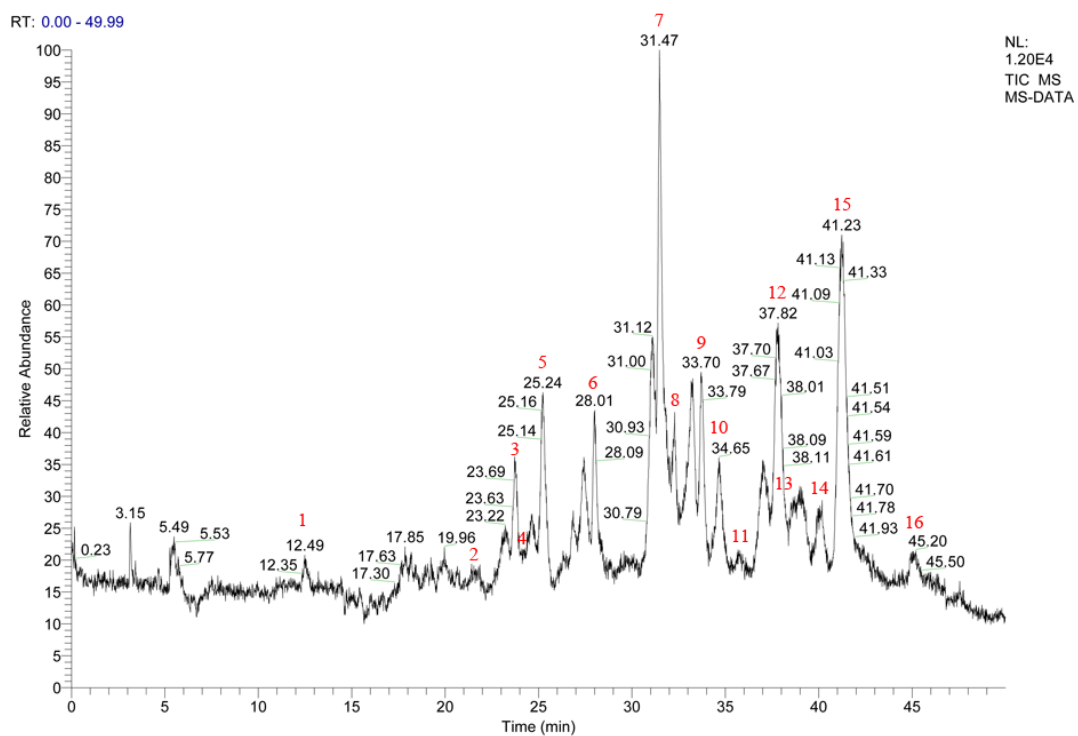


Table S1. Primer sequences used for real time-PCR analysis

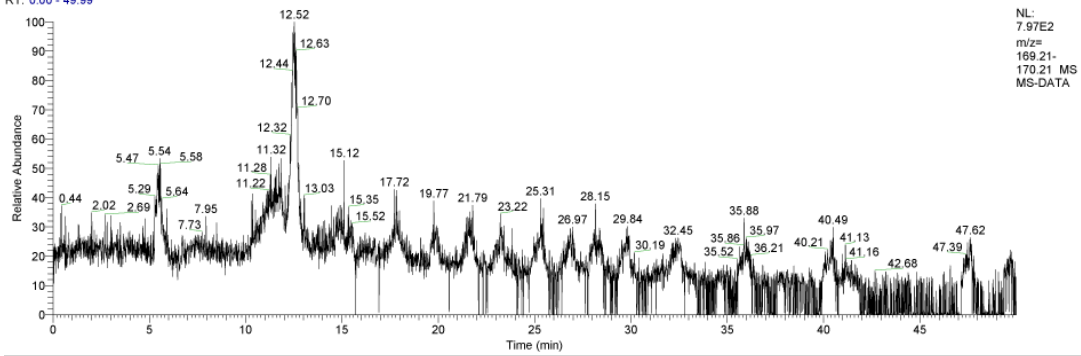
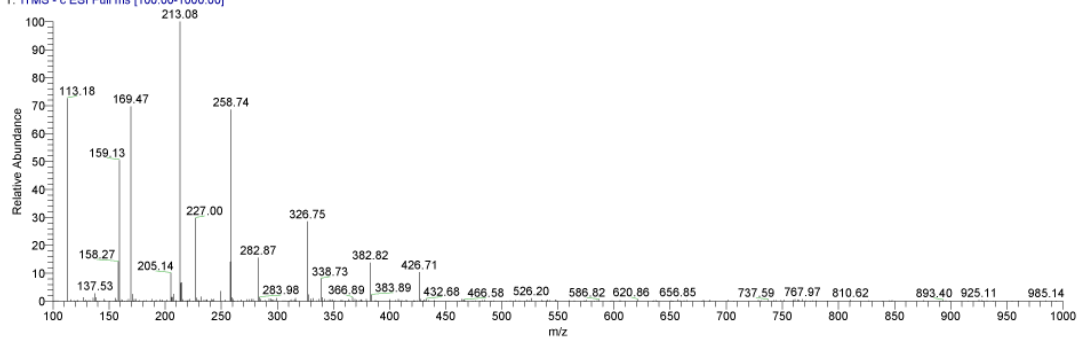
Gene	Direction	Sequence
GAPDH	Forward	GGCATCGTGGAAGGGCTCAT
	Reverse	AGGGATGATGTTCTGGGCTGC
PDGFR	Forward	CATTGGGGACAGGGAAGTGGAC
	Reverse	CCTGATGGTGATGCTCTCGC
CTGF	Forward	CTGAAAGAATAGCTGGCTTCA
	Reverse	CAGGTACTAGCTGAGGTCAT
NF- κ B	Forward	AACACTGCCGAGCTCAAGAT
	Reverse	CATCGGCTTGAGAAAAGGAG
MMP-2	Forward	CTTGCTGGTGGCCACATTC
	Reverse	CTCATTCCCTGCGAAGAACAC
TIMP-1	Forward	GCCTACACCCCAGCCAT
	Reverse	ATGCCAGGGAACCAGGAAGC
TIMP-2	Forward	GGCAACCCCATCAAGAGGATTCAAT
	Reverse	CACACTGCTGAGGAGGGG
α -SMA	Forward	TTCGTGACTACTGCTGAGCGTGAGA
	Reverse	AAAGATGGCTGGAAGAGGGTC
Bcl-2	Forward	TCCTTCCAGCCTGAGAGCAAC
	Reverse	GCGACGGTAGCGACGAGAG
Bax	Forward	ATGGGCTGGACACTGGACTTC
	Reverse	GAGTGAGGCAGTGAGGAC



Supplemental Figure 1. The total ion current (TIC) output from the ESI-MS in negative mode.

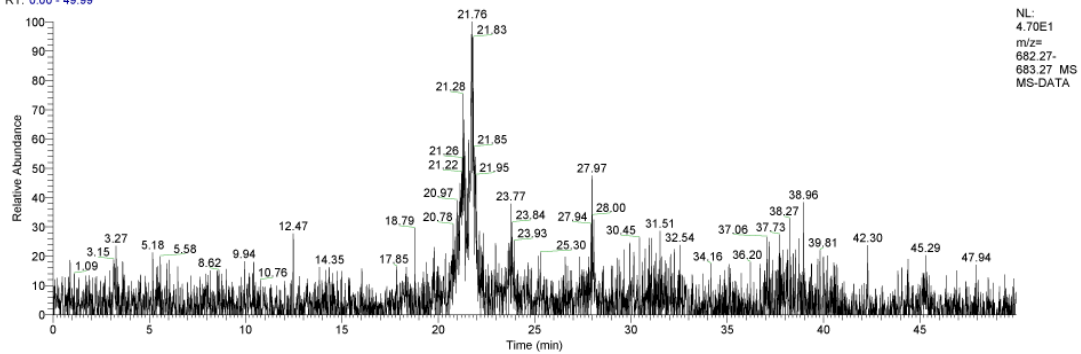
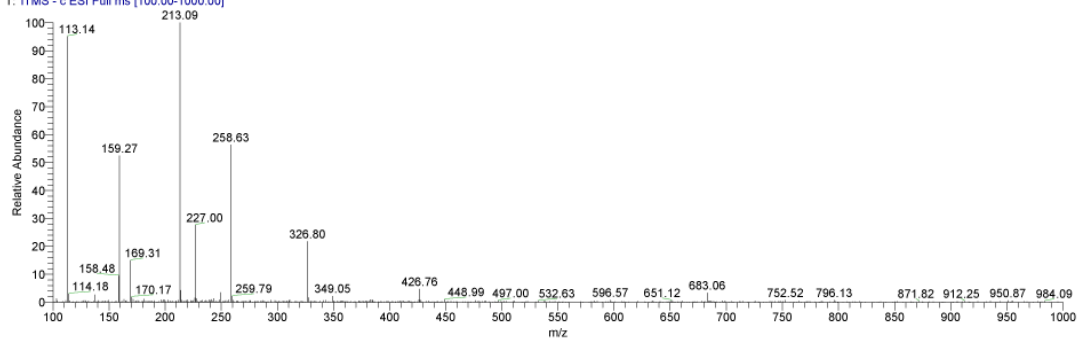
1

RT: 0.00 - 49.99

MS-DATA #989 RT: 12.52 AV: 1 NL: 1.14E3
T: ITMS - c ESI Full ms [100.00-1000.00]

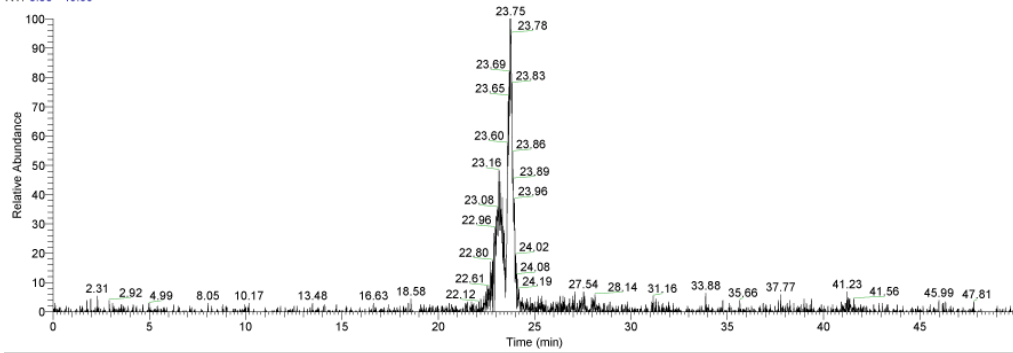
2

RT: 0.00 - 49.99

MS data #1718 RT: 21.76 AV: 1 NL: 1.79E3
T: ITMS - c ESI Full ms [100.00-1000.00]

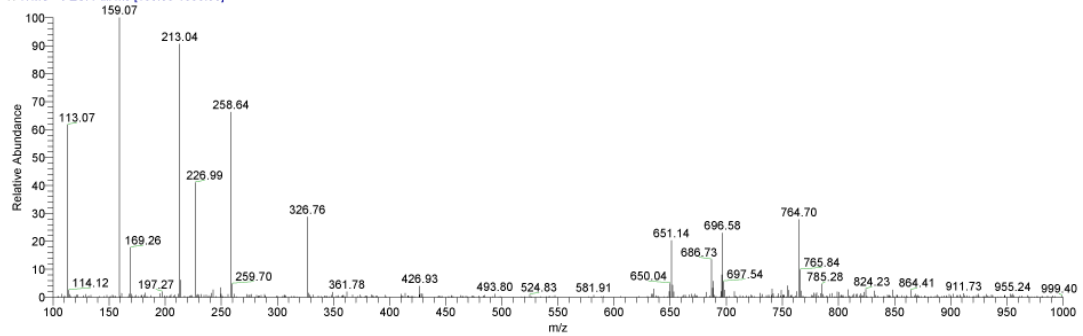
3

RT: 0.00 - 49.99



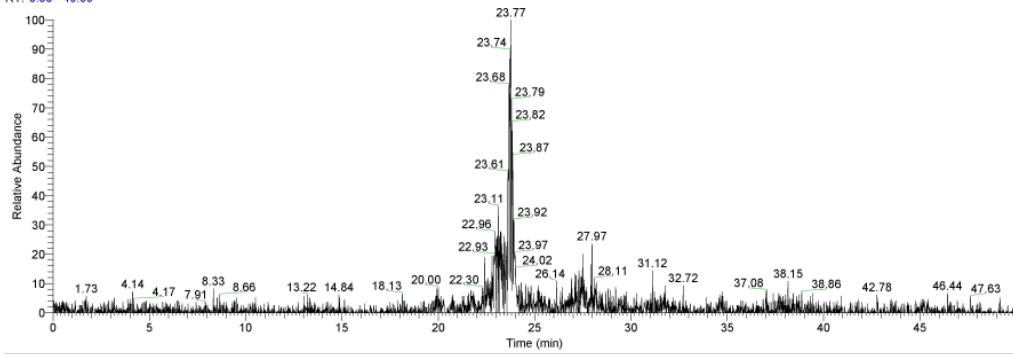
NL:
2.05E2
m/z=
764.51-
765.51 MS
MS-DATA

MS-DATA #1875 RT: 23.75 AV: 1 NL: 7.40E2
T: ITMS - c ESI Full ms [100.00-1000.00]



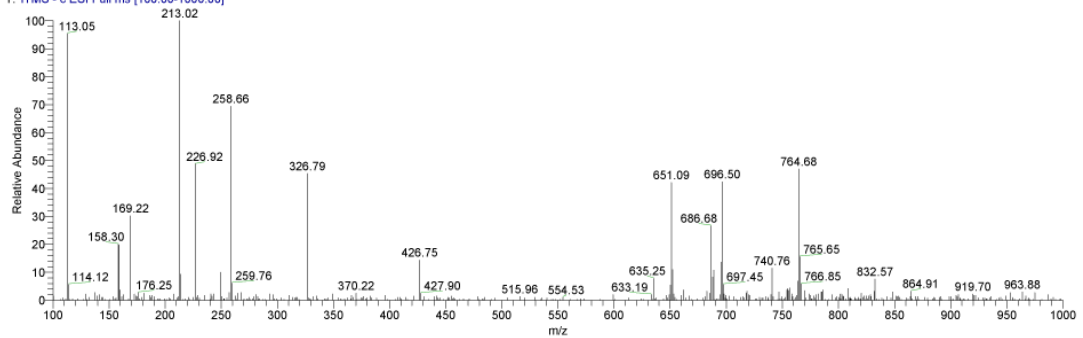
4

RT: 0.00 - 49.99



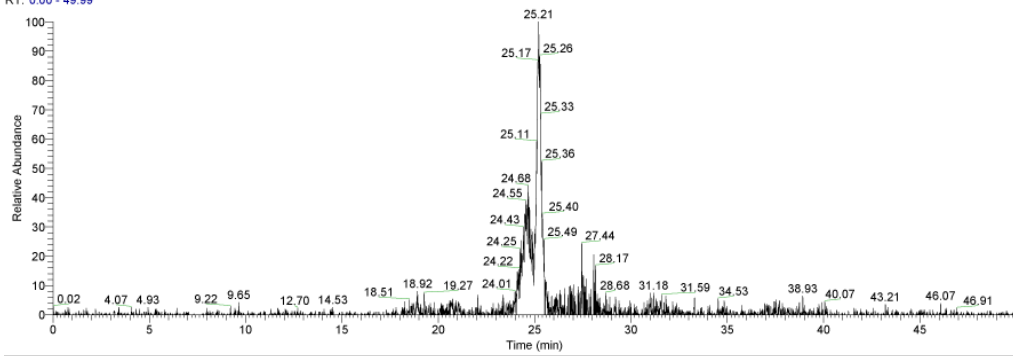
NL:
1.95E2
m/z=
650.15-
651.15 MS
MS-DATA

MS-DATA #1876 RT: 23.77 AV: 1 NL: 4.11E2
T: ITMS - c ESI Full ms [100.00-1000.00]



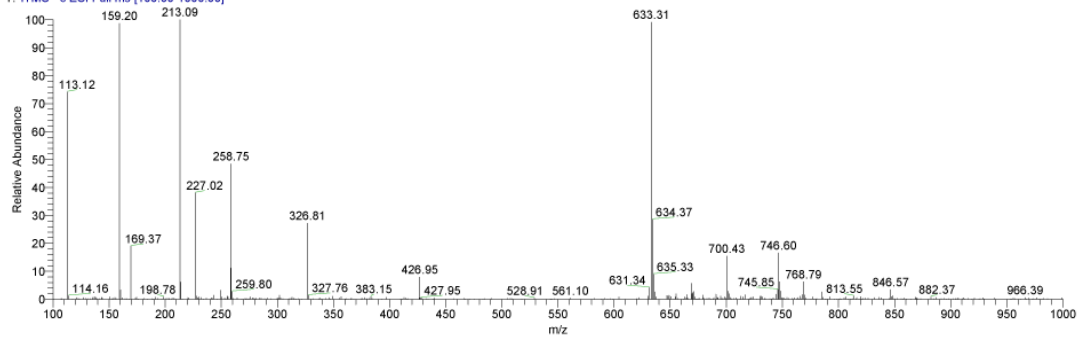
5

RT: 0.00 - 49.99



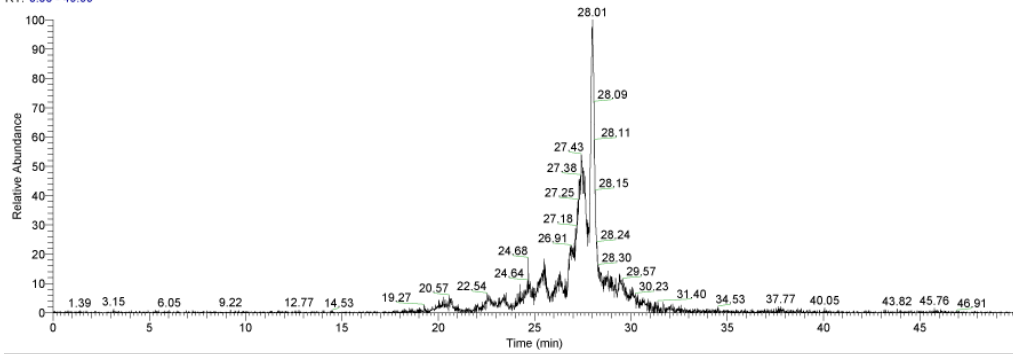
NL:
3.11E2
m/z=
633.70-
634.70 MS
MS-DATA

MS-DATA #1990 RT: 25.21 AV: 1 NL: 1.09E3
T: ITMS - c ESI Full ms [100.00-1000.00]



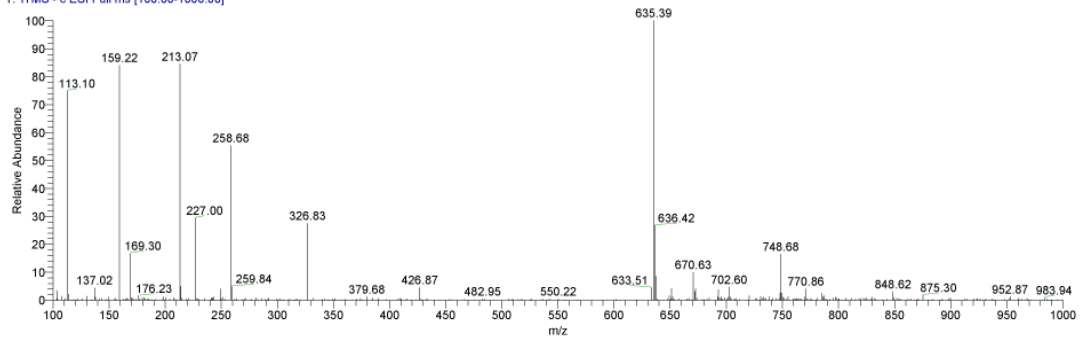
6

RT: 0.00 - 49.99



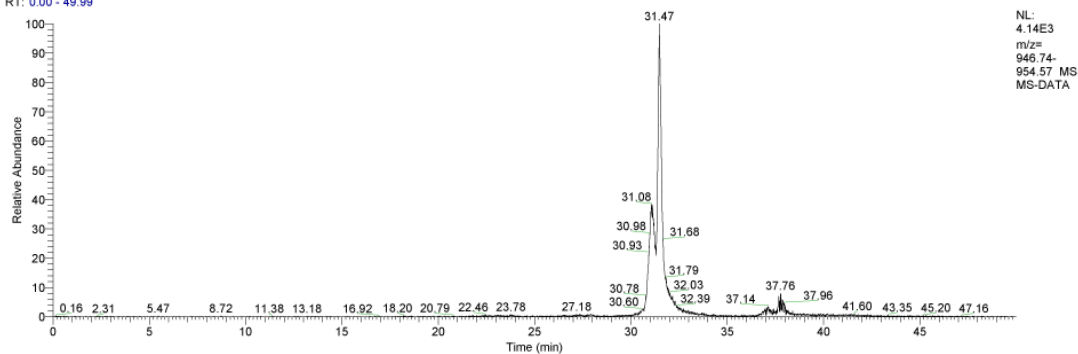
NL:
1.03E3
m/z=
634.49-
635.49 MS
MS-DATA

MS-DATA #2211 RT: 28.01 AV: 1 NL: 1.03E3
T: ITMS - c ESI Full ms [100.00-1000.00]



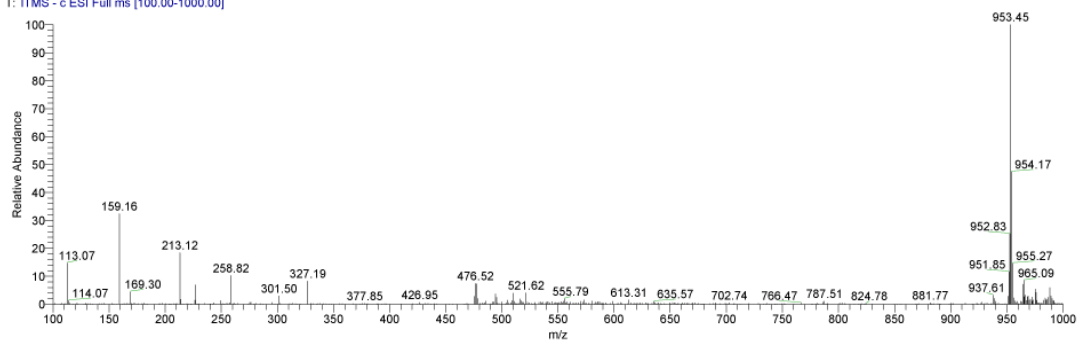
7

RT: 0.00 - 49.99



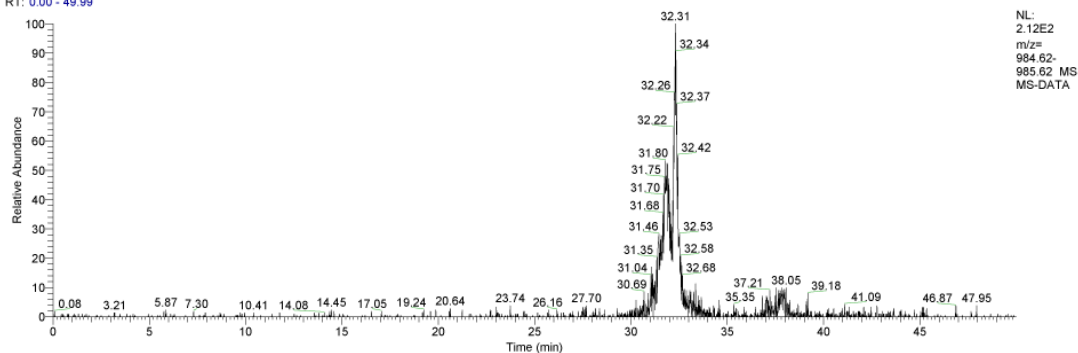
NL:
4.14E3
m/z=
946.74-
954.57 MS
MS-DATA

MS-DATA #2484 RT: 31.47 AV: 1 NL: 2.21E3
T: ITMS - c ESI Full ms [100.00-1000.00]



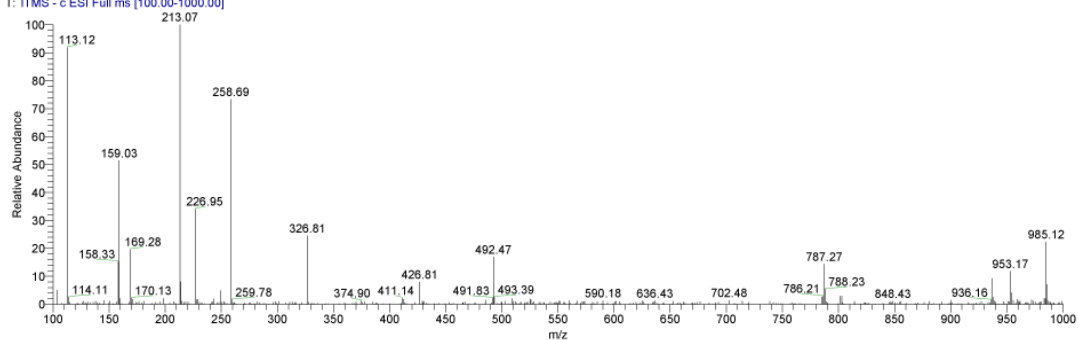
8

RT: 0.00 - 49.99

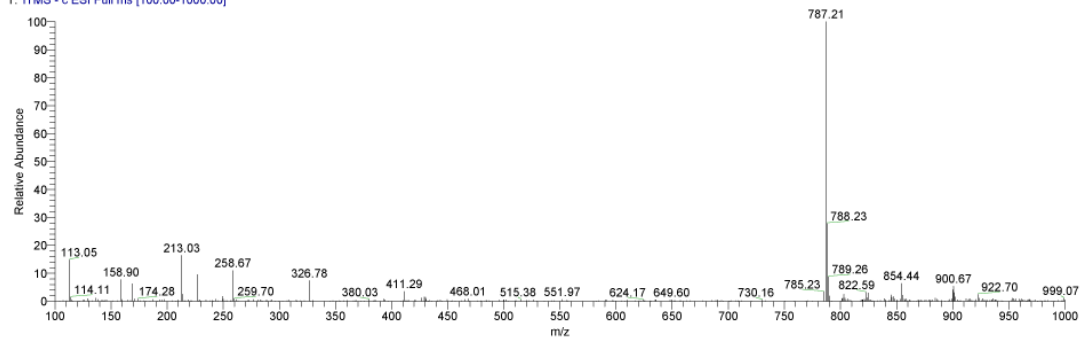
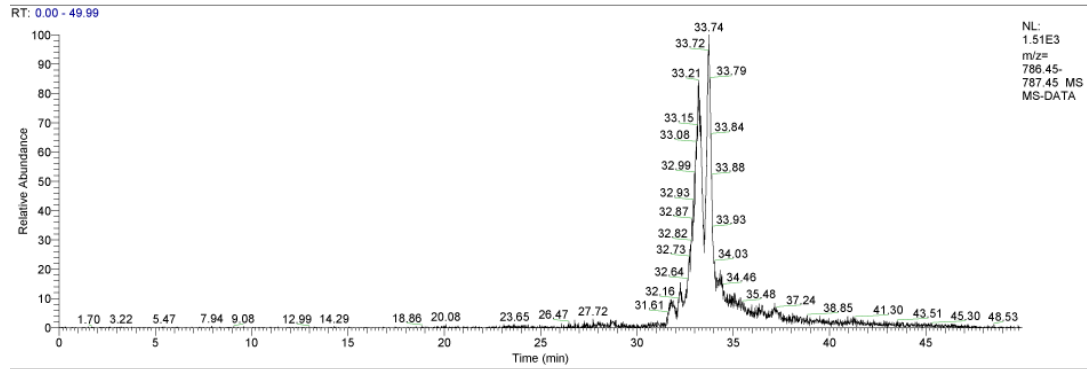


NL:
2.12E2
m/z=
984.62-
985.62 MS
MS-DATA

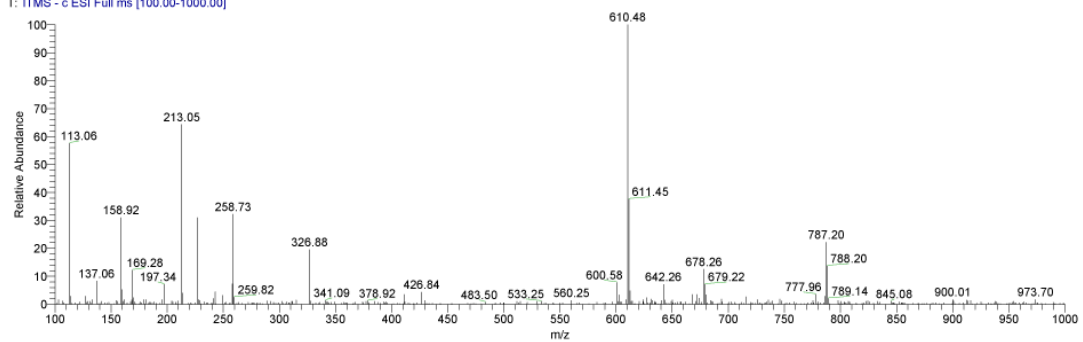
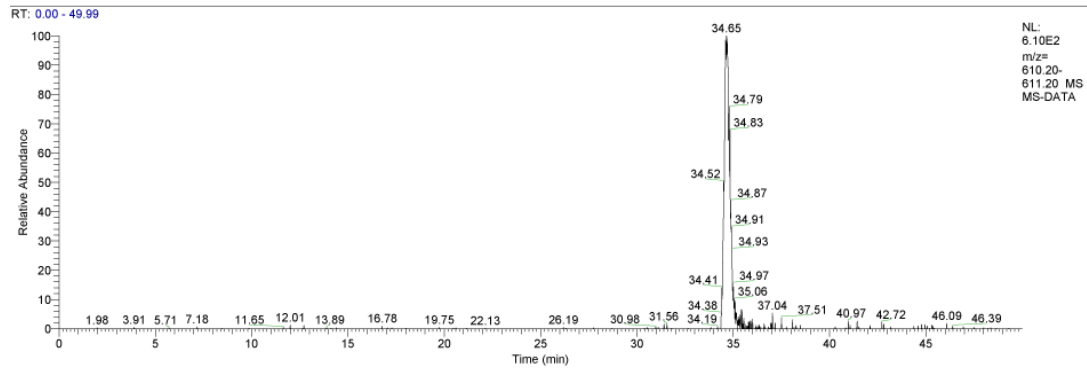
MS-DATA #2550 RT: 32.31 AV: 1 NL: 9.54E2
T: ITMS - c ESI Full ms [100.00-1000.00]



9

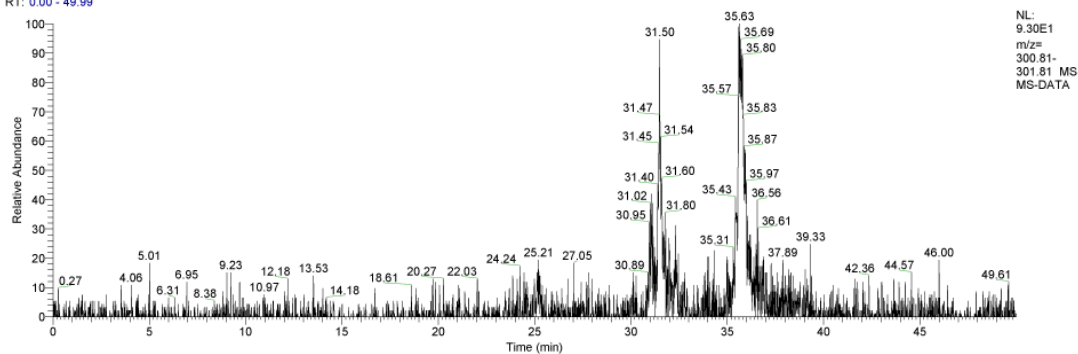


10



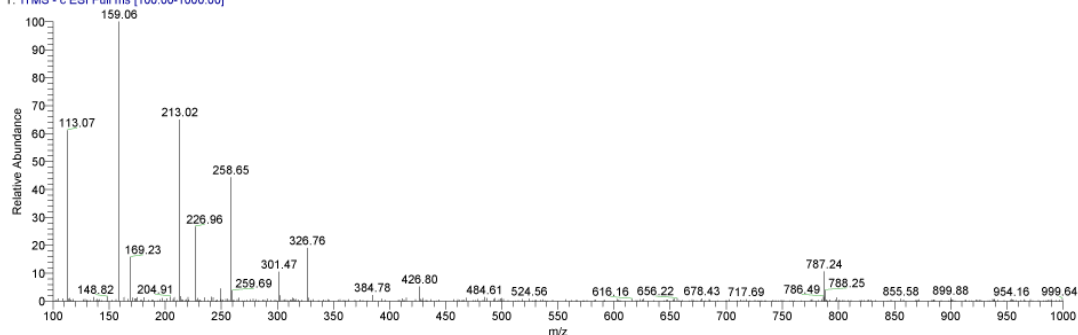
11

RT: 0.00 - 49.99



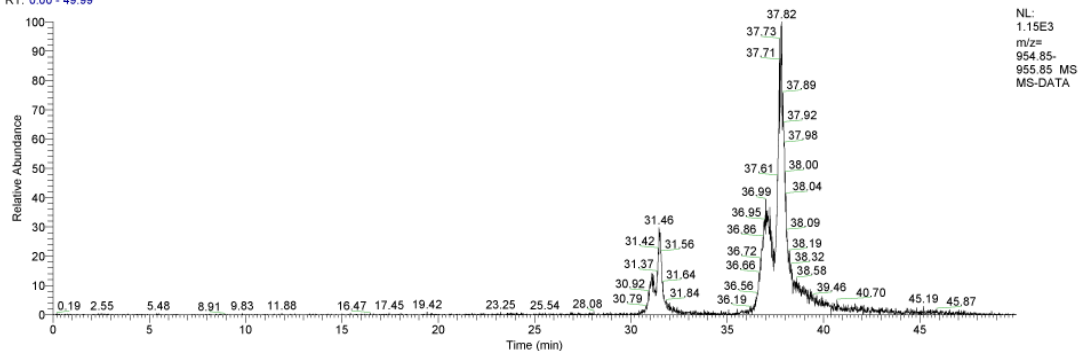
MS-DATA #2812 RT: 35.63 AV: 1 NL: 8.90E2

T: ITMS - c ESI Full ms [100.00-1000.00]



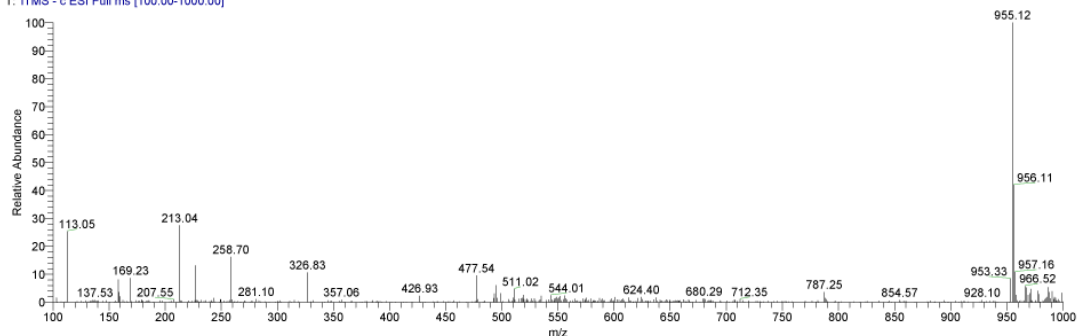
12

RT: 0.00 - 49.99



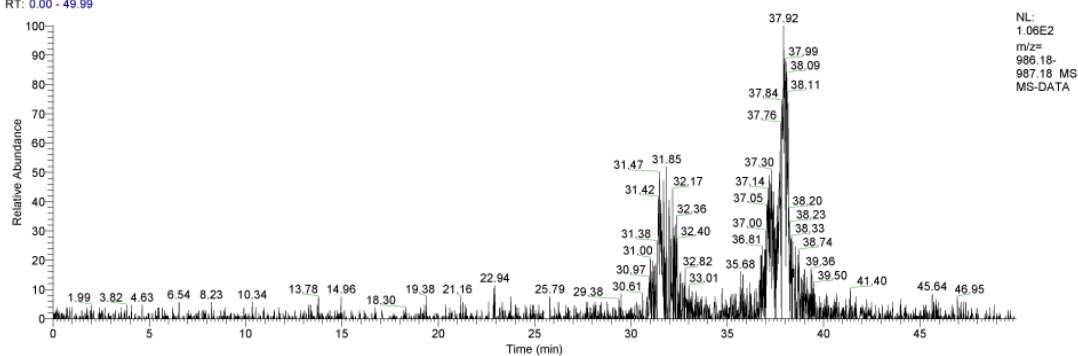
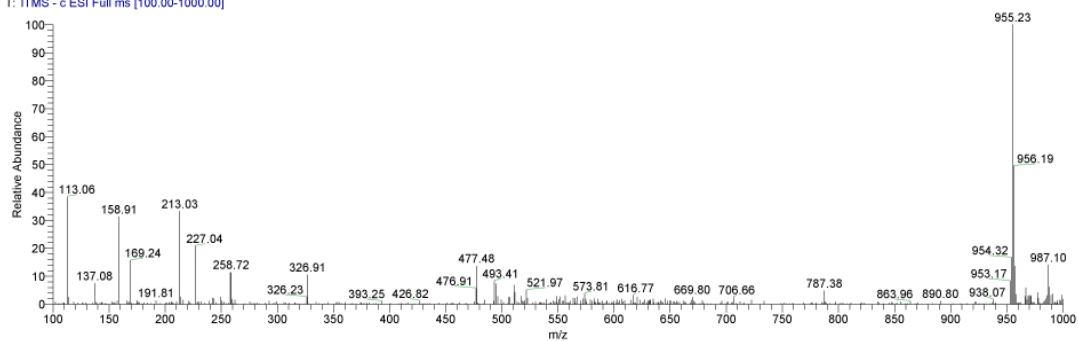
MS-DATA #2985 RT: 37.82 AV: 1 NL: 1.15E3

T: ITMS - c ESI Full ms [100.00-1000.00]



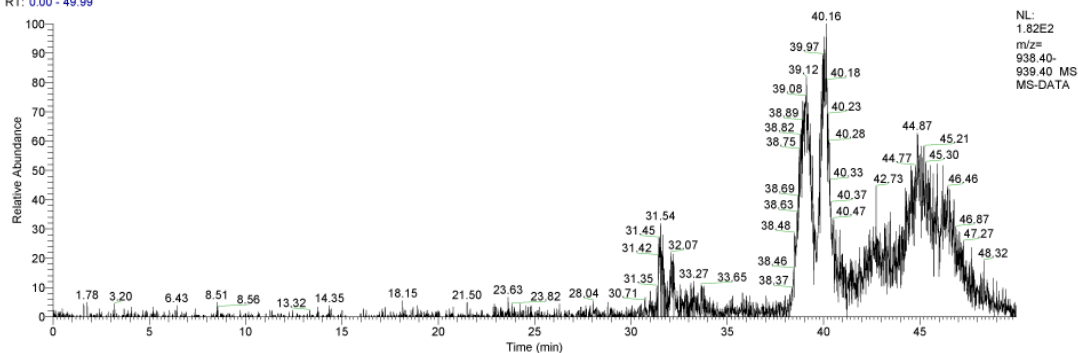
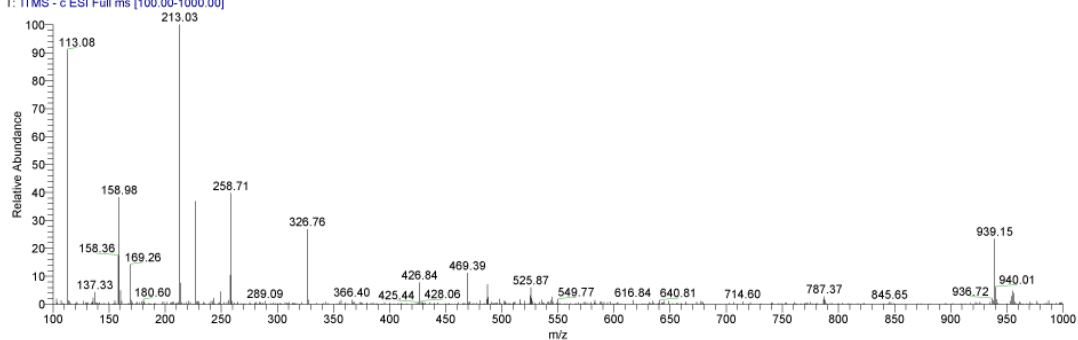
13

RT: 0.00 - 49.99

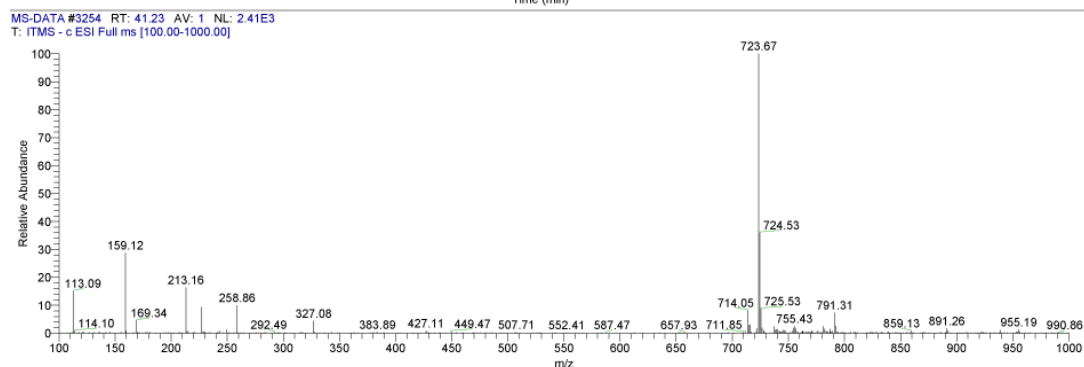
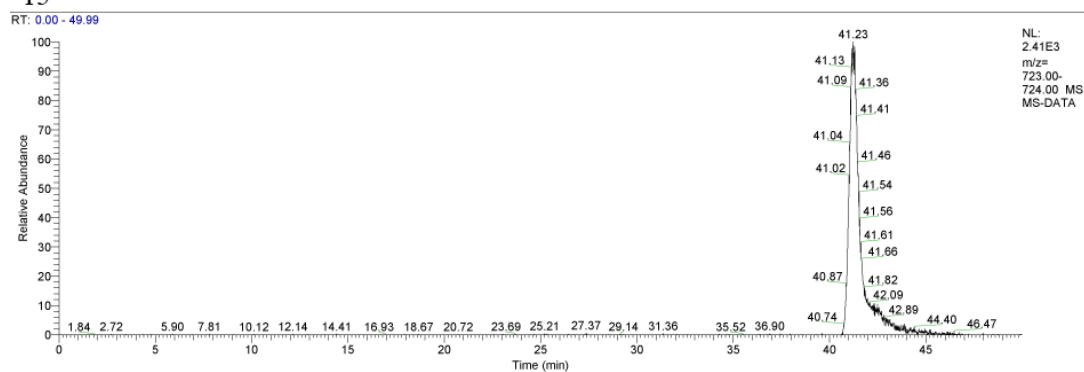
MS-DATA #2993 RT: 37.92 AV: 1 NL: 7.56E2
T: ITMS - c ESI Full ms [100.00-1000.00]

14

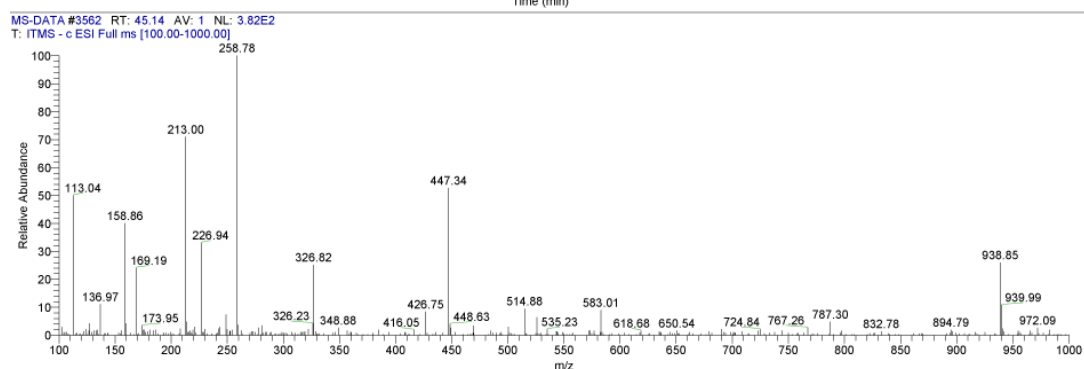
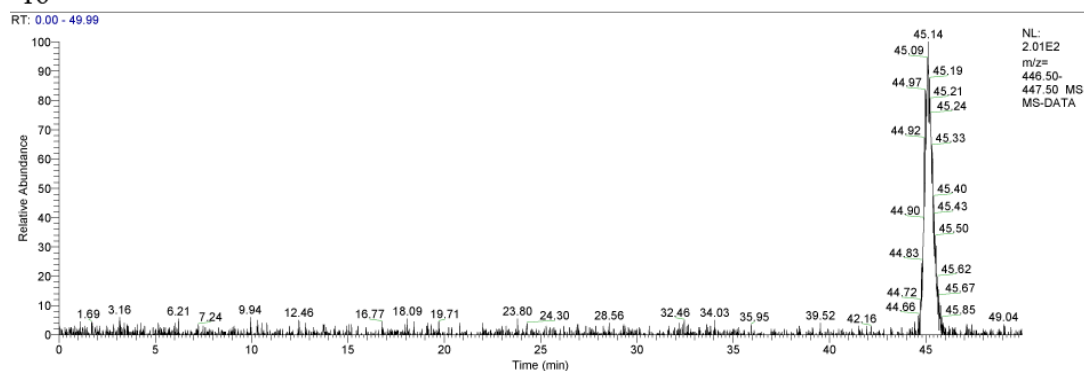
RT: 0.00 - 49.99

MS-DATA #3169 RT: 40.16 AV: 1 NL: 7.80E2
T: ITMS - c ESI Full ms [100.00-1000.00]

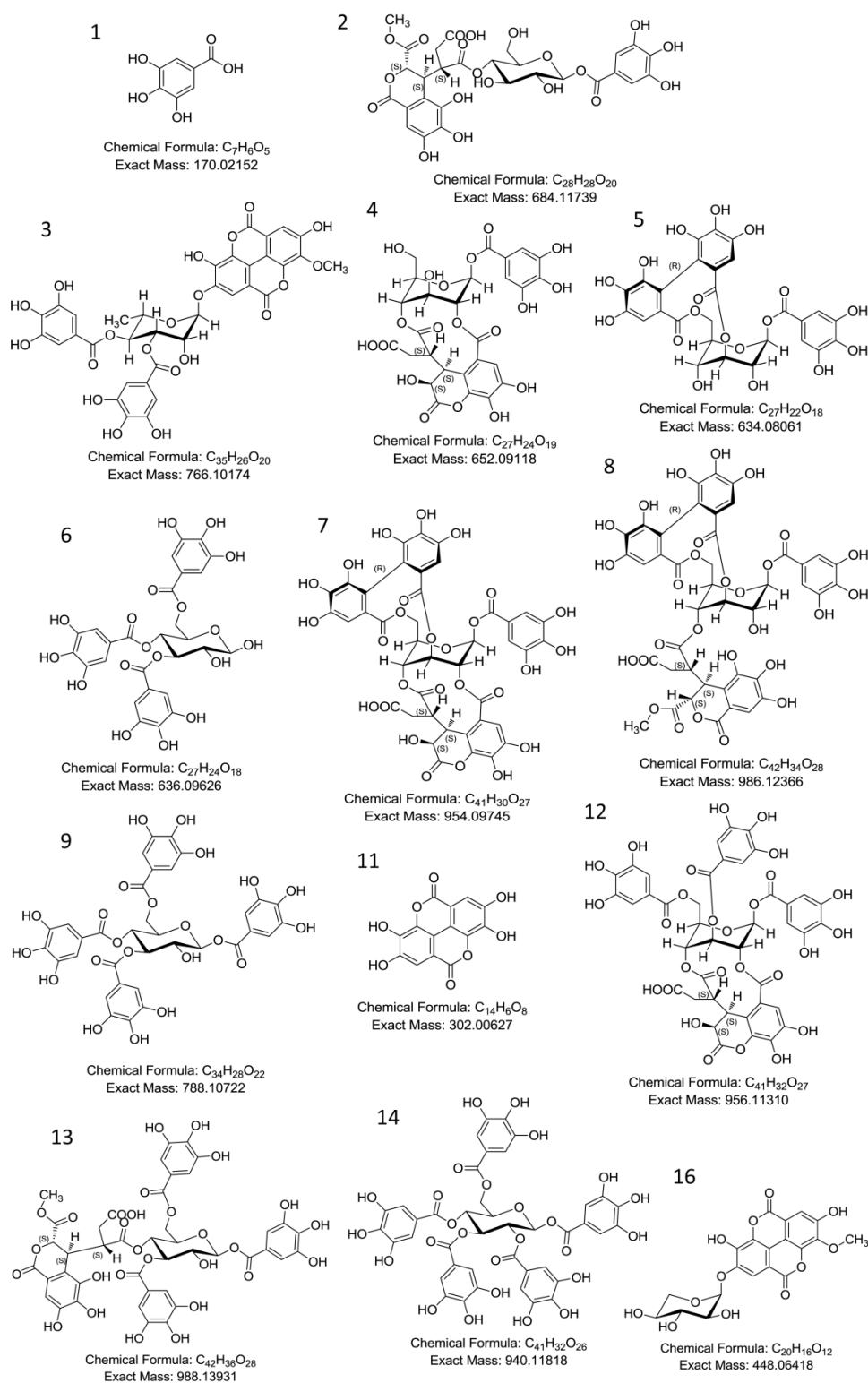
15



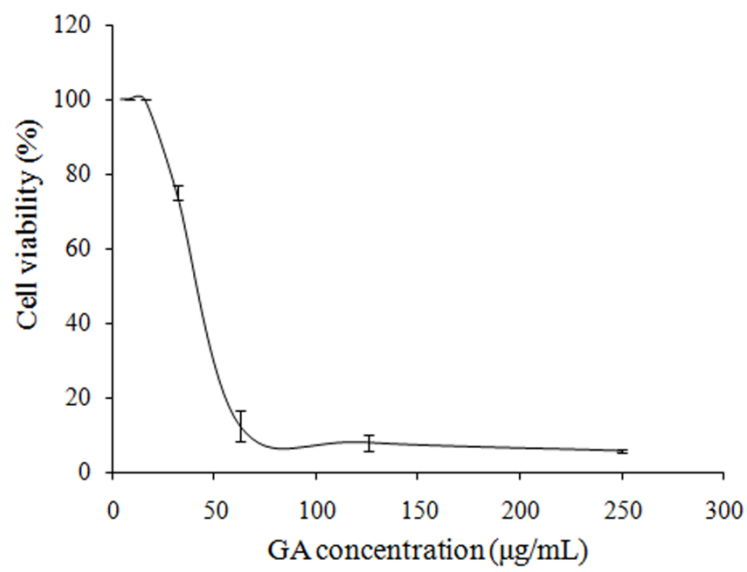
16



Supplemental Figure 2. MS spectra of the putative 16 compounds in EF. The number of each spectrum corresponds to the peak number in Table 2, Figure 1, Supplemental Figure 1 and Supplemental Figure 3.



Supplemental Figure 3. Structures of the 14 phenolic compounds identified from EF based on the HPLC-MS data and the known structures of reference compounds. Compound numbers, chemical formula and exact mass are presented.



Supplemental Figure 4. Effect of GA on HSC-T6 cell viability. The viability rate (%) of HSC-T6 cells manifest decreased with increased GA concentration.