

Supporting information

for

**Profiling of Aldehyde-containing Compounds by Stable
Isotope Labelling- Assisted Mass Spectrometry Analysis**

Lei Yu, Ping Liu, Ya-Lan Wang, Qiong-Wei Yu, Bi-Feng Yuan, Yu-Qi Feng*

Key Laboratory of Analytical Chemistry for Biology and Medicine (Ministry of Education),
Department of Chemistry, Wuhan University, Wuhan 430072, P.R. China

*To whom correspondence should be addressed. Tel: +86-27-68755595; fax: +86-27-68755595.

E-mail address: yqfeng@whu.edu.cn

Synthesis of 4-APC and 4-APC-*d*₄

N-(4-(2-Bromoethoxy)phenyl)acetamide p-Hydroxyacetanilide (2.0 g) and 1,2-dibromoethane (24.7 g) were added to saturated sodium carbonate solution (20 mL). Tetrabutylammonium perchlorate (0.1 g) was added as a catalyst. The mixture was vigorously stirred for 28 h at 80 °C after which water and the excess of 1,2-dibromoethane were evaporated under high vacuum. The residue was extracted with chloroform (250 mL) to remove inorganic salts and amounts of disubstituted products. The chloroform extract was washed with a saturated sodium carbonate solution (2×). The chloroform layer was dried over MgSO₄, filtered, and concentrated. Recrystallization from ethanol/water yielded a white solid. ¹H NMR (CD₃Cl/drop DMSO-*d*₆, 200 MHz) δ 7.5 (d, 2H, CHCNHAc), 6.8 (d, 2H, CHCOCH₂), 4.2 (t, 2H, OCH₂), 3.6 (t, 2H, CH₂Br), 2.0 (s, 3H, COCH₃).

(2-(4-Acetamidophenoxy)ethyl)trimethylammonium Bromide *N*-(4-(2-Bromoethoxy)phenyl)acetamide (0.5 g) was mixed with a solution of Me₃N in EtOH (5 mL). Cosolvent methylethylketone (5 mL) was added, and the mixture was stirred at room temperature for 4 days, during which time the product precipitated. It was filtered and recrystallized from hot EtOH/water to afford C as pinkish/beige transparent crystals. ¹H NMR (D₂O, 200 MHz) δ 7.3 (d, 2H, CHCNHAc), 7.0 (d, 2H, CHCOCH₂), 4.4 (m, 2H, OCH₂), 3.7 (m, 2H, CH₂NMe₃), 3.2 (s, 9H, NMe₃), 2.1 (s, 3H, COCH₃).

(4-(2-(trimethylammonio)ethoxy)benzenaminium halide (2-(4-Acetamidophenoxy)ethyl)trimethylammonium (0.2 g) was heated at reflux with hydrochloric acid (5 N, 3 mL) for 30 min. After cooling, the solvent was evaporated from the solution leaving a solid residue, which was recrystallized from hot EtOH/water. Filtering, washing, and drying afforded shiny beige crystals with excellent purity. ¹H NMR (D₂O, 200 MHz) δ 7.4 (d, 2H, CHCNH₃), 7.1 (d,

2H, CHCOCH₂), 4.5 (m, 2H, OCH₂), 3.8 (m, 2H, CH₂NMe₃), 3.2 (s, 9H, NMe₃).

4-APC-*d*₄ was prepared in a similar procedure with 1,2-dibromoethane-*d*₄ instead of 1,2-dibromoethane. ¹HNMR (D₂O, 200 MHz) δ 7.2 (d, 2H, CHCNH₃), 7.0 (d, 2H, CHCOCH₂), 3.1 (s, 9H, NMe₃).

Table S1. The slopes of regression lines of the measured peak intensity ratios versus the mean concentration ratios (1:20, 1:10, 1:5, 1:1, 5:1, 10:1 and 20:1) of 4-APC/4-APC-*d*₄ labelled standards of hexanal, benzaldehyde and heptanal using QTRAP mass spectrometer.

Analytes	Linear range (light / heavy)	Regression line		
		Slope	Intercept	R ²
Hexanal	0.05-20	1.064	0.0229	0.9914
Benzaldehyde	0.05-20	0.951	0.0122	0.9973
Heptanal	0.05-20	0.985	-0.0009	0.9972

Table S2. The measured concentration of creatinine in pooled urine sample of healthy controls and lung cancer patients using LC-UV spectrometer.

Analytes	Concentration (mmol/L)
Healthy controls	7.1 ± 0.4
Lung cancer patients	4.8 ± 0.3

Table S3. List of the measured peak intensity ratios (pooled urine sample of lung cancer patients / pooled urine sample of healthy controls) of 16 aldehyde-containing compounds using QTRAP mass spectrometer.

Number	Retention Time (min)	4-APC Labelled (m/z)	4-APC-<i>d</i>₄ Labelled (m/z)	Forward Labelling	Reverse Labelling
1	17.46/17.42	279.2496	283.1369	1.6±0.1	1.7±0.1
2	14.96/14.89	285.1660	289.1653	2.6±0.1	2.8±0.1
3	20.38/20.31	293.2646	297.1595	1.8±0.4	1.9±0.2
4	23.19/23.10	307.2795	311.2139	2.1±0.3	1.9±0.2
5	25.67/25.68	321.2946	325.3206	1.5±0.2	1.7±0.3
6	14.19/14.08	265.2419	269.1554	2.1±0.3	2.3±0.2
7	14.00/13.97	281.1802	285.2436	0.3±0.0	0.3±0.1
8	13.25/13.14	267.1811	271.1567	0.5±0.1	0.3±0.0
9	16.80/16.69	295.1655	299.1372	1.6±0.2	1.8±0.3
10	26.63/26.60	335.3114	339.3357	1.6±0.2	1.8±0.3
11	18.16/18.13	351.2148	355.2672	1.3±0.2	1.3±0.0
12	16.05/16.02	369.1517	373.2130	1.6±0.2	1.8±0.0
13	20.79/20.81	217.1268	221.1210	1.3±0.0	1.4±0.1
14	6.48/6.38	273.1210	277.1547	1.5±0.2	1.6±0.0
15	25.53/25.24	337.0839	341.1340	1.4±0.1	1.7±0.2
16	21.99/21.43	395.9088	400.0807	1.8±0.3	1.9±0.2

Table S4. The measured m/z of 4-APC and 4-APC- d_4 labelled aldehyde-containing compounds from white wine and their prospective molecular formulas using QTOF-MS spectrometer.

No.	RT (min) (4-APC/4-APC- d_4 Labelled)	4-APC Labelled (m/z)	4-APC- d_4 Labelled (m/z)	Prospective Formulas	Compound Name
1	17.35/17.28	279.2496	283.2743	C ₆ H ₁₂ O	Hexanal
2	14.77/14.72	285.1384	289.0646	C ₇ H ₆ O	Benzaldehyde
3	20.33/20.29	293.2646	297.2895	C ₇ H ₁₄ O	Heptanal
4	23.22/23.17	307.2795	311.3049	C ₈ H ₁₆ O	Octanal
5	2.76/2.74	223.1709	227.1902	C ₂ H ₄ O	Acetaldehyde
6	6.42/6.26	251.2132	255.2390	C ₄ H ₈ O	Butanal
7	17.07/17.02	265.2419	269.1560	C ₅ H ₁₀ O	Pentanal
8	17.10/17.03	261.2134	265.2422	C ₂ H ₄ NO	Aminoacetaldehyde
9	8.80/8.74	275.1780	279.2035	C ₅ H ₄ O ₂	Furfural
10	16.37/16.34	299.2190	303.2082	C ₈ H ₈ O	Phenylacetaldehyde
11	27.80/27.76	335.3114	339.3357	C ₁₀ H ₂₀ O	Decanal
12	16.76/16.63	342.1907	346.2894	C ₉ H ₉ NO ₂	p-Acetaminobenzaldehyde
13	16.73/16.76	343.2091	347.2346	C ₉ H ₈ O ₃	Caffeic aldehyde
14	24.32/24.36	349.3255	353.3519	C ₁₁ H ₂₂ O	Undecanal
15	29.04/28.92	383.3114	387.2120	C ₁₄ H ₂₀ O	9Z-13-Tetradecadien-11-ynal
16	19.54/19.56	393.3190	397.3447	C ₁₂ H ₂₂ O ₃	12-Oxo-dodecanoic acid
17	2.77/2.73	224.2809	228.3021		
18	18.39/18.32	237.2011	241.2217		
19	17.07/17.04	262.2011	266.2270		

No., number; RT, retention time.

Figure S1. The pathway for the synthesis of 4-APC and 4-APC- d_4

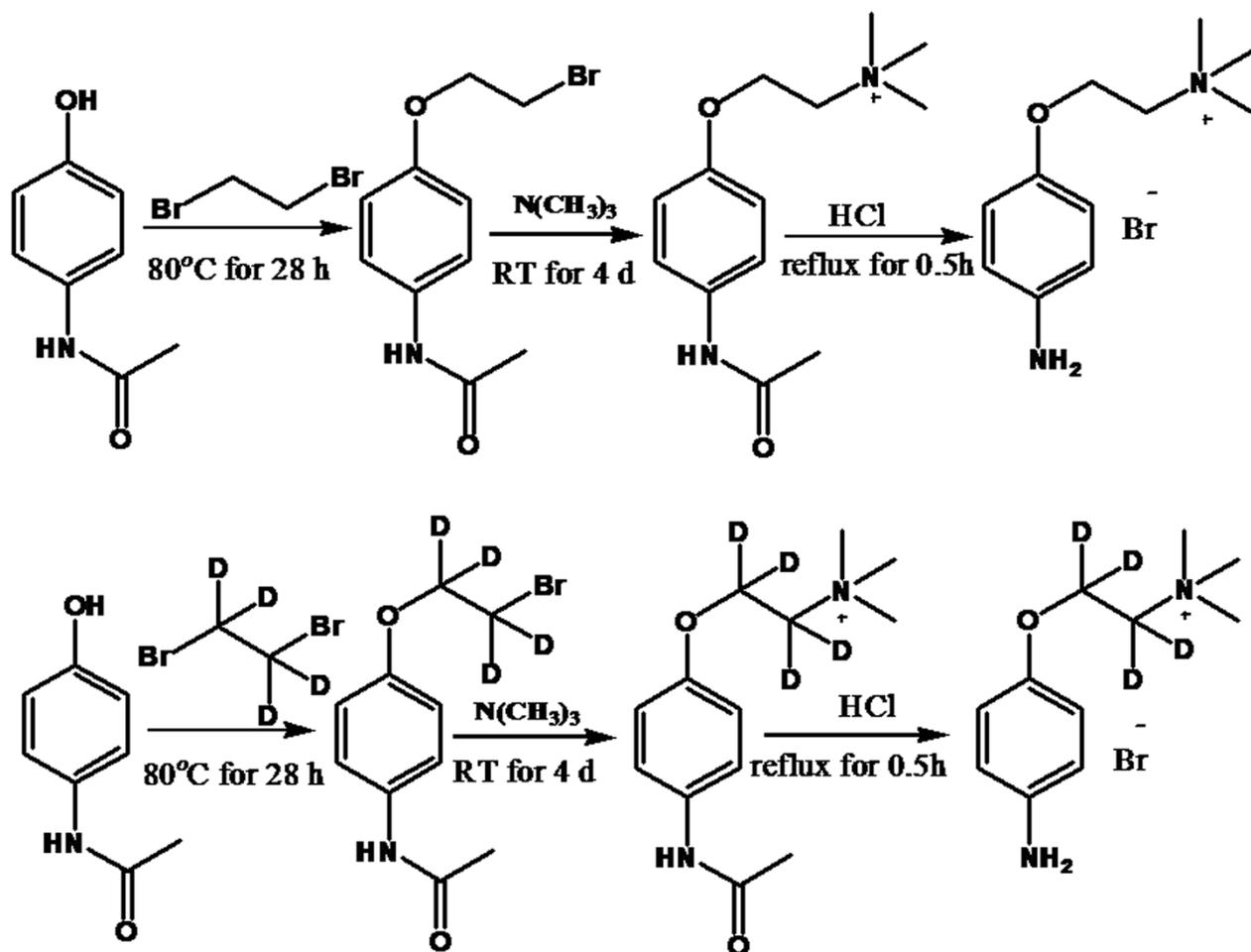


Figure S2. The proposed fragmentation pathway of 4-APC labelled aldehydes.

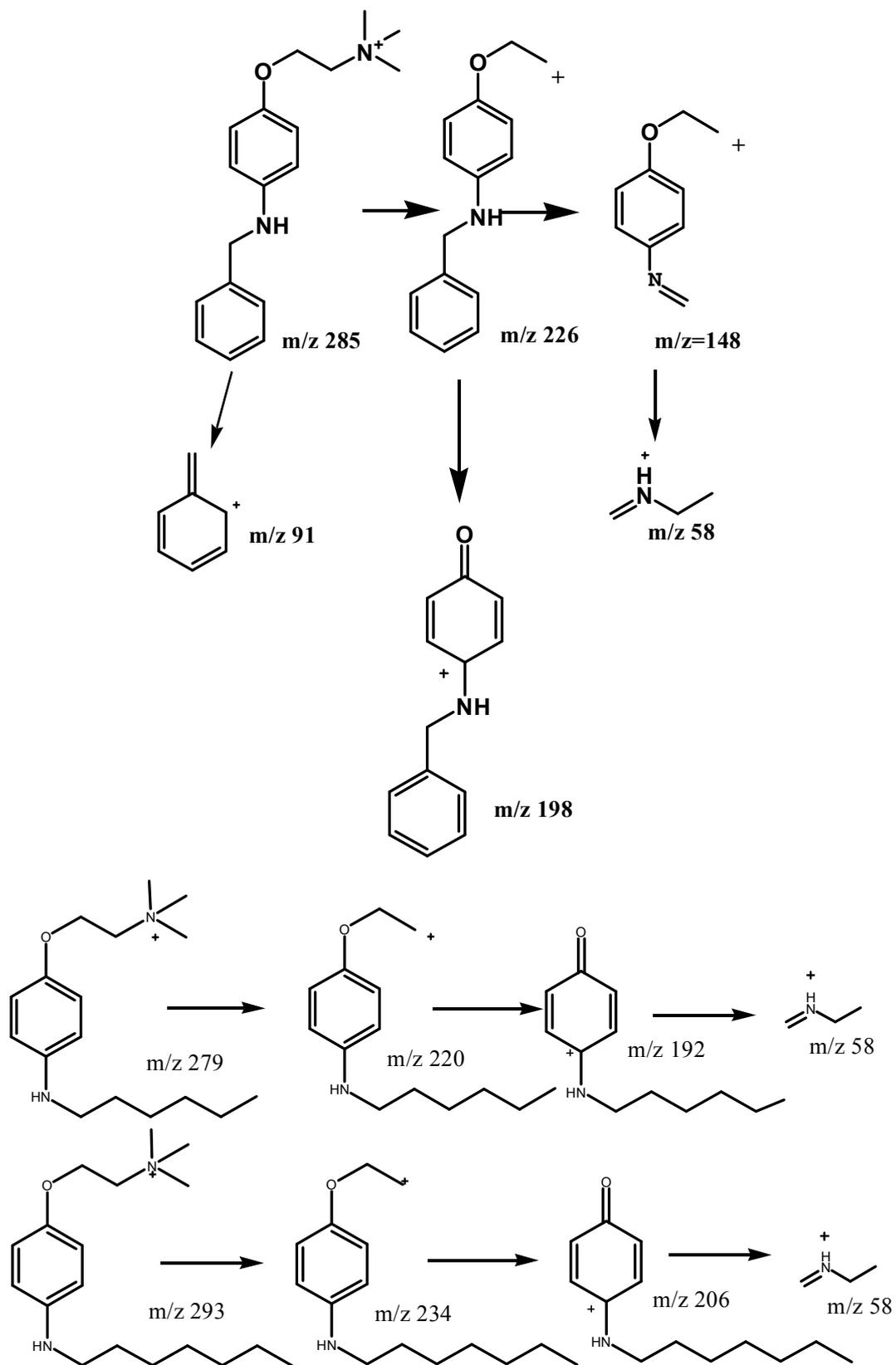
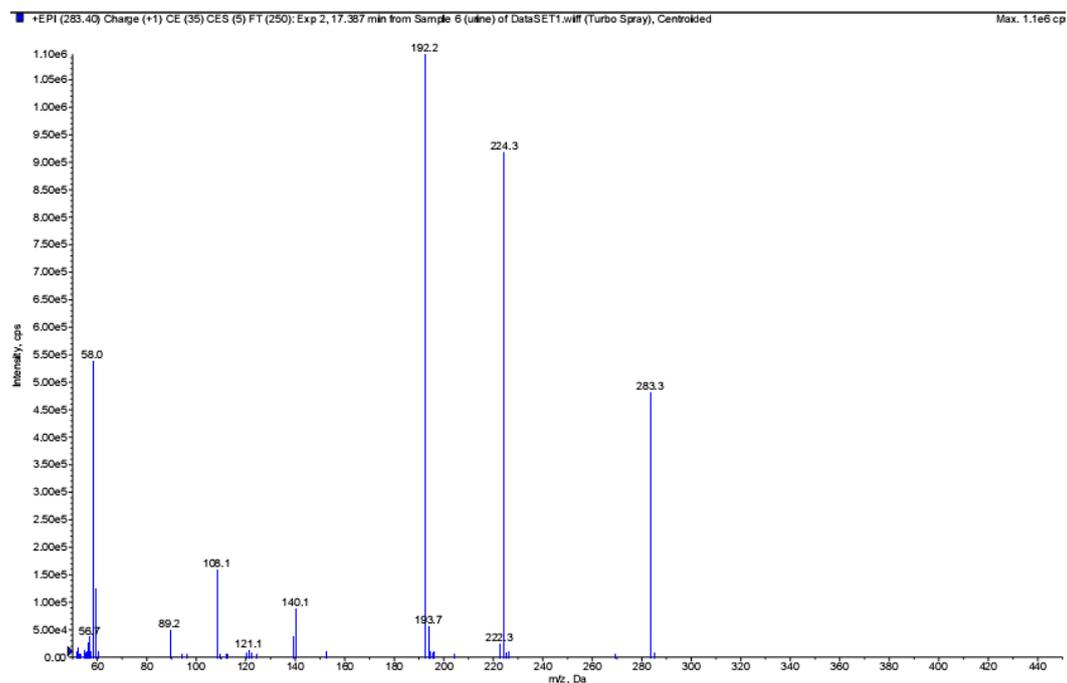
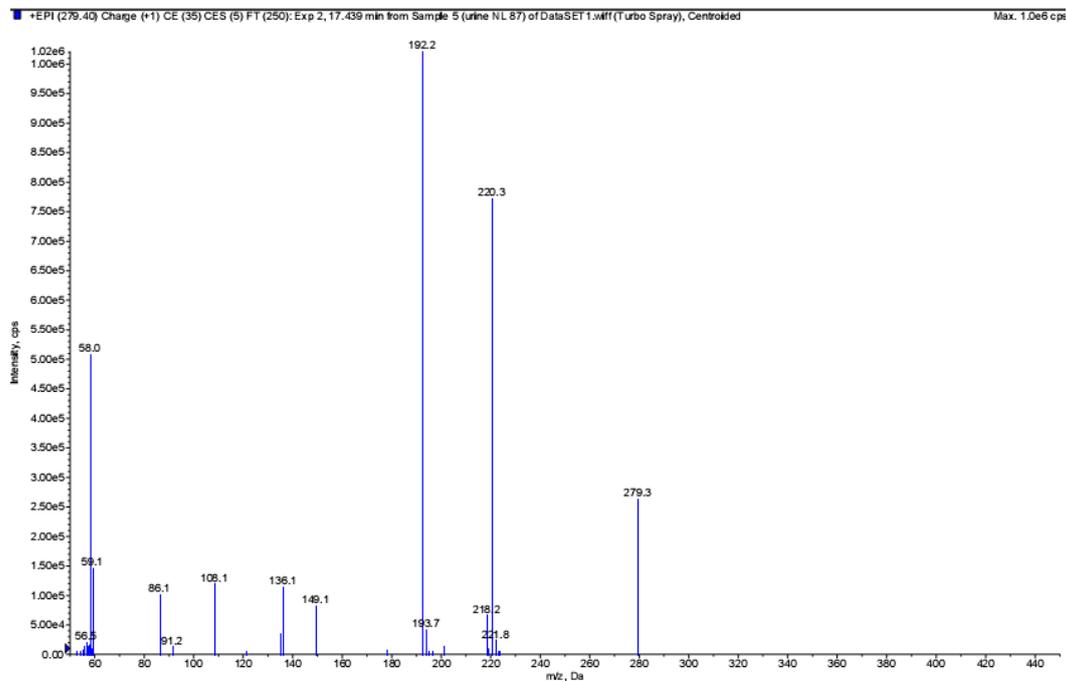
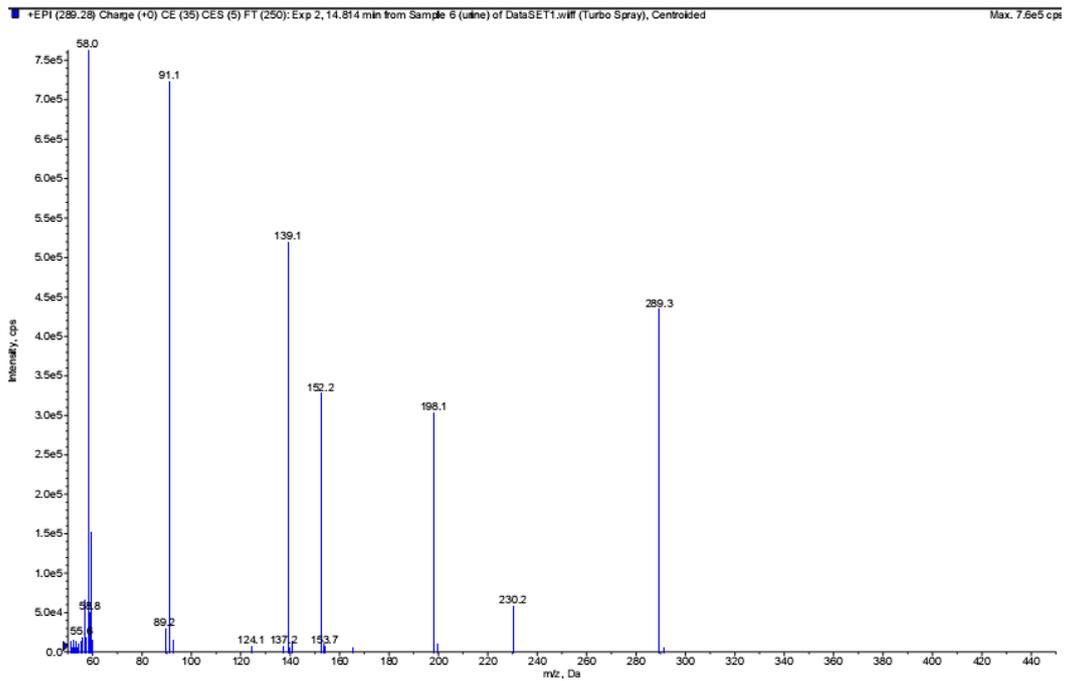
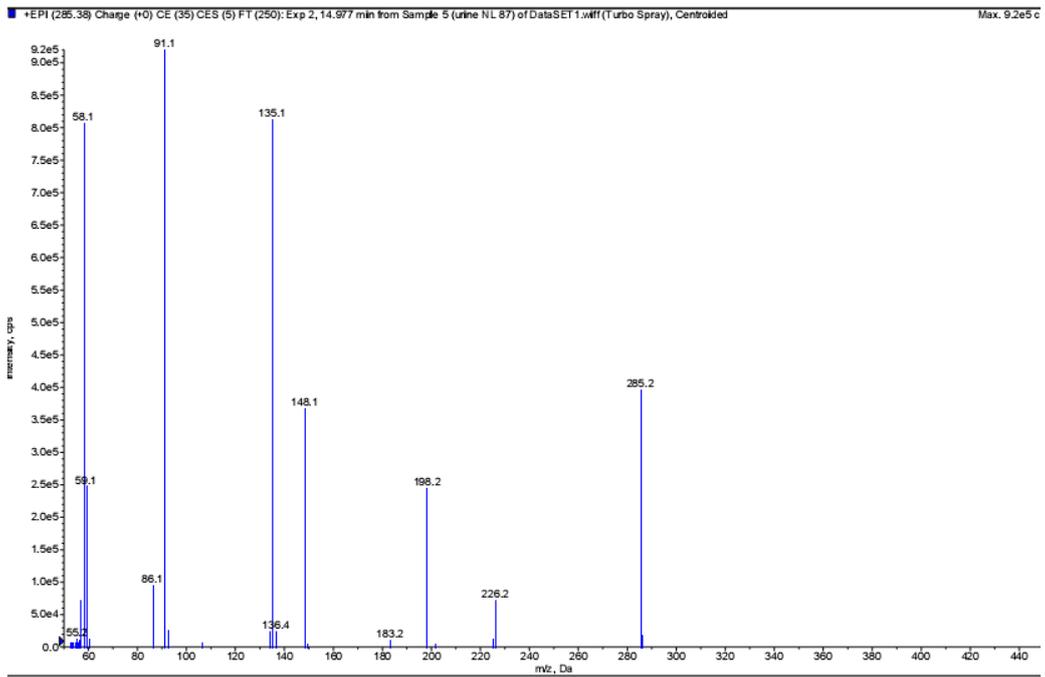


Figure S3. The product ion mass spectra of 4-APC and 4-APC- d_4 labelled compounds in human urine using QTRAP mass spectrometer.

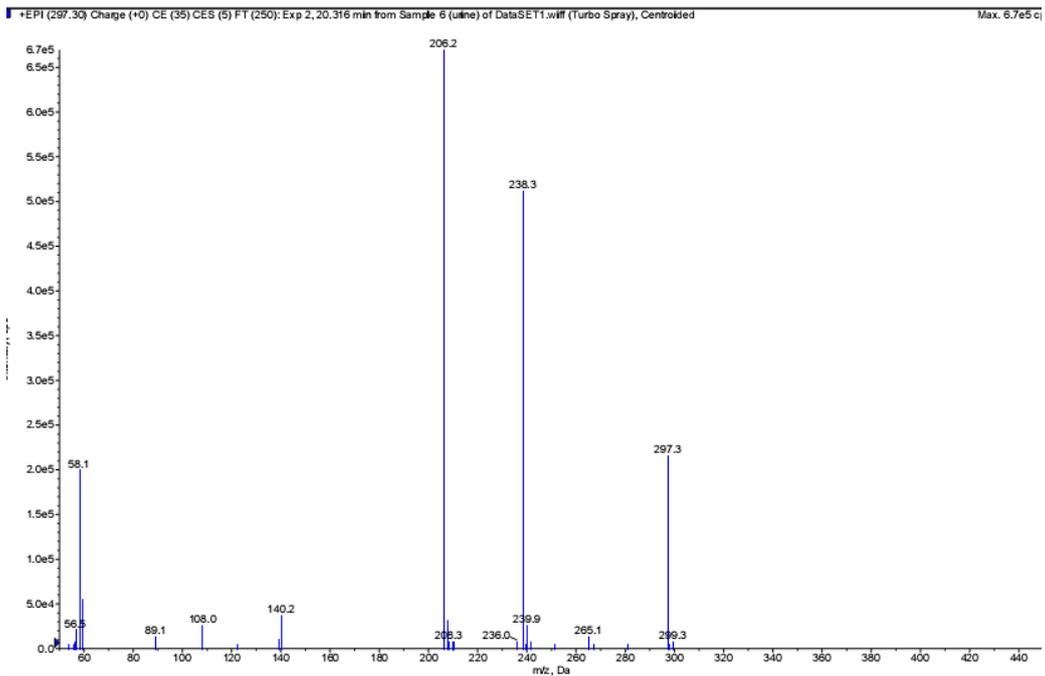
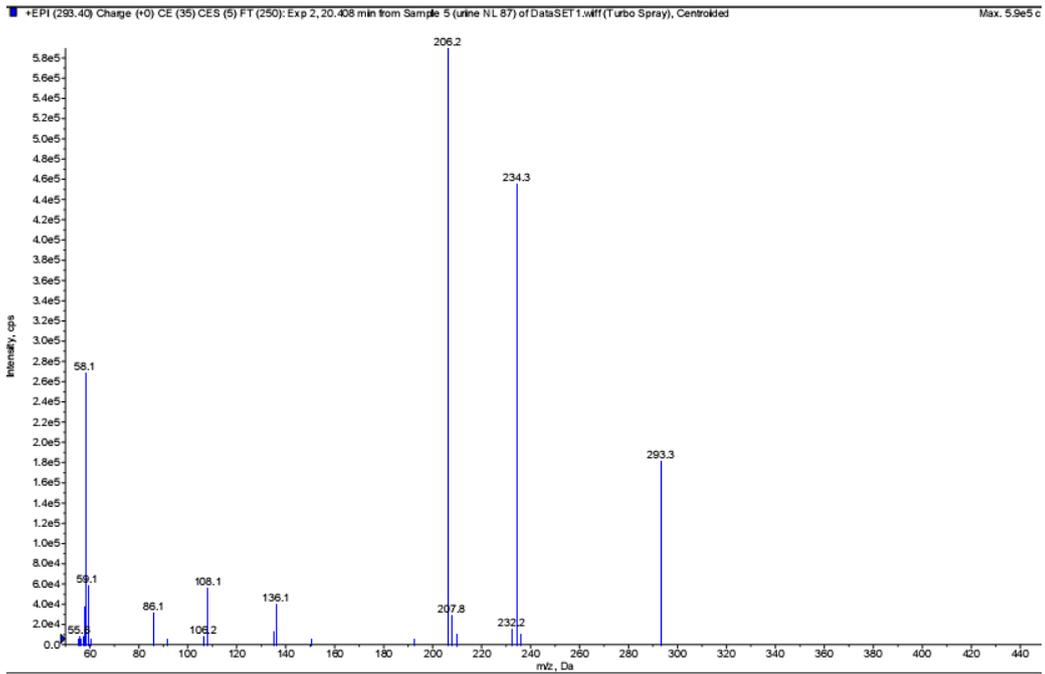
Compound 1, m/z 279/283, $t_r=17.46$ min



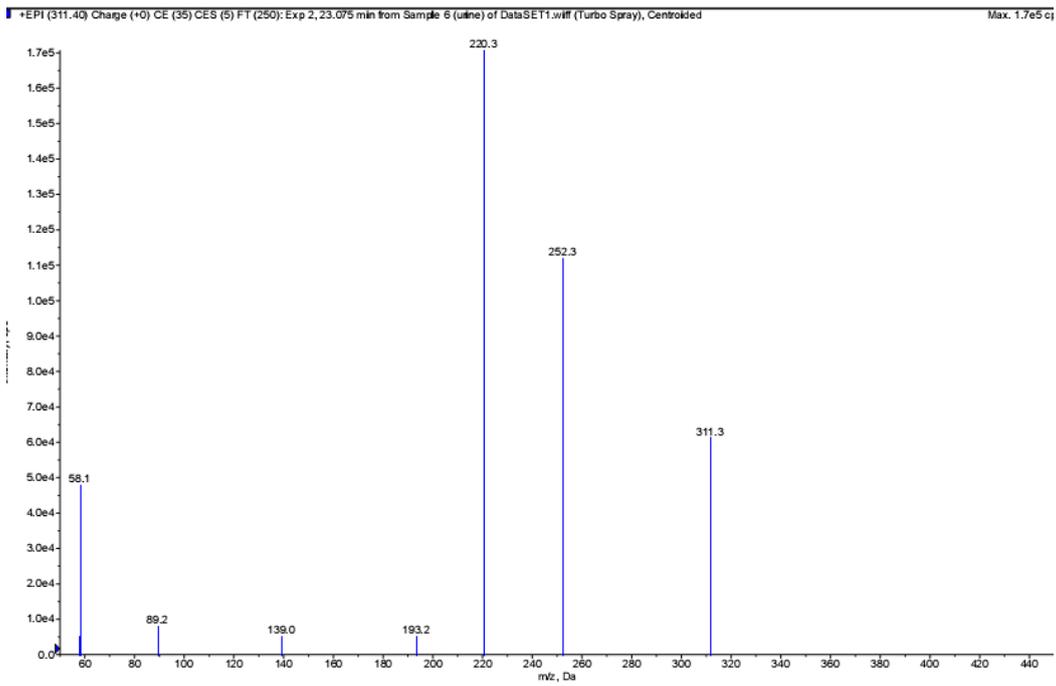
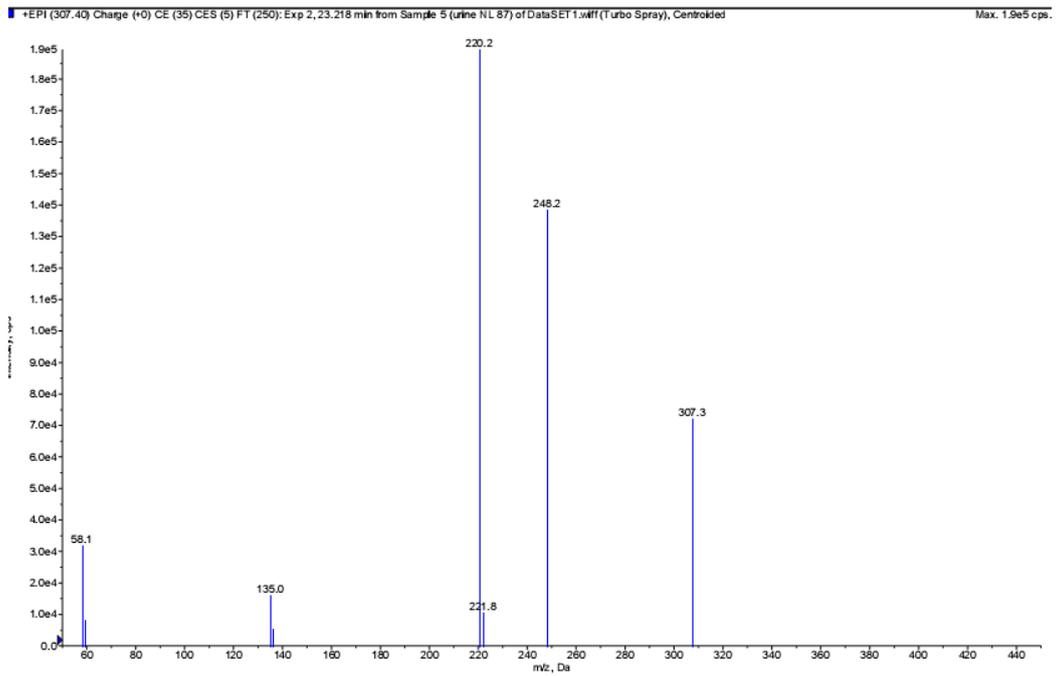
Compound 2, m/z 285/289, $t=14.96$ min



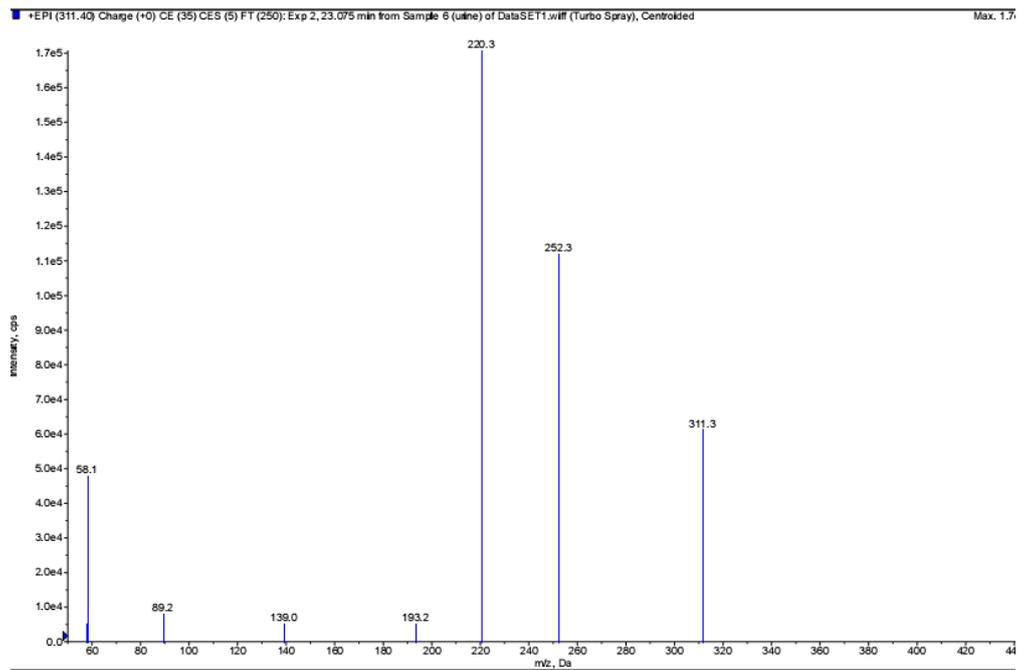
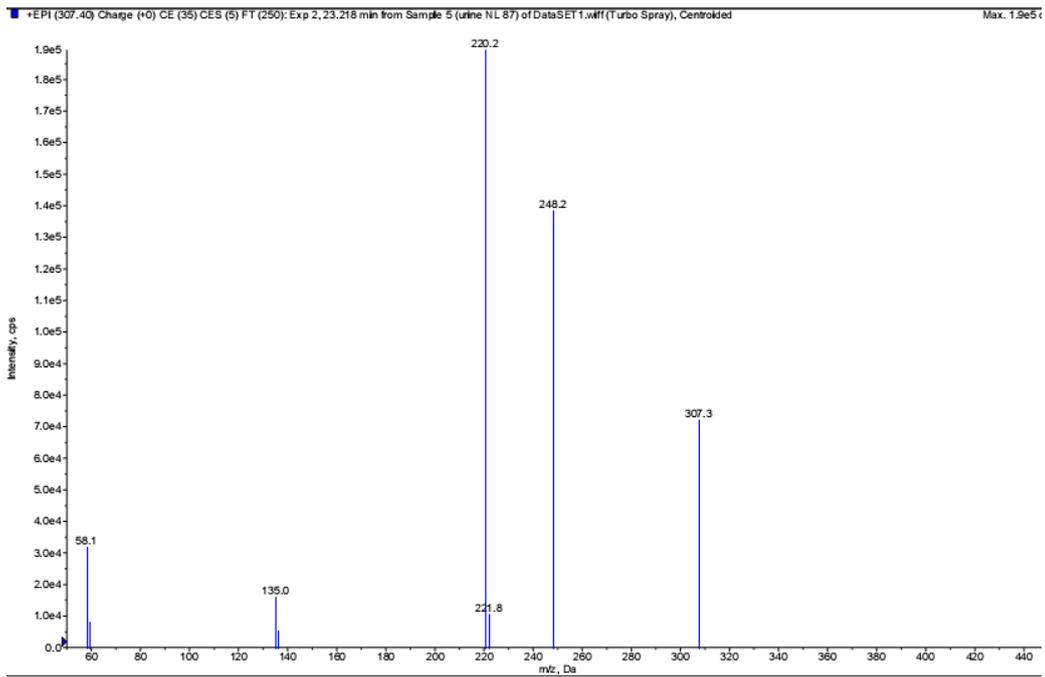
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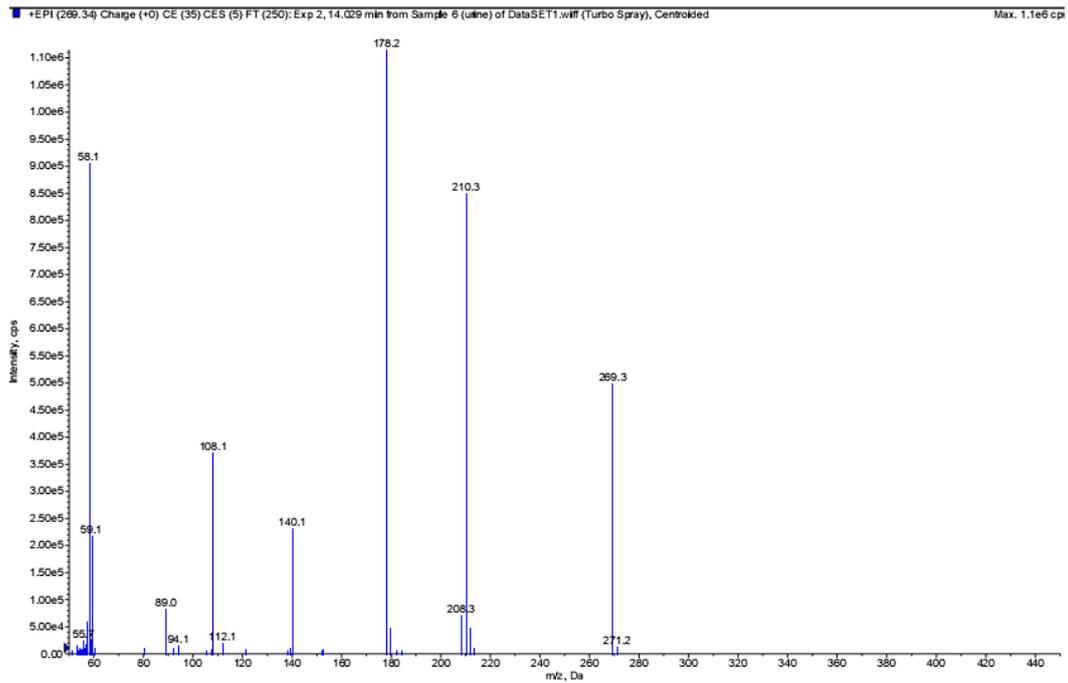
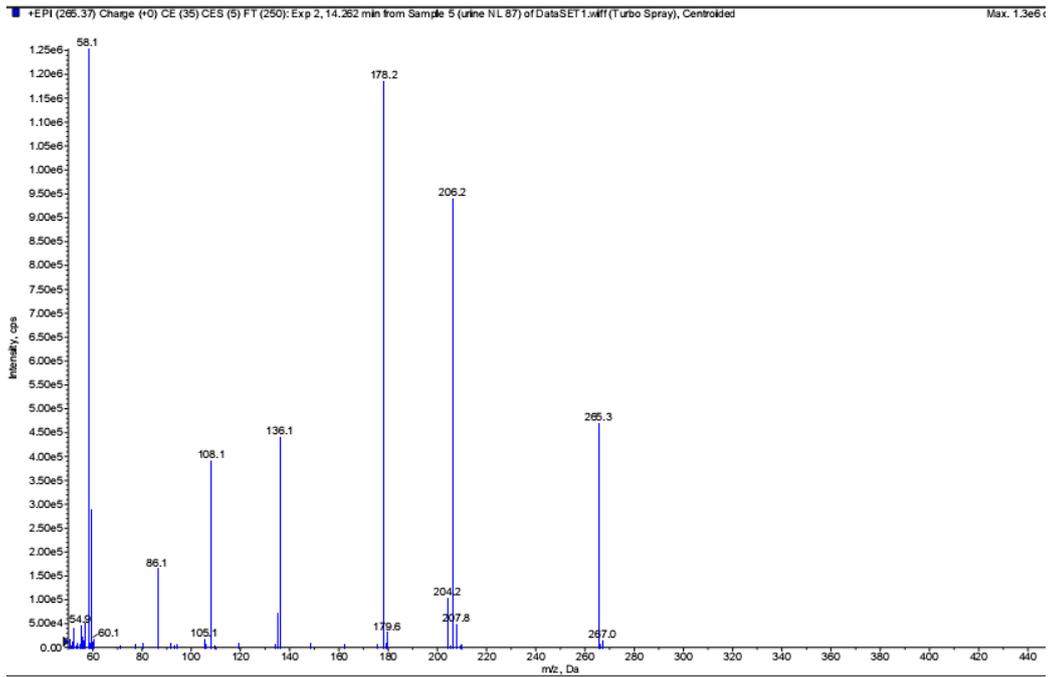
Compound 4, m/z 307/311, $t=23.19$ min



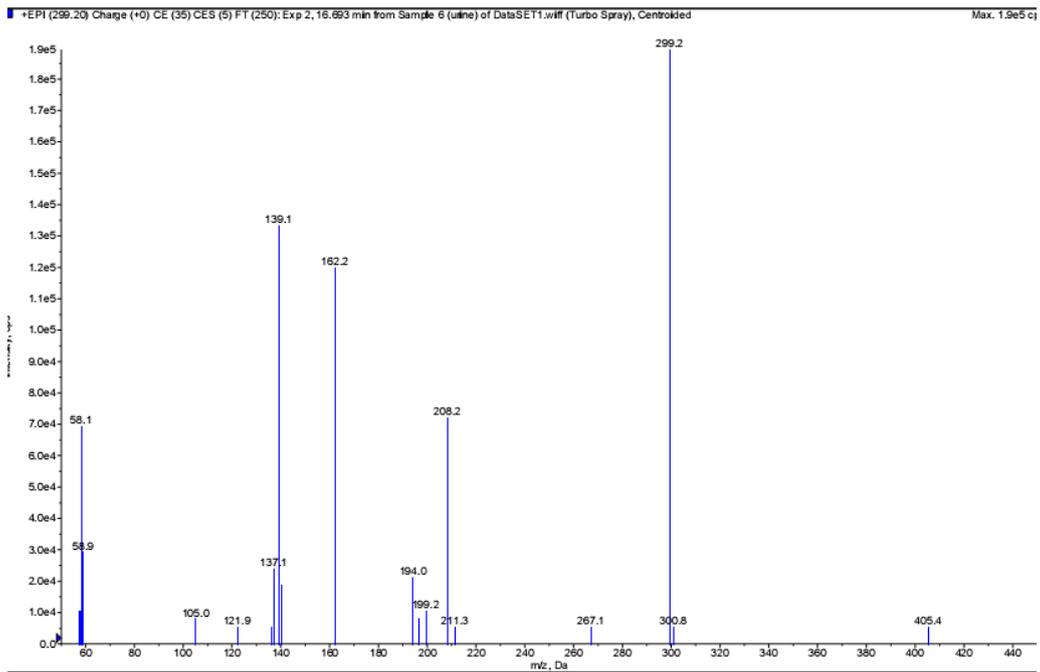
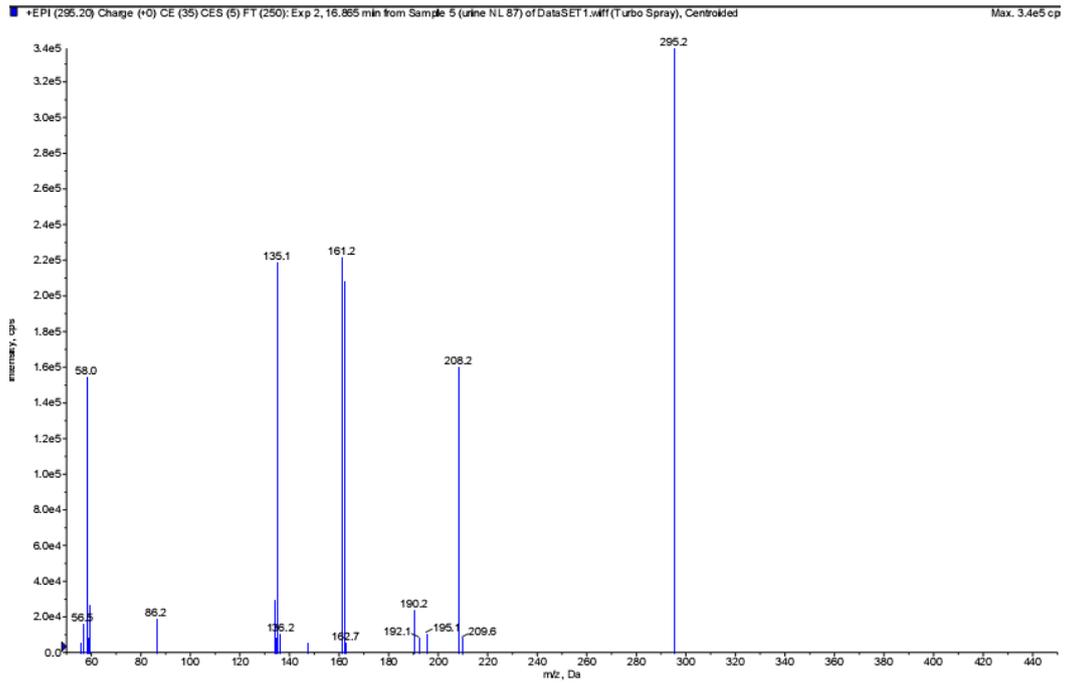
Compound 5, m/z 321/325, $t=25.67$ min



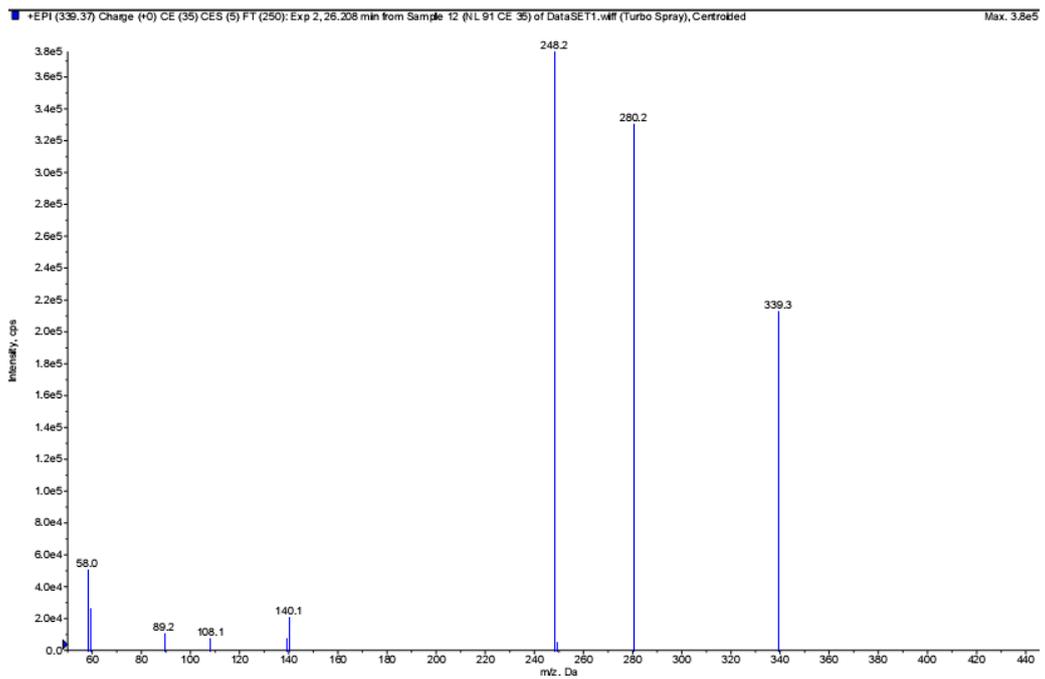
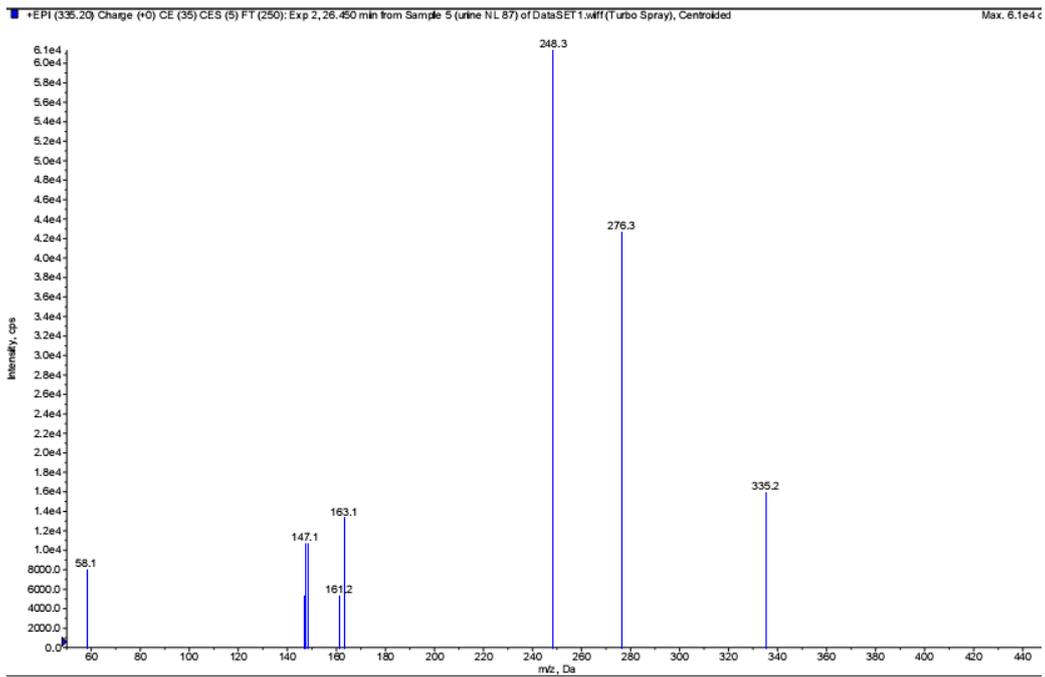
Compound 6, m/z 265/269, $t=14.19$ min



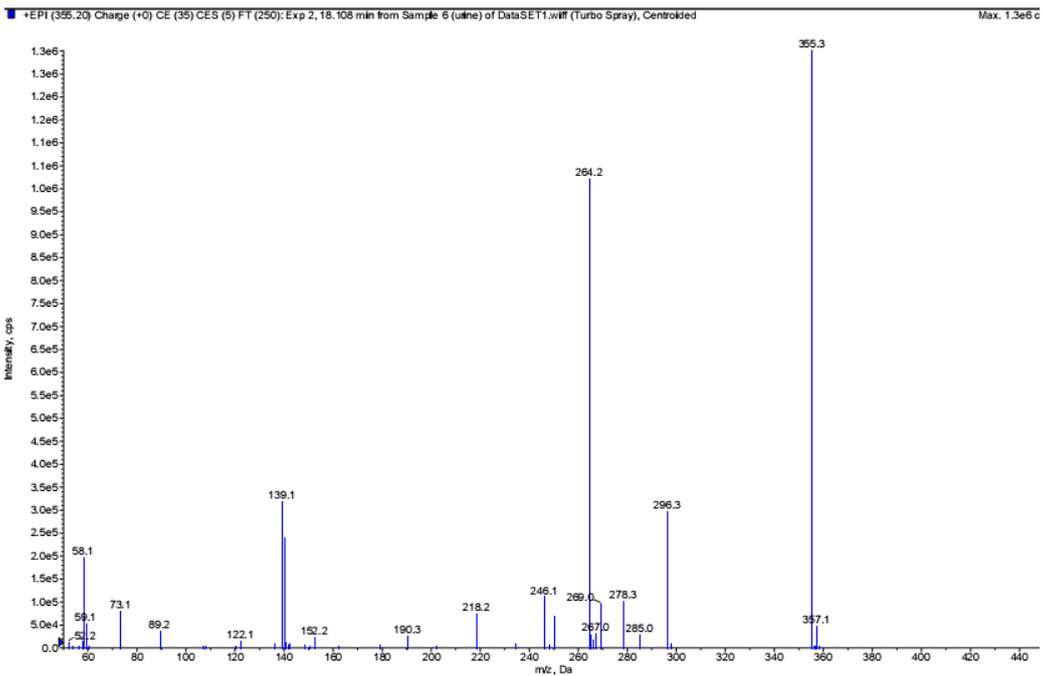
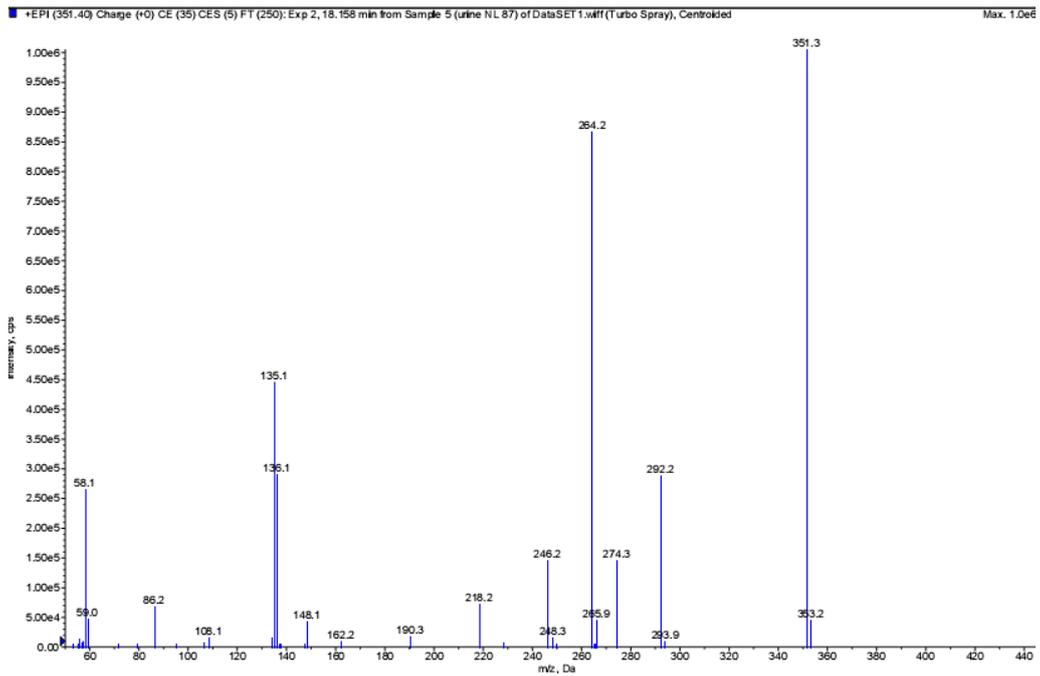
Compound 9, m/z 295/299, $t=16.80$ min



Compound 10, m/z 335/339, $t=26.63$ min



Compound 11, m/z 351/355, t=18.16 min



Compound 12, m/z 369/373, t=16.05 min

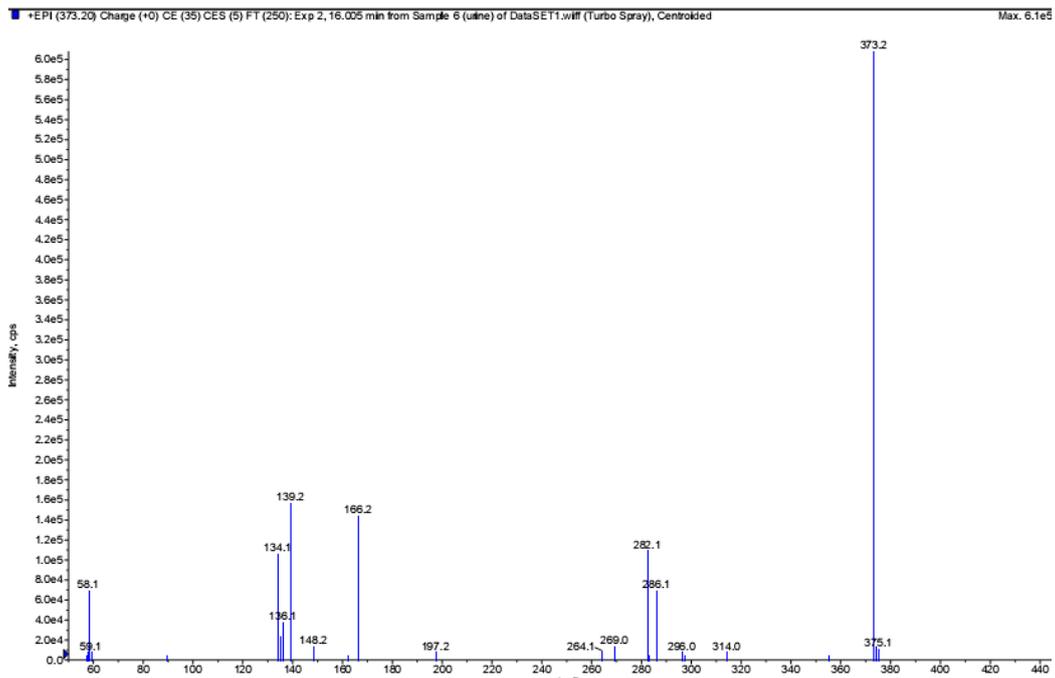
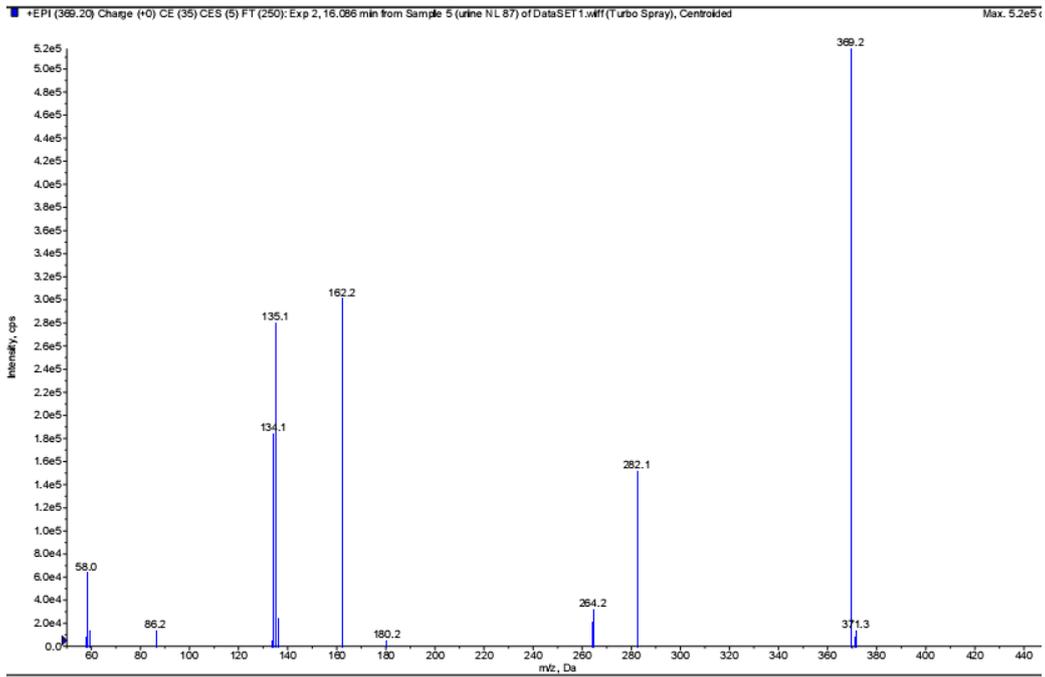


Figure S4. The flow chart illustrating the structure and individual steps of the approach.

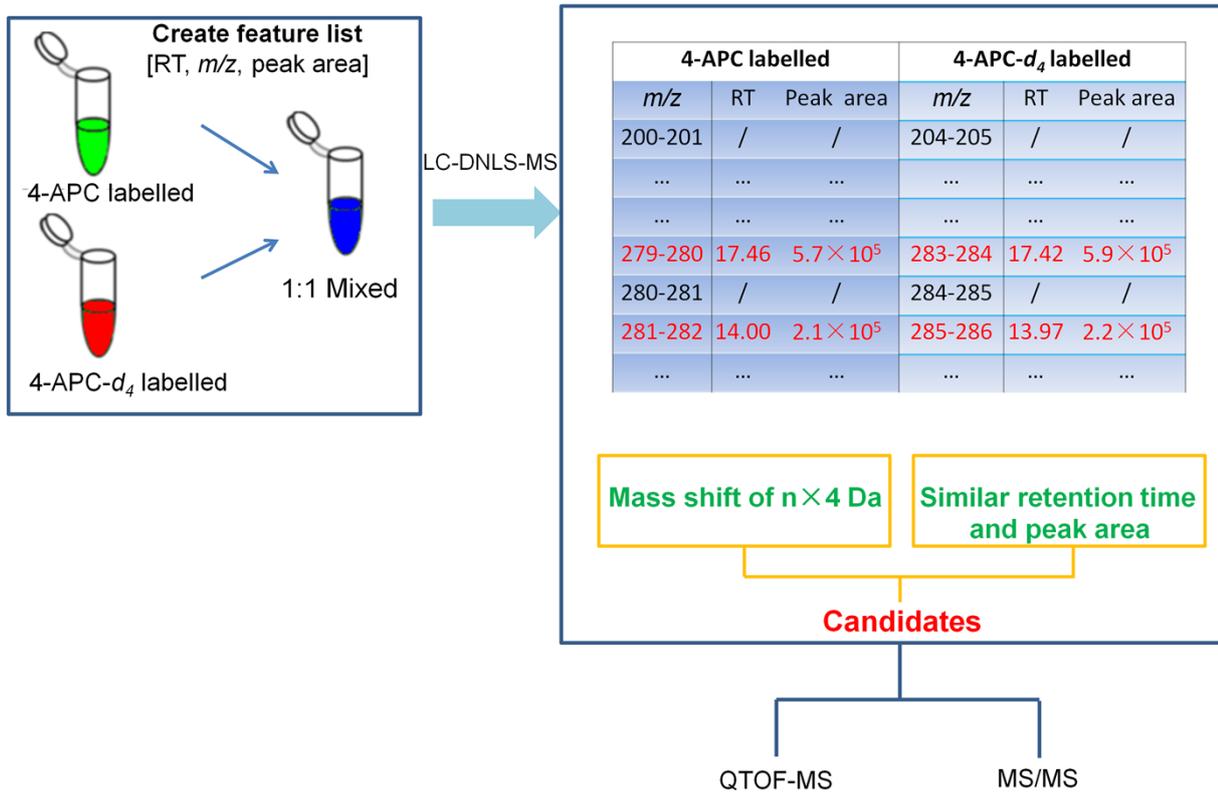
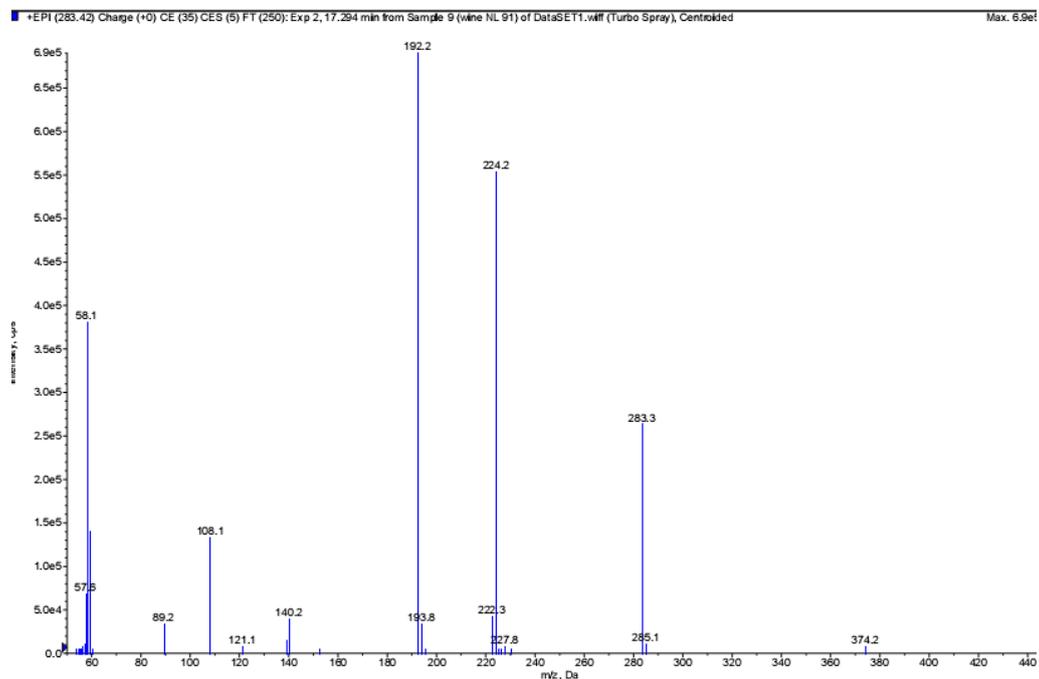
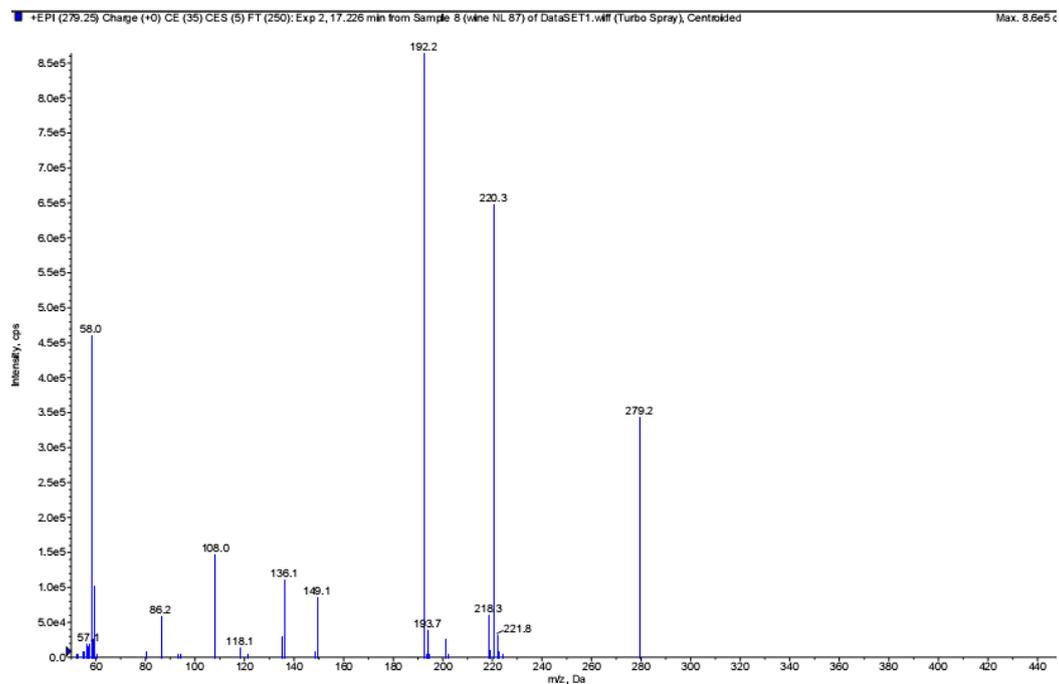
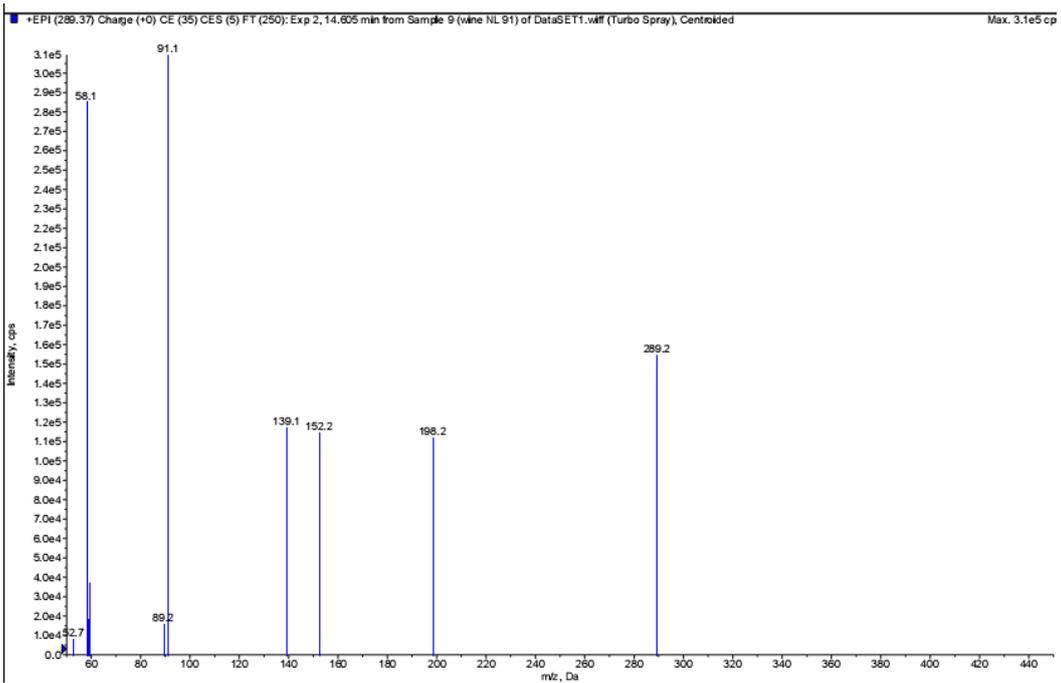
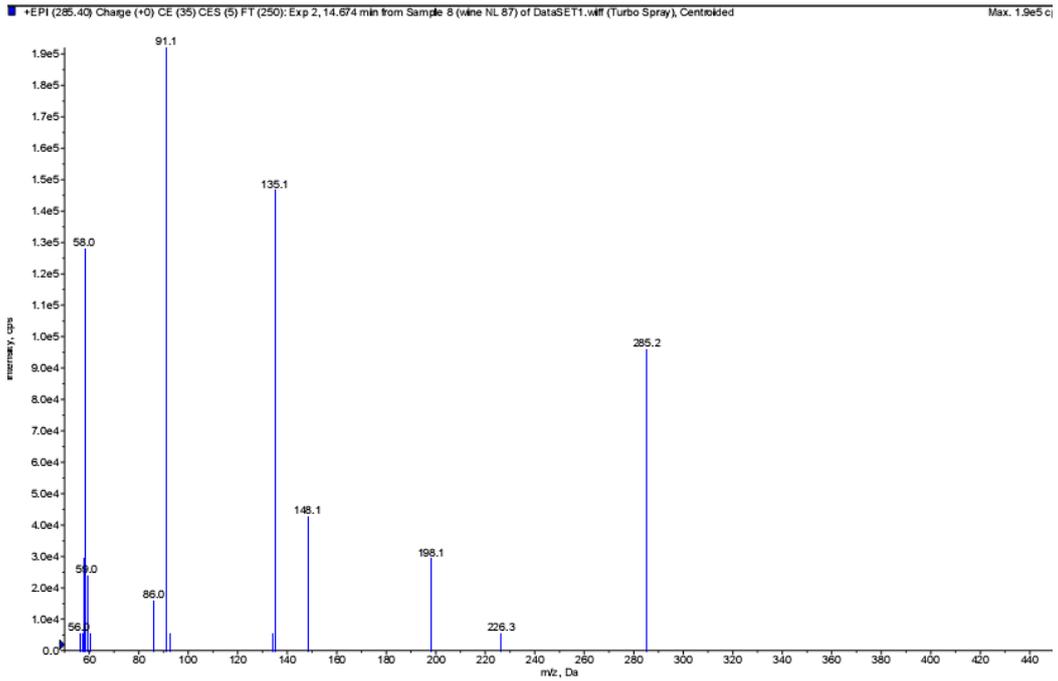


Figure S5. The product ion mass spectra of 4-APC and 4-APC- d_4 labelled compounds in white wine using QTRAP mass spectrometer.

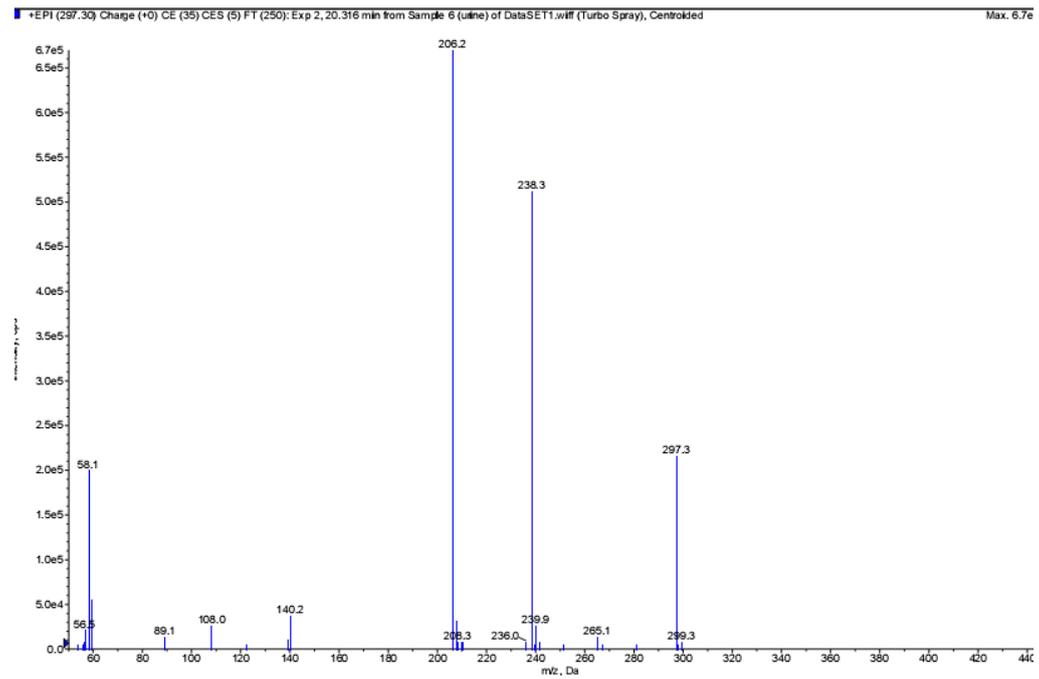
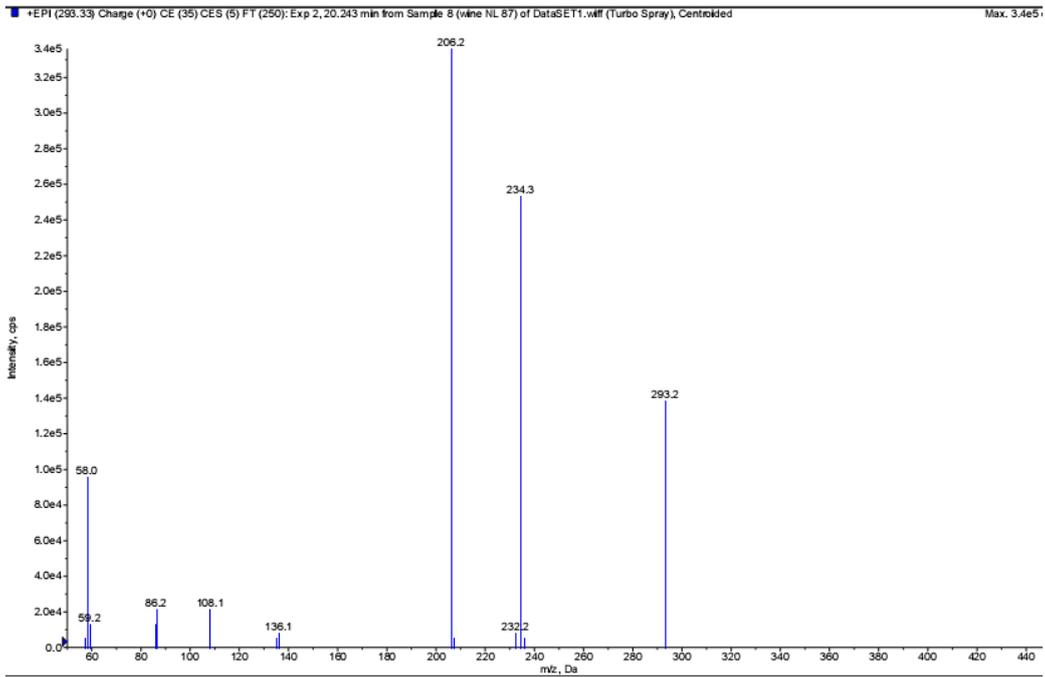
Compound 1, m/z 279/283, $t=17.35$ min



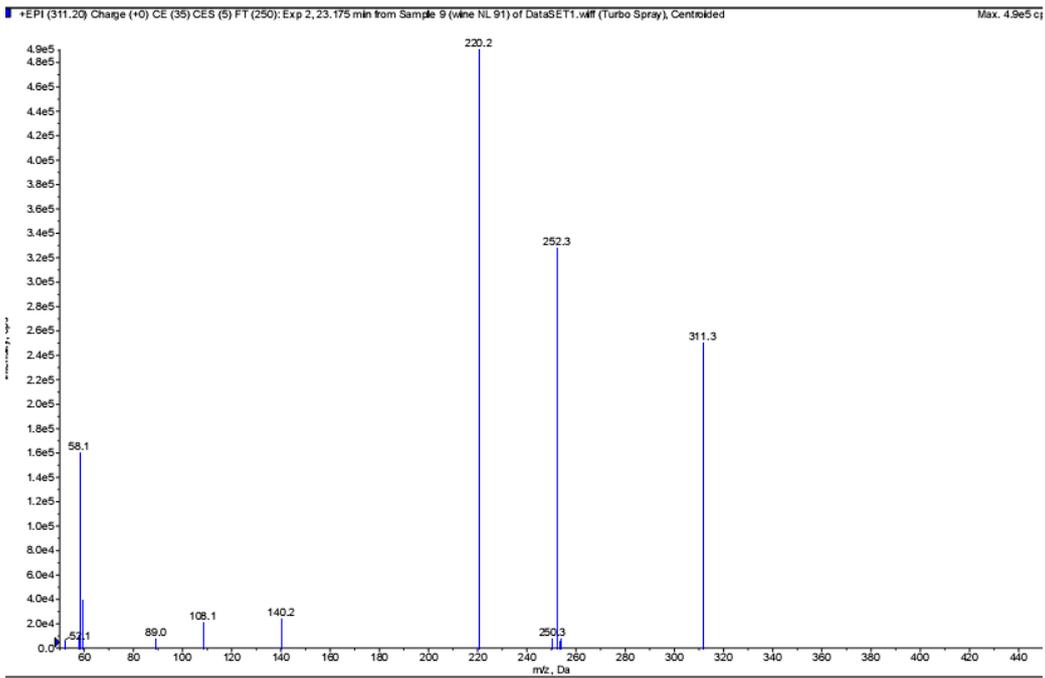
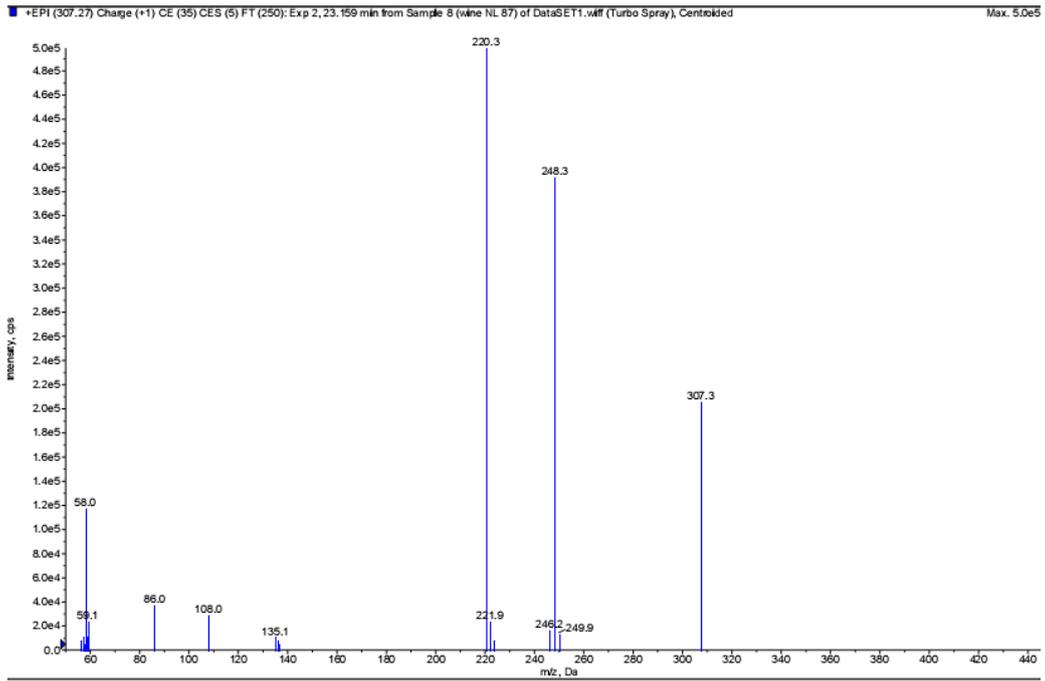
Compound 2, m/z 285/289, $t=14.77$ min



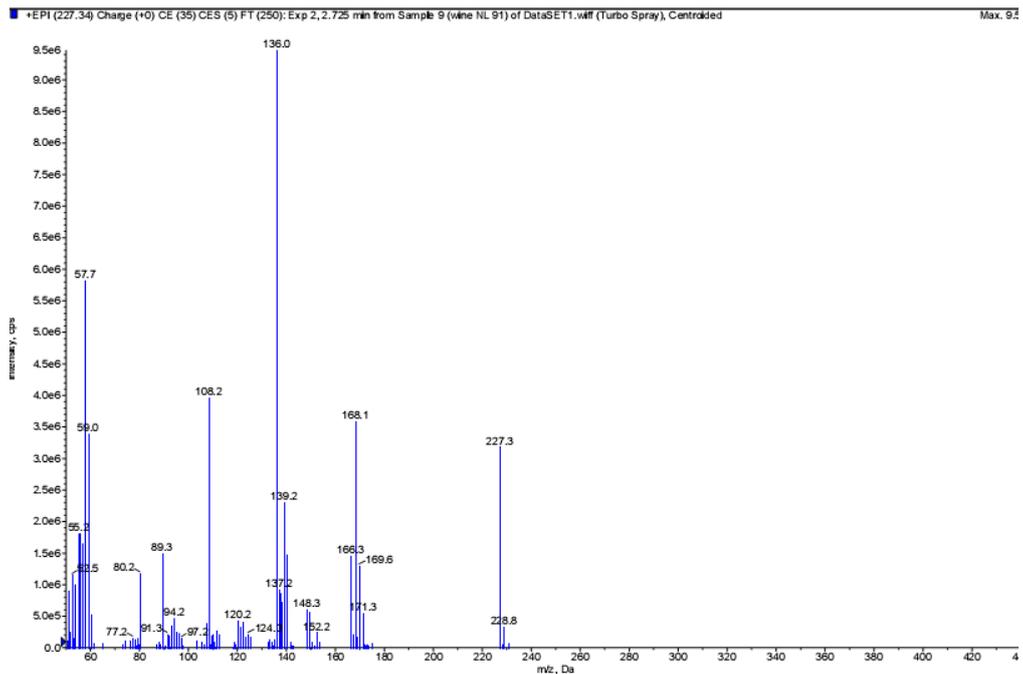
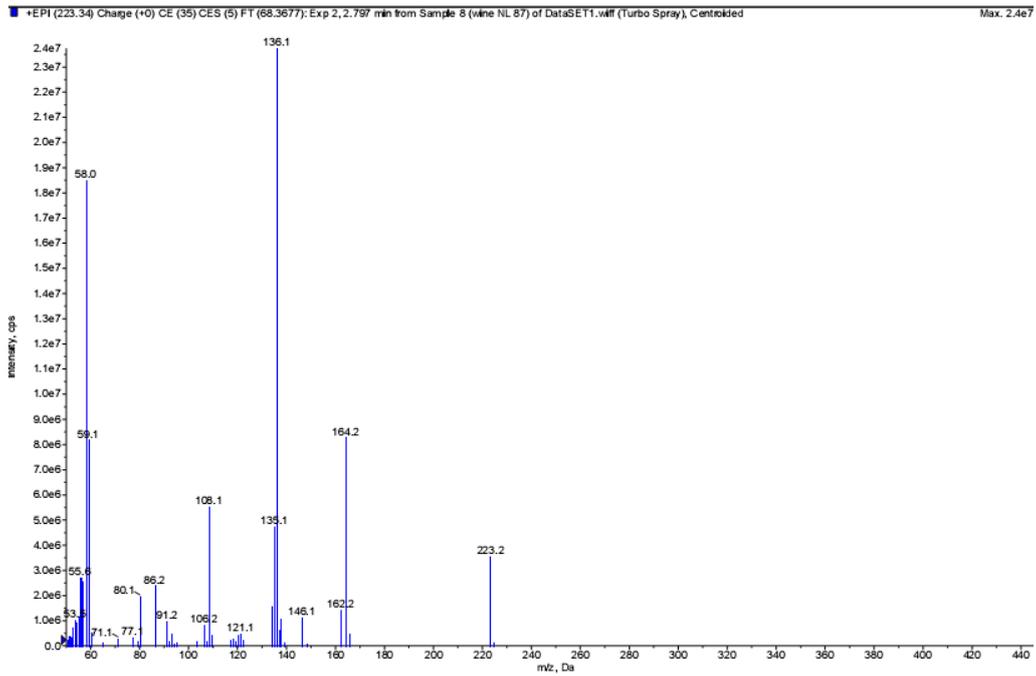
Compound 3, m/z 293/297, $t=20.33$ min



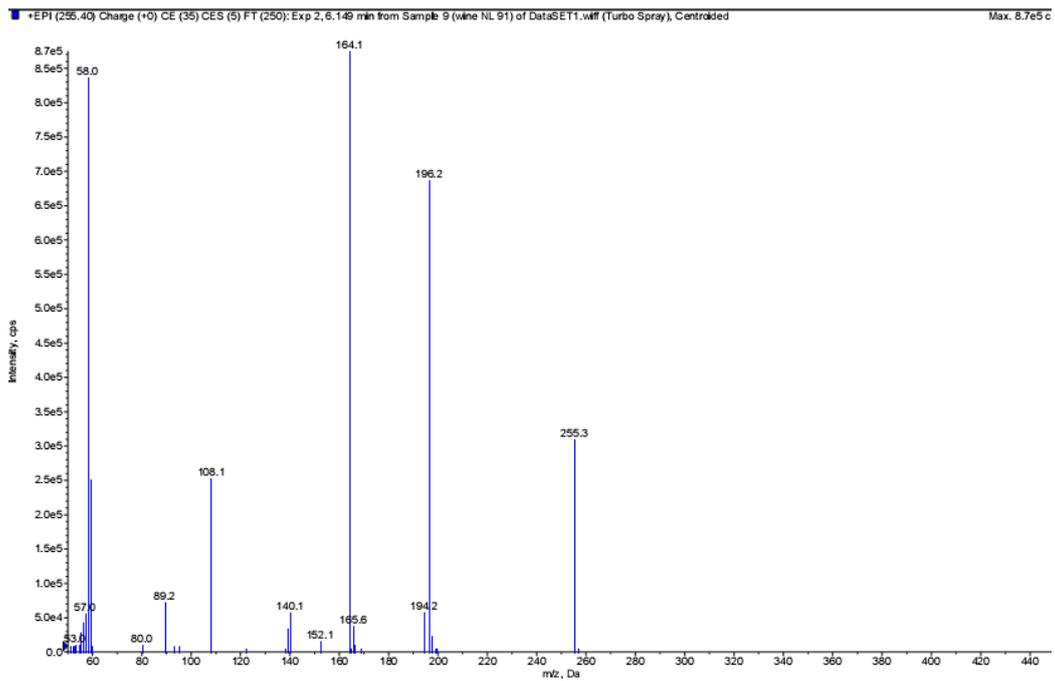
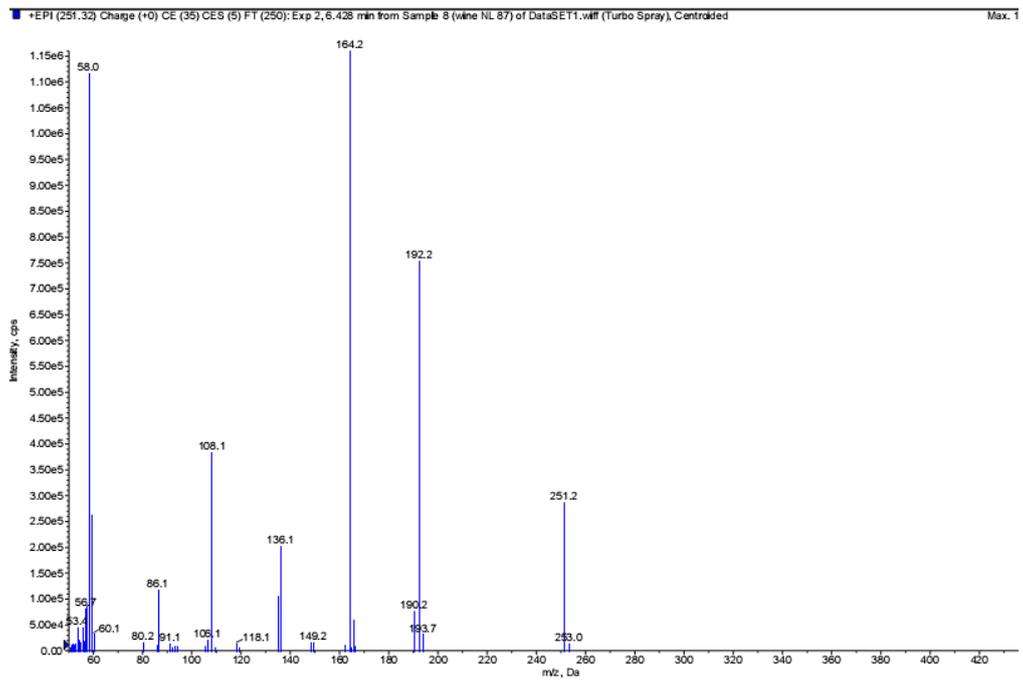
Compound 4, m/z 307/311, $t=23.22$ min



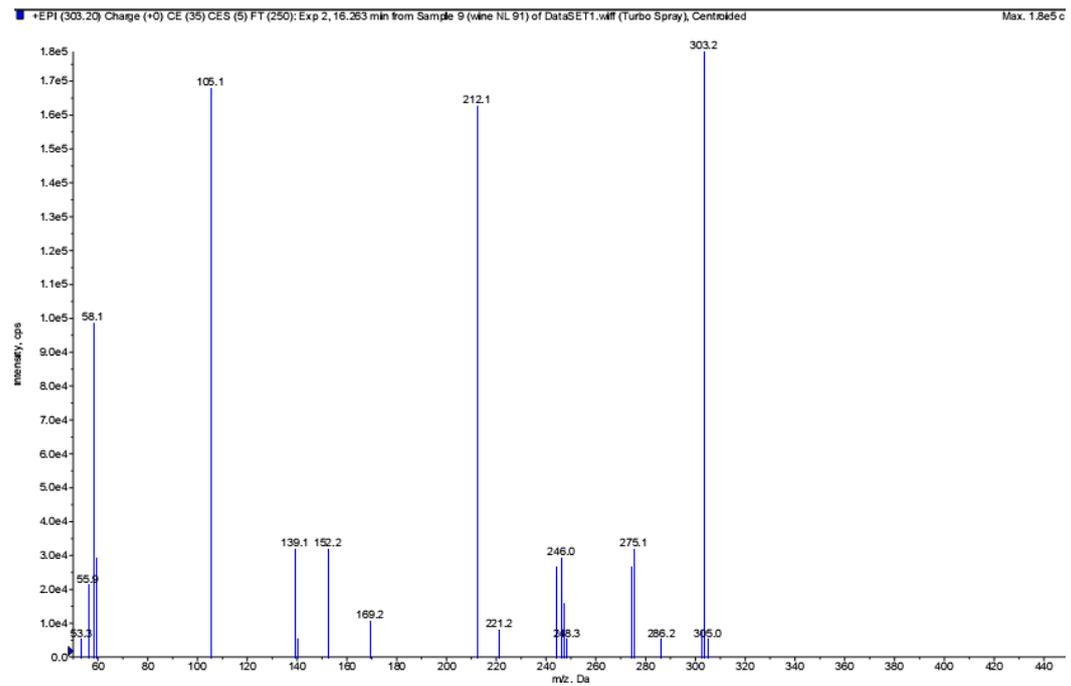
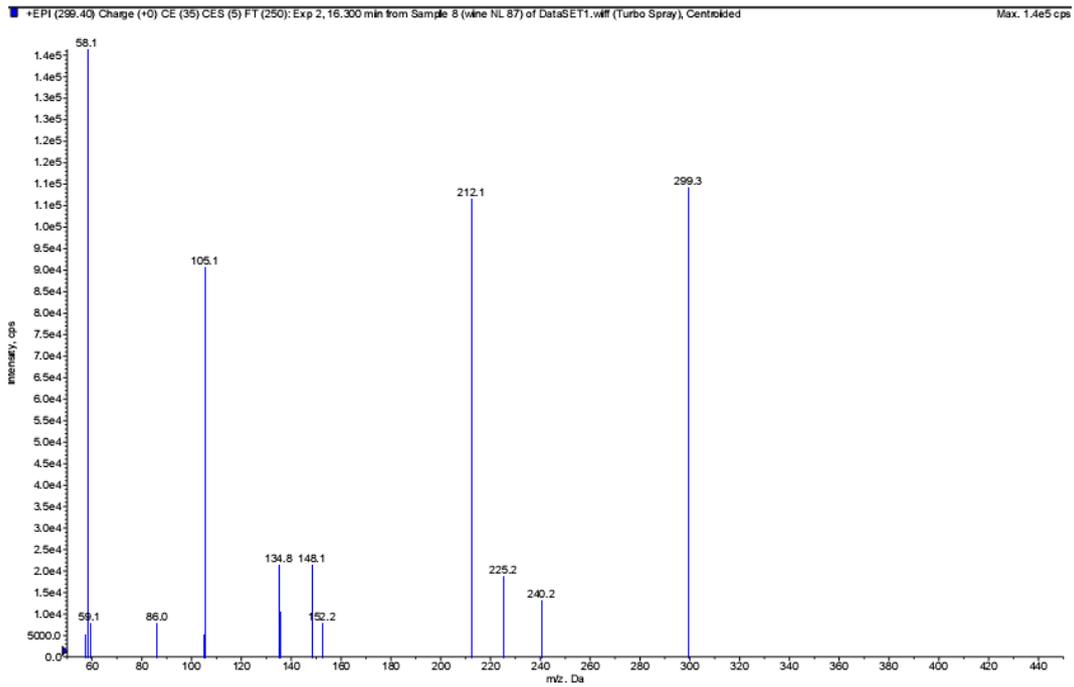
Compound 5, m/z 223/227, t=2.76 min



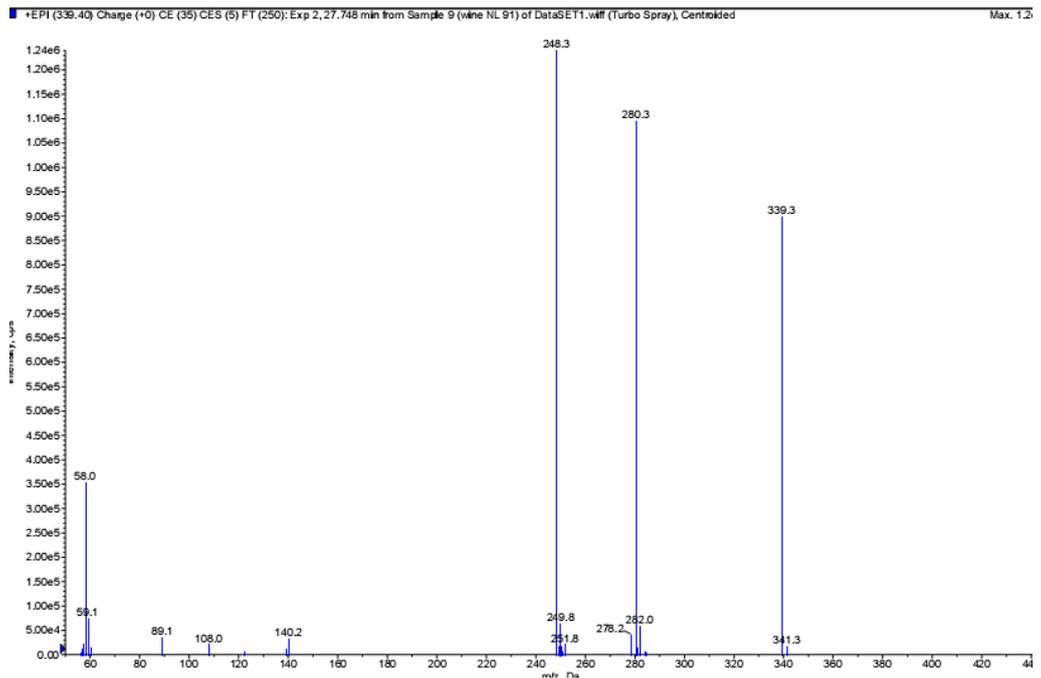
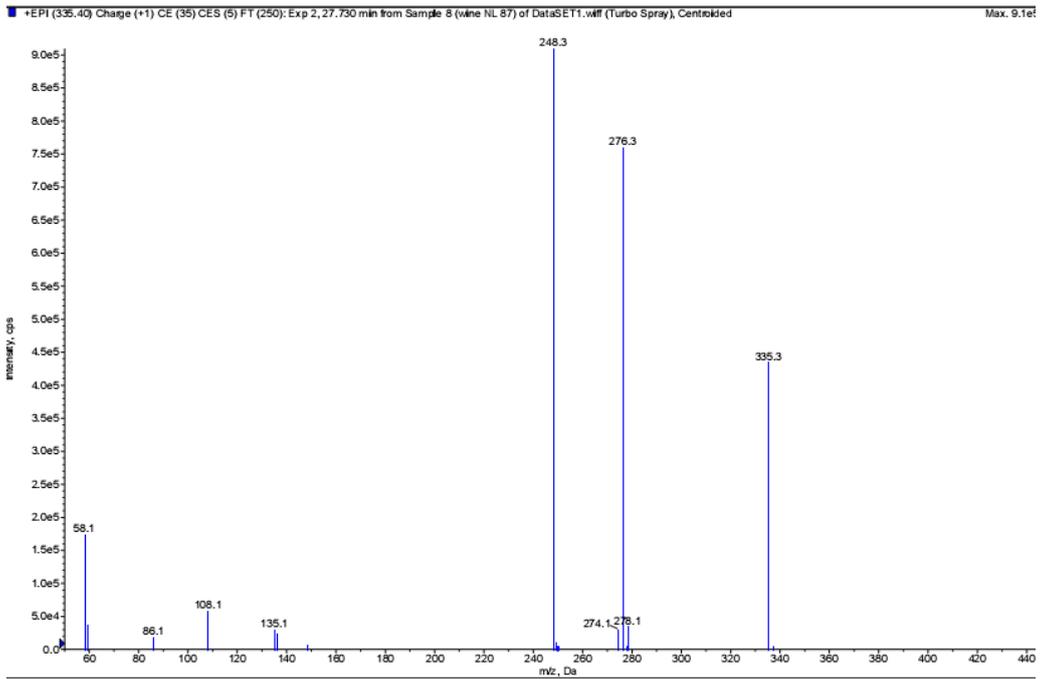
Compound 6, m/z 251/255, $t=6.42$ min



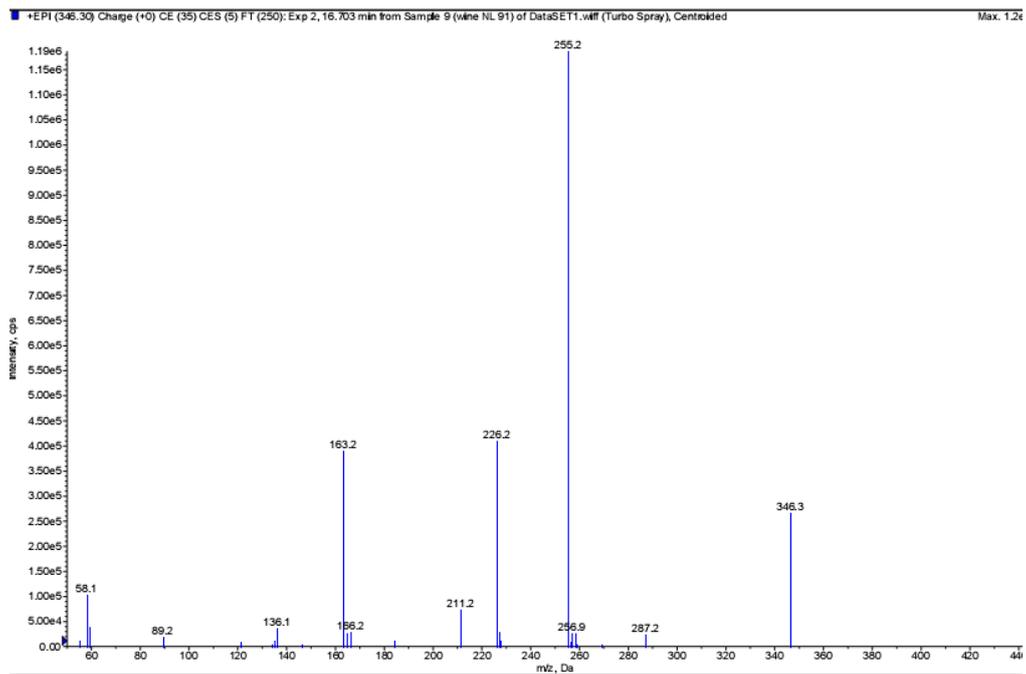
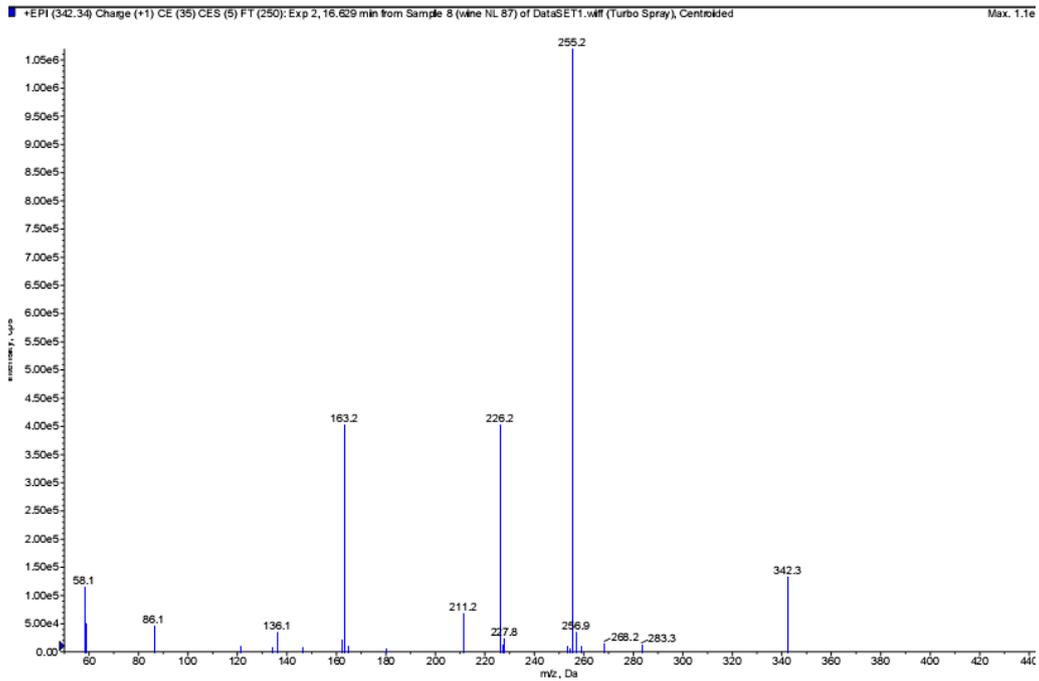
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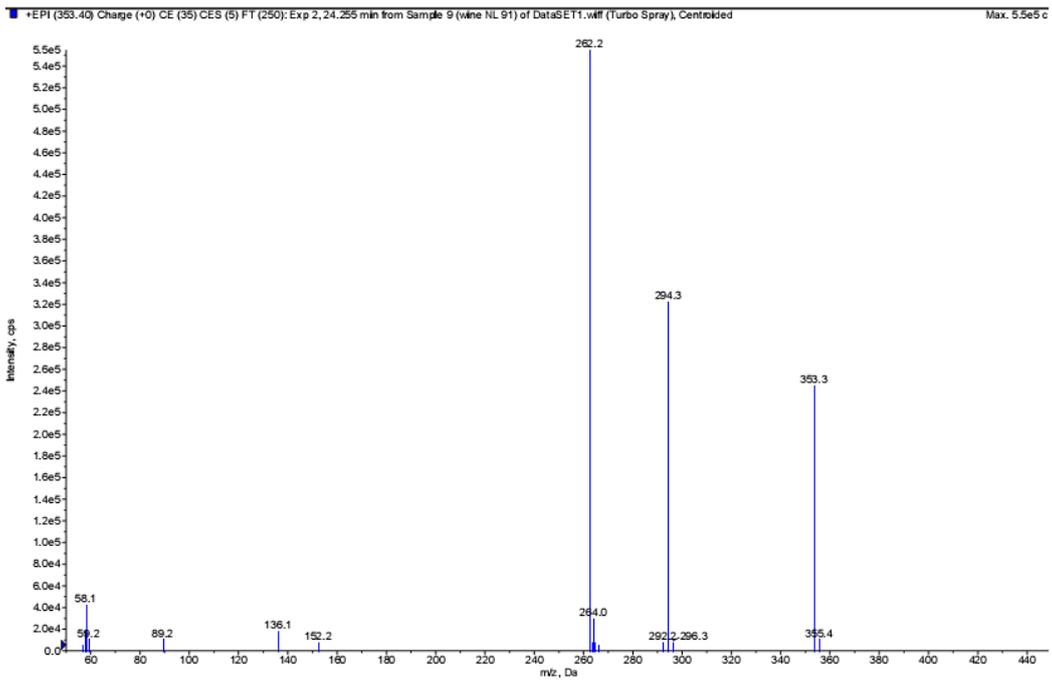
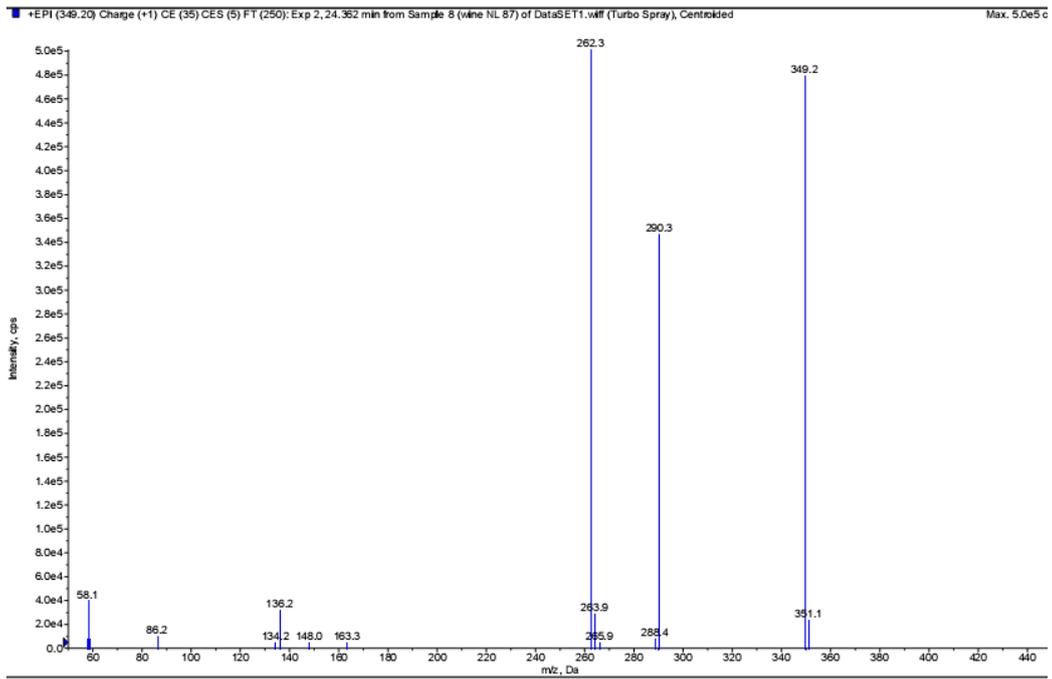
Compound 11, m/z 335/339, $t=27.80$ min



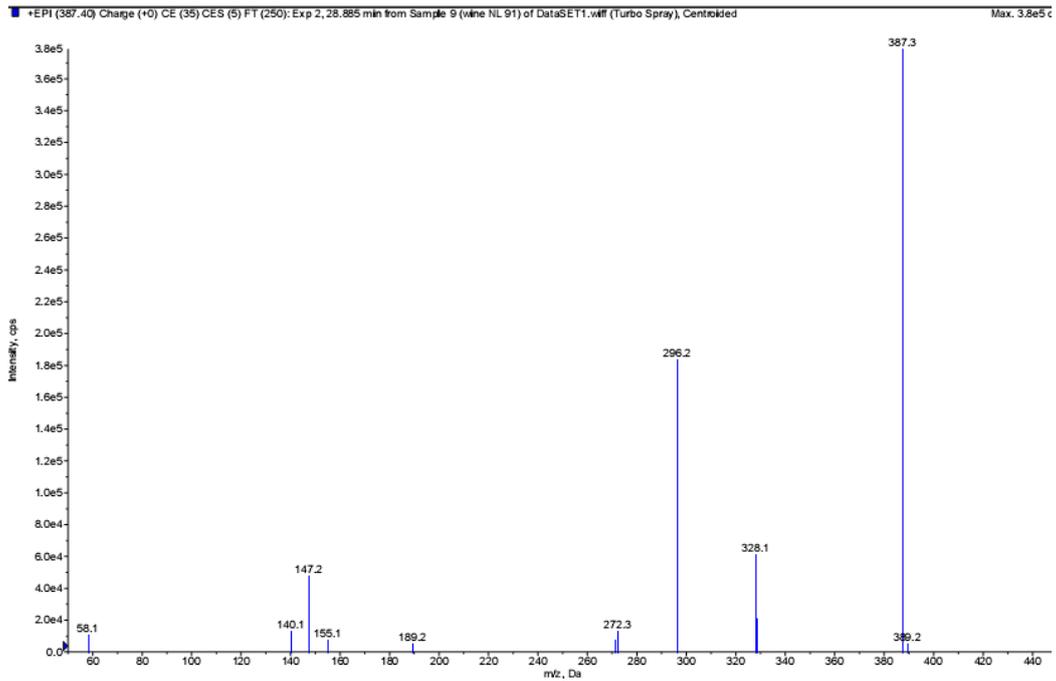
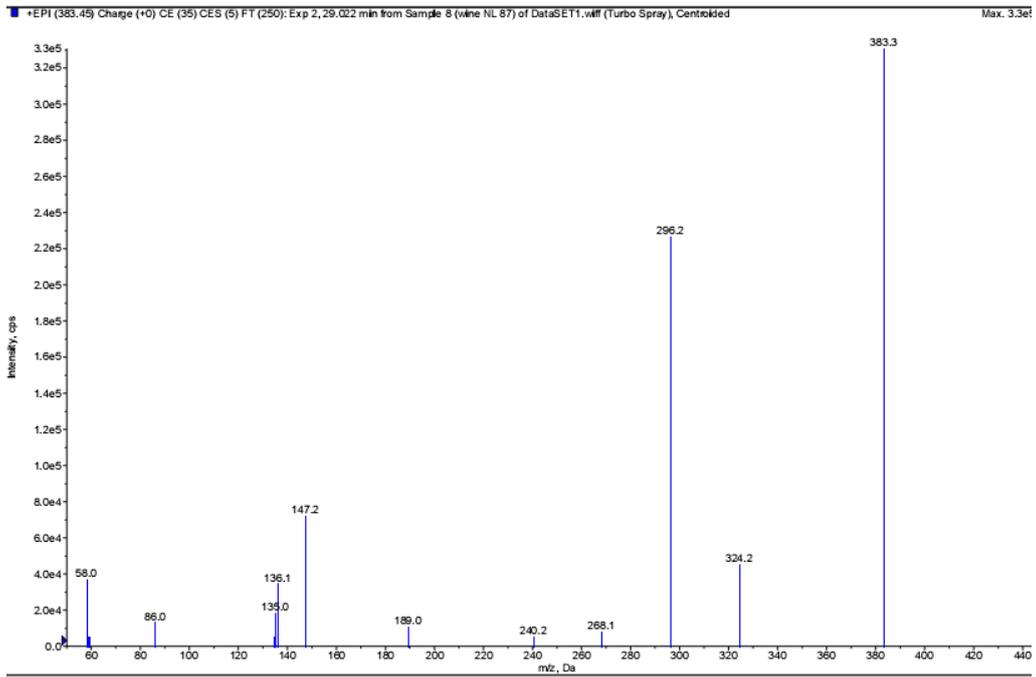
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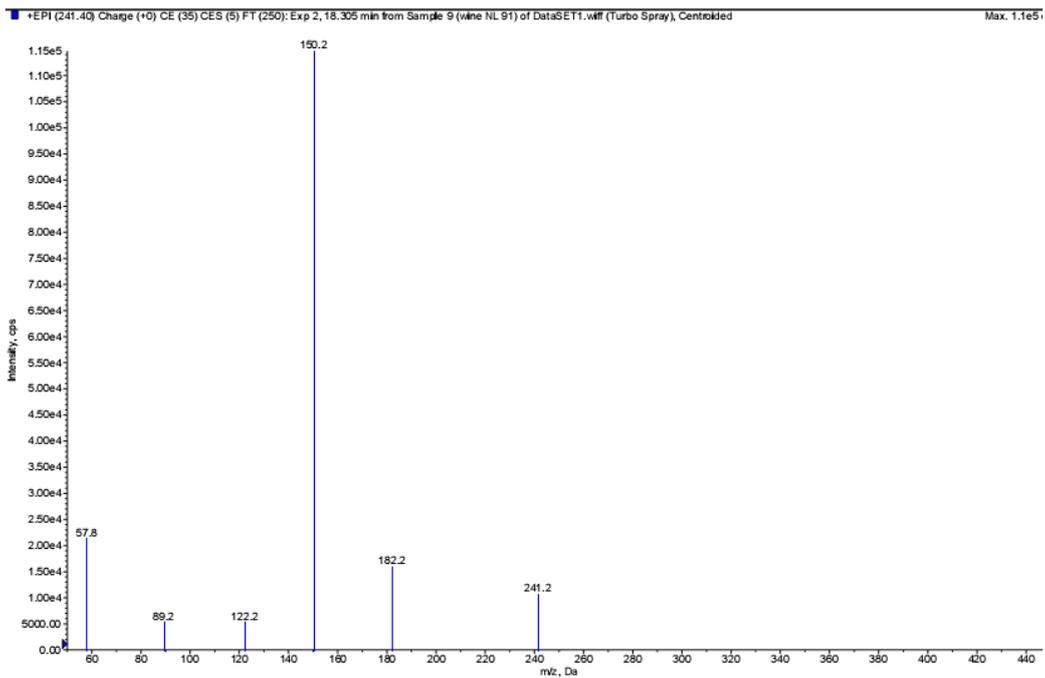
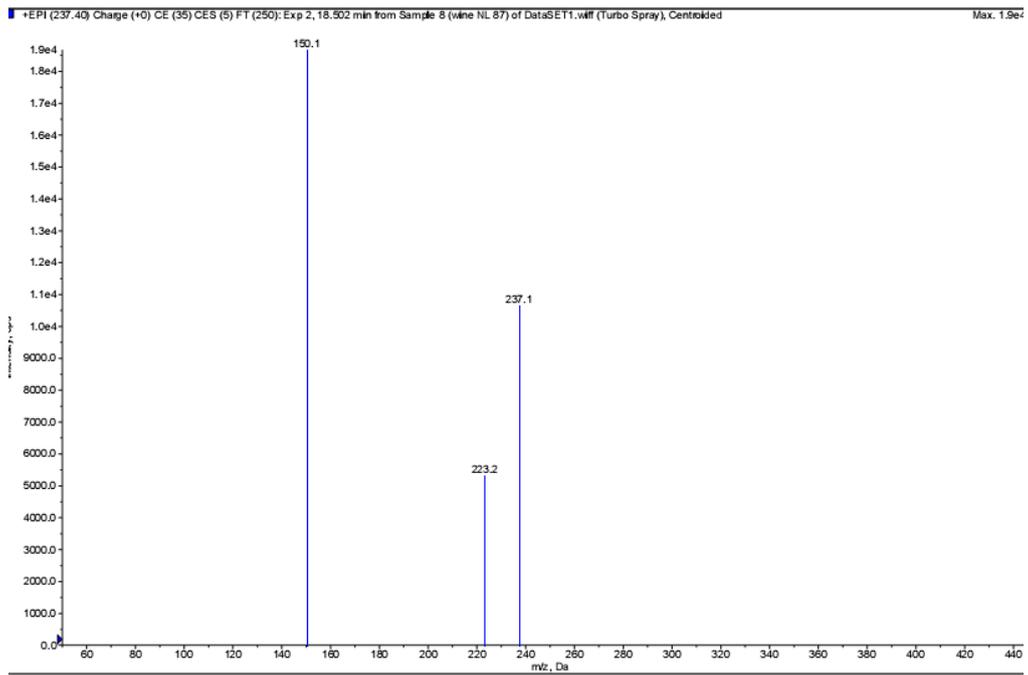
Compound 14, m/z 349/353, $t=24.32$ min



Compound 15, m/z 383/387, $t=29.04$ min



Compound 18, m/z 237/241, $t=18.39$ min



Compound 19, m/z 262/266, $t=17.07$ min

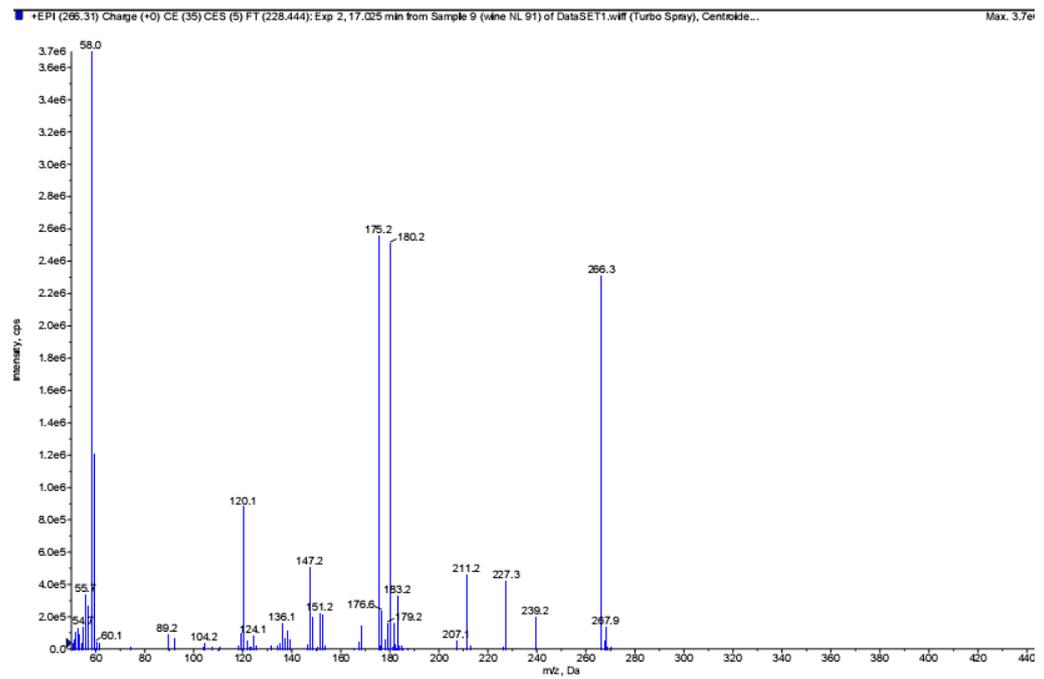
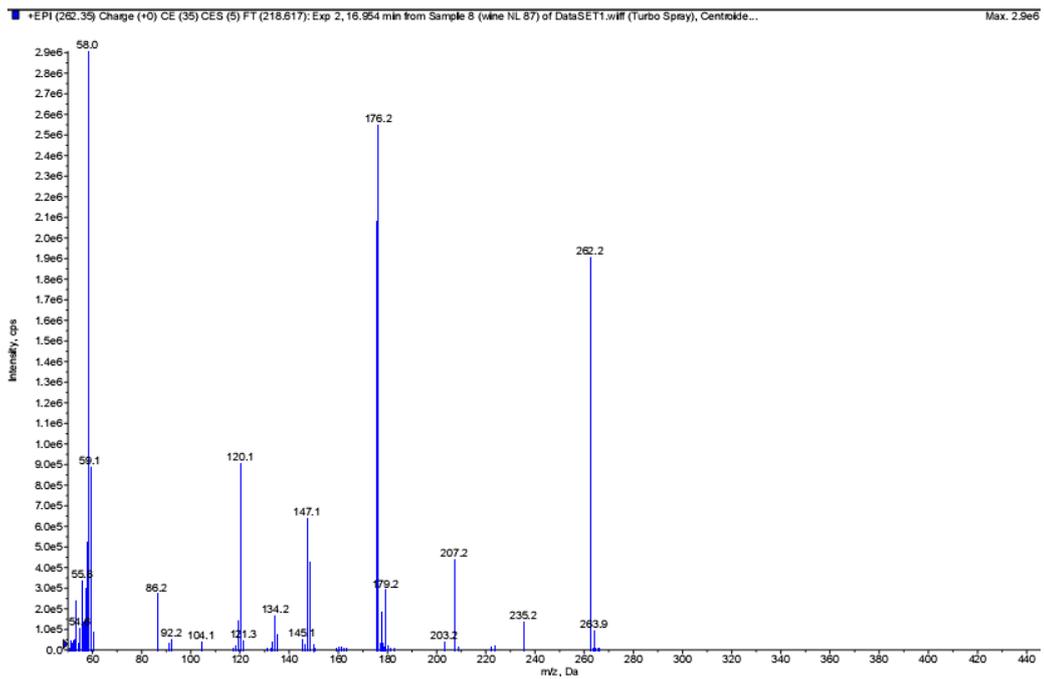


Figure S6. Comparison of the retention time of 4-APC and 4-APC- d_4 labelled hexanal, benzaldehyde, heptanal and octanal from the white wine with standards spiked in white wine by DNLS analysis using QTRAP mass spectrometer. (A) 4-APC and 4-APC- d_4 labelled hexanal; (B) 4-APC and 4-APC- d_4 labelled benzaldehyde; (C) 4-APC and 4-APC- d_4 labelled heptanal; (D) 4-APC and 4-APC- d_4 labelled octanal.

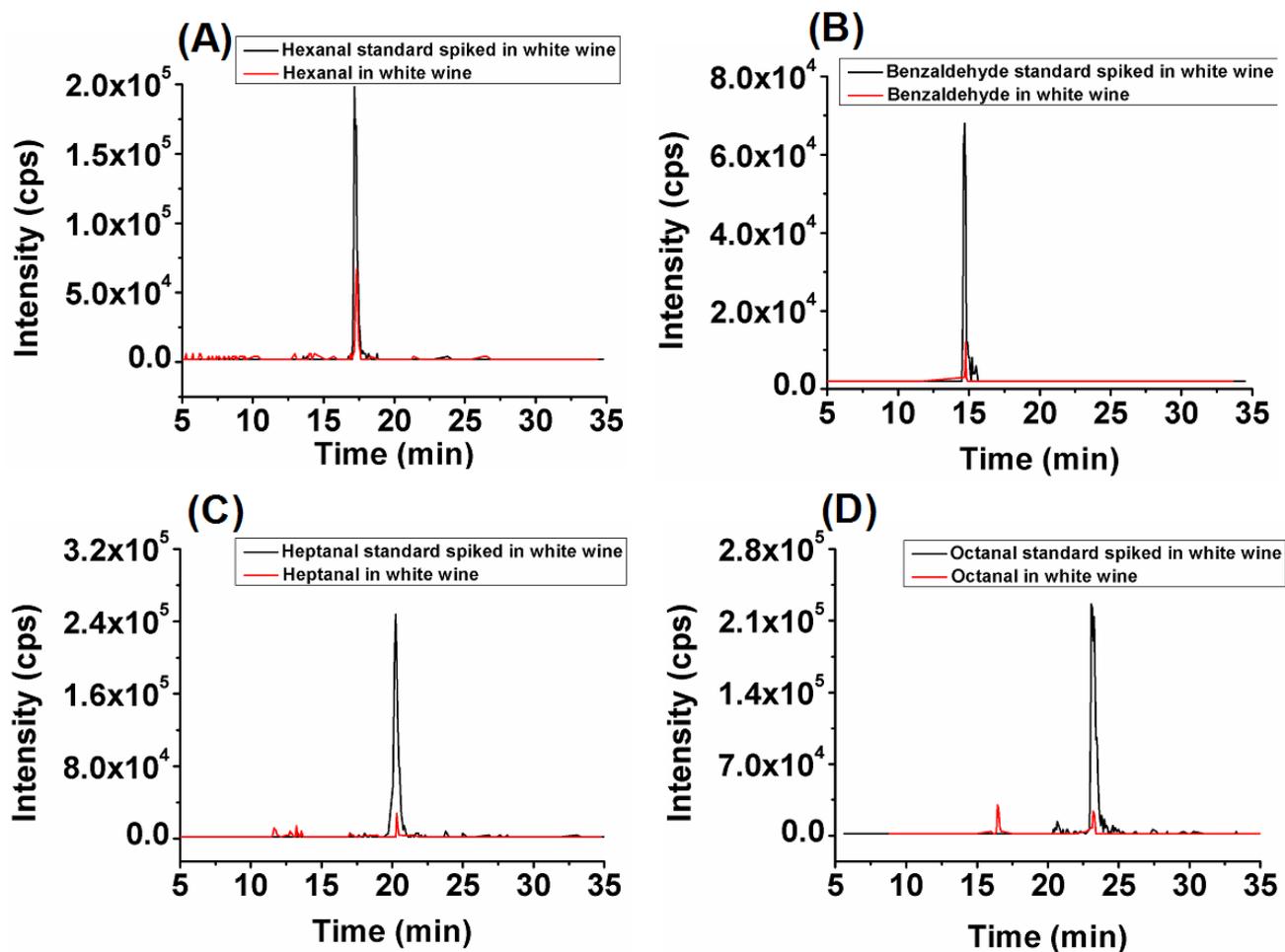


Figure S7. Comparison of the MS/MS spectrum of 4-APC labelled hexanal, benzaldehyde, heptanal, octanal and nonanal from the urine with the standards using QTRAP mass spectrometer. (A) 4-APC labelled hexanal standard; (B) 4-APC labelled hexanal in white wine; (C) 4-APC labelled benzaldehyde standard; (D) 4-APC labelled benzaldehyde in white wine; (E) 4-APC labelled heptanal standard; (F) 4-APC labelled heptanal in white wine; (G) 4-APC labelled octanal standard; (H) 4-APC labelled octanal in white wine.

