

*Supplementary Materials*

# **Antihyaluronidase and Antioxidant Potential of *Atriplex sagittata* Borkh. in Relation to Phenolic Compounds and Triterpene Saponins**

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**Table S1.** Optimized parameters for the quantitative analysis of phenolic acids.

Compound	Retention time [min]	MRM (Q1/Q3) [m/z]	DP [V]	EP [V]	CEP [V]	CE [eV]	CXP [V]
gallic acid	1.01	168.7/78.9	-35	-3	-12	-36	0
		168.7/124.9	-35	-3	-12	-14	0
protocatechuic	2.25	152.9/80.9	-55	-1	-10	-26	0
		152.9/107.8	-55	-1	-10	-38	0
gentisic acid	3.34	352.9/80	-70	-4	-16	-110	0
		352.9/96.9	-70	-4	-16	-52	0
4-hydroxybenzoic acid	4.24	136.8/92.9	-30	-7	-10	-18	0
vanillic acid	5.52	166.8//107.9	-35	-4	-12	-18	0
		166.8/123	-35	-4	-12	-12	0
caffeic acid	5.75	178.7/88.9	-30	-6,5	-12	-46	0
		178.7/134.9	-30	-6,5	-12	-16	0
syringic acid	6.34	196.9/122.8	-30	-9	-12	-24	0
		196.9/181.9	-30	-9	-12	-12	-2
4-hydroxycynamic acid	6.70	162.8/93	-30	-8	-12	-44	0
		162.8/119	-30	-8	-12	-14	0
ferulic acid	6.84	192.8/133.9	-25	-11,5	-14	-16	0
		192.8/177.9	-25	-11,5	-14	-12	-2
salicylic acid	6.86	136.9/75	-35	-4	-10	-48	0
		136.9/93	-35	-4	-10	-16	-2
sinapic acid	6.87	222.8/121	-35	-8,5	-10	-36	0
		222.8/148.9	-35	-8,5	-10	-20	0
veratric acid	6.88	180.7/121.9	-35	-6	-14	-18	0
		180.7/136.9	-35	-6	-14	-12	0
3-hydroxycynamic acid	6.89	162.8/91	-35	-4,5	-12	-36	0
		162.8/119	-35	-4,5	-12	-14	0
rosmarinic acid	7.01	358.7/132.6	-50	-5	-26	-44	0
		358.7/160.8	-50	-5	-26	-20	-2

Abbreviations: Q1 – precursor ion, Q3 – product ion, <sup>a</sup>DP- Declustering Potential; - <sup>b</sup>EP- Entrance Potential; <sup>c</sup>CEP- Cell Entrance Potential; <sup>d</sup>CE- Collision Energy; <sup>e</sup>CXP- Collision Cell Exit Potential.

**Table S2.** Analytical parameters of the quantitative LC-MS/MS method for determination of phenolic acids.

Compound	LOD [ng/mL]	LOQ [ng/mL]	R <sup>2</sup>	Linear range [ng/mL]
gallic acid	50	100	0.9995	100-50000
protocatechuic	10	60	0.9993	60-12000
gentisic acid	8	15	0.9982	30-7500
4-hydroxybenzoic acid	30	50	0.9990	50-50000
vanillic acid	60	120	0.9994	120-30000
caffeic acid	60	315	0.9976	313-6250
syringic acid	50	155	0.9986	155-7750
4-hydroxycynamic acid	20	30	0.9980	30-7500
ferulic acid	15	25	0.9982	25-5000
salicylic acid	5	10	0.9979	44-1000
sinapic acid	7	30	0.9992	30-6000
veratric acid	100	500	0.9975	500-25000
3-hydroxycynamic acid	25	100	0.9994	100-25000
rosmarinic acid	20	25	0.9995	25-25000

Abbreviations: LOD -limit of detection. LOQ -limit of quantification. R<sup>2</sup> - determination coefficient.

**Table S3.** Optimized parameters for the quantitative analysis of flavonoids.

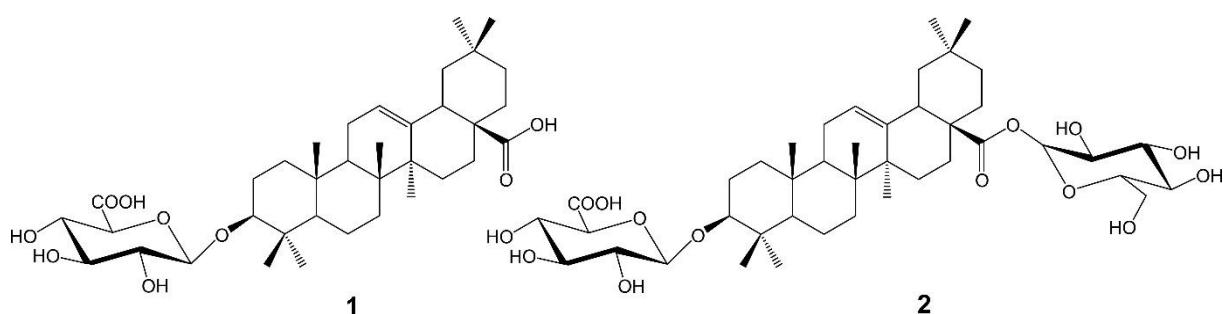
Compound	Retention time [min]	Q1 [m/z]	Q3 [m/z]	DP <sup>a</sup> [V]	EP <sup>b</sup> [V]	CEP <sup>c</sup> [V]	CE <sup>d</sup> [eV]	CXP <sup>e</sup> [V]
kaempferol-3-gluco-side-7-rhamnoside	11.25	592.818	284.9 430.9	-145 -145	-4.5 -4.5	-30 -30	-42 -36	-4 -16
rutin	11.99	608.745	299.6 270.9	-90 -90	-8 -8	-30 -30	-46 -60	-4 -4
isovitexin/vitexin	12.4	430.821	310.9 340.9	-65 -65	-4.5 -4.5	-18 -18	-28 -26	-4 -14
isoquercetin	13	462.739	299.7 270.7	-85 -85	-1.5 -1.5	-20 -20	-30 -44	-4 -4
kaempferol-3-rutinoside	13.31	592.697	284.8 226.7	-65 -65	-12 -12	-30 -30	-38 -68	-2 -2
narcisoside	13.52	622.817	314.9 298.8	-90 -90	-4.5 -4.5	-30 -30	-40 -52	-4 -4
naringin	14.50	579.109	151 271	-80 -80	-4 -4	-26 -26	-54 -42	-2 -4
astragalin	14.66	446.72	226.8 254.8	-75 -75	-9 -9	-20 -20	-54 -40	-2 -2
isorhamnetin-3-gluco-side	14.76	476.807	313.9 270.9	-95 -95	-10 -10	-22 -22	-30 -44	-4 -4

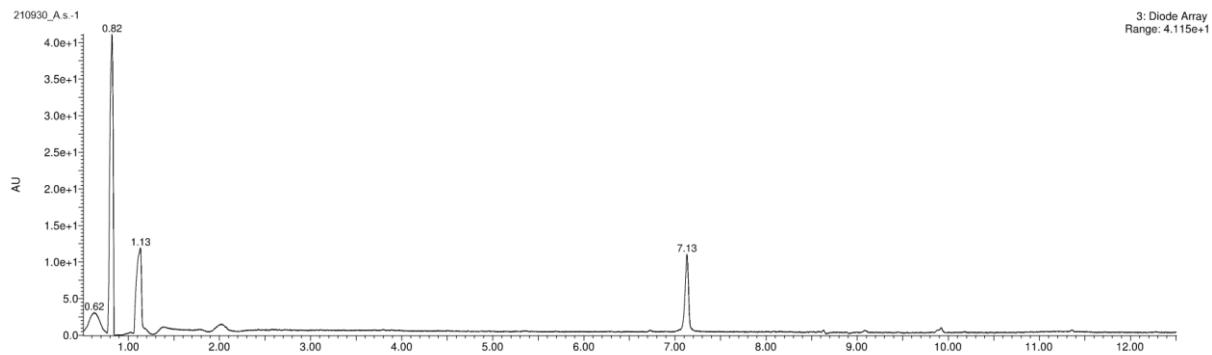
Abbreviations: Q1 – precursor ion, Q3 – product ion <sup>a</sup>DP- Declustering Potential; - <sup>b</sup>EP- Entrance Potential; <sup>c</sup>CEP- Cell Entrance Potential; <sup>d</sup>CE- Collision Energy; <sup>e</sup>CXP- Collision Cell Exit Potential.

**Table S4.** Analytical parameters of the quantitative LC-MS/MS method for determination of flavonoids.

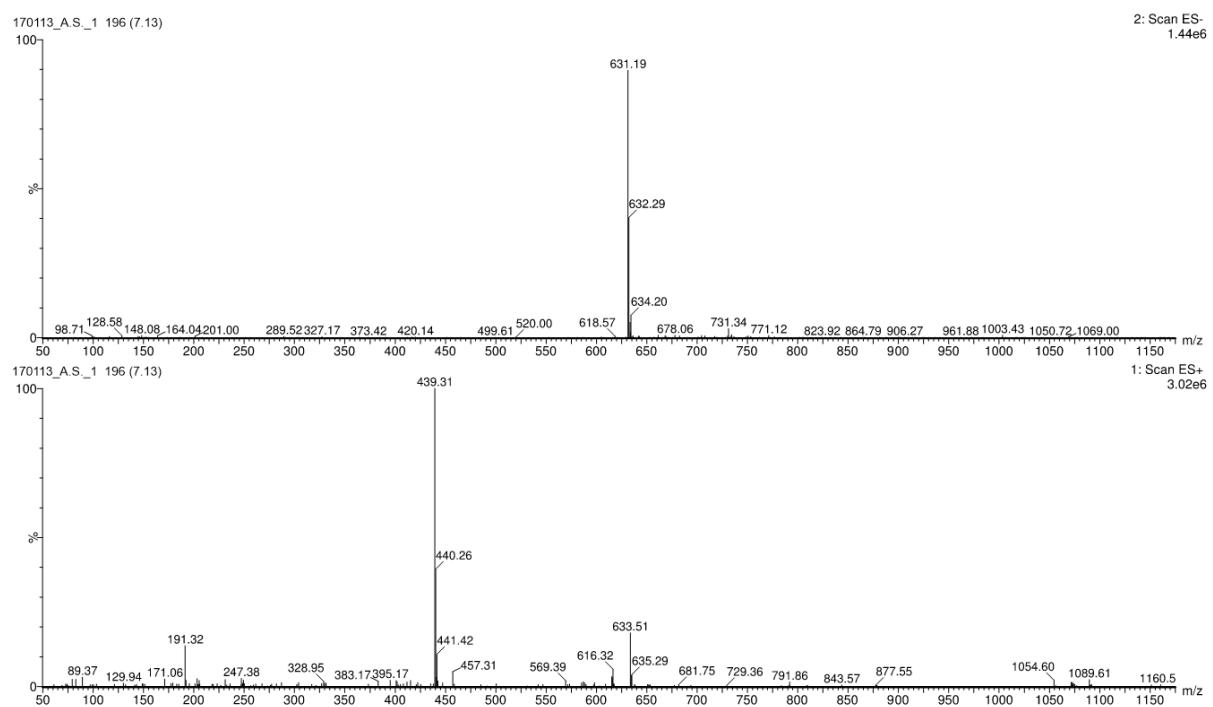
Compound	LOD [ng/mL]	LOQ [ng/ mL]	R <sup>2</sup>	Linear range [ng/ mL]
kaempferol-3-glucoside-7-rhamnoside	90	150	0.9980	400-40000
rutin	100	250	0.9973	2000-28000
isovitexin/vitexin	100	250	0.9989	2500-80000
isoquercetin	167	250	0.9987	2000-25000
kaempferol-3-rutinoside	50	100	0.9991	200-20000
narcisoside	50	100	0.9975	200-20000
naringin	50	100	0.9985	1000-25000
astragalin	120	240	0.9978	1200-24000
isorhamnetin-3-glucoside	100	200	0.9985	2000-20000

Abbreviations: LOD -limit of detection. LOQ -limit of quantification. R<sup>2</sup> - determination coefficient.

**Figure S1.** Structures of compounds 1 (calenduloside E) and 2 (chikusetsusaponin IVa).

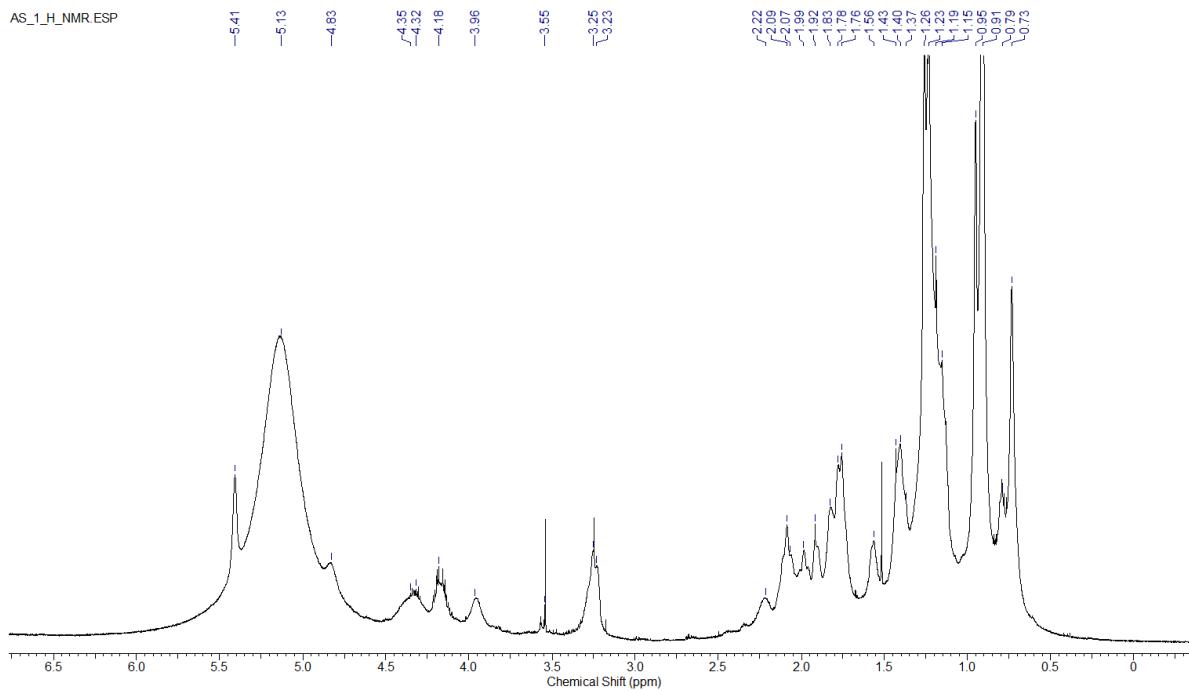


**Figure S2.** UPLC-PDA chromatogram of compound 1.

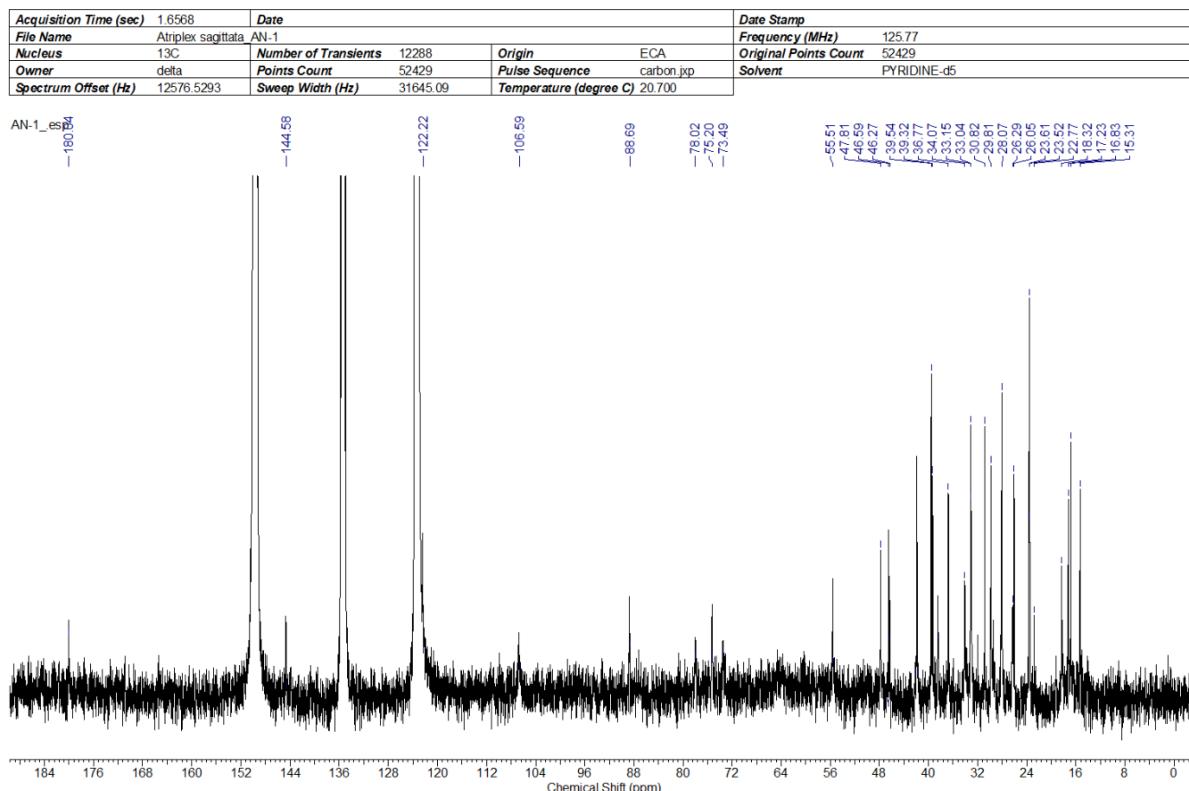


**Figure S3.** ESI-QTOF-MS (negative and positive ion mode) of compound 1.

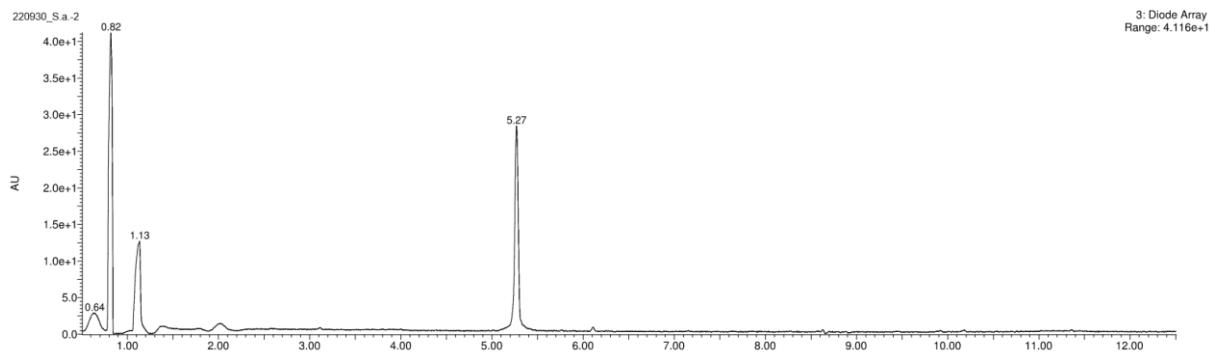
<b>Acquisition Time (sec)</b>	3.4919	<b>Date</b>		<b>Date Stamp</b>	
<b>File Name</b>	Atriplex sagittata/AS-1			<b>Frequency (MHz)</b>	500.16
<b>Nucleus</b>	<sup>1</sup> H	<b>Number of Transients</b>	8	<b>Original Points Count</b>	26214
<b>Owner</b>	delta	<b>Points Count</b>	26214	<b>Pulse Sequence</b>	proton.jpx
<b>Spectrum Offset (Hz)</b>	3251.0396	<b>Sweep Width (Hz)</b>	7507.11	<b>Temperature (degree C)</b>	20.700



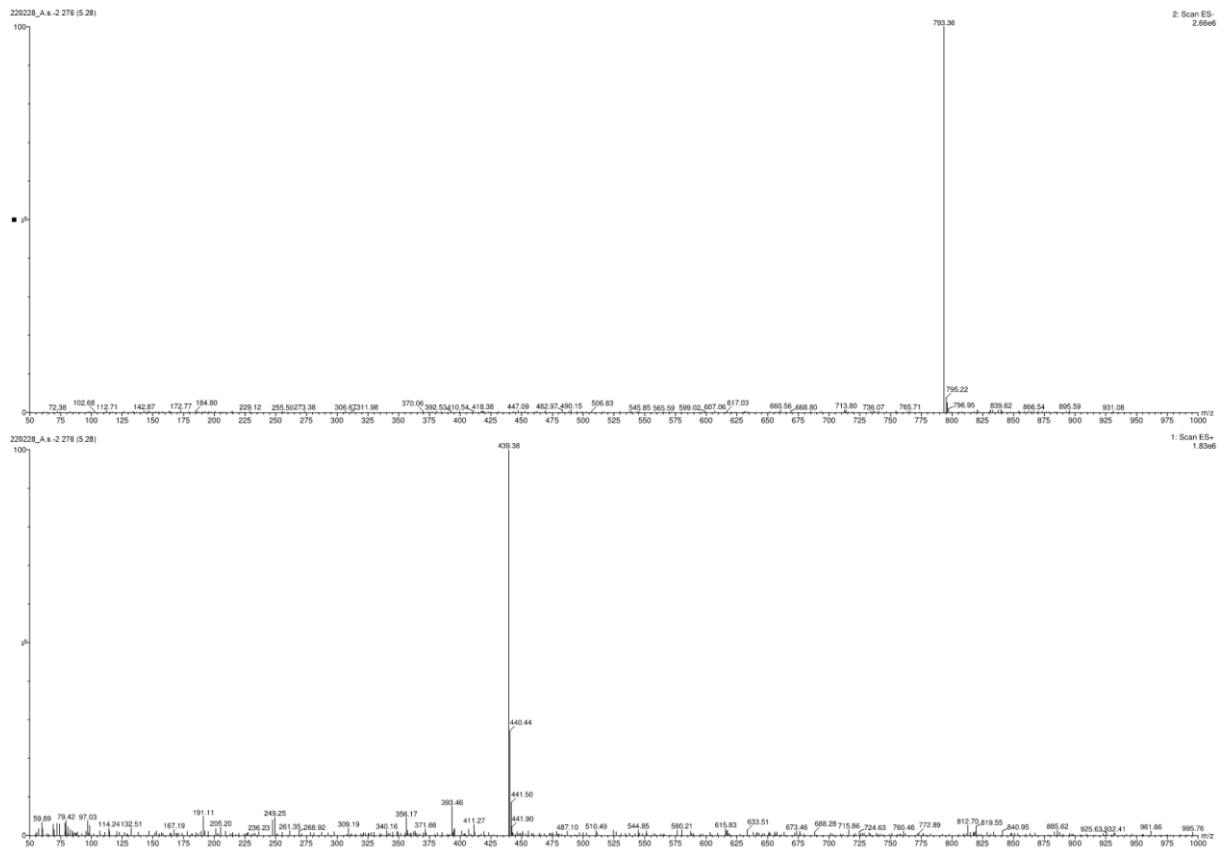
**Figure S4.** The <sup>1</sup>H NMR (500 MHz, pyridine-d5) spectrum of compound 1.



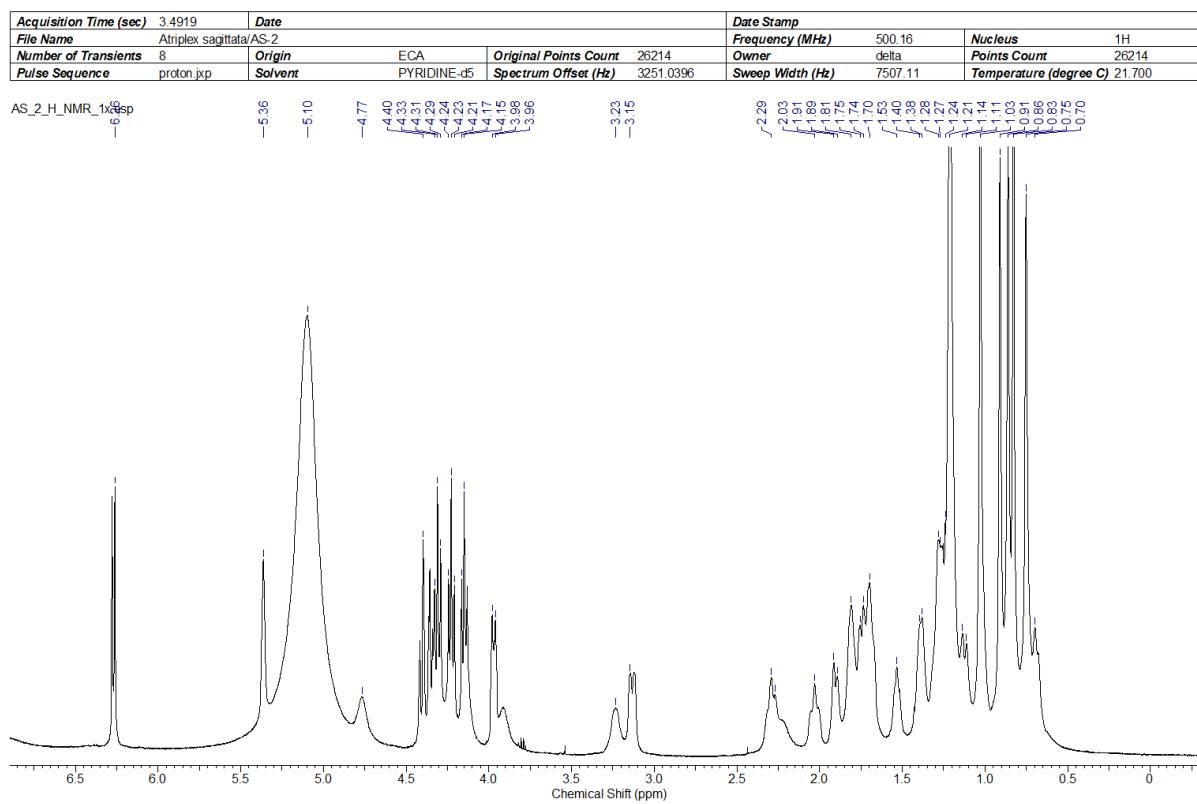
**Figure S5.** The <sup>13</sup>C NMR (125 MHz, pyridine-d5) spectrum of compound 1.



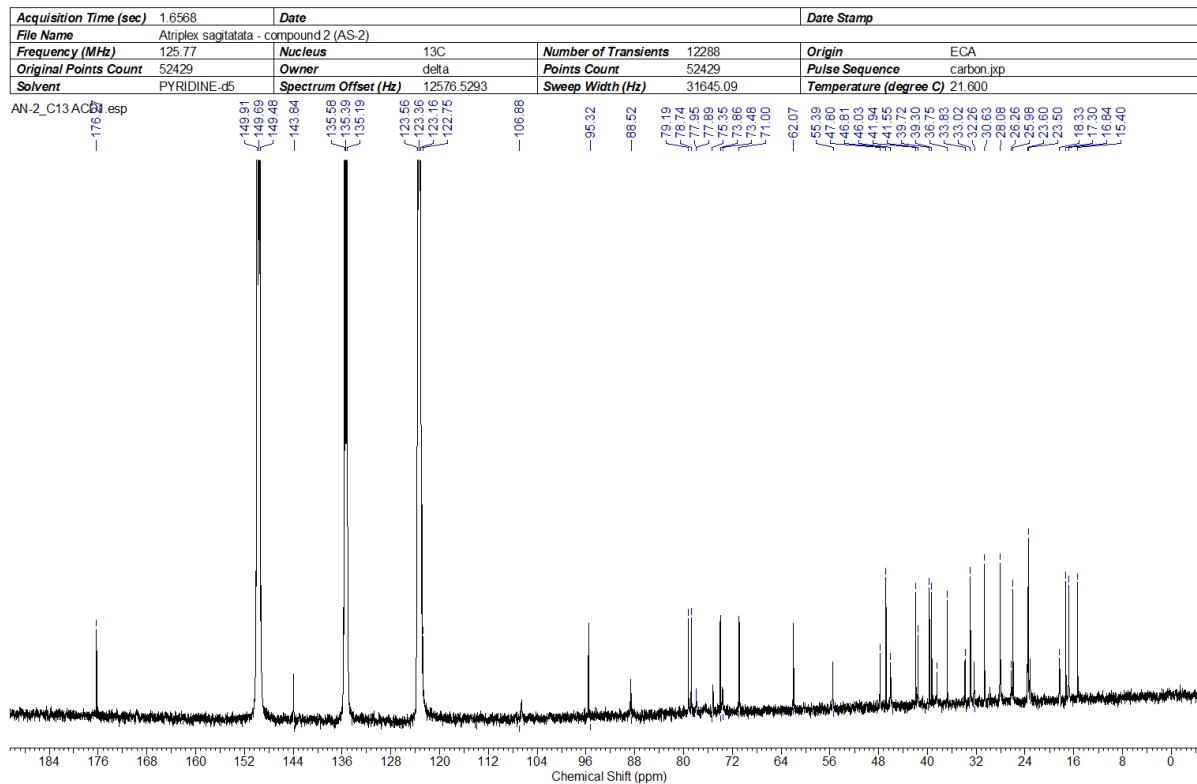
**Figure S6.** UPLC-PDA chromatogram of compound 2.



**Figure S7.** ESI-QTOF-MS (negative and positive ion mode) of compound 2.



**Figure S8.** The <sup>1</sup>H NMR (500 MHz, pyridine-d5) spectrum of compound 2.



**Figure S9.** The <sup>13</sup>C NMR (125 MHz, pyridine-d5) spectrum of compound 2.

**Table S5.**  $^1\text{H}$  (500 MHz) and  $^{13}\text{C}$  (125 MHz) NMR spectral data ( $\delta$  ppm) for compound **1** and **2** (pyridine-d5).

No.	Compound 1		Compound 2	
	$\delta\text{C}$	$\delta\text{H (J in Hz)}^*$	$\delta\text{C}$	$\delta\text{H (J in Hz)}^*$
<b>1</b>	38.41	1.37; 0.79	38.48	1.28; 0.75
<b>2</b>	26.29	2.22; 1.78	26.26	2.29; 1.81
<b>3</b>	88.69	3.26	88.52	3.23
<b>4</b>	39.32		39.30	
<b>5</b>	55.51	0.73	55.53	0.70
<b>6</b>	18.32	1.43; 1.23	18.33	1.38; 1.21
<b>7</b>	33.04	1.40; 1.23	32.92	1.40; 1.24
<b>8</b>	39.54		39.72	
<b>9</b>	47.81	1.56	47.81	1.53
<b>10</b>	36.77		36.75	
<b>11</b>	23.52	1.83	23.49	1.81
<b>12</b>	122.22	5.41	122.75	5.36
<b>13</b>	144.58		143.96	
<b>14</b>	42.05		41.94	
<b>15</b>	29.81	2.09; 1.19	29.81	2.27; 1.14
<b>16</b>	22.77	2.07; 1.92	23.23	2.03; 1.91
<b>17</b>	46.59		46.81	
<b>18</b>	41.72	3.23	41.55	3.15
<b>19</b>	46.27	1.76; 1.26	46.03	1.74; 1.21
<b>20</b>	30.82		30.62	
<b>21</b>	34.07	1.37; 1.15	33.83	1.27
<b>22</b>	33.02	1.99; 1.19	32.37	1.80; 1.70
<b>23</b>	28.07	1.32	28.07	1.21
<b>24</b>	16.83	0.91	16.84	0.91
<b>25</b>	15.31	0.78	15.40	0.75
<b>26</b>	17.23	0.91	17.30	1.03
<b>27</b>	26.05	1.26	25.98	1.21
<b>28</b>	180.04		176.27	
<b>29</b>	33.16	0.91	33.02	0.86
<b>30</b>	23.61	0.95	23.60	0.83
<i>3-O-<math>\beta</math>-D-GlcA</i>				
1	106.59	4.83	106.57	4.77
2	75.24	3.96	75.35	3.98
3	78.04	4.19	77.95	4.17
4	73.44	4.32	73.48	4.23
5	77.98	4.35	77.89	4.29
6	nd		nd	
<i>28-O-<math>\beta</math>-D-Glc</i>				
1			95.32	6.26 d (8.0)
2			73.86	4.15
3			78.74	4.21
4			71.00	4.24
5			79.19	3.96
6			62.07	4.40; 4.33

\* Overlapping signals are reported without designated multiplicity.