

Supplementary Information

**UPLC-ESI-Q-TOF-MS^E -based metabolomics analysis of *Acer mono* Sap and evaluation
for osteogenic activity in mouse osteoblast cells**

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Table S1 List of forward and reverse primers used in the gene expression studies by qRT-PCR analysis

Gene	Forward primer	Reverse primers	Tm
Alkaline Phosphatase	CACAGATTCCCAAAGCACCT	GGGATGGAGGAGAGAAGGTC	55.45/59.5
Osteocalcin	CCTGAGTCTGACAAAGCCTCA	GCCGGAGTCTGTTCACTACCTT	58.4/60.0
Collagen1 α 1	GCTTCACCTACAGCACCC TTGT	TGACTGTCTGCCCAAGTTC	60.0/58.20
Osteopontin	CCCGGTGAAAGTGACTGATTCT	GATCTGGGTGCAGGCTGTAAA	58.4/58.2
Runx2	GGCACAGACAGAACGCTTGATGA	GAATGCGCCCTAAATCACTGA	58.4/56.2
Osterix	GGAGGTTCACTCCATTCCA	TAGAAGGAGCAGGGACAGA	55.4/57.4

Fig. S1

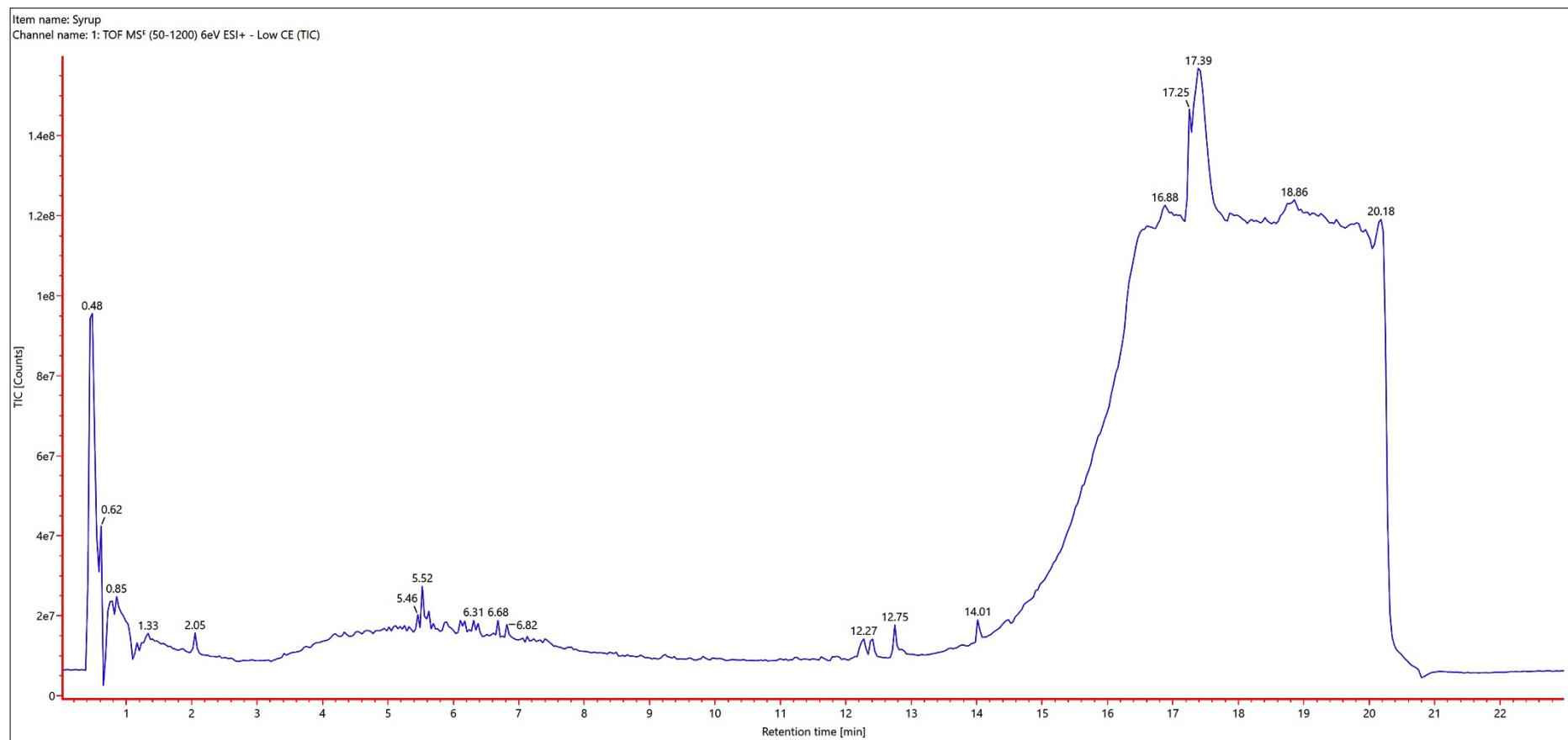
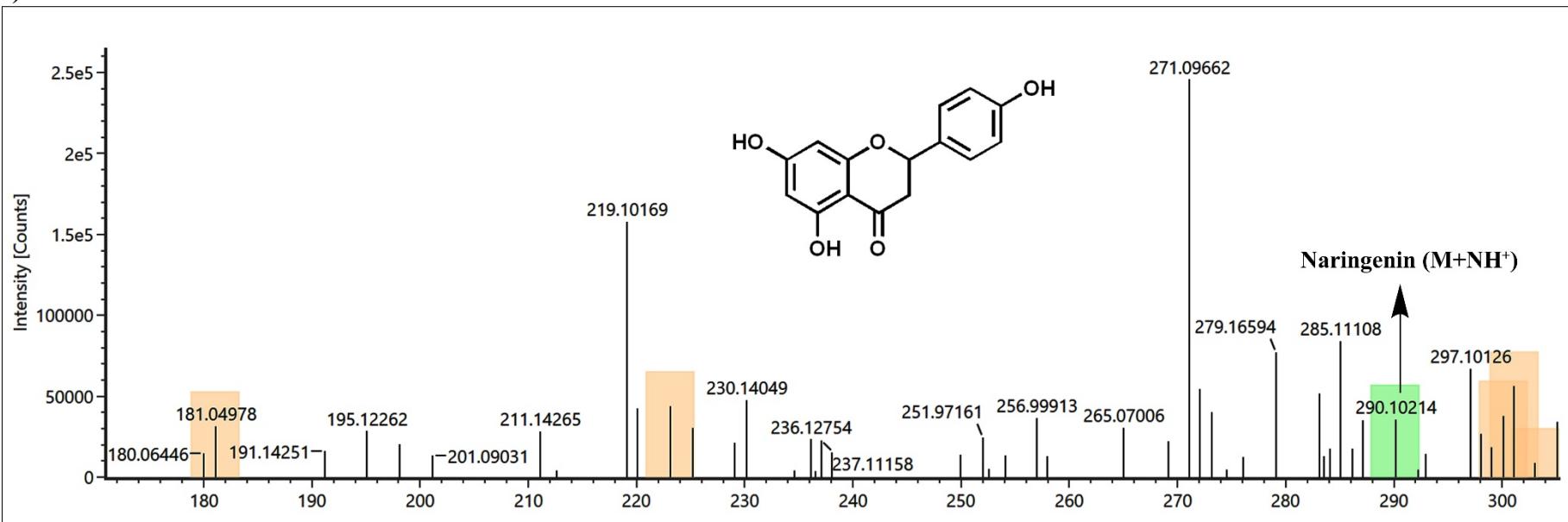


Fig. S2

a)



b)

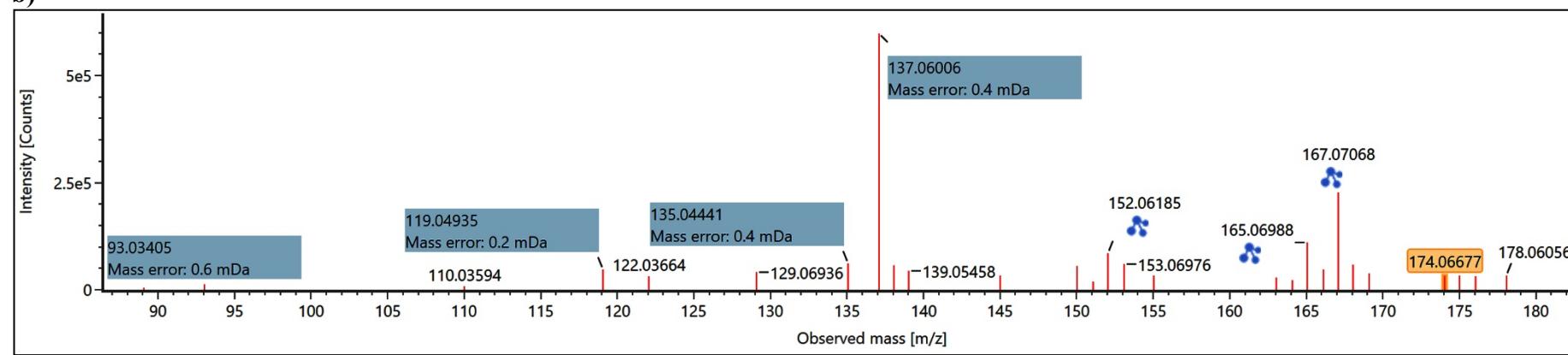


Fig. S3

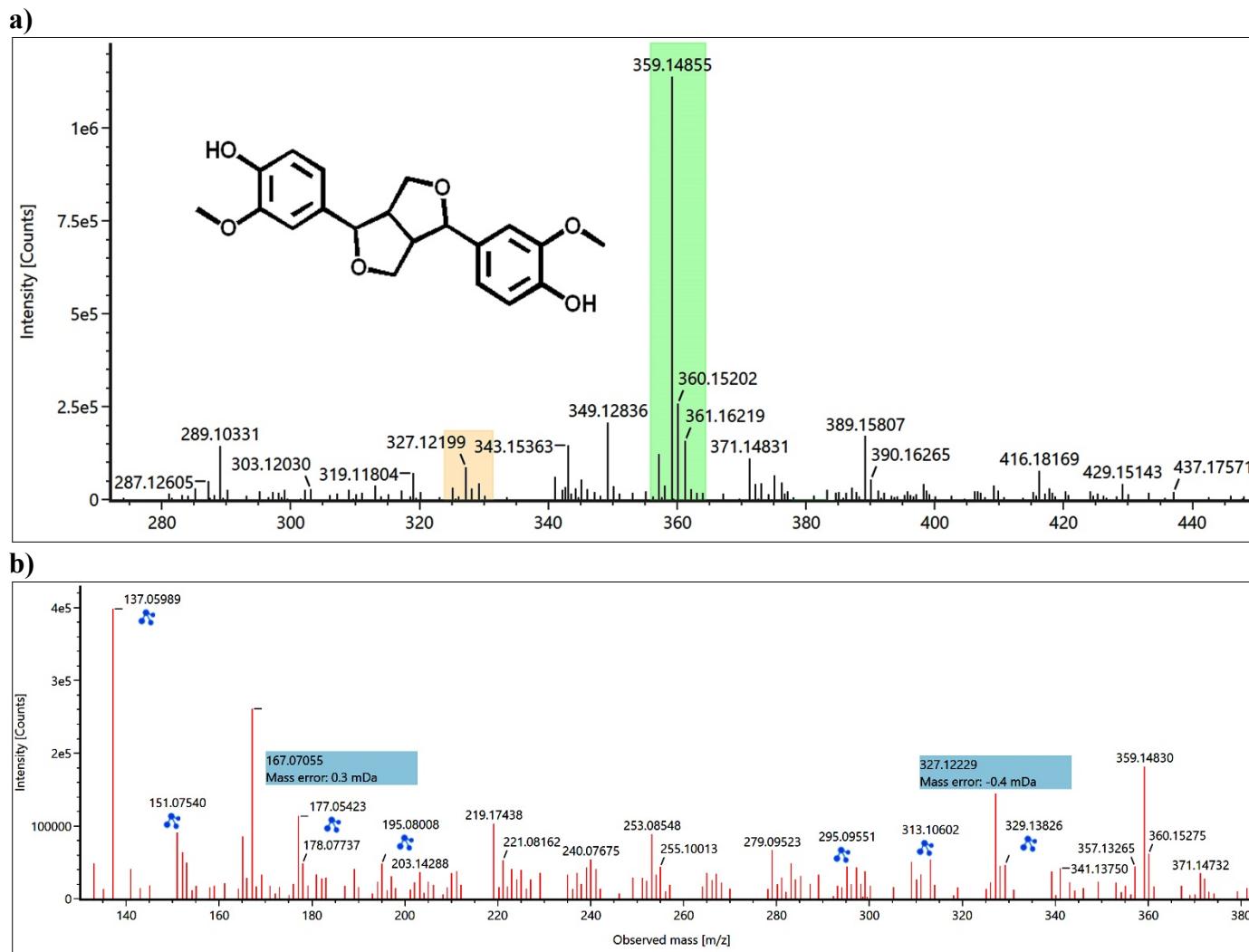


Fig. S4 Structures of the compounds identified from SAM using MS/MS fragmentation analysis

