

Natural deep eutectic solvents for the extraction of triterpenoid saponins from *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen

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Tables

Table S1. The list of triterpene saponins predicted in different parts of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen based on literature mining

Compound Name	Elemental composition	Molar mass	Theoretical m/z	Ref
7- <i>O</i> - α -rhamnopyranosyl-(S)-quercetin-3- <i>O</i> - β -6-caffeoyl-glucopyranosyl-(1 \rightarrow 2)- α -rhamnopyranoside	C ₄₂ H ₄₆ O ₂₃	918.8134	917.8061	[1]
7- <i>O</i> - α -rhamnopyranosyl-(S)-quercetin-3- <i>O</i> - β -glucopyranosyl-(1 \rightarrow 2)- α -rhamnopyranoside	C ₃₃ H ₄₀ O ₂₀	756.6686	755.6613	[1]
Acanthopanaxoside E	C ₄₂ H ₆₆ O ₁₅	810.4402	809.4329	[2]
Acanthoside D	C ₄₈ H ₇₈ O ₁₈	942.5188	941.5115	[3]
Acutoside A	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[4]
Anchusosid 2	C ₄₈ H ₇₈ O ₁₈	942.5188	941.5115	[4]
Aradecoside D	C ₅₉ H ₉₆ O ₂₇	1237.395	1236.3877	[5]
Araliasaponin I	C ₄₇ H ₇₆ O ₁₈	928.5032	927.4959	[6]
Araliasaponin II	C ₄₇ H ₇₆ O ₁₉	944.4981	943.4908	[6]
Araliasaponin III	C ₅₃ H ₈₆ O ₂₃	1090.5559	1089.5486	[6]
Araliasaponin IV	C ₅₄ H ₈₈ O ₂₃	1104.5716	1103.5643	[6]
Araliasaponin XVII	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[7]
Araliaarmoside	C ₅₉ H ₉₄ O ₂₈	1250.5932	1249.5859	[8]
Araliachinosides I	C ₃₆ H ₅₆ O ₈	616.3975	615.3902	[9]
Araliasaponin IV	C ₅₄ H ₈₈ O ₂₄	1120.5666	1119.5593	[10]
Araliasaponin IX	C ₅₃ H ₈₆ O ₃₂	1090.5560	1089.5487	[6]
Araliasaponin V	C ₅₄ H ₈₈ O ₂₃	1104.5716	1103.5643	[6]
Araliasaponin VI	C ₅₄ H ₈₈ O ₂₄	1120.5666	1119.5593	[6]
Araliasaponin VII	C ₅₄ H ₈₈ O ₂₄	1120.5666	1119.5593	[6]
Araliasaponin VIII	C ₆₀ H ₉₈ O ₃₀	1298.6143	1297.6070	[6]
Araloside A	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[11]
Araloside A methyl ester	C ₄₈ H ₇₆ O ₁₈	940.5032	939.4959	[3]
Araloside B	C ₅₂ H ₈₂ O ₂₂	1058.5298	1057.5225	[11]

Araloside C	C ₅₃ H ₈₄ O ₂₃	1088.5403	1087.5330	[3]
Araloside D	C ₄₆ H ₇₄ O ₁₆	882.4977	881.4904	[12]
Araloside G	C ₅₄ H ₈₈ O ₂₃	1104.5716	1103.5643	[3]
Araloside H	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[13]
Araloside J	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[13]
Armatoside A	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[14]
Armatoside B	C ₄₇ H ₇₄ O ₁₉	942.4824	941.4751	[14]
Calendulaglycoside A	C ₅₄ H ₈₆ O ₂₄	1118.5509	1117.5436	[15]
Calendulaglycoside C	C ₄₈ H ₇₆ O ₁₉	956.4980	955.4907	[15]
Calenduloside B	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[16]
Calenduloside E	C ₃₆ H ₅₆ O ₉	632.3924	631.3851	[16]
Calenduloside G	C ₄₁ H ₆₄ O ₁₃	764.4347	763.4274	[16]
Chikusetsusaponin (1)	C ₃₆ H ₅₈ O ₈	618.4131	617.4058	[3]
Chikusetsusaponin IV	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[3]
Chikusetsusaponin Ib	C ₄₁ H ₆₄ O ₁₃	764.4347	763.4274	[17]
Chikusetsusaponin IVa	C ₄₂ H ₆₆ O ₁₄	808.4609	807.4536	[18]
Chikusetsusaponin V	C ₄₈ H ₇₆ O ₁₉	956.4981	955.4908	[18]
Collinsonidin	C ₄₁ H ₆₆ O ₁₃	766.4503	765.4430	[4]
Congmunoside IX	C ₅₃ H ₈₆ O ₂₃	1088.5403	1087.5330	[19]
Congmunoside VIII	C ₆₀ H ₉₈ O ₃₀	1298.6143	1297.6070	[2]
Congmunoside X	C ₆₀ H ₉₈ O ₂₈	1266.6245	1265.6172	[20]
Congmunoside XV	C ₅₄ H ₈₈ O ₂₄	1120.5666	1119.5593	[21]
Congmuyanoside A	C ₄₁ H ₆₆ O ₁₄	782.4453	781.4380	[4]
Congmuyanoside B	C ₄₈ H ₇₈ O ₁₉	958.5137	957.5064	[22]
Congmuyanoside D	C ₄₂ H ₆₈ O ₁₄	764.4609	763.4536	[23]
Congmuyanoside E	C ₄₈ H ₇₆ O ₁₉	956.4981	955.4908	[4]
Congmuyanoside G	C ₆₀ H ₉₈ O ₂₉	1282.6193	1281.6120	[4]
Congmuyenoside A	C ₄₁ H ₆₆ O ₁₃	766.4503	765.4430	[24]
Congmuyenoside III	C ₆₀ H ₉₈ O ₂₉	1282.6193	1281.6120	[4]
Congmuyenoside IV	C ₄₂ H ₆₈ O ₁₅	812.4558	811.4485	[23]
Durupcoside C	C ₄₇ H ₇₆ O ₁₈	928.5032	927.4959	[25]
Echinocystic acid-3- <i>O</i> -β- <i>D</i> -glucopyranosyl-(1-3)- β-	C ₄₆ H ₄₇ O ₁₅	839.2915	838.2842	[1]

<i>D</i> -glucuronopyranoside-6'- <i>O</i> -butyl ester				
Eclalbasaponin I	C ₄₂ H ₆₈ O ₁₄	764.4609	763.4536	[23]
Eclalbasaponin III	C ₄₈ H ₇₈ O ₁₉	958.5137	957.5064	[26]
Ecliptasaponin B	C ₄₁ H ₆₆ O ₁₄	958.5137	957.5064	[27]
Elatoside A	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[18]
Elatoside B	C ₄₈ H ₇₆ O ₁₉	956.4981	955.4908	[18]
Elatoside D	C ₅₄ H ₈₆ O ₂₄	1118.5509	1117.5436	[28]
Elatoside G	C ₃₆ H ₅₆ O ₁₁	664.3824	663.3751	[29]
Elatoside H	C ₄₂ H ₆₆ O ₁₅	810.4402	809.4329	[29]
Ginsenin R0	C ₄₈ H ₇₆ O ₁₉	956.4981	955.4908	[18]
Guaiacin B	C ₄₇ H ₇₆ O ₁₇	912.5083	911.5010	[30]
Hemsloside G2	C ₅₅ H ₈₈ O ₂₄	1132.5666	1131.5593	[7]
Kalopanax-saponin F	C ₅₃ H ₈₄ O ₂₃	1088.5403	1087.5330	[31]
Kalopanax-saponin F methyl ester	C ₅₃ H ₈₆ O ₂₃	1088.5403	1087.5330	[31]
Lucynoside E	C ₄₂ H ₆₈ O ₁₄	764.4609	763.4536	[23]
Lucynoside H	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[4]
Momordin Ic	C ₄₁ H ₆₄ O ₁₃	764.4347	763.4274	[32]
Olean-12-en-28-oic acid, 3- <i>O</i> -β-d-glucopyranosyl(1 → 4)-β-d-glucopyranosyl-olean acid	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[23]
Pseudoginsenoside RT1	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[33]
Pseudoginsenoside RT1 butyl ester	C ₅₁ H ₈₂ O ₁₈	982.5501	981.5428	[33]
Quinoasaponin 2	C ₄₈ H ₇₈ O ₁₉	958.5137	957.5064	[34]
Quinoasaponin B	C ₄₇ H ₇₆ O ₁₈	928.5032	927.4959	[10]
Randianin	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[4]
Salsoloside C	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[35]
Silphioside A	C ₄₃ H ₆₈ O ₁₄	808.4609	807.4536	[3]
Silphioside B	C ₄₂ H ₆₈ O ₁₃	780.4660	779.4587	[23]
Silphioside E	C ₄₈ H ₇₈ O ₁₈	942.5188	941.5115	[4]
Spinasaponin A	C ₄₂ H ₆₆ O ₁₄	808.4609	807.4536	[36]
Spinasaponin A ₂₈ - <i>O</i> -glycoside	C ₄₈ H ₇₆ O ₁₉	956.4981	955.4908	[36]
Stipuleanoside R1	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[18]
Stipuleanoside R2	C ₅₃ H ₈₄ O ₂₃	1088.5403	1087.5330	[28]
Taibaienoside I	C ₅₁ H ₈₂ O ₁₈	982.5501	981.5428	[37]

Taibaienoside II	C ₄₉ H ₇₈ O ₁₈	954.5188	953.5115	[37]
Taibaienoside III	C ₅₅ H ₈₈ O ₂₃	1116.5716	1115.5643	[37]
Taibaienoside IV	C ₄₆ H ₇₄ O ₁₄	850.5078	849.5005	[37]
Taibaienoside V	C ₅₂ H ₈₆ O ₁₄	934.6018	933.5945	[37]
Taibaienoside VI	C ₄₇ H ₇₄ O ₁₈	926.4875	925.4802	[37]
Taibaienoside VII	C ₅₅ H ₈₈ O ₂₃	1116.5716	1115.5643	[37]
Taibaienoside VIII	C ₅₇ H ₉₂ O ₂₃	1144.6029	1143.5956	[37]
Taibaienoside IX	C ₄₇ H ₇₆ O ₁₇	912.5083	911.5010	[4]
Tarasaponin III	C ₄₈ H ₇₆ O ₁₉	882.4977	881.4904	[2]
Tarasaponin VI	C ₄₁ H ₆₄ O ₁₃	764.4347	763.4274	[2]
Tragopogonoside B	C ₄₂ H ₆₆ O ₁₅	810.4402	809.4329	[38]
Yuzhizioside IV	C ₅₂ H ₈₄ O ₂₁	1044.5505	1043.5432	[17]
Zingibroside R1 dimethyl ester	C ₄₄ H ₇₀ O ₁₄	822.4766	821.4693	[36]
α -Hederin	C ₄₁ H ₆₆ O ₁₂	750.4554	749.4481	[25]
Congmuyenoside I	C ₄₂ H ₆₈ O ₁₄	764.4609	763.4536	[23]
Congmuyenoside II	C ₄₈ H ₇₈ O ₁₉	958.5137	957.5064	[23]
Congmuyenoside III	C ₆₀ H ₉₈ O ₂₉	1282.6193	1281.6120	[23]
Congmuyenoside IV	C ₄₂ H ₆₈ O ₁₅	812.4558	811.4485	[23]

Table S2. Triterpene saponins annotated in the extracts of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots by reversed ultra phase high performance liquid chromatography-mass spectrometry (RP-UHPLC-QqTOF-MS) in untargeted SWATH experiments (with the inclusion list containing the *m/z* of all predicted [M-H]⁻ ions using published data)

No	t _R (min)	<i>m/z</i> [M-H] ⁻ - observed	<i>m/z</i> [M-H] ⁻ - calculated	Elemental composition [M-H] ⁻	MS2 fragmentation patterns - product ions, <i>m/z</i> (rel. intensity)	Δ <i>m</i> (ppm)	Assignment
3	3.7	749.4481	749.4482	C ₄₁ H ₆₅ O ₁₂ ⁻	455.3532 (60%), 485.1508 (75%), 617.4060 (35%), 749.4481 (40%)	0.13	α-Heredin
4	3.4	765.4443	765.4431	C ₄₁ H ₆₅ O ₁₃ ⁻	471.3479 (30%), 959.3343 (20%), 633.4011 (15%), 757.3878 (100%), 765.4443 (17%)	- 1.57	Collinsonidin
5	3.5	778.3300	778.4509	C ₄₂ H ₆₆ O ₁₃ ^{•-}	425.1316 (10%), 455.3541 (20%), 485.1516 (65%), 778.3300 (60%)	155	Acutoside A
6	4.9	781.4171	781.4380	C ₄₁ H ₆₅ O ₁₄ ⁻	387.2855 (90%), 401.2492 (20%), 619.3331 (10%), 665.3183 (15%), 779.4215 (10%), 777.4070 (9%), 780.4209 (8%), 781.4171 (5%)	27	Congmuyenoside A
7	4.6	809.4333	809.4329	C ₄₂ H ₆₅ O ₁₅ ⁻	556.3175 (40%), 751.4649 (30%), 767.4594 (35%), 779.4564 (15%), 781.4732 (100%), 793.4362 (90%), 809.4333 (60%)	-0.49	Elatoside
8	3.3	811.4519	811.4485	C ₄₂ H ₆₇ O ₁₅ ⁻	345.2268 (100%), 471.5677 (15%), 485.1509 (30%), 721.3301 (20%), 777.3617 (28%), 809.4022 (45%), 811.4519 (48%)	-4.2	Congmuyenoside IV
9	3.8	911.5015	911.5010	C ₄₇ H ₇₅ O ₁₇ ⁻	455.3515 (20%), 617.4057 (15%), 749.4479 (20%), 77.4590 (18%), 911.5015 (100%)	- 0.5	Guaiacin B
10	4.5	925.4814	925.4802	C ₄₇ H ₇₃ O ₁₈ ⁻	455.3541 (3%), 569.3852 (30%), 613.3746 (10%), 775.4288 (4%), 794.4439 (2%), 925.4814 (100%)	-1.3	Araloside A
11	3.5	927.4955	927.4959	C ₄₇ H ₇₅ O ₁₈ ⁻	471.3419 (20%), 633.4008 (20%), 747.4391 (18%), 765.4456 (20%), 795.4537 (40%), 927.4955 (100%)	0.5	Araliasaponin I
12	5.1	939.4618	939.4959	C ₄₈ H ₇₅ O ₁₈ ⁻	569.3852 (10%), 631.3857 (26%), 793.4388 (40%), 847.4473 (20%), 865.4611 (25%), 939.4618 (100%)	36	Araloside A methyl ester
13	4.6	941.4753	941.4752	C ₄₇ H ₇₃ O ₁₉ ⁻	455.3509 (5%), 471.3465 (5%), 631.3852 (7%), 747.4331 (35%), 793.4388 (42%), 809.4331 (3%), 927.4957 (36%), 941.4753 (100%)	-0.1	Armatoside B
14	3.8	941.5046	941.5115	C ₄₈ H ₇₇ O ₁₈ ⁻	456.3556 (3%), 585.3772 (5%), 617.4064 (7%), 779.4526 (20%), 925.4833 (16%), 941.5046 (100%)	7.3	

15	3.5	943.4905	943.4908	C ₄₇ H ₇₅ O ₁₉ ⁻	765.4419 (18%), 779.4282 (17%), 927.4974 (60%), 943.4905 (100%)	0.3	Araliasaponin II
16	3.9	953.5104	953.5115	C ₄₉ H ₇₇ O ₁₈ ⁻	455.3523 (15%), 617.4082 (12%), 749.4462 (18%), 795.4536 (9%), 822.4728 (9%), 893.4927 (10%), 911.5016 (30%), 925.4814 (26%), 953.5104 (100%)	1.2	Taibaienoside II
17	4.9	955.4890	955.4908	C ₄₈ H ₇₄ O ₁₉ ⁻	454.3310 (3%), 594.3418 (5%), 761.4049 (6%), 793.4257 (10%), 823.3948 (5%), 913.4877 (9%), 925.4790 (25%), 955.4890(100%)	1.9	Clendulagluconide C
18	4.4	957.4942	957.5067	C ₄₈ H ₇₇ O ₁₉ ⁻	791.4209 (10%), 925.4787 (13%), 955.4912 (100%), 957.4942 (15%)	13	Congmuyanoside B
19	4.5	1043.5432	1043.5432	C ₅₂ H ₈₃ O ₂₁ ⁻	569.3822 (15%), 613.3765 (8%), 731.4388 (9%), 775.4326 (12%), 925.4788 (100%), 1043.5432 (9%)	0	Yuzhizioside IV
20	4.4	1057.5363	1057.5225	C ₅₂ H ₈₁ O ₂₂ ⁻	455.3532 (2%), 551.3732 (3%), 585.3641 (3%), 701.4259 (7%), 745.4197 (3%), 895.4713 (15%), 911.5030 (5%), 955.4884 (3%), 1057.5363 (100%)	13	Araloside B
21	4.2	1087.5337	1087.5335	C ₅₃ H ₈₃ O ₂₃ ⁻	455.3501 (2%), 551.3726 (2%), 702.42875 (2%), 775.4270 (2%), 793.4383 (1.5%), 925.4798 (8%), 956.4949 (5%), 1087.5337 (100%)	-0.6	Araloside C
22	4.5	1089.5476	1089.5487	C ₅₃ H ₈₅ O ₂₃ ⁻	455.3521 (2%), 587.3969 (5%), 719.4372 (20%), 749.4493 (13%), 881.4925 (100%), 925.4795 (12%), 1043.5444 (35%), 1089.5476 (30%)	1	Araliasaponin III
23	5.1	1117.5419	1117.5436	C ₅₄ H ₈₅ O ₂₄ ⁻	959.3599 (15%), 749.4479 (10%), 793.4357 (35%), 866.4649 (15%), 919.4631 (7%), 937.4834 (50%), 955.4900 (51%), 1013.5322 (47%), 1057.5603 (36%), 1099.5340 (55%), 1117.5419 (100%)	1.5	Calendulaglycoside A
24	4.0	1119.5585	1119.5593	C ₅₄ H ₈₇ O ₂₄	609.3126 (10%), 775.4263 (5%), 955.4899 (7%), 1087.5365 (3%), 1119.5585 (100%)	0.7	Araliasaponin IV
25	3.4	1249.5862	1249.5859	C ₅₉ H ₉₃ O ₂₈	763.4331 (10%), 795.4525 (11%), 925.4817 (100%), 1249.5862 (95%)	-0.2	Araliaarmoside

Table S3. List of compounds confirmed by MS² analysis in the roots of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen

Compound Name	t _R	<i>m/z</i> [M-H] ⁻ observed	<i>m/z</i> [M-H] ⁻ of aglycone
Guaiacin B	3.8	911.4993	455.3507
Kalopanax-Saponin F	3.9	1087.5317	455.3509
Oleanolic acid-3- <i>O</i> -(triglucopyranosyl-1-3-arabinopyranosyl)-28-1-glucopyranosyl	3.8	1235.6107	455.3522
Calendulaglycoside A	3.9	1117.5490	455.3499
Araliaarmoside	4.0	1249.5869	455.3487
Calendulaglycoside C	4.3	955.4859	455.3511
Araloside B	4.4	1057.549	455.3517
Oleanolic acid-3- <i>O</i> -(diglucopyranosyl-1-3-arabinopyranosyl)-28-1-glucopyranosyl ester	4.3	1073.5569	455.3518
Oleanolic acid-3- <i>O</i> -(methyldioxy-trihexopyranosyl-1-3-pentopyranosyl)-28-1-hexopyranosyl ester	4.3	1119.5606	455.3521
Oleanolic acid-3- <i>O</i> -(hexosyl)-28-1-hexouronide ester	4.6	793.4312	455.3471
Araliasaponin III	4.5	1089.5493	455.3501
Araloside A	5.3	925.4805	455.3508
Oleanolic acid 3- <i>O</i> -hexuronide-(1-3-penta-furanoside)	5.7	763.4260	455.3502

Table S4. Triterpene saponins annotated in the extracts of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in SWATH and data-dependent acquisition (DDA) experiments

Nº	t _R (min)	m/z [M-H] ⁻ observed	m/z [M-H] ⁻ calculated	Elemental composition [M-H] ⁻	MS2 fragmentation patterns - product ions, m/z (rel. intensity)	Δm (ppm)	Assignment	Ref	Suppl. spectra
1	7.8	911.4993	911.501	C ₄₇ H ₇₅ O ₁₇ ⁻	455.3507 (20), 617.4041 (15), 749.4472 (15), 911.4993 (100)	1.9	Guaiacin B isomer 1	[30]	S1-1
2	7.9	1087.5317	1087.5331	C ₅₃ H ₈₃ O ₂₃ ⁻	455.3509 (10), 701.4265 (5), 925.4814 (15), 1087.5317 (100)	1.3	Kalopanax-Saponin F isomer 1	[31]	S1-2
3	8.0	1235.6107	1235.6066	C ₅₉ H ₉₅ O ₂₇ ⁻	455.3522 (50), 617.4059 (15), 749.4492 (100), 911.5035 (10), 1235.6107 (10)	-3.3	Oleanolic acid-3-O- (triglucopyranosyl-1- 3-arabinopyranosyl)- 28-1-glucopyranosyl		S1-3
4	8.2	1117.539	1117.5436	C ₅₄ H ₈₅ O ₂₄ ⁻	455.3499 (3), 731.4347 (5), 955.4898 (10), 1117.5390 (100)	4.1	Calendulaglycoside A	[15]	S1-4
5	8.4	1249.5869	1249.5859	C ₅₉ H ₉₃ O ₂₈ ⁻	455.3487 (5), 701.4254 (15), 925.4743 (7), 1057.5219 (80), 1087.5337 (50), 1153.5538 (100), 1249.5869 (70)	-0.8	Araliaarmoside	[8]	S1-5
6	8.4	1087.5336	1087.5331	C ₅₃ H ₈₃ O ₂₃ ⁻	455.3509 (10), 701.4265 (5), 925.4814 (15), 1087.55336 (100)	-0.5	Kalopanax-Saponin F isomer 2	[31]	S1-2
7	8.6	955.4859	955.4908	C ₄₈ H ₇₅ O ₁₉ ⁻	455.3511 (5), 569.379 (5), 793.4317 (15), 955.4859 (100)	5.1	Calendulaglycoside C isomer 1	[15]	S1-6
8	8.7	1057.5249	1057.5225	C ₅₂ H ₈₁ O ₂₂ ⁻	455.3517 (5), 701.4286 (7), 763.4309 (5), 895.4698 (7), 1057.5249 (100)	-2.3	Araloside B isomer 2	[11]	S1-7
9	8.9	1073.5569	1073.5538	C ₅₃ H ₈₅ O ₂₂ ⁻	455.3518 (50), 617.4052 (20), 749.4485 (30), 911.5022 (100), 1073.5569 (15)	-2.9	Oleanolic acid-3-O- (diglucopyranosyl-1- 3-arabinopyranosyl)- 28-1-glucopyranosyl ester		S1-8
10	8.9	1119.5606	1119.5616	C ₅₄ H ₈₇ O ₂₄ ⁻	455.3521 (5), 617.4057 (5), 749.4490 (7), 911.5019 (100), 1073.5551 (15), 1119.5606 (13)	0.9	Oleanolic acid-3-O- (ethyl- trihexopyranosyl-1- 3-pentopyranosyl)-		S1-9

							28-1-hexopyranosyl ester		
11	9.0	793.4312	793.438	C ₄₂ H ₆₅ O ₁₄ ⁻	455.3471 (10), 631.3785 (15), 793.4312 (100)	8.6	Oleanolic acid-3-O-(hexosyl)-28-1-hexouronide ester isomer 1		S1-10
12	9.3	1089.5493	1089.5487	C ₅₃ H ₈₅ O ₂₃ ⁻	455.3501 (5), 719.4360 (10), 881.4901 (100), 1043.5437 (12), 1089.5493 (12)	-0.6	Araliasaponin III	[6]	S1-11
13	10.1	955.4899	955.4908	C ₄₈ H ₇₅ O ₁₉ ⁻	455.3511 (5), 569.379 (5), 793.4317 (15), 955.4899 (100)	0.9	Calendulaglycoside C isomer 2	[15]	S1-6
14	10.3	925.4805	925.4802	C ₄₇ H ₇₃ O ₁₈ ⁻	455.3508 (2), 569.3831 (5), 731.4366 (20), 925.4803 (100)	0.3	Araloside A isomer 1	[11]	S1-12
15	10.4	925.4807	925.4802	C ₄₇ H ₇₃ O ₁₈ ⁻	455.3508 (2), 569.3831 (5), 731.4366 (20), 925.4803 (100)	0.5	Araloside A isomer 2	[11]	S1-12
16	10.8	895.4696	-	-	455.3508 (5), 551.3731 (5), 895.4696 (100)	-	Oleanolic acid unknown derivatives		S1-13
17	10.8	793.436	793.438	C ₄₂ H ₆₅ O ₁₄ ⁻	455.3471 (10), 631.3785 (15), 793.4312 (100)	2.5	Oleanolic acid-3-O-(hexosyl)-28-1-hexouronide ester isomer 2		S1-10
18	10.9	763.426	763.4274	C ₄₁ H ₆₃ O ₁₃ ⁻	455.3502 (3), 631.3822 (5), 763.4260 (100)	1.8	Oleanolic acid 3-O-hexuronide-(1-3-penta-furanoside)		S1-14
19	11.0	911.5016	911.501	C ₄₇ H ₇₅ O ₁₇ ⁻	455.3507 (20), 617.4041 (15), 749.4472 (15), 911.5016 (100)	-0.7	Guaiacin B isomer 2	[30]	S1-1
20	11.0	911.4949	911.501	C ₄₇ H ₇₅ O ₁₇ ⁻	455.3507 (20), 617.4041 (15), 749.4472 (15), 911.4949 (100)	6.7	Guaiacin B isomer 3	[30]	S1-1

The extracts were prepared with conventional (water or ethanol) or natural deep eutectic solvents (NADES) and analyzed by a Waters ACQUITY UPLC I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled online to a hybrid quadrupole-time of flight mass spectrometer (QqTOF-MS) AB Sciex TripleTOF 6600 (AB Sciex, Darmstadt, Germany)

Table S5. The conditions of reverse phase-ultra high performance liquid chromatographic (RP-UHPLC) separation and the settings for electrospray ionization-quadrupole-time of flight mass spectrometry (ESI-QqTOF-MS) applied for the profiling (untargeted analysis) of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen root semi-polar secondary metabolites with QqTOF hybride mass spectrometer SHIMADZU LCMS-9030 System (SHIMADZU Corporation, Kyoto, Japan).

Chromatography	
SIL-30AC Autosampler	
Injection mode	Partial Loop
Injection volume	5 μ L
Wash solvent	50% MeOH
Wash solvent volume	300 μ L
Cooler temperature	4.0 C
Rinse type	Internal & external
Rinse mode	Before and after aspiration
Needle overfill flush	Rinse port + rinse pump
Rinse time	2 sec
Column conditions	
Separation column	Phenomenex Kinetex C18 Column (100 x 2.1 mm, particle size 1.7 μ m)
Column oven temperature	40.0 C
LC separation parameters	
Eluent A	aqueous 0.1% (v/v) FA
Eluent B	0.1% (v/v) FA in acetonitrile
Flow rate	0.5 mL/min

Elution program	gradient to 100% eluent B – 10 min
	100% eluent B isocratic – 2 min
	gradient to 5% eluent B – 0.1 min
	5% eluent B isocratic – 0.9 min (re-equilibration)

Mass spectrometry

General

Mass analyzer type	quadrupole-time of flight (QqTOF-MS)
Ionsource	ESI
Experiment type	Sequential Windowed Acquisition of All Theoretical Fragment Ion Mass Spectra (DIA)
Operatinon mode	positive, negative
Cycle time (s)	0.996
Duration	10.0 min

Ion source settings

Nebulizer gas (L/min)	3
Drying gas (L/min)	10
Ion spray voltage (kV)	4.0/-3.0 (negative mode)
Ion source temperature (°C)	180

MS settings

Experiment type	TOF-MS
<i>m/z</i> range	240 - 1350
Accumulation time (ms)	100
ID function	ON

MS/MS Setting

Experiment type	SWATH
Collision gas	Ar
MS/MS experiment type	DIA
SWATH window number	28 u (650-1350 <i>m/z</i>)
SWATH window width (<i>m/z</i>)	25.0
CE	18-52 (35 ± 17 V)
Accumulation time (ms)	21
ID function	OFF
Collision potential (V)	35/-35 (negative mode)
Collision energy spread (V)	17 (negative mode)
SWATH, sequential window acquisition of all theoretical fragment ion spectra	

Table S6. The conditions of reversed phase-ultrahigh performance liquid chromatographic (RP-UHPLC) separation and the settings for electrospray ionization-quadrupole-time of flight mass spectrometry (ESI-QqTOF-MS) applied for the SWATH and DDA MS/MS analysis of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen root semi-polar secondary metabolites with Waters ACQUITY UPLC I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled online to a hybrid quadrupole-time of flight mass spectrometer (QqTOF-MS) AB Sciex TripleTOF 6600 (AB Sciex, Darmstadt, Germany).

Chromatography	
ACQUITY Sample Manager (SM)	
Injection mode	Partial Loop
Injection volume	5 μ L
Weak wash solvent	0.3 mmol/L aq. ammonium formate, pH 3.5 (adjusted using formic acid)
Weak wash volume	800 μ L
Strong wash solvent	acetonitrile
Strong wash volume	400 μ L
Target sample temperature	4.0 C
Needle overfill flush	automatic
Column conditions	
Separation column	EC 150/2 NUCLEOSHELL RP 18 (150 x 2 mm, particle size 2.7 μ m)
Target column temperature	40.0 C
ACQUITY Binary Solvent Manager (BSM)	
Eluent A	0.3 mmol/L aq. ammonium formate, pH 3.5 (adjusted using formic acid)
Eluent B	acetonitrile

Seal wash duration	5 min
Flow rate	0.4 mL/min
Elution program	5% eluent B isocratic - 2 min gradient to 95% eluent B – 17 min 95% eluent B isocratic – 2 min gradient to 5% eluent B – 0.1 min 5% eluent B isocratic – 3 min (re-equilibration)

Mass spectrometry

General

Mass analyzer type	quadrupole-time of flight (QqTOF-MS)
Ionsource	DuoSpray™ ion source
Experiment type	Sequential Windowed Acquisition of All Theoretical Fragment Ion Mass Spectra (SWATH)
Operatinon mode	negative
Cycle time (s)	1.1
Pause between ranges (ms)	1.049
Auto adjust with mass	on
Settling time (s)	0
Time bins to sum	4
Duration	23 min

Ion source settings

Nebulizer gas (psig)	60
Drying gas (psig)	70
Curtain gas (psig)	55

Ion spray voltage (kV)	-4.5
Ion source temperature (°C)	450
MS settings	
Experiment type	TOF-MS
<i>m/z</i> range	65 - 1250
Accumulation time (ms)	50/75 (SWATH/DDA)
Declustering potential (V)	-35
Collision potential (V)	-10
MS/MS Setting	
Fragmentation mode	collision-activated dissociation (CAD)
MS/MS experiment type	DDA
For ions greater than	100 Da
Charge state	1-2
With intensity greater than	3000
Maximum number of candidate ions to monitor per cycle	5
Exclude former target ions for	4 sec
Fragment intensity multiplier	2
Maximum accumulation	2 sec
Analyte type	small molecules
Accumulation time (ms)	175
Declustering potential (V)	-35
Collision potential (V)	-45
Collision energy spread (V)	35
Ion release delay (V)	-30

Ion release width (V)	-15
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MS/MS Setting	
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Fragmentation mode	collision-activated dissociation (CAD)
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MS/MS experiment type	SWATH
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SWATH window number	15
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SWATH window width (<i>m/z</i>)	80
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SWATH window overlap (<i>m/z</i>)	1
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Rolling collision energy	off
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Analyte type	small molecules
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Accumulation time (ms)	60
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Declustering potential (V)	-35
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Collision potential (V)	-45
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Collision energy spread (V)	35
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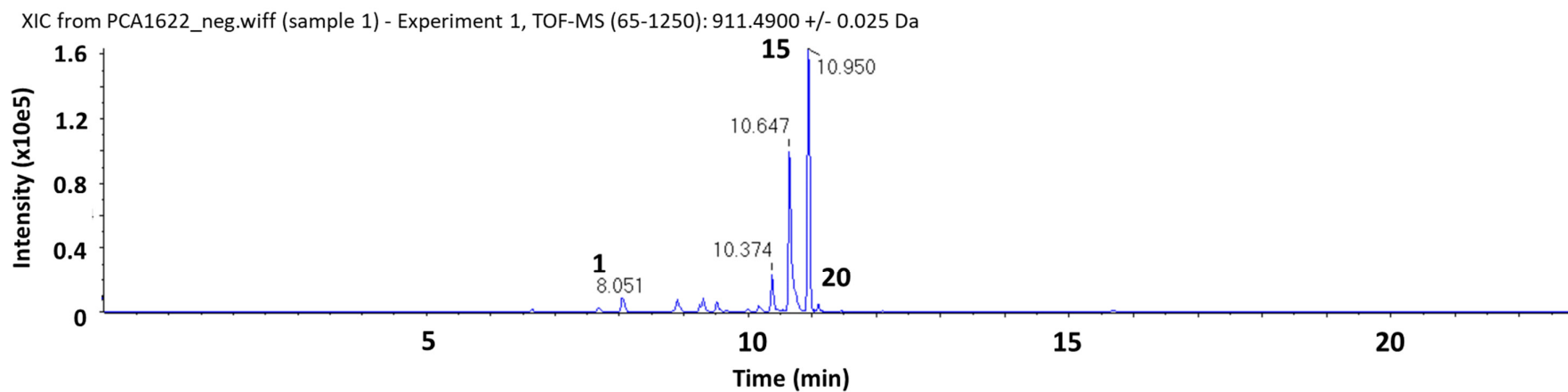
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Ion release width (V)	-15
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Figures

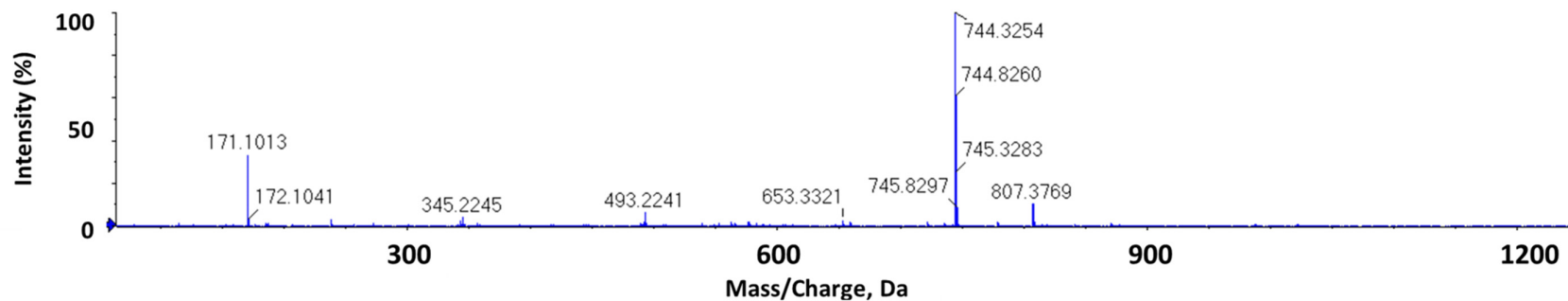
Chromatographic and mass spectral data of triterpene saponins annotated in the extracts of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in SWATH and data-dependent acquisition (DDA) experiments

A

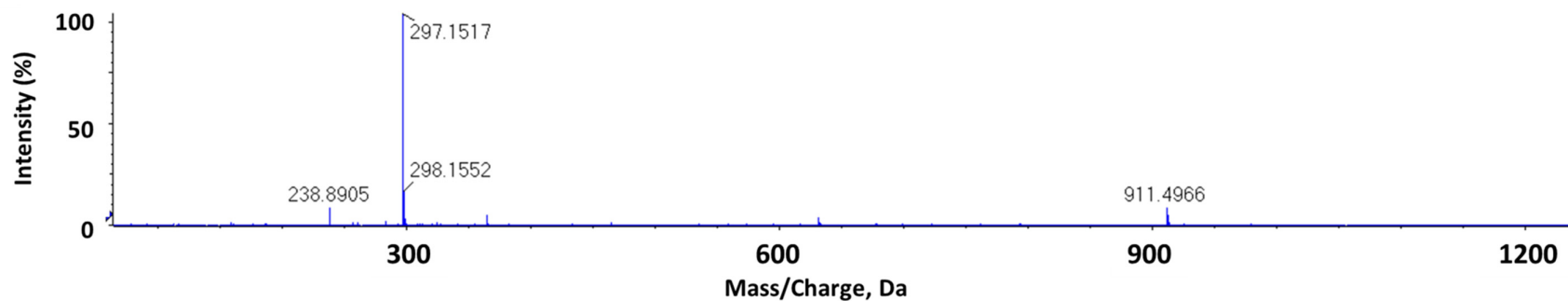


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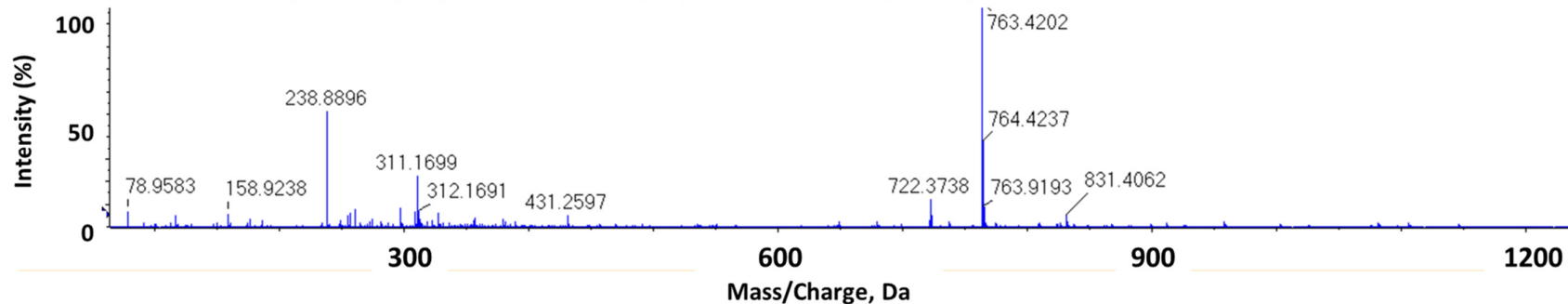
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Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.950 min



Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 11.093 min



C

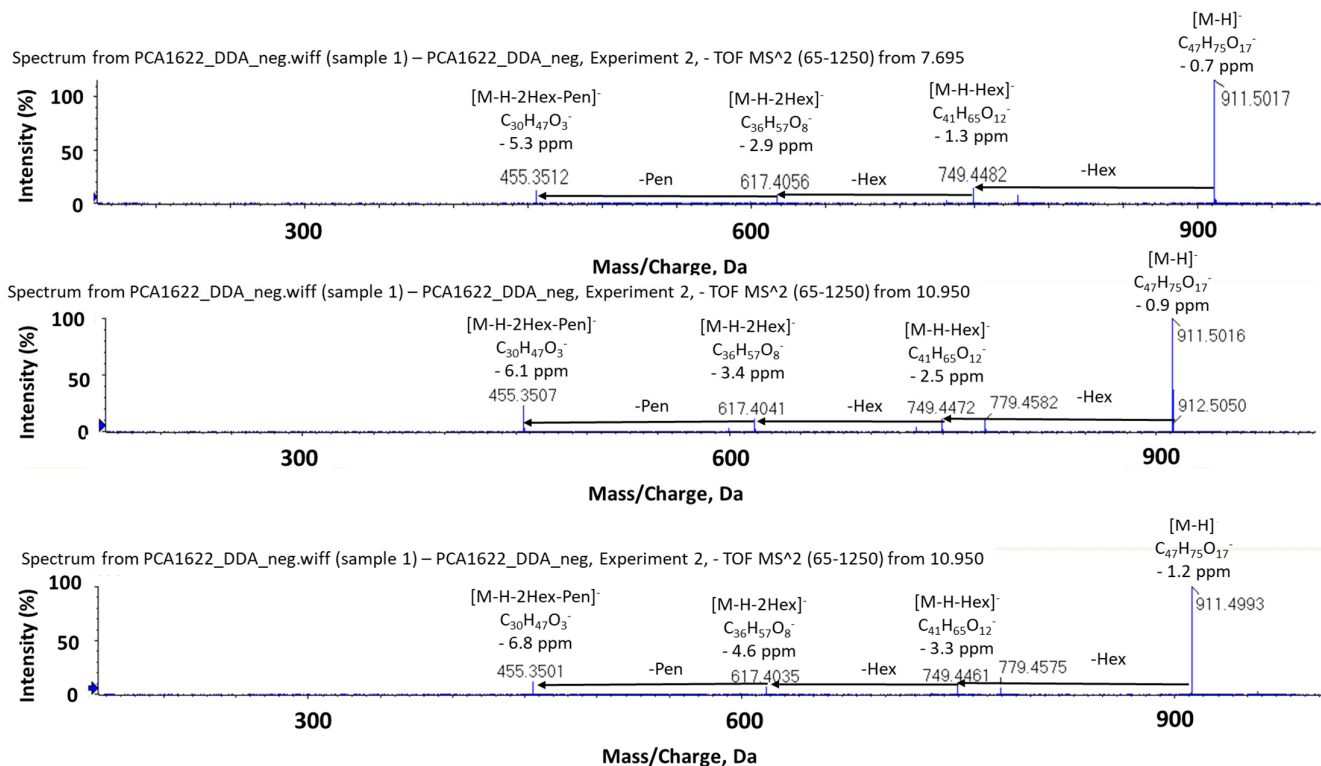
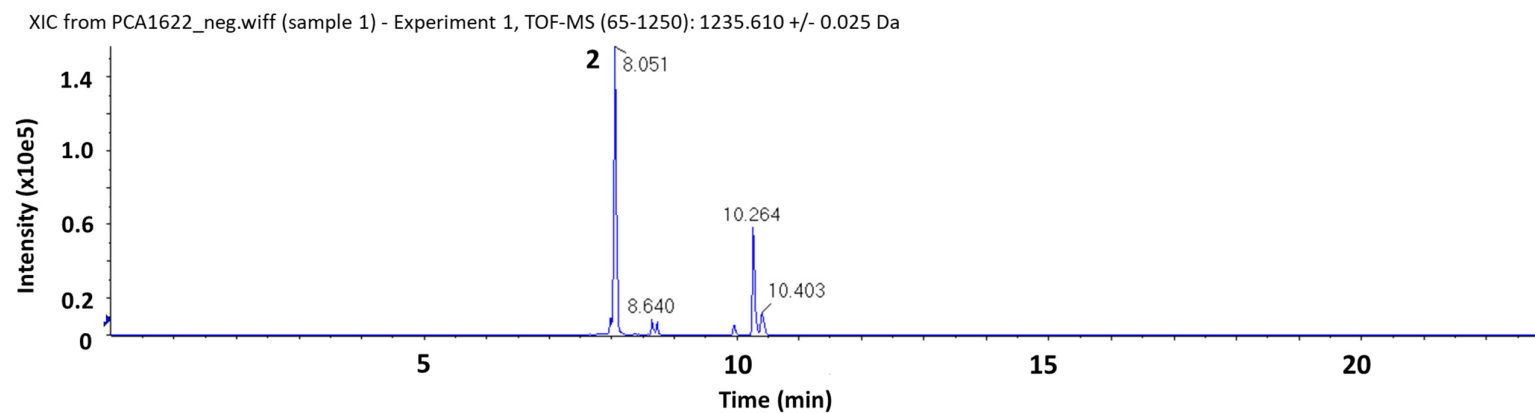
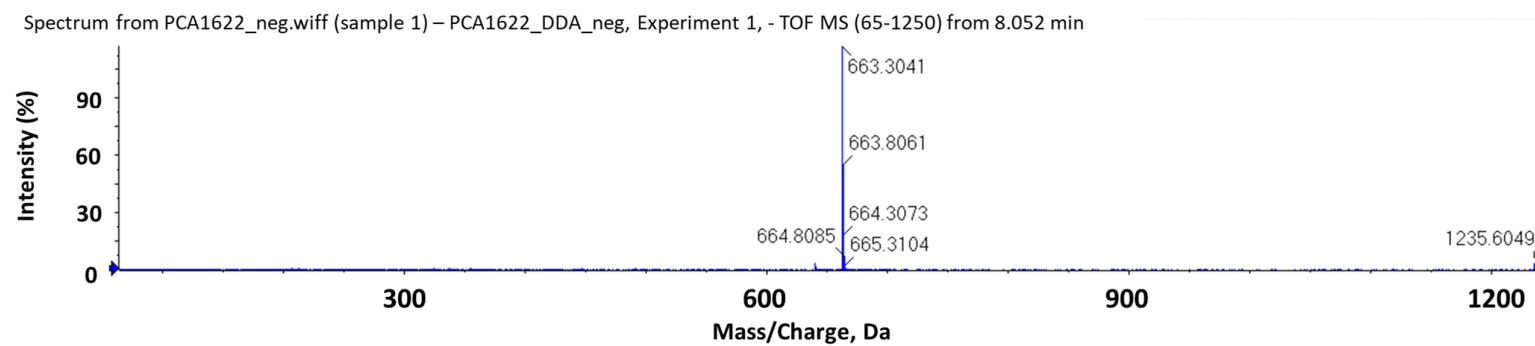


Figure S1. Extracted ion chromatogram (XIC) of m/z 911.5000 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 1, 15, 20 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as guaiacin B isomers at t_R 7.77, 10.98 and 11.01, respectively. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted [M-H]⁻ ions annotated at the MS1 level).

A**B**

C

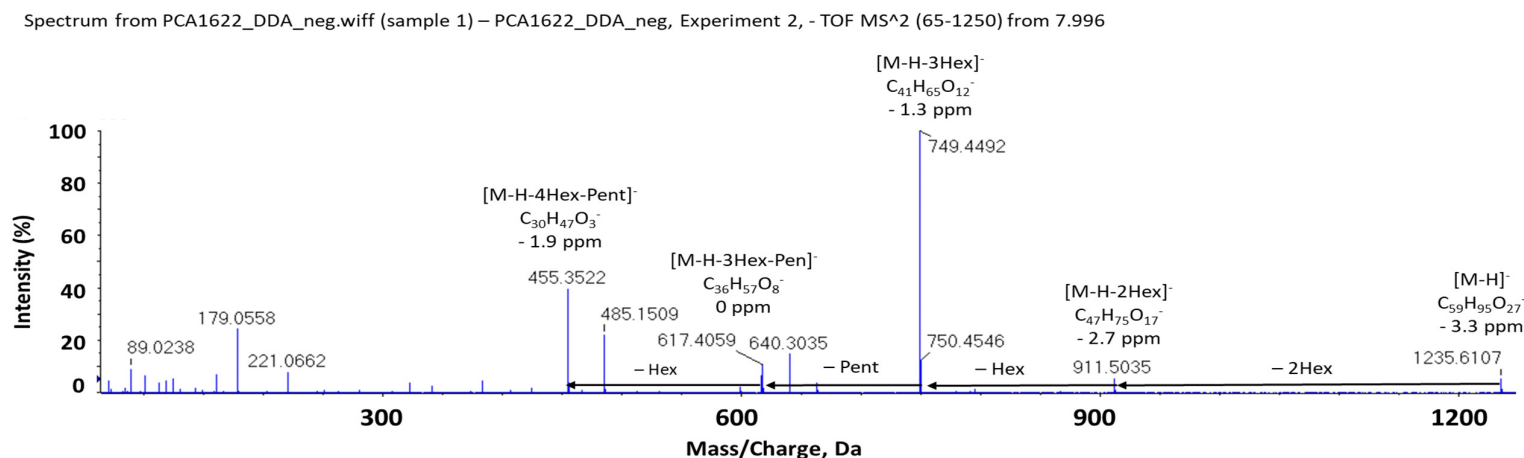
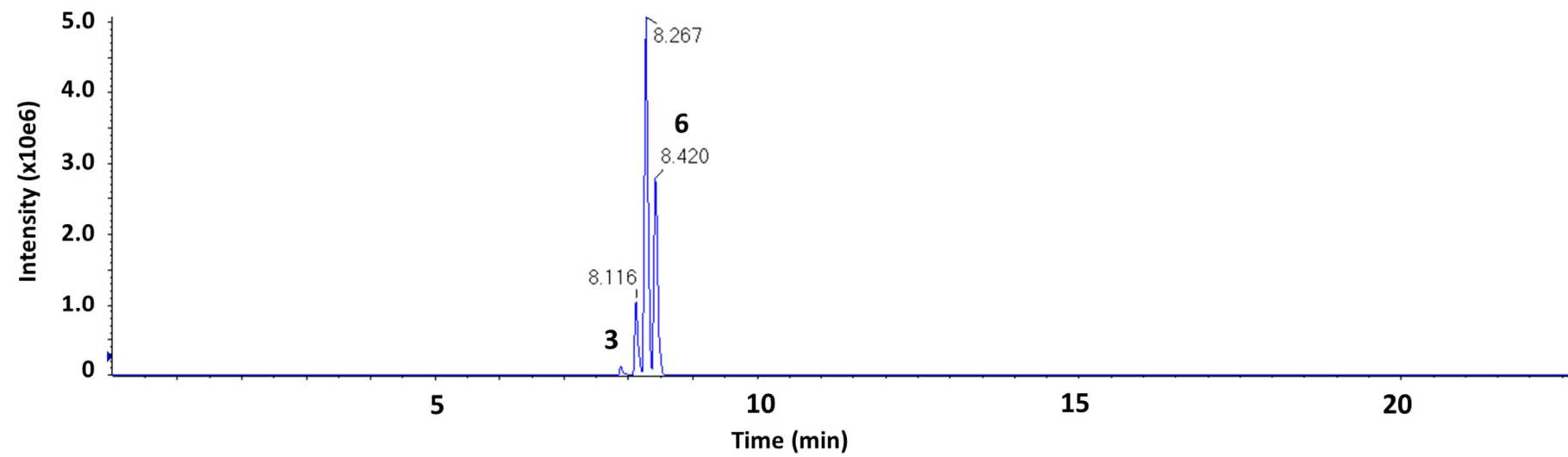


Figure S2. Extracted ion chromatogram m/z 1235.6100 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 2 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid-3-*O*-(triglucopyranosyl-1-3-arabinopyranosyl)-28-1-glucopyranosyl. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

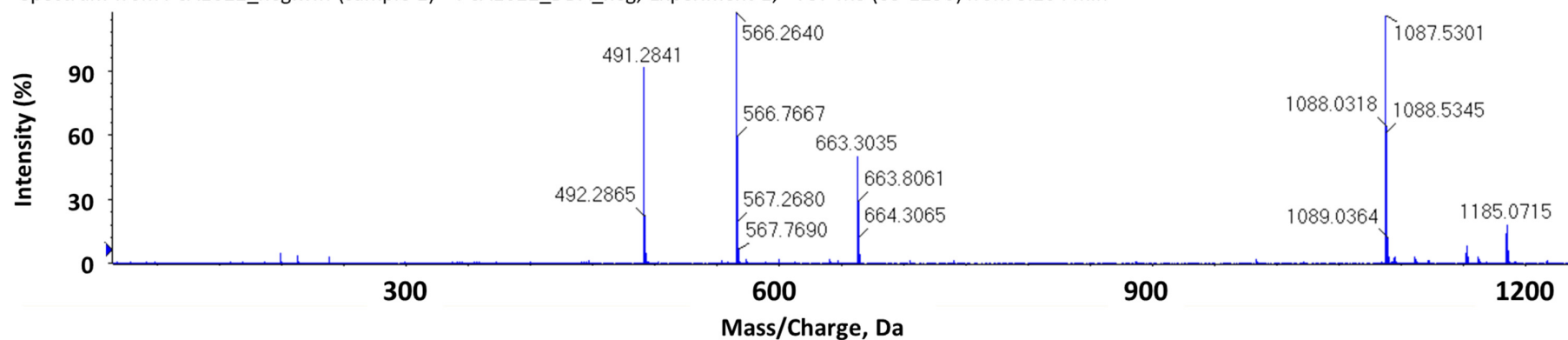
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1087.5400 +/- 0.025 Da

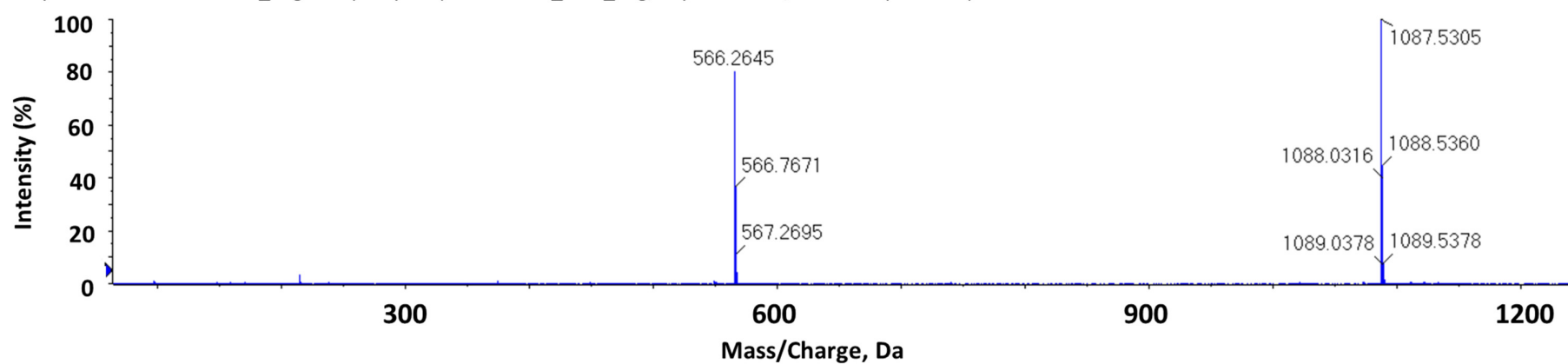


B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.104 min



Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.425 min



C

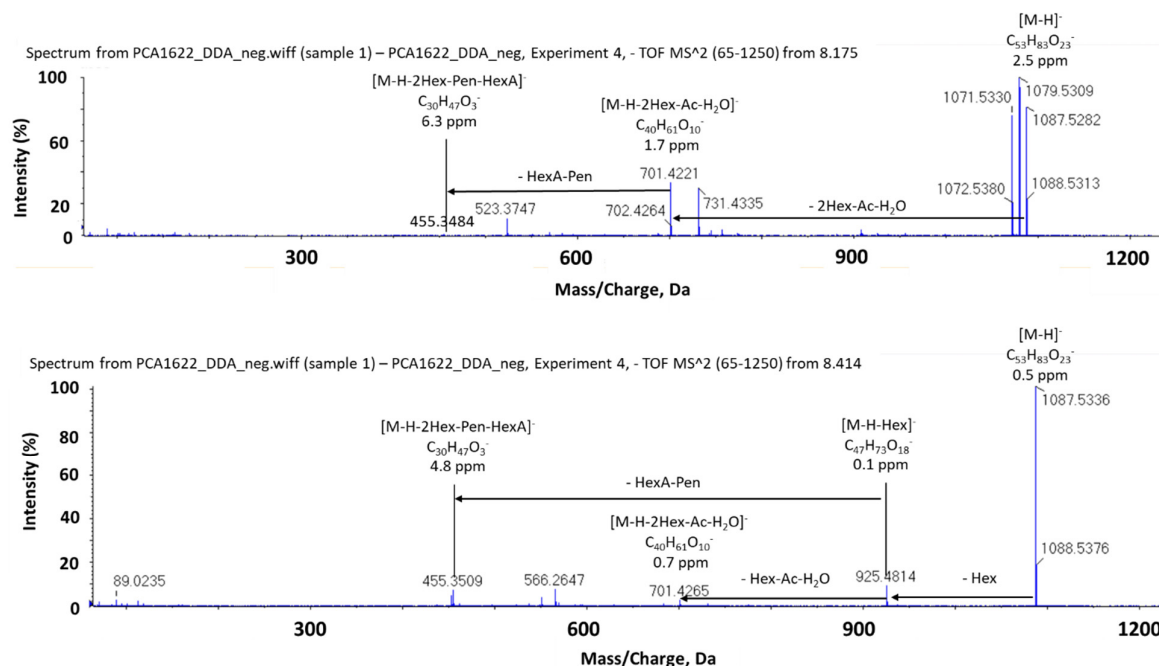
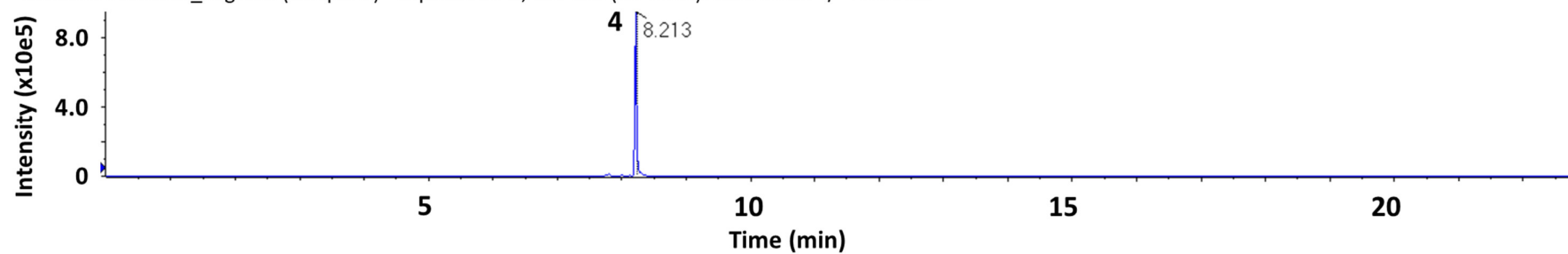


Figure S3. Extracted ion chromatogram m/z 1087.5400 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 3, 6 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as kalopanax-saponin F isomers at t_R 7.89 and 8.41, respectively. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

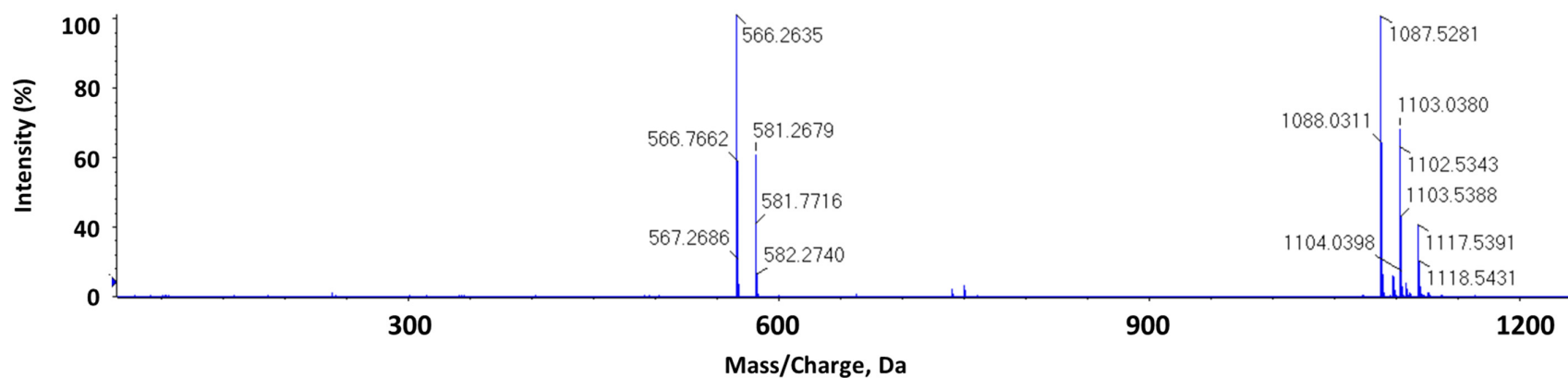
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1117.5400 +/- 0.025 Da



B

Spectrum from PCA1622_neg.wiff (sample 1) - PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.232 min



C

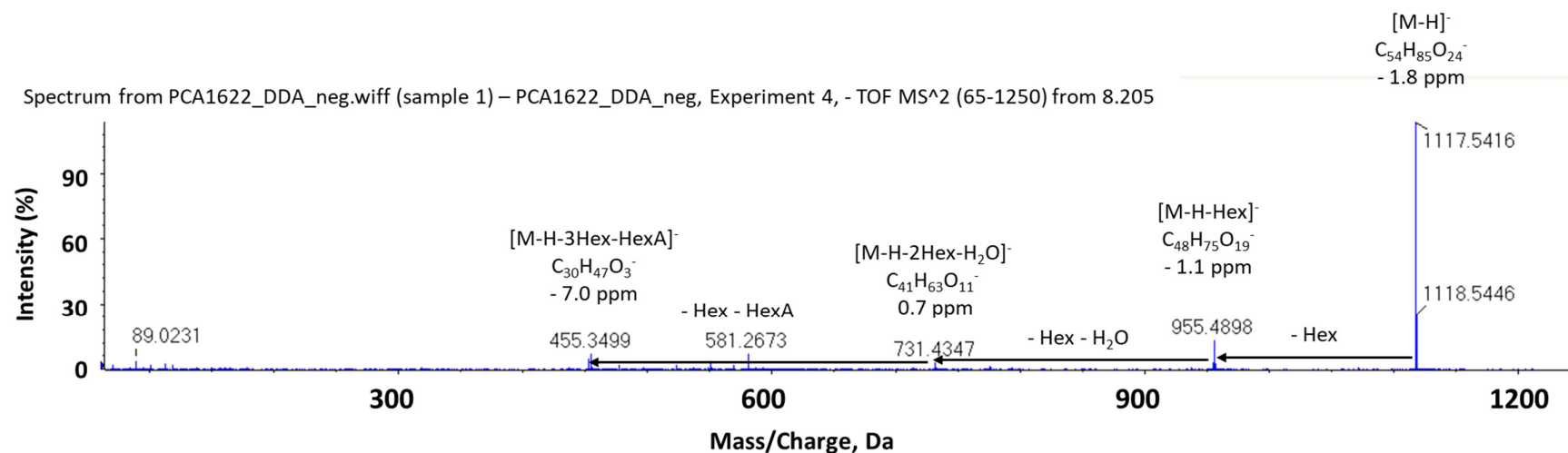
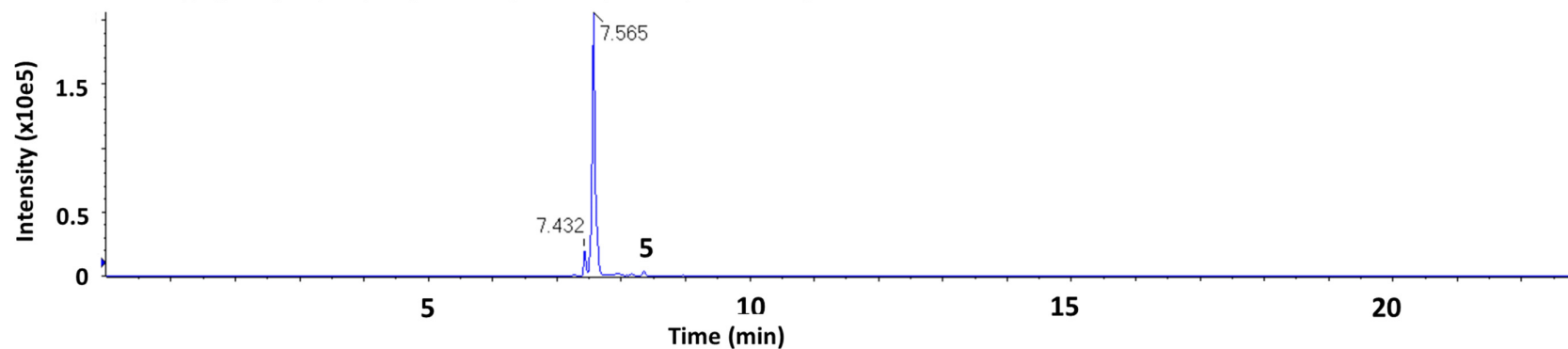


Figure S4. Extracted ion chromatogram m/z 1117.5400 ± 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 4 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as calendulaglycoside A. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted [M-H]⁻ ions annotated at the MS1 level).

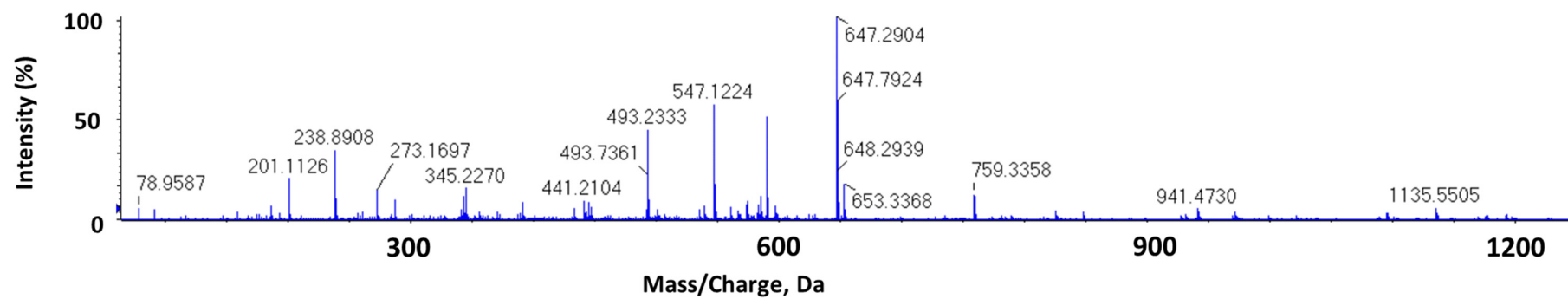
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1249.580 +/- 0.025 Da



B

Spectrum from PCA1622_neg.wiff (sample 1) - PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 7.552 min



C

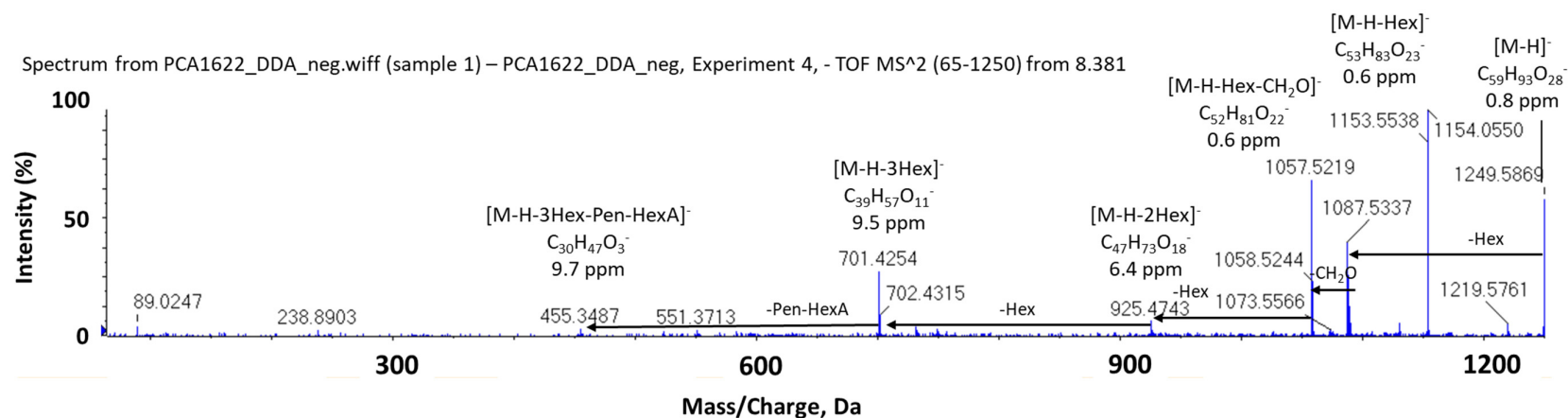
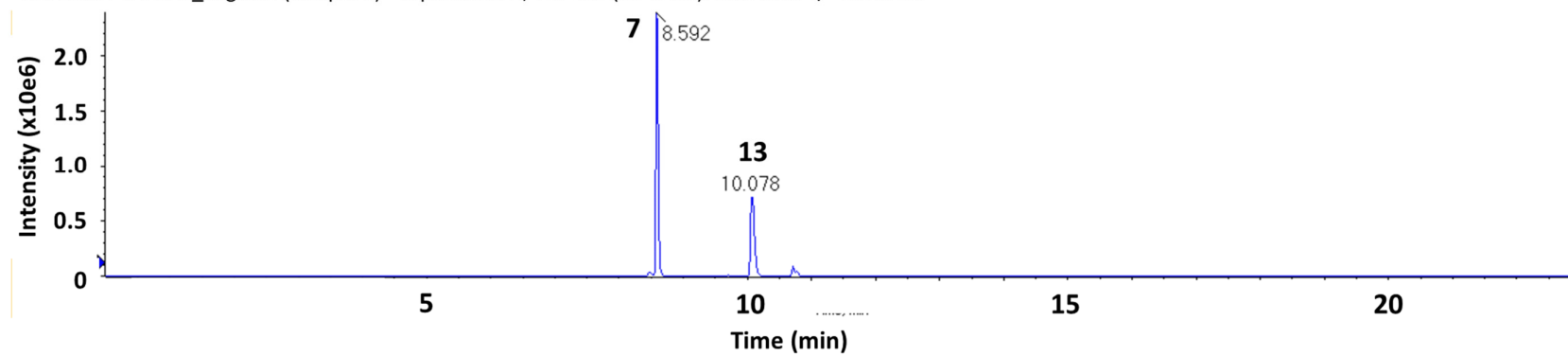


Figure S5. Extracted ion chromatogram m/z 1249.5900 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 5 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as araliaarmoside. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

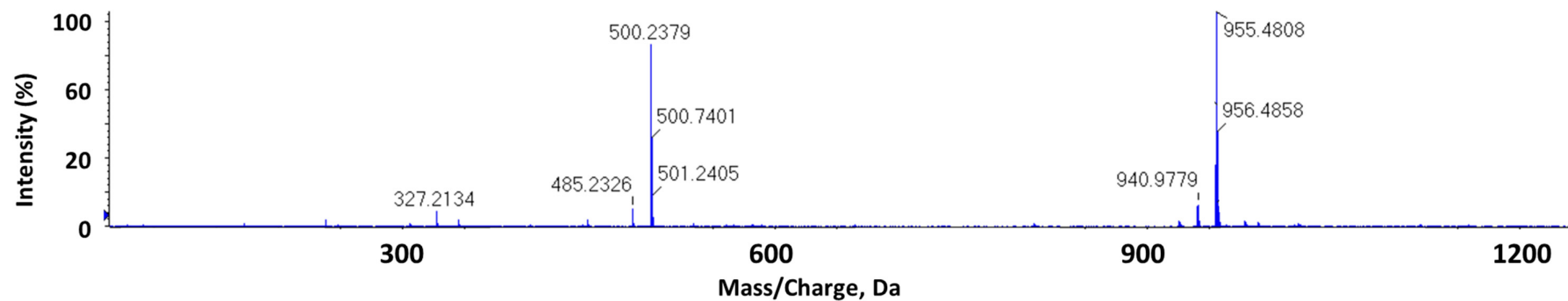
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 955.4800 +/- 0.025 Da

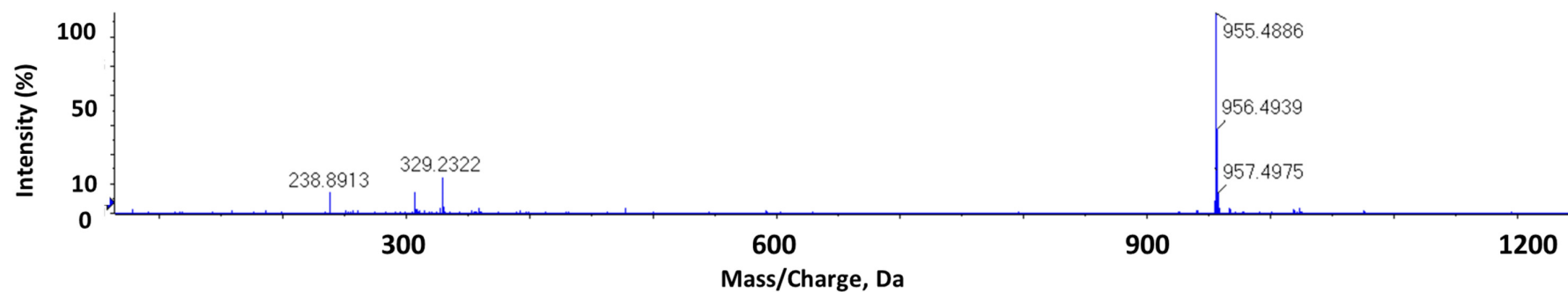


B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.592 min



Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.091 min



C

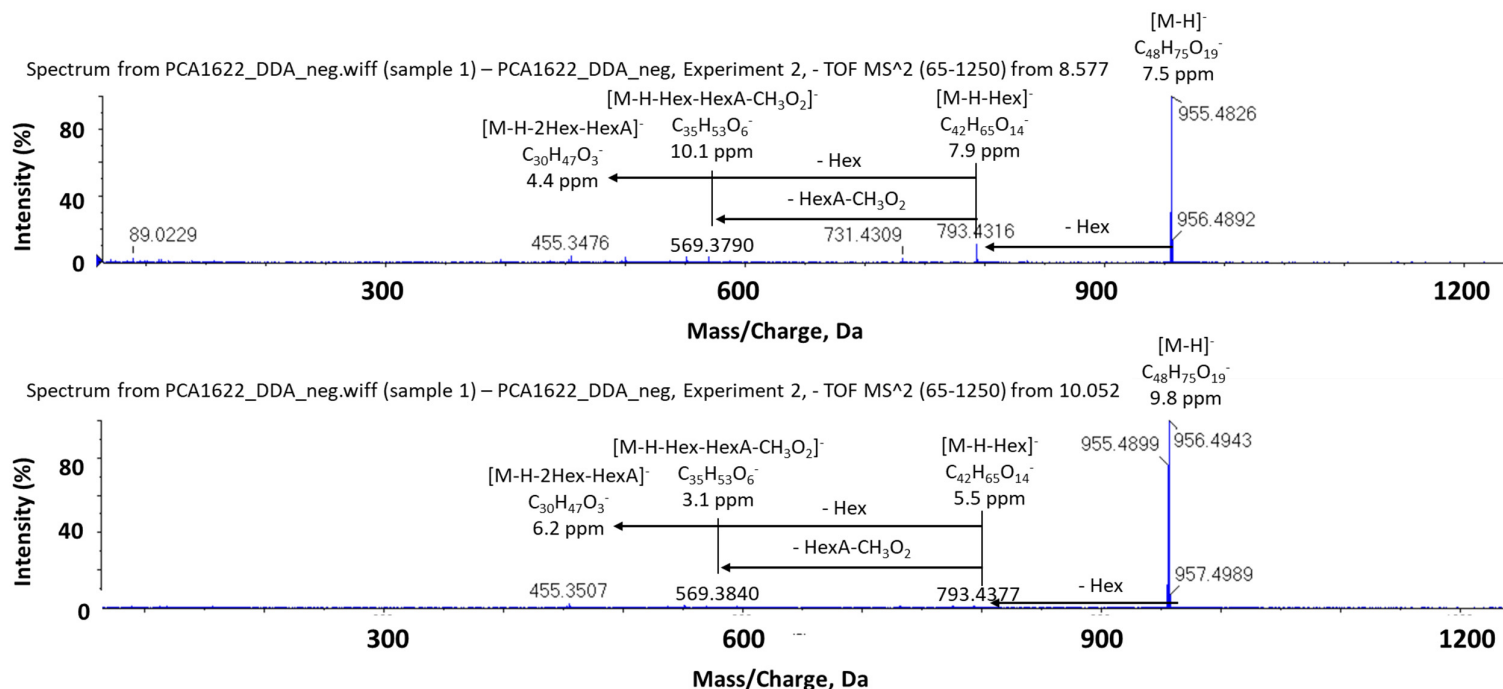
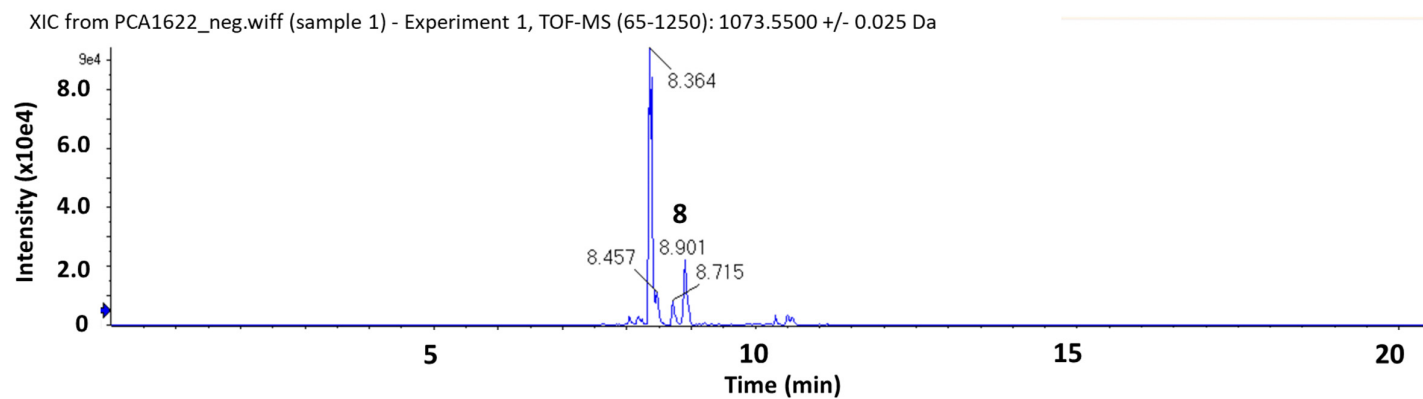
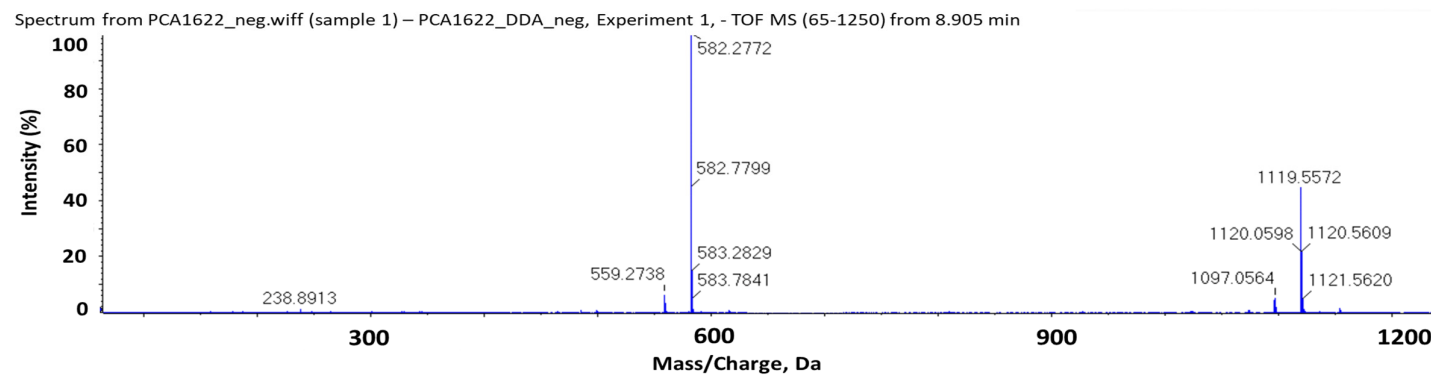


Figure S6. Extracted ion chromatogram m/z 955.4900 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 7, 13 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as calendulaglycoside C isomers at t_R 8.59 and 10.07, respectively. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

A



B



C

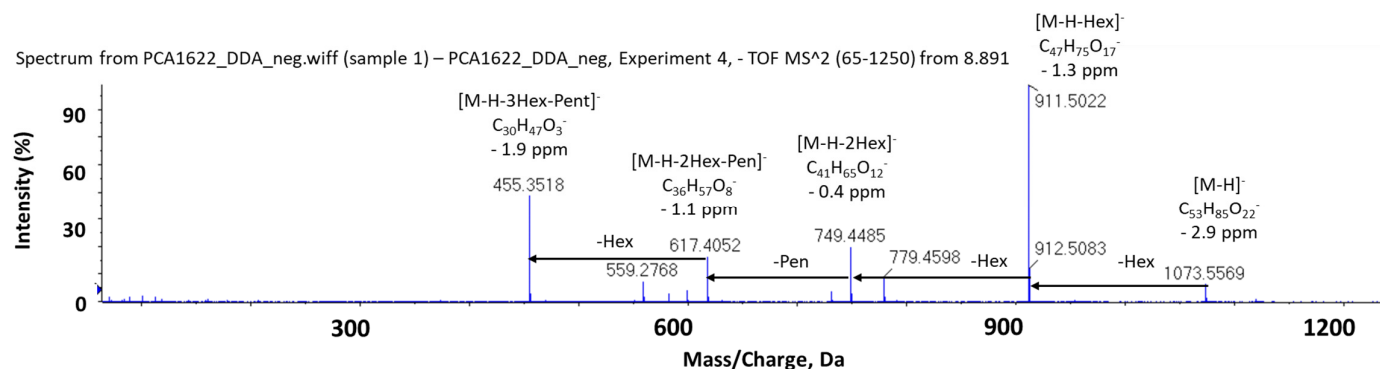
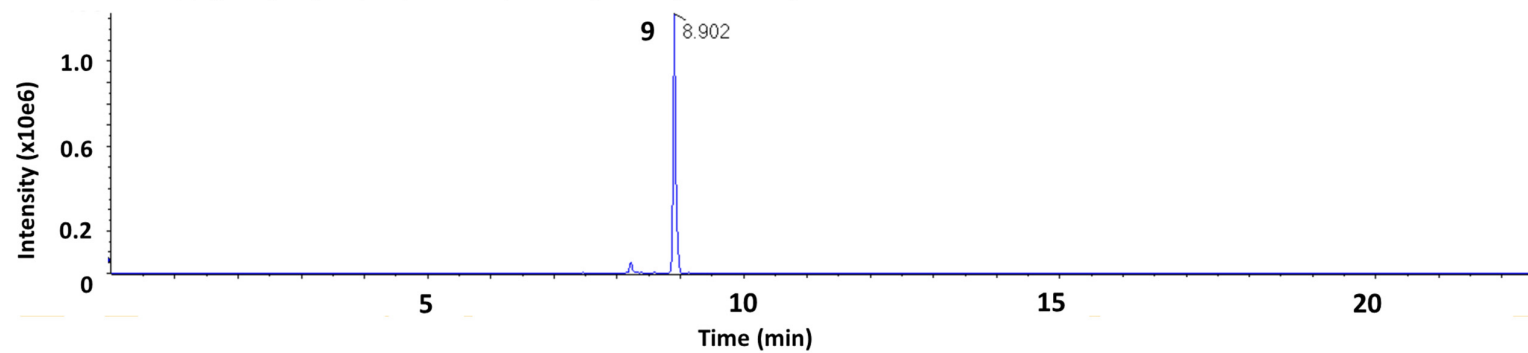


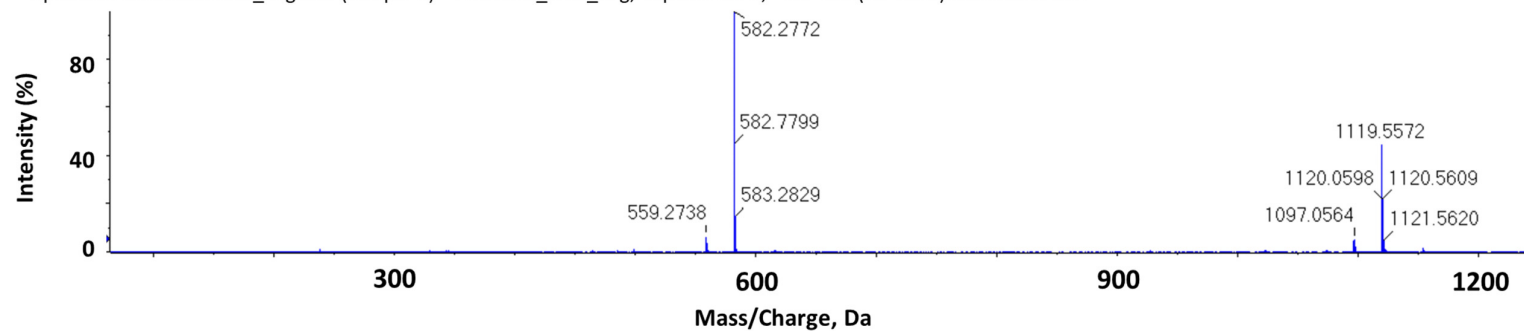
Figure S7. Extracted ion chromatogram m/z 1073.5600 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 8 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid-3-*O*-(diglucopyranosyl-1-3-arabinopyranosyl)-28-*O*-glucopyranosyl ester. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted [M-H]⁻ ions annotated at the MS1 level).

A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1119.550 +/- 0.025 Da

**B**

Spectrum from PCA1622_neg.wiff (sample 1) - PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.710 min



C

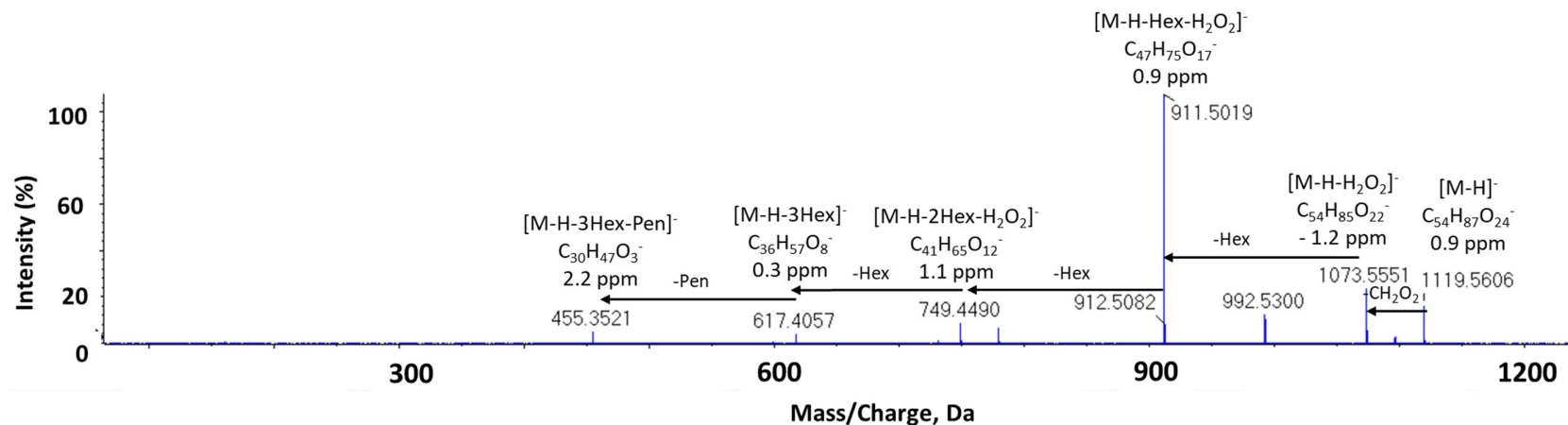
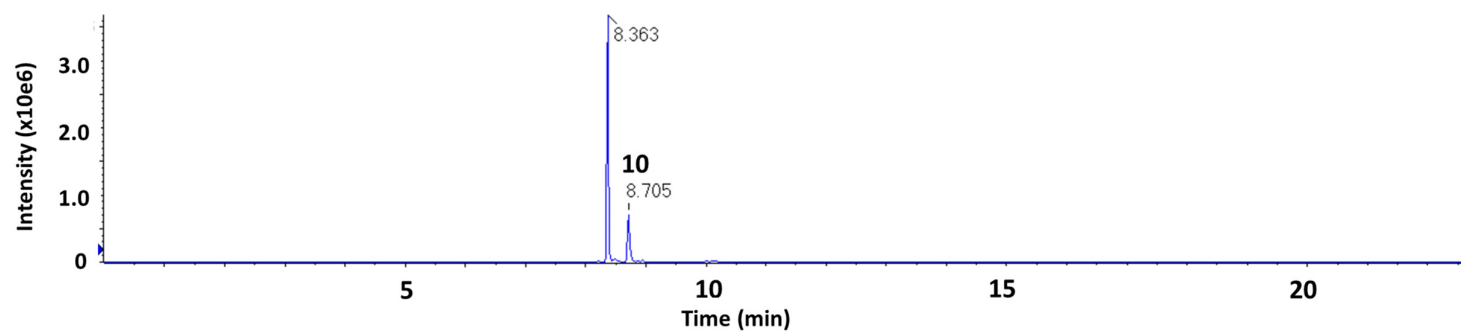
Spectrum from PCA1622_DDA_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 4, - TOF MS² (65-1250) from 8.719

Figure S8. Extracted ion chromatogram m/z 1119.5600 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 9 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid-3-*O*-(methyldioxy-trihexopyranosyl-1-3-pentopyranosyl)-28-1-hexopyranosyl ester. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

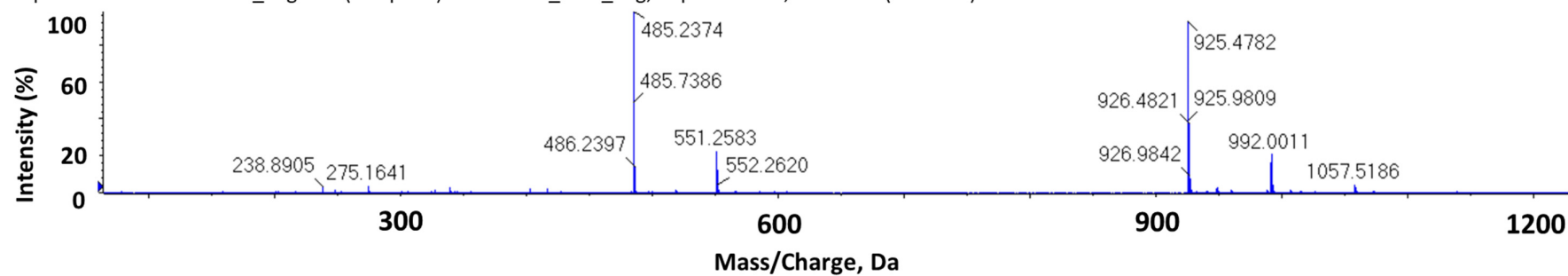
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1057.5200 +/- 0.025 Da



B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 8.710 min



C

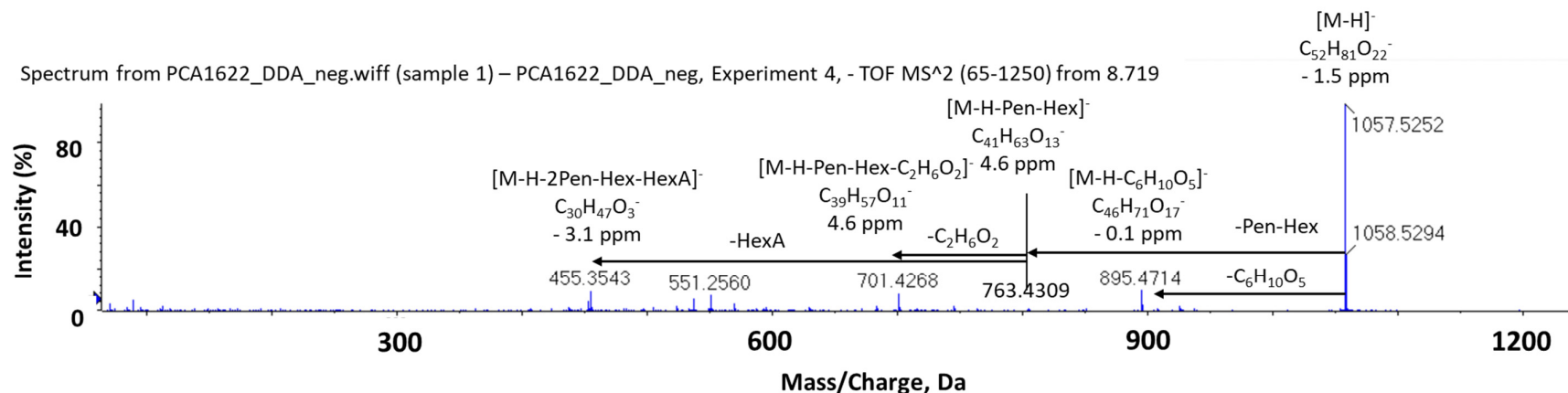
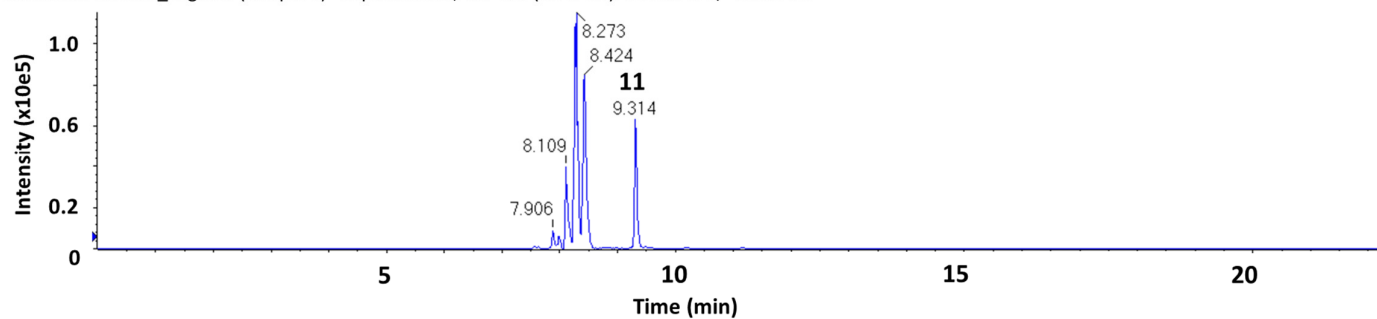


Figure S9. Extracted ion chromatogram m/z 1057.5300 ± 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 10 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as araloside B. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

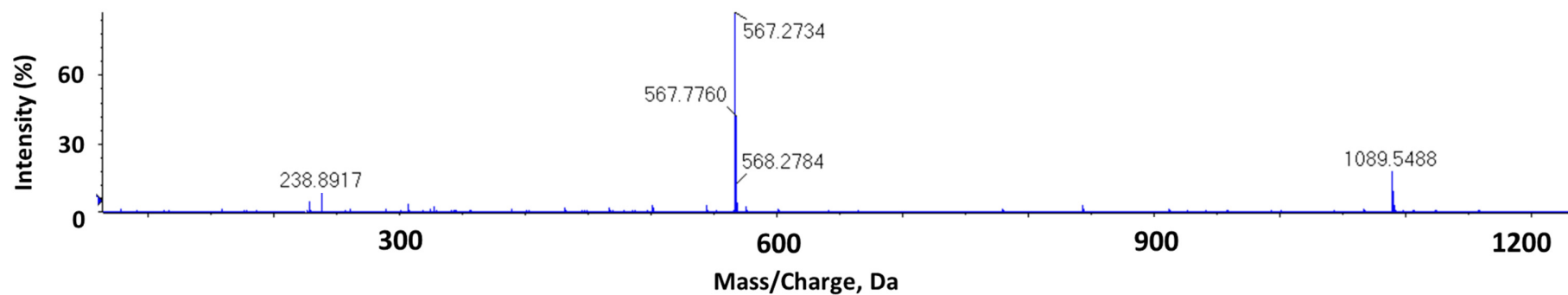
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 1089.540 +/- 0.025 Da



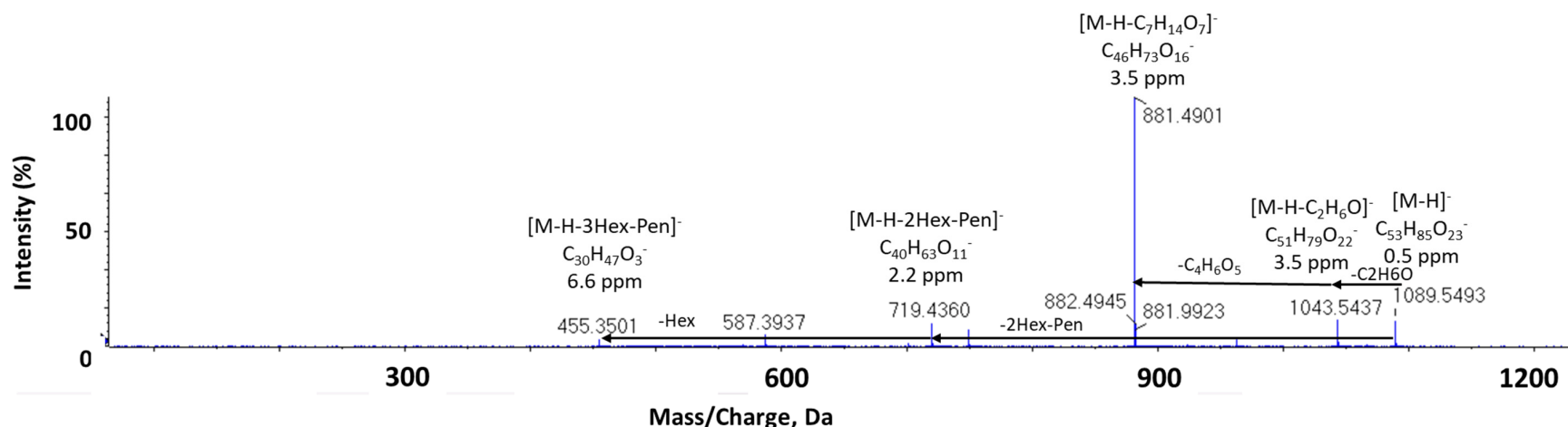
B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 9.307 min



C

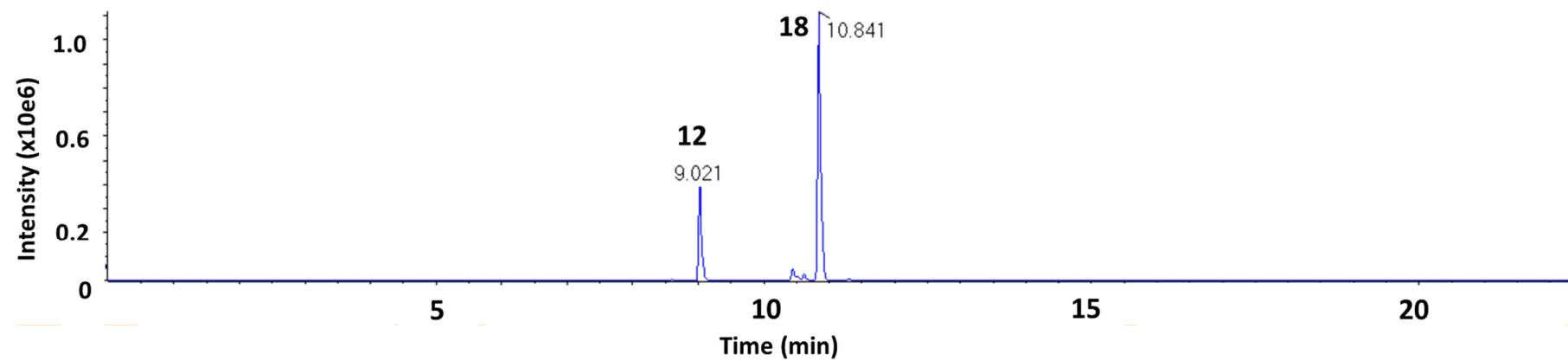
Spectrum from PCA1622_DDA_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 4, - TOF MS² (65-1250) from 9.317



Figures S10. Extracted ion chromatogram m/z 1089.5500 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 11 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as araliasaponin III. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

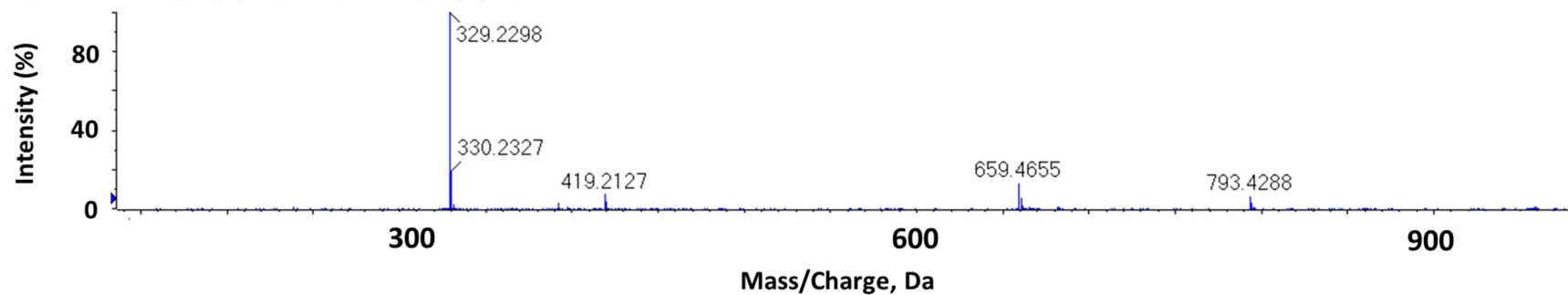
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 793.430 +/- 0.025 Da

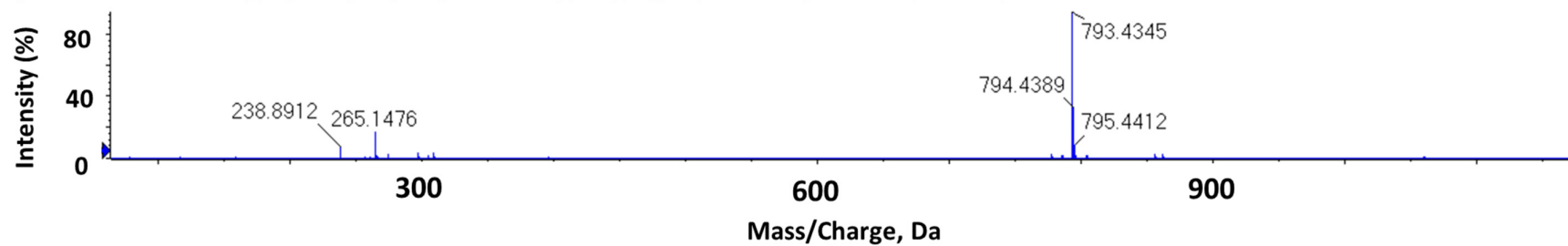


B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 9.021 min



Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.838 min



C

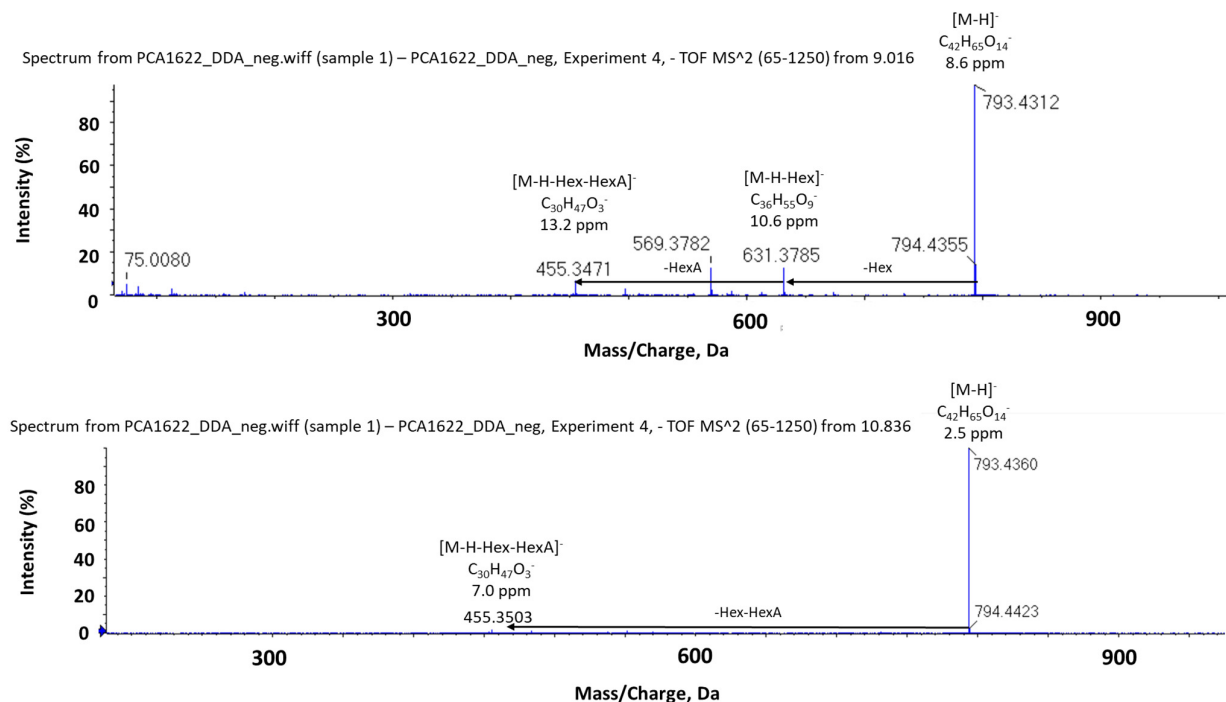
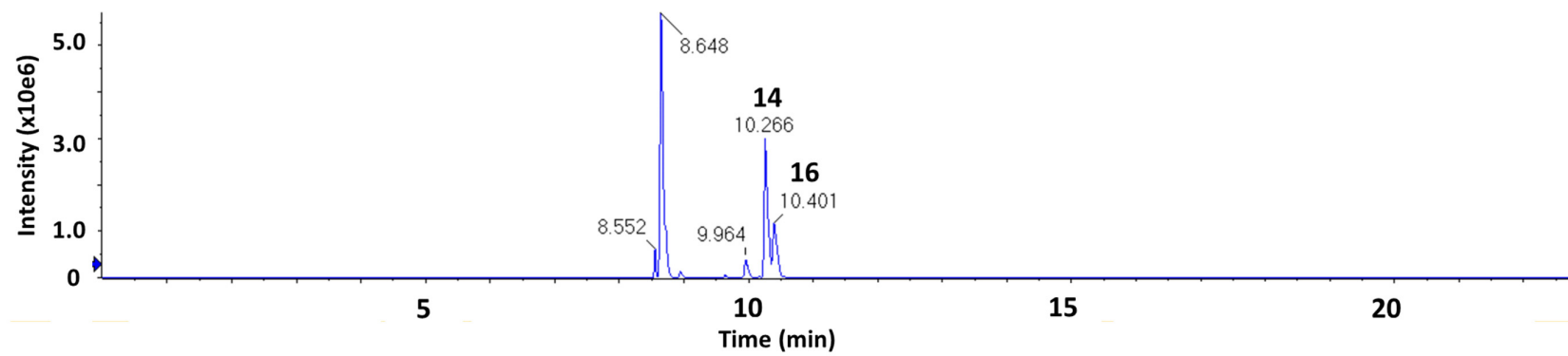


Figure S11. Extracted ion chromatogram m/z 793.4400 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 12, 18 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid-3-*O*-(hexosyl)-28-1-hexouronide ester isomers t_R 9.02 and 10.84, respectively. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

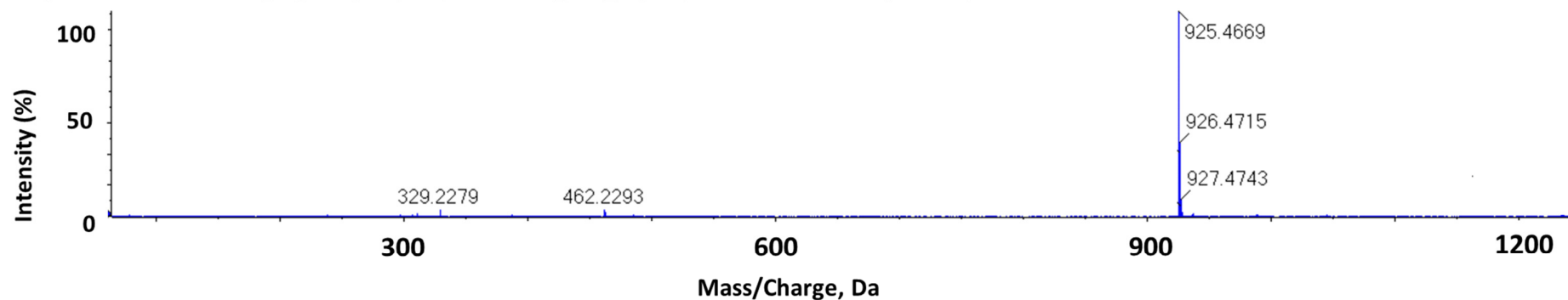
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 925.4700 +/- 0.025 Da

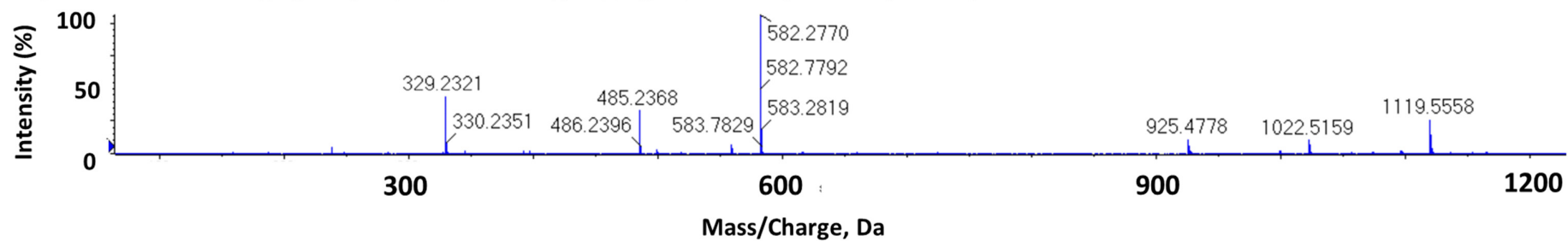


B

Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.266 min



Spectrum from PCA1622_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.401 min



C

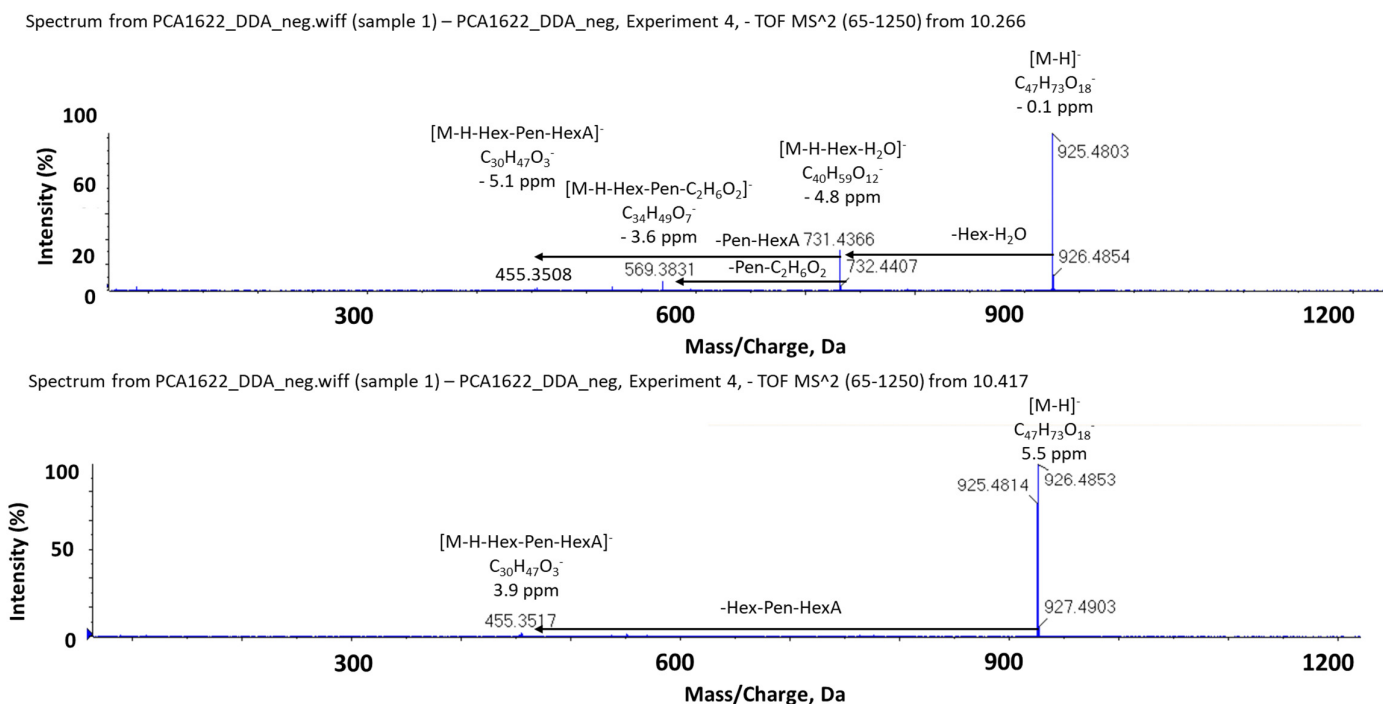
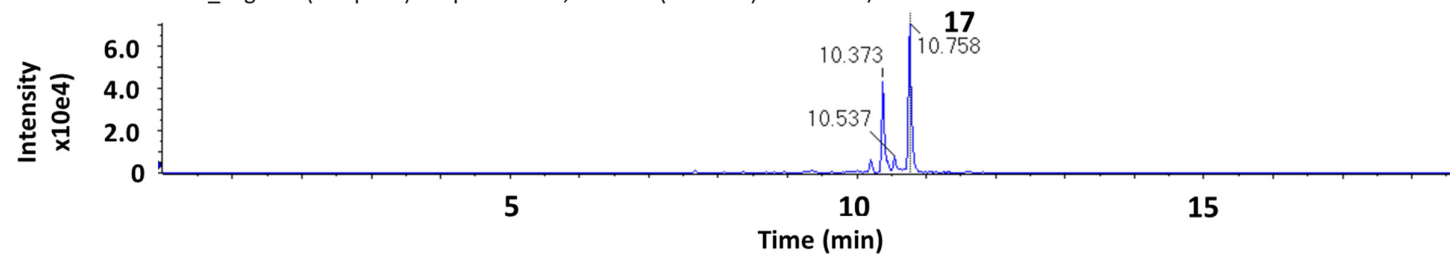


Figure S12. Extracted ion chromatogram m/z 925.4800 ± 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 14, 16 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as araloside A isomers t_R 10.266 and 10.401, respectively. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted $[M-H]^-$ ions annotated at the MS1 level).

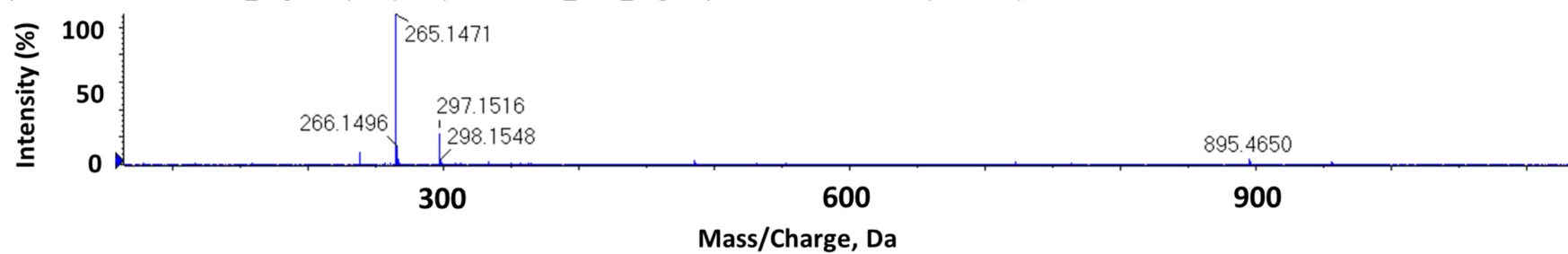
A

XIC from PCA1622_neg.wiff (sample 1) - Experiment 1, TOF-MS (65-1250): 895.460 +/- 0.025 Da



B

Spectrum from PCA1622_neg.wiff (sample 1) - PCA1622_DDA_neg, Experiment 1, - TOF MS (65-1250) from 10.763 min



C

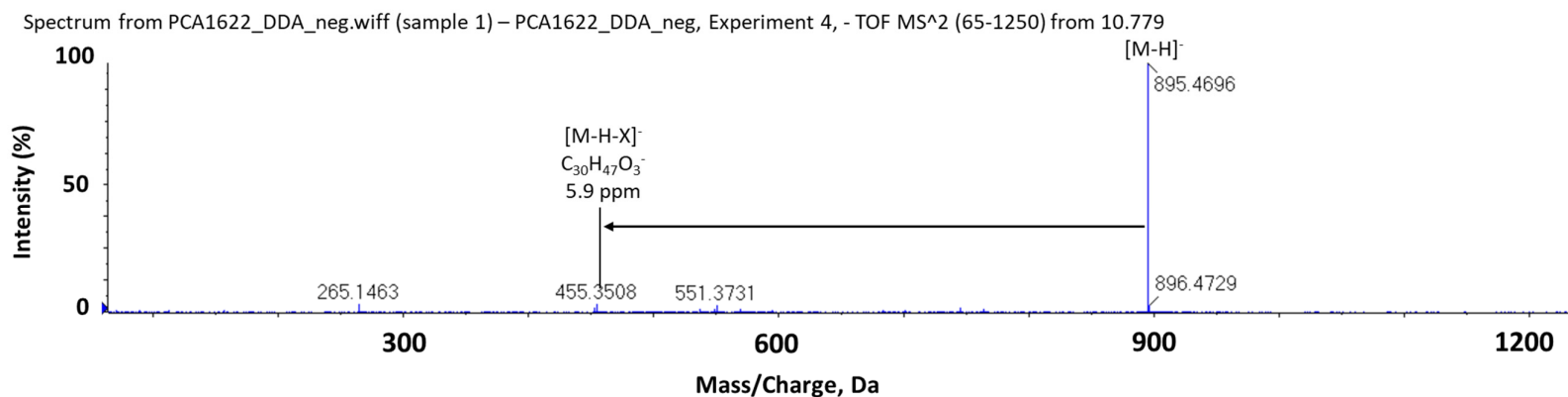
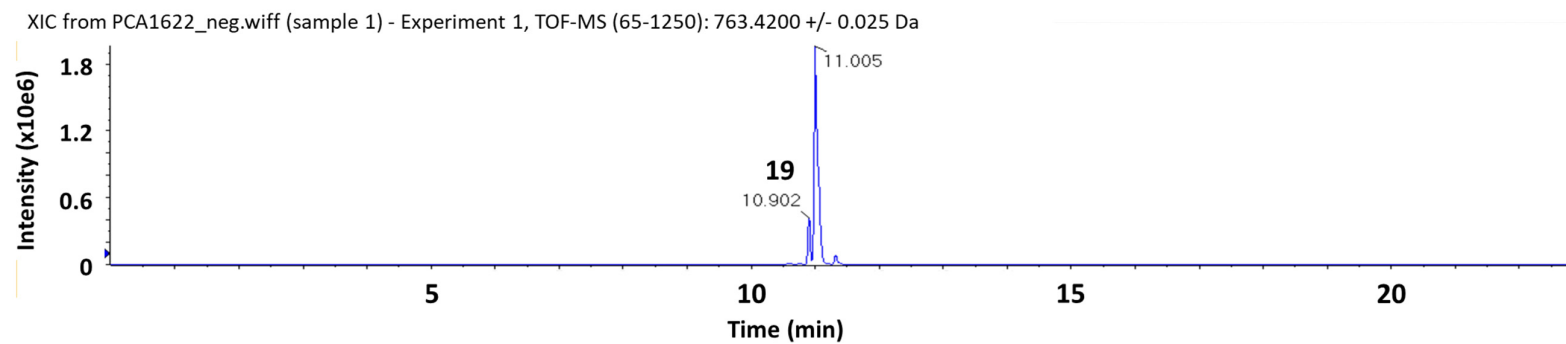
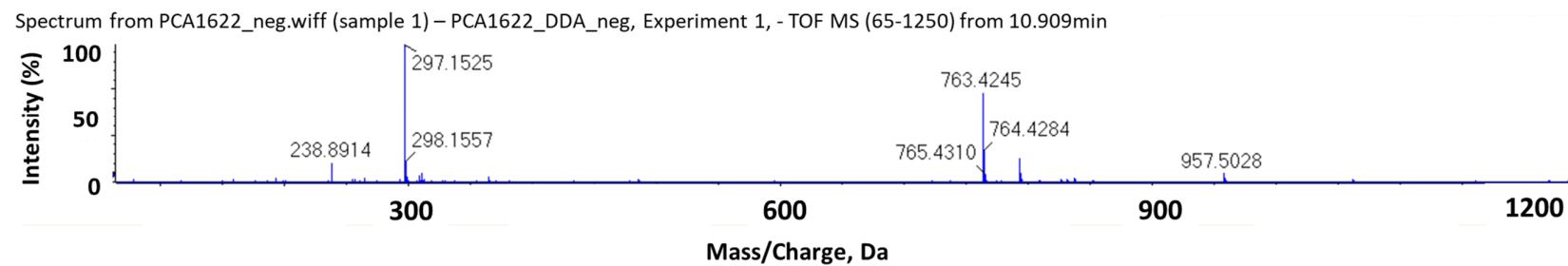


Figure S13. Extracted ion chromatogram m/z 895.4500 ± 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 17 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid unknown derivatives. The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted [M-H]⁻ ions annotated at the MS1 level).

A



B



C

Spectrum from PCA1622_DDA_neg.wiff (sample 1) – PCA1622_DDA_neg, Experiment 4, - TOF MS² (65-1250) from 11.059

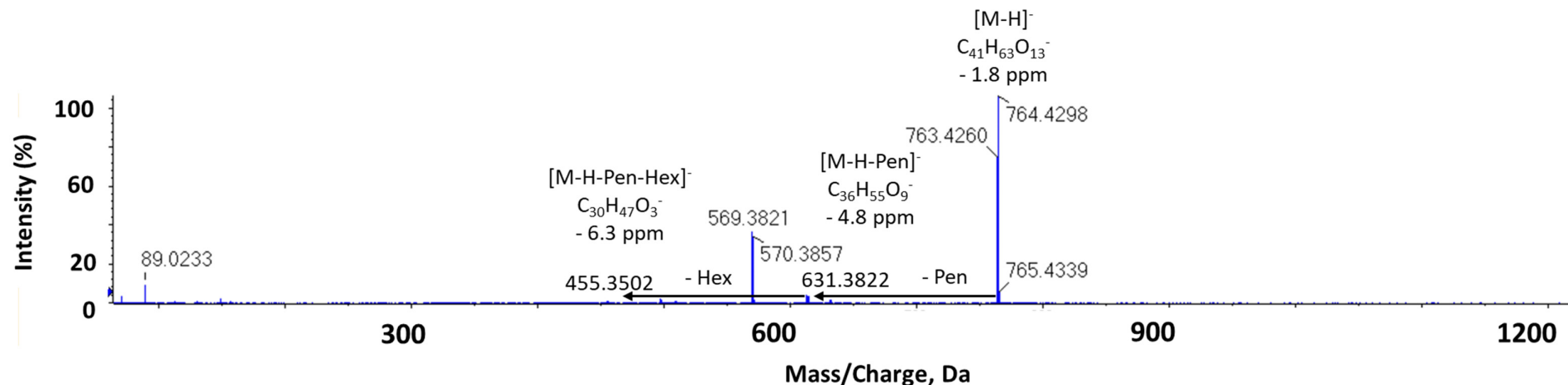


Figure S14. Extracted ion chromatogram m/z 763.4300 \pm 0.02 (A), the MS spectra (B) and MS/MS fragmentation patterns (C) of the compounds 19 annotated in the total ethanolic extract of *Aralia elata* var. *mandshurica* (Rupr. & Maxim.) J. Wen roots as oleanolic acid 3-*O*-hexuronide-(1-3-pentafuranoside). The analysis relied on RP-UHPLC-QqTOF-MS accomplished with a Waters ACQUITY I-Class UPLC System (Waters GmbH, Eschborn, Germany) coupled on-line to a Triple-TOF6600 hybrid mass spectrometer (Sciex, Darmstadt, Germany) in the negative ion mode. Metabolites were annotated by reversed phase ultra-high performance liquid chromatography-mass spectrometry and tandem mass spectrometry (RP-UHPLC-QqTOF-MS and MS/MS) in targeted data-dependent acquisition (DDA) experiments (with the inclusion list containing the m/z of all predicted [M-H]⁻ ions annotated at the MS1 level).

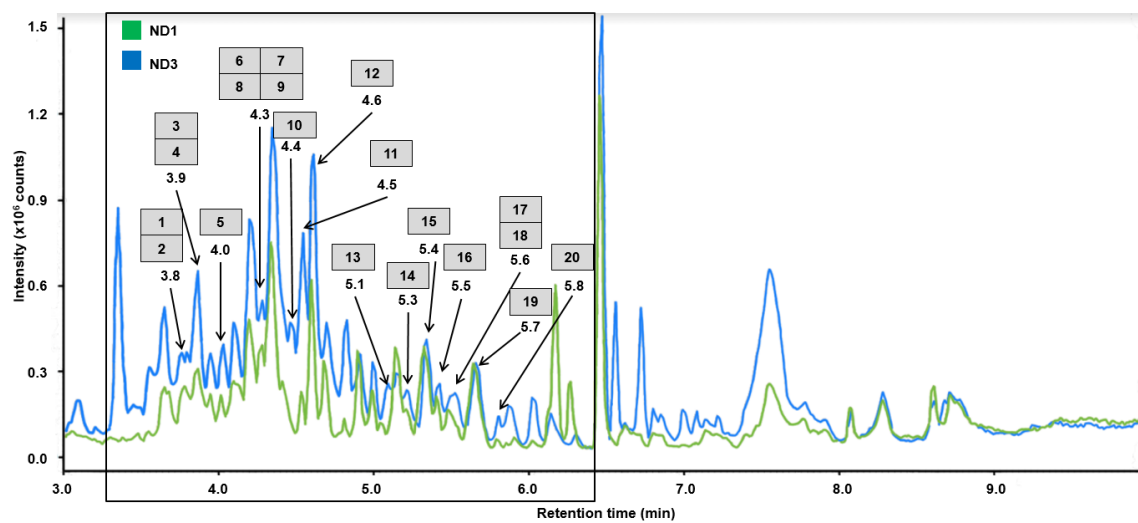


Figure S15. The full t_R range obtained in the chromatograms of ND1 and ND3 extracts of whole roots of *A. elata* (peak numbers correspond to compounds listed in Table 1).

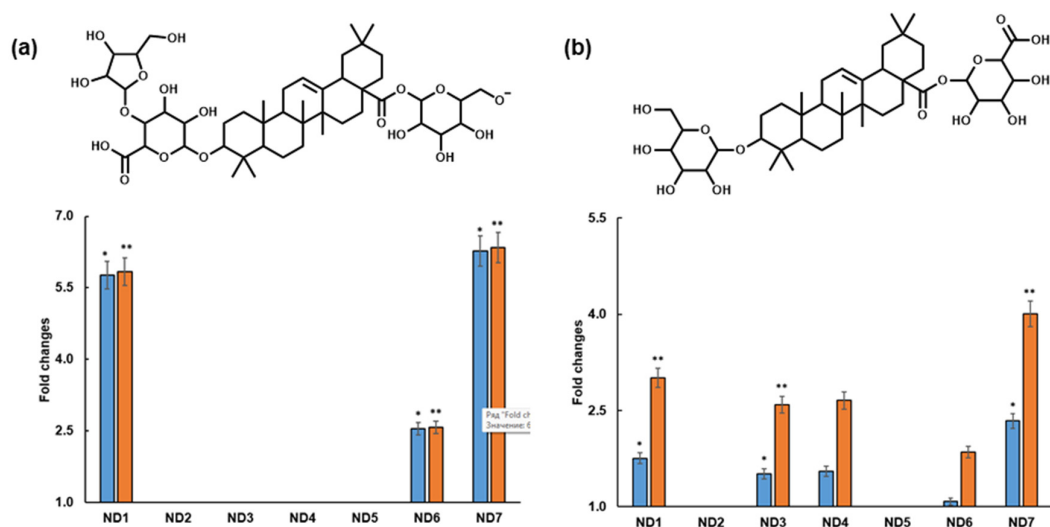


Figure S16. Structures and relative recoveries of **14** (araloside isomer 1 (a)), **18** (oleanolic acid-3-*O*-(hexosyl)-28-1-hexouronide ester isomer 1 (b)), **13**, expressed as the difference (fold) in comparison to those observed in aqueous and ethanolic extracts. Fold changes comparison with water (blue), fold changes comparison with ethanol (orange), * (for recoveries in relation to water) or ** (for recoveries in relation to ethanol) - $p \leq 0.05$ vs. control. The compounds are numbered as in Table 1. ND1 – NADES with choline chloride and malic acid (molar ratio 1:1), ND2 – NADES with the molar ratio of choline chloride and malic acid of 1:2, ND3 – NADES with the molar ratio of choline chloride and lactic acid of 1:3, ND4 - NADES with the molar ratio of choline chloride and lactic acid of 1:3+ 30% (v/v) water, ND6 – NADES with the molar ratio of sorbitol and malic acid of 1:1 + 10% (v/v) water, ND7 - NADES with the molar ratio of sorbitol and malic acid of 1:2+20% (v/v) water.

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