0.0431	98.5935	98.5935	Methadone	150.1277	1.5506	99.6178	1.3213	99.5427
3-Fluorofentanyl		355.2180	0.4399	99.5225	0.4362	95.1465	Methamphetamine	164.1070	1.1943	99.6563	0.9424	99.5560	99.5560	Methcathinone	156.0841	1.0051	99.4840	0.9507	99.0218
3-Fluorophenmetazine		196.1322	1.6006	99.4387	1.0015	99.5006	Methiopropamine	248.1645	1.6191	99.4144	1.1553	99.2055	99.2055	Methoxetamine	353.2224	0.0704	98.6097	0.7161	98.3199
4/5-APDB		178.1226	1.1267	99.6459	1.0252	97.9841	Methoxyacetifentanyl	296.2009	1.1864	99.5226	1.0278	99.0897	99.0897	Methoxyphenidine	208.0968	1.1695	99.2013	1.1699	99.2699
4-Chloro-Alpha-PVP		266.1306	1.4243	98.4546	0.8829	99.0547	Methylenedione	234.1489	0.9005	99.5964	0.3965	99.0849	99.0849	Methylenephedate	300.1473	1.5702	99.0772	0.9129	99.0097
4-Chloromethcathinone		198.0680	1.5290	99.5521	1.4755	98.7582	Metoclopramide	208.1332	1.5558	99.3208	1.5388	98.9703	98.9703	Mexedrone	326.0855	1.0985	99.2322	0.6859	98.9765
4-Methoxybutrylfentanyl		381.2537	0.1689	99.3459	0.9730	97.9662	Mildazolam	266.1652	0.8574	99.5545	0.8258	98.9637	98.9637	Mirtazapine	399.2278	1.3999	99.2904	0.8436	99.3310
5-DBFPV		274.1802	1.1097	99.4470	1.0646	99.1194	Mitragynine	296.0716	1.0768	97.5970	1.0625	95.8292	95.8292	Modafinil	300.1473	1.5702	99.0772	0.9129	92.9154
5-Fluoro-PB-22		377.1660	1.3315	98.9360	1.1144	98.9819	Morphine	286.1438	0.2784	99.5352	0.9745	92.9154	92.9154	Naloxone	328.1543	0.4403	98.8440	0.7928	96.4197
6/5-MAPB		190.1226	1.0261	99.6256	1.5285	99.3207	N-Desalkylflurazepam	289.0538	1.1746	99.6046	1.2005	98.9066	98.9066	N-Desalkylflurazepam	470.2317	1.1216	96.8872	0.7430	97.1506
6-Acetylmorphine		328.1543	0.4403	98.8440	0.7928	96.4197	Nefazodone	250.1437	2.2353	97.0007	1.0438	98.5456	98.5456	N-Ethylpentylene/N,N-Dimethylpentylene	305.1859	0.3910	99.5322	0.4108	97.5489
7/4/3-MAPB		190.1226	1.0261	99.6256	1.5285	99.3207	N-Methyl-Norcarfentanil	271.0633	1.1876	99.3032	0.7893	98.9215	98.9215	Nordiazepam	266.1539	1.4257	98.9884	1.2922	99.0696
7-Aminoclonazepam		286.0742	1.1568	99.3415	0.8375	98.9995	Nordoxepin	296.1257	0.7822	99.5385	0.9228	98.6686	98.6686	Norfluoxetine	324.1387	0.0979	85.4136	0.9646	93.8497
7-Aminofunitrazepam		284.1194	2.0733	97.7982	0.6751	99.0319	Norketamine	224.0837	1.3322	99.3716	1.2579	99.2282	99.2282	Normeperidine	234.1489	0.9005	99.5964	0.3965	99.0849
7-APDB		178.1226	1.1267	99.6459	1.0252	97.9841	Norpropoxyphene	326.2115	1.0246	99.2507	0.6611	98.1907	98.1907	Oxazepam	287.0582	-0.0699	97.9141	0.8636	93.8947
AB-PINACA		331.2129	1.2961	98.7074	0.3699	98.9559	Oxycodone	316.1543	0.8103	99.7328	1.2426	98.7149	98.7149	Oxymorphone	302.1387	0.0979	85.4136	0.9646	93.8497
Acetaminophen		152.0706	1.3173	96.1147	-0.0152	95.2489	Paracetamol	369.2337	0.0421	98.5065	0.7074	93.1344	93.1344	Paracetamol	353.2224	0.1671	99.7847	0.5610	98.9805
Acetylcodeine		342.1700	1.2909	99.3449	0.7533	99.2765	Paracetamol	353.2024	0.1671	99.7847	0.5610	98.9805	98.9805	Paracetamol	353.2024	0.1671	99.7847	0.5610	98.9805
Acetylentanyl/Benzylfentanyl		323.2118	-0.2557	86.9129	-0.3048	76.5476	Paracetamol	353.2024	0.1671	99.7847	0.5610	98.9805	98.9805	Paracetamol	353.2024	0.1671	99.7847	0.5610	98.9805
Acrylfentanyl		335.2118	0.0969	99.3141	0.6229	95.0780	PCP	244.2060	1.2132	99.4043	1.0048	99.3979	99.3979	Pentazocine	286.2165	1.3908	99.4933	1.1017	99.4688
AH-7921		329.1182	0.3177	99.4097	0.6871	97.5881	Pentylone	236.1281	1.4895	99.3200	1.1191	99.0859	99.0859	Phenazepam</td					