

Qualitative and Quantitative Analysis of Organic Acids in Urine by Gas Chromatography-High Resolution Mass Spectrometry with Hard Ionization and Soft Ionization

Product: Mass spectrometer (MS)

Introduction

The test of urine organic acids is an essential component of the study of inherited metabolic diseases, such as inborn error of metabolism. Urine contains a variety of organic acids that forms through diverse sources, including normal and abnormal metabolism, drugs and drug metabolism, or even xenobiotics and dietary supplements. The gas chromatography - mass spectrometry (GC-MS), especially GC - single quadrupole MS(GC-QMS) is generally chosen for the analysis of urine organic acids. However, since GC-QMS lacks the ability to the accurate identification of organic acids in urine due to its complicated mixture especially when creating a new method, GC-high resolution MS (GC-HRMS) is a more qualified system instead.

JMS-T2000GC, a GC-HRMS system that incorporates a combination ion source of electron ionization (EI) and field ionization (FI), is capable of switching between the EI and FI modes without breaking the vacuum. The analysis of data acquired by the 2 ionization methods through msFineAnalysis AI software leads to the estimation of chemical structures as well as compositions, which is practical for the determination of compounds in organic acids.

In this work, the qualitative analysis of the chemical composition and structure of organic compounds in urine and the quantitative analysis of target organic acid using JMS-T2000GC with msFineAnalysis AI was executed.

Measurement

The organic acids as sample solutions were extracted from acidified urines by liquid-liquid extraction with ethyl acetate, followed by trimethylsilyl (TMS) derivatization to increase the volatility of organic acids. 1 μ L of the sample solution was then injected in JMS-T2000GC by an autosampler. The msFineAnalysis AI software was used for the qualitative analysis and the Escrime™ software was used for quantitative analysis. The detailed GC-HRMS measurement conditions are presented in Table 1.

Table 1. Measurement conditions



JMS-T2000GC "AccuTOF™ GC Alpha"

GC-HRMS	JMS-T2000GC (JEOL Ltd.)
GC inlet mode	Splitless
GC inlet temperature	280 °C
GC Column	DB-5MS, 30m x 0.25mm, 0.25 μ m
GC Oven	60°C (1min) → 10°C/min → 300°C (3min)
Carrier gas	He, 1.0mL/min
MS Ionization	EI+: 70eV, 300 μ A
	FI+: -10 kV with JEOL 5 μ m emitter (10mA, 6msec)
MS Monitor ion range	<i>m/z</i> 45-800
Analysis software	Qualitative: msFineAnalysis AI (JEOL Ltd.) Quantitative: Escrime™ (JEOL Ltd.)

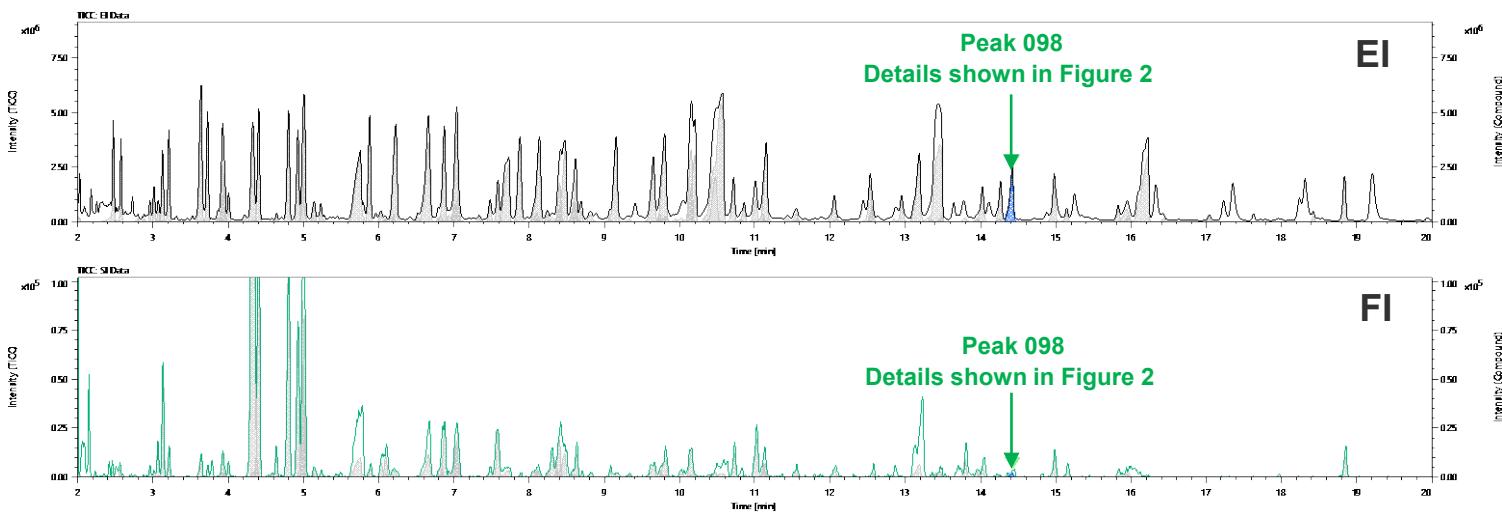


Figure 1. TICCs of a urine organic acid from sample (top: EI, bottom: FI)

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Qualitative Analysis Results

Figure 1 shows both the EI and FI total ion current chromatograms (TICCs) of urine organic acids from the sample, resulting in the detection of 108 compounds including 29 target organic acids, whose msFineAnalysis AI analysis results are shown in red columns in Table 2. Peak [33] was assigned as 5-Methyl-3-isoxazolepropanoic acid trimethylsilyl ester, a TMS derivative compound from oximated succinyl acetone. Some co-eluted components due to the failure of chromatographically separation such as peak [023] and [024] were however detected separately by the deconvolution function of msFineAnalysis AI.

The results of both integrated analysis and AI structural analysis of peak[098] acquired by msFineAnalysis AI were shown in Figures 2 and 3 respectively. Molecular ion was not detected in the EI mass spectrum of Figure 2 while the $[M+H]^+$ ion was detected in the FI mass spectrum, which implies the importance of the combination analysis of soft ionization methods. The chemical formula was estimated to be $C_{16}H_{34}NO_5Si_2$ from the exact mass of the detected $[M+H]^+$ ion, which did not match the estimated chemical composition from the molecular ion or the candidate compound from the NIST database search results. However, msFineAnalysis AI obtained an AI library that contains plenty of EI mass spectra of compounds which are not registered in the NIST database, and compounds that match the chemical composition estimated from the molecular ion and have a matching EI fragment ions pattern were shown as candidates in the search result of AI library (Figure 3).

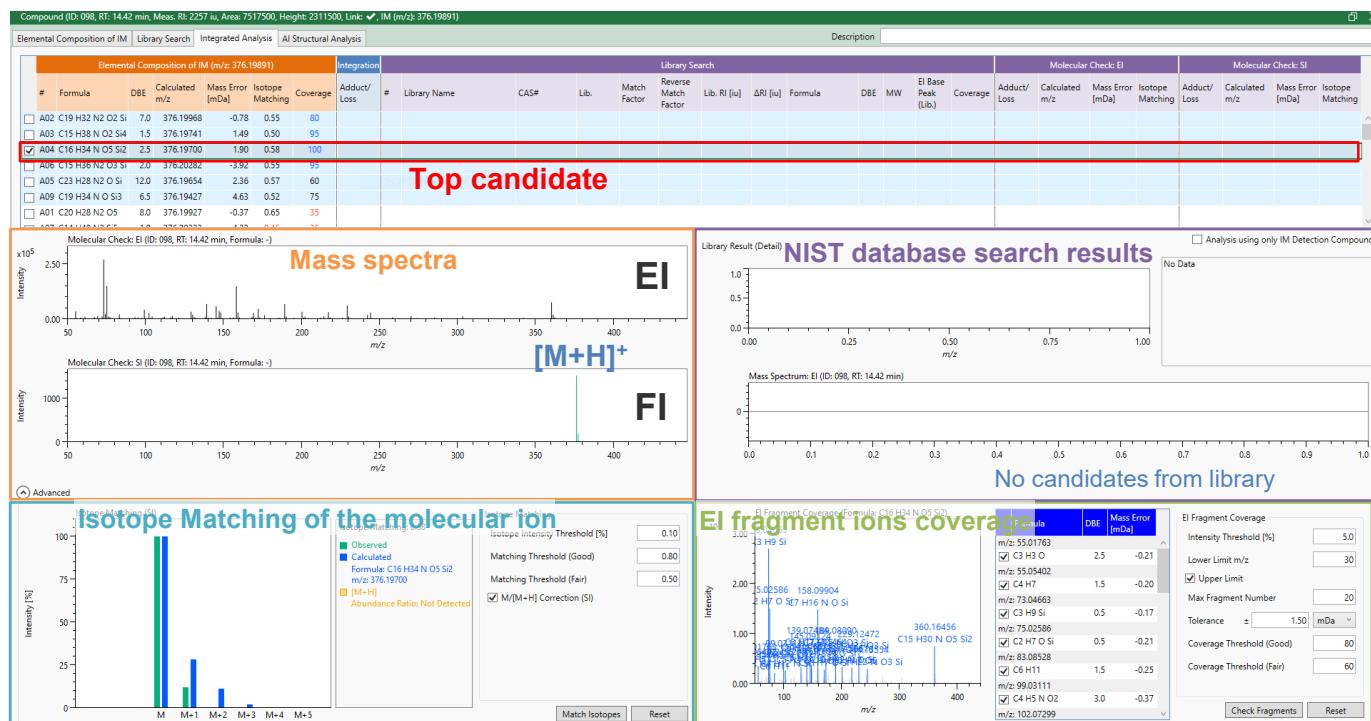


Figure 2. Integrated Analysis results of peak 098

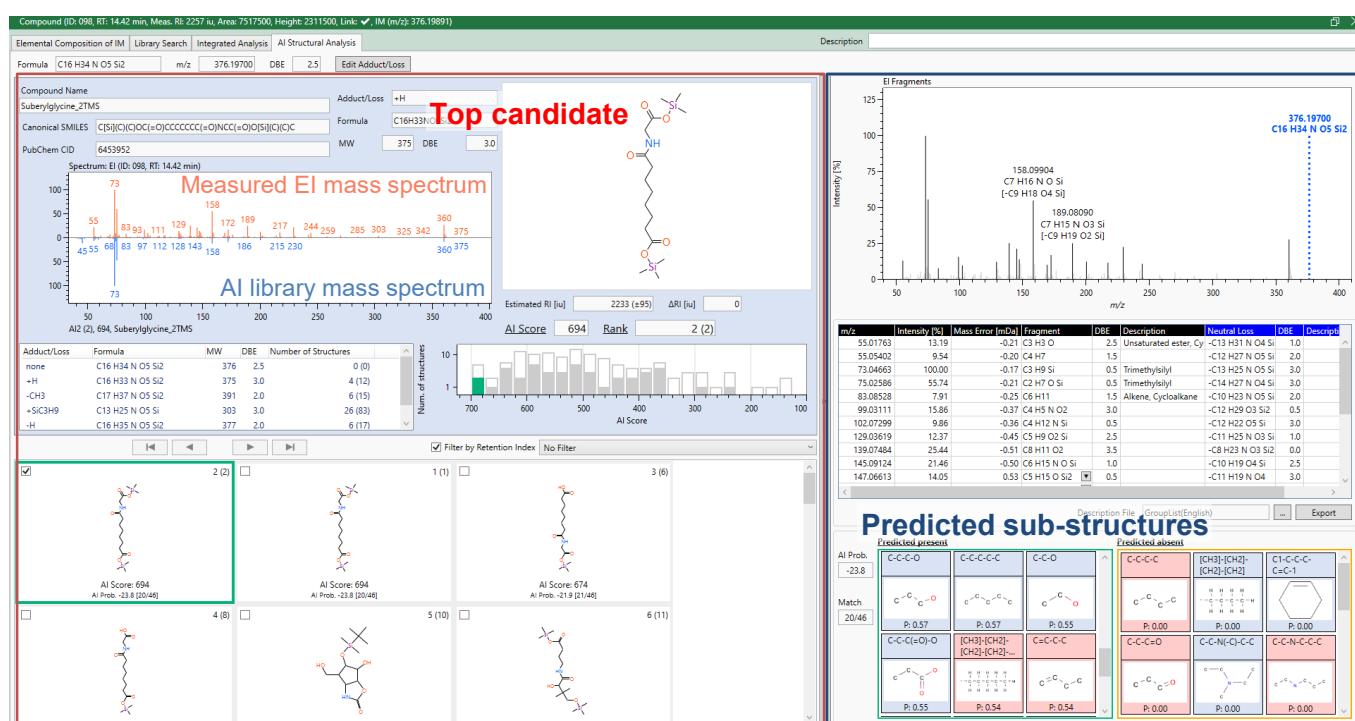


Figure 3. AI Structural Analysis results of peak 098

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Table 2. msFineAnalysis AI results

ID	RT [min]	General			Compound Name	Total Result							
		Meas. RI [i.u]	Area	IM m/z		Lib.	Match Factor / AI Score	Lib. RI [i.u]	ΔRI [i.u]	Formula	DBE	Adduct/ Loss	Calculated m/z
001	2.00	1096	840854	159.10743	EI	trimethylsilyl N-methylpropanimidate	AI	793	929 (±95)	C7H17N O Si	1.0	none	159.10739
002	2.48	1139	6417982	219.08602	EI	Lactic Acid, 2TMS derivative	mainlib	918	1066	C8H19 N O3 Si2	1.5	-CH3	219.08672
003	2.50	1140	153422	233.10237	EI	α-Hydroxyisobutyric acid, 2TMS derivative	repplib	717	1071	C9 H21 O3 Si2	1.5	-CH3	233.10237
004	2.58	1147	4160738	205.07132	EI	Glycolic acid, 2TMS derivative	repplib	953	1080	C7H17 O3 Si2	1.5	-CH3	205.07107
005	2.80	1167	257103	187.15102	EI	2-Octanol, TMS derivative	repplib	872	1108	C10 H23 O Si	0.5	-CH3	187.15127
006	2.88	1174	250064	143.08833	EI	trimethyl-[{E}]-2-methylbut-1-enoylsilane	AI	738	984 (±95)	C7 H15 O Si	1.5	-CH3	143.08867
007	2.96	1181	586136	233.09056	EI	Glyoxylic oxime acid, bis(trimethylsilyl)-	mainlib	785	1135	C8H19 N O3 Si2	2.0	none	233.08980
008	3.02	1186	1871002	307.12302	EI	Oxalic acid, 2TMS derivative	mainlib	767	1134	C11 H27 O4 Si3	1.5	+SiC3H9	307.12117
009	3.07	1191	778960	180.09790	EI	p-Cresol, TMS derivative	mainlib	779	1153	C10 H16 O Si	4.0	none	180.09649
010	3.13	1196	4614950	247.10678	EI	Pyruvic acid oxime, bis(trimethylsilyl)- deriv.	mainlib	923	1157	C9 H21 N O3 Si2	2.0	none	247.10545
011	3.22	1203	5653093	233.10260	EI	3-Hydroxybutyric acid, 2TMS derivative	repplib	889	1167	C9 H21 O3 Si2	1.5	-CH3	233.10237
012	3.64	1237	13610480	321.13698	EI	Propanedioic acid, 2TMS derivative	repplib	931	1216	C12 H29 O4 Si3	1.5	+SiC3H9	321.13682
013	3.73	1244	7871900	262.10558	EI	Methylmalonic acid, 2TMS derivative	repplib	933	1223	C10 H22 O4 Si2	2.0	none	262.10511
014	3.93	1260	6156020	261.13359	EI	2-Hydroxyisocaprylic acid, 2TMS derivative	repplib	858	1248	C11 H25 O3 Si2	1.5	-CH3	261.13367
015	3.95	1262	5469197	233.09926	EI	4-Hydroxybutanoic acid, 2TMS derivative	repplib	833	1238	C9 H21 O3 Si2	1.5	-CH3	233.10237
016	4.01	1267	1771635	258.09575	EI	2-Propanoic acid, 2-(acetylamo)-_2TMS	AI	700	1302 (±95)	C10 H20 N O3 Si2	3.5	-CH3	258.09762
017	4.23	1284	274853	189.08699	EI	Urea, 2TMS derivative	mainlib	771	1246	C6 H17 N2 O2 Si2	1.5	-CH3	189.08739
018	4.33	1292	12007097	315.10364	EI	Silanol, trimethyl-, phosphate (3:1)	mainlib	916	1285	C9 H28 O4 P Si3	-0.5	+H	315.10275
019	4.41	1298	9735226	349.16825	EI	Ethylmalonic acid, 2TMS derivative	mainlib	908	1286	C14 H33 O4 Si3	1.5	+SiC3H9	349.16812
020	4.64	1317	548152	254.11738	EI	Catechol, 2TMS derivative	mainlib	871	1324	C12 H22 O2 Si2	4.0	none	254.11528
021	4.80	1330	11317113	335.15279	EI	Butanedioic acid, 2TMS derivative	mainlib	947	1321	C13 H31 O4 Si3	1.5	+SiC3H9	335.15247
022	4.93	1340	7172905	349.16846	EI	Methylsuccinic acid, 2TMS derivative	mainlib	928	1330	C14 H33 O4 Si3	1.5	+SiC3H9	349.16812
023	4.99	1345	4213726	329.15392	EI	Uracil, 2TMS derivative	repplib	872	1345	C13 H29 N2 O2 Si3	3.5	+SiC3H9	329.15313
024	5.02	1347	10038423	395.19272	EI	Glyceric acid, 3TMS derivative	repplib	925	1342	C15 H39 O4 Si4	0.5	+SiC3H9	395.19199
025	5.12	1355	336850	337.17194	EI	2,3-Dihydroxybutanoic acid tritms	mainlib	714	1362	C13 H33 O4 Si3	0.5	+H	337.16812
026	5.14	1357	1678911	337.17750	EI	serine hydroxamate,_3TMS	AI	745	1452 (±95)	C10 H33 N2 O3 Si3	0.5	+H	337.17935
027	5.24	1364	1227728	409.20702	EI	(R*,S*)-2,3-Dihydroxybutanoic acid, tris(trimethylsilyl) deriv.	mainlib	910	1234 (±381)	C16 H41 O4 Si4	0.5	+SiC3H9	409.20764
028	5.75	1406	16077539	290.16031	EI	Isobutyrylglycine, TMS derivative	mainlib	886	1372	C12 H28 N O3 Si2	1.5	+SiC3H9	290.16022
029	5.88	1417	10130214	349.16790	EI	Pentanedioic acid, 2TMS derivative	mainlib	931	1409	C14 H33 O4 Si3	1.5	+SiC3H9	349.16812
030	6.03	1429	1000163	362.19938	EI	Glycine, N-(2-methyl-1-propoxyl)-N-(trimethylsilyl), trimethylsilyl ester	mainlib	809	1414	C15 H36 N O3 Si3	1.5	+SiC3H9	362.19975
031	6.09	1434	286530	291.14075	EI	bis(trimethylsilyl) 2-methylpentanedioate	AI	718	1406 (±95)	C10 H27 O4 Si2	1.5	+H	291.14424
032	6.19	1443	572961	300.14389	EI	trimethylsilyl 3-(3-methyl-1,2-oxazol-5-yl)propanoate	AI	647	1400 (±95)	C13 H26 N O3 Si2	3.5	+SiC3H9	300.14457
033	6.23	1446	14930674	227.09678	EI	5-Methyl-3-isoxazolepropanoic acid trimethylsilyl ester	mainlib	789	1344 (±381)	C10 H17 N O3 Si	4.0	none	227.09722
034	6.52	1470	431613	365.16214	EI	D(-)-Citramalic acid, 3TMS derivative	mainlib	729	1487	C14 H33 O5 Si3	1.5	+H	365.16303
035	6.66	1482	14654733	232.13616	EI	N-(2-methylbutanoyl)glycine,_TMS	AI	725	1492 (±95)	C10 H22 N O3 Si	1.5	+H	232.13635
036	6.79	1492	349385	289.13156	EI	bis(trimethylsilyl) 3-methylpent-2-enoate	AI	662	1474 (±95)	C12 H25 O4 Si2	2.5	+H	289.12859
037	6.83	1496	859327	304.17513	EI	N-2-Methylbutyrylglycine, 2TMS derivative	mainlib	737	1491	C13 H30 N O3 Si2	1.5	+H	304.17587
038	6.85	1497	242023	355.15705	EI	trimethylsilyl 2-trimethylsilyloxybenzoate	AI	621	1510 (±95)	C16 H31 O3 Si3	4.5	+SiC3H9	355.15755
039	6.87	1500	10697228	282.11085	EI	Salicylic acid, 2TMS derivative	repplib	940	1522	C13 H22 O3 Si2	5.0	none	282.11020
040	6.89	1501	503265	349.16897	EI	trimethyl-[4-trimethylsilyloxy-5-(trimethylsilyloxyethyl)oxolan-2-yl]oxysilane	AI	633	1495 (±95)	C14 H33 O4 Si3	1.5	-H	349.16812
041	6.94	1505	907439	236.03786	EI	6-Methyl-1,3-oxathiazin-4(3H)-one 2,2-dioxide, TMS	mainlib	811	1523	C17 H4 N O4 Si	2.5	+H	236.04073
042	7.03	1513	12796141	291.14529	EI	Hexanedioic acid, 2TMS derivative	repplib	933	1512	C12 H27 O4 Si2	1.5	+H	291.14424
043	7.05	1515	4517844	346.16920	EI	DL-Pyroglutamic acid, 2TMS	mainlib	880	1523	C14 H32 N O3 Si3	2.5	+SiC3H9	346.16845
044	7.33	1539	360644	289.12803	EI	3-Methyladipic acid, 2TMS derivative	mainlib	822	1544	C12 H25 O4 Si2	2.5	-CH3	289.12859
045	7.48	1552	1776015	286.10543	EI	5-Hydroxymethyl-2-furoic acid, 2TMS derivative	mainlib	897	1547	C12 H22 O4 Si2	4.0	none	286.10511
046	7.59	1561	3556196	296.07581	EI	1,2-Dithiane-4,5-diol, (Z)-, 2TMS derivative	repplib	889	1601	C10 H24 O2 S2 Si2	1.0	none	296.07508
047	7.73	1573	12249784	230.11963	EI	Glycine, N-(2-methyl-1-oxo-butene)-, trimethylsilyl ester, (E)-	mainlib	915	1579	C10 H20 N O3 Si	2.5	+H	230.12070
048	7.88	1587	11201975	365.16051	EI	Pentanedioic acid, 2-[(trimethylsilyl)oxy]-, bis(trimethylsilyl) ester	repplib	920	1583	C14 H33 O5 Si3	1.5	+H	365.16303
049	8.05	1602	456369	296.12605	EI	3-Hydroxyphenylacetic acid, 2TMS derivative	mainlib	755	1618	C14 H24 O3 Si2	5.0	none	296.12585
050	8.10	1606	1308053	379.17960	EI	Pentanedioic acid, 3-methyl-3-[(trimethylsilyl)oxy]-, bis(trimethylsilyl) ester	mainlib	739	1614	C15 H35 O5 Si3	1.5	+H	379.17868
051	8.14	1610	12976144	305.15935	EI	Pimelic acid, 2TMS derivative	mainlib	909	1610	C13 H29 O4 Si2	1.5	+H	305.15989
052	8.24	1619	827860	282.11082	EI	4-Hydroxybenzoic acid, 2TMS derivative	repplib	843	1635	C13 H22 O3 Si2	5.0	none	282.11020
053	8.31	1625	9707076	295.14316	EI	Paracetamol, 2TMS derivative	repplib	818	1625	C14 H25 N O2 Si2	5.0	none	295.14183
054	8.41	1634	4823139	296.12605	EI	4-Hydroxybenzeneacetic acid, 2TMS derivative	repplib	905	1648	C14 H24 O3 Si2	5.0	none	296.12585
055	8.48	1641	12537944	377.14935	EI	A-ketoglutaric acid oxime,_3TMS	AI	702	1656 (±95)	C14 H31 N O5 Si3	3.0	none	377.15045
056	8.60	1652	314578	242.08656	EI	trimethylsilyl 2-(furan-2-carbonylamino)acetate	AI	760	1657 (±95)	C10 H16 N O4 Si	4.5	+H	242.08431
057	8.62	1653	5215354	318.19184	EI	Hexanoyl glycine, bis(trimethylsilyl)-	mainlib	911	1663	C14 H32 N O3 Si2	1.5	+H	318.19152
058	8.69	1660	1029228	378.15546	EI	2-Ketoglutaric acid, tri-TMS	mainlib	718	1645	C16 H32 N O5 Si3	2.5	+H	378.15628
059	8.81	1671	1069656	237.02884	EI	2-(Formylamino)benzoic acid, TMS derivative	mainlib	855	1693	C11 H15 N O3 Si	6.0	none	237.08157
060	9.07	1694	3961868	309.12433	EI	2-(Formylamino)benzoic acid, 2TMS derivative	mainlib	797	1707	C14 H23 N O3 Si2	6.0	none	309.12110
061	9.74	1777	329413	295.08909	EI	Terephthalic acid, 2TMS derivative	mainlib	911	1707	C14 H13 O4 Si2	6.5	-CH3	295.08164
062	10.02	1784	715977	384.15999	EI	4-Hydroxymandelic acid, 3TMS derivative	mainlib	811	1738	C17 H32 O4 Si3	5.0	none	384.16029
063	10.15	1797	10465545	324.14308	EI	Hippuric acid, 2TMS derivative	repplib	890	1821	C15 H26 N O3 Si2	5.5	+H	324.14457
064	10.73	1757	1138142	312.12172	EI	Vanilllic Acid, 2TMS derivative	repplib	785	1776	C14 H24 O4 Si2	5.0	none	312.12076
065	9.80	1763	13581808	391.14231	EI	Aconitic acid, (Z)-, 3TMS derivative	repplib	939	1741	C15 H31 O6 Si3	3.5	+H	391.14229
066	9.82	1765	970253	325.13607	EI	Homovanillic Acid, 2TMS derivative	repplib	806	1785	C15 H26 O4 Si2	5.0	none	326.13641
067	9.94	1777	329413	295.08909	EI	805	1799	C13 H19 O4 Si2	6.5	-CH3	295.08164		
068	10.02	1784	715977	384.15999	EI	4-Hydroxymandelic acid, 3TMS derivative	mainlib	734	1797	C17 H32 O4 Si3	5.0	none	384.16029
069	10.86	1867	1865334	351.17974	EI	2-Hydroxybiphenyl, O-bis-TMS	mainlib	898	1888	C18 H31 O3 Si2	5.5	-CH3	351.18062
070	10.21	1802	405.23020	312.12172	EI	Azelaic acid, 2TMS derivative	mainlib	876	1806	C14 H24 O4 Si2	1.5	+SiC3H9	405.23072
071	10.47	1829											

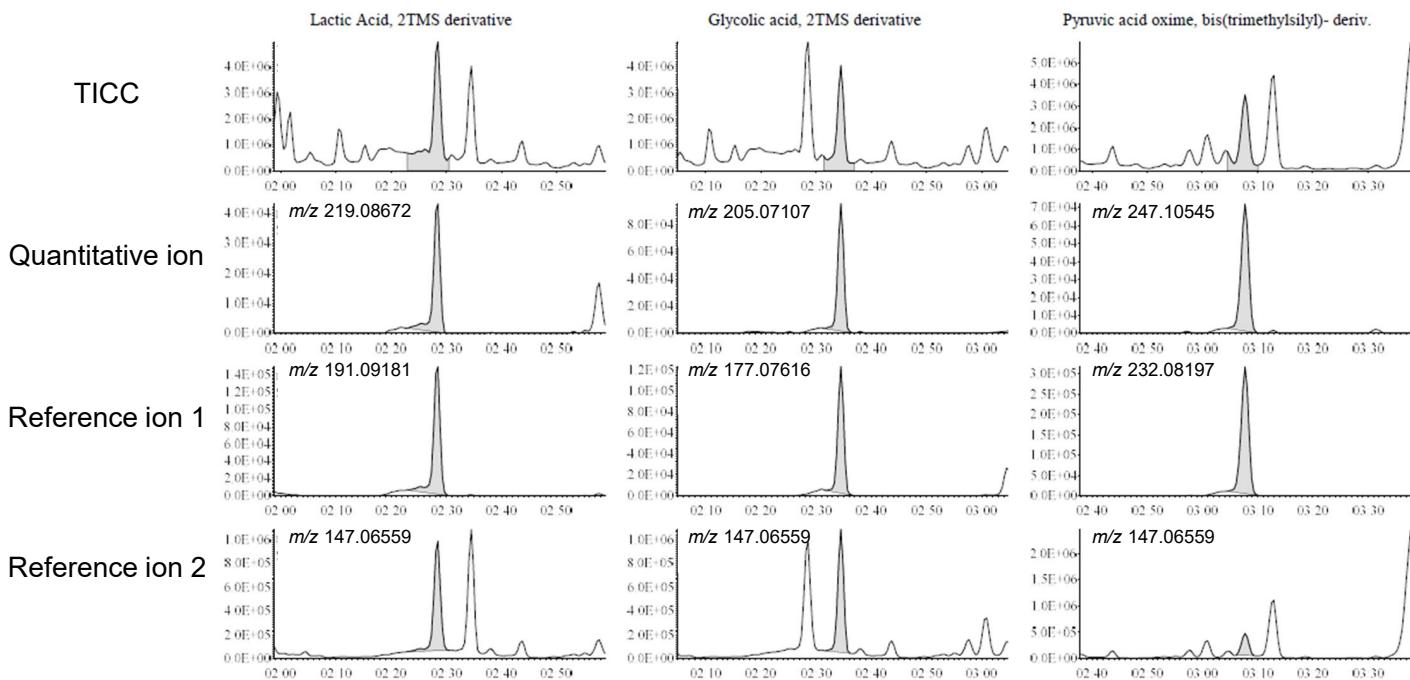
Quantitative Analysis Results

Figure 4 shows the EI TICC and extracted ion chromatograms (EICs) of three target organic acids: 2TMS-lactic acid, 2TMS-glycolic acid and 2TMS-pyruvic acid oxime. Multiple peaks were detected in TICC for the complicated components in urine. However, the EICs created by the accurate m/z signal manages to detect the target compound while limiting the influence of other compounds to the minimum.

All organic acids, including co-eluted organic acids, show detectable peak shapes for quantitation analysis (partially presented in Figure 4). As an example of quantitative analysis, the quantitative values of target organic acids in several urine samples shown in Table 3 were calculated from the area ratio with reference samples at known concentrations.

Table 3. Quantitative values of target organic acids (Unit: μM)

Target Compounds	Reference sample	Negative control	Sample 1	Sample 2	Sample 3
Lactic Acid, 2TMS derivative	110.39	22.125	47.965	21.408	106.486
Glycolic acid, 2TMS derivative	101.82	21.094	296.258	95.486	68.736
Pyruvic acid oxime, bis(trimethylsilyl)- deriv.	69.32	3.29	68.008	17.624	99.834
3-Hydroxybutyric acid, 2TMS derivative	85.26	24.108	189.822	62.896	140.502
Malonic acid (bis-TMS)	597.39	N.D.	52.678	N.D.	N.D.
Methylmalonic acid, 2TMS derivative	16.71	N.D.	2.226	N.D.	4.476
2-Hydroxyisocaproic acid, 2TMS derivative	26.07	N.D.	N.D.	N.D.	0.947
Ethylmalonic acid, 2TMS derivative	34.06	1.178	6.897	1.399	8.493
Succinic acid (2TMS)	58.99	9.027	11.01	3.012	19.784
Methylsuccinic acid, 2TMS derivative	37.46	0.848	4.966	0.578	3.38
Uracil, 2TMS derivative	120.47	0.366	5.711	N.D.	0.446
Glyceric acid, 3TMS derivative	215.37	N.D.	3.335	N.D.	6.28
Isobutyrylglycine, TMS derivative	32.89	0.056	1.301	0.248	0.581
Glutaric acid (2TMS)	31.39	0.679	3.483	1.149	2.107
5-Methyl-3-isoxazolepropanoic acid trimethylsilyl ester	130.87	0.037	N.D.	N.D.	0.016
2-Methylbutyrylglycine, TMS ester	21.03	0.116	0.755	0.023	0.23
DL-Pyroglutamic acid, 2TMS	101.79	15.708	96.916	16.772	47.851
Adipic acid, (2TMS)	31.54	1.259	12.437	1.074	4.637
Tiglyl glycine, 2TMS derivative	117.44	0.258	32.114	0.315	15.06
2-Hydroxyglutaric acid, tri-TMS	72.16	1.863	13.407	1.767	17.665
3-Hydroxy-3-methylglutaric acid, tri-TMS	124.06	N.D.	9.917	43.703	8.687
Pimelic acid, 2TMS derivative	42.46	1.165	4.868	3.24	7.039
Hexanoyl glycine, bis(trimethylsilyl)-	24.07	N.D.	0.061	N.D.	N.D.
Suberic acid, 2TMS derivative	29.27	0.695	2.989	2.026	5.466
Orotic Acid, 3TMS derivative	48.76	0.108	0.621	0.116	0.902
Azelaic acid, 2TMS derivative	27.62	0.983	4.021	2.833	7.592
Sebacic acid, 2TMS derivative	22.87	0.057	0.355	0.153	0.102
N-Acetyltyrosine, 2TMS	25.41	0.068	0.19	0.012	0.372
Suberylglycine-TMS	12.12	N.D.	N.D.	N.D.	N.D.



**Figure 4. TICC and EICs of extracted urine sample.
(Right: 2TMS-lactic acid, Center: 2TMS-glycolic acid, Left: 2TMS-pyruvic acid oxime)**

Summary

The JMS-T2000GC with msFineAnalysis AI is a powerful tool for the identification of the chemical composition and chemical structure of components such as TMS derivatives of organic acids in urine, and for the quantitative analysis by EICs which are created by accurate m/z . The JMS-T2000GC with msFineAnalysis AI is expected to have a wide range of applications for metabolomics and clinical analysis in the future.

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