

A New Phenylpropanoid Glycoside: Serratumoside A from *Clerodendrum serratum*

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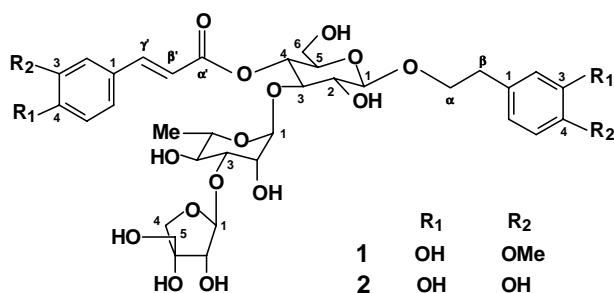
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Abstract: A new phenylpropanoid glycoside, serratumoside A, was isolated from the aerial parts of *Clerodendrum serratum* var. *amplexifolium* Moldenke. Its structure was determined by spectral and chemical methods.

Keywords: *Clerodendrum serratum*, Verbenaceae, phenylpropanoid glycoside, serratumoside A.

In the previous papers^{1,2,3,4}, we have reported some chemical constituents from *Clerodendrum serratum* var. *amplexifolium* Moldenke. A continuation of our studies on the same plant led to the isolation of a new phenylpropanoid, serratumoside A (**1**), which is reported in this paper.

Figure 1



Serratumoside A (**1**) with three known phenylpropanoid glycosides identified as acteoside⁵, martinoside⁶ and myricoside⁷, respectively, were isolated from the *C. serratum*. Compound **1** was isolated as a brown amorphous powder, $[\alpha]_D^{19} + 89.42$ (c 0.261, MeOH), the negative FABMS gave quasimolecular ion peak at m/z 783 [M-1]⁻, suggesting the molecular formula as C₃₆H₄₈O₁₉, which was confirmed by the high resolution negative FABMS (found [M-1]⁻ 783.2640, calcd. 783.2711) and the NMR

spectral data of **1**. Its UV [λ_{\max} (log ϵ): 214.5 (5.15), 258 (4.73), 282.0 (4.59) and 325.5 (5.21) nm] and IR (ν : 3417 br., 1705, 1630 and 1595 cm^{-1}) spectra showed the presence of hydroxyl groups, α , β -unsaturated ester and aromatic rings. The ^1H and ^{13}C NMR spectral data (**Table 1** and **2**) of **1** showed that the signals were in good agreement with those of myricoside (**2**) except for the differences in the aglycone and acyl moieties, *i.e.* the existence of two additional methoxy groups [δ_{H} 3.75 (3H, s); δ_{C} 55.8 (q) and δ_{H} 3.83 (3H, s); δ_{C} 55.8 (q)] in the aglycone and acyl moiety, respectively.

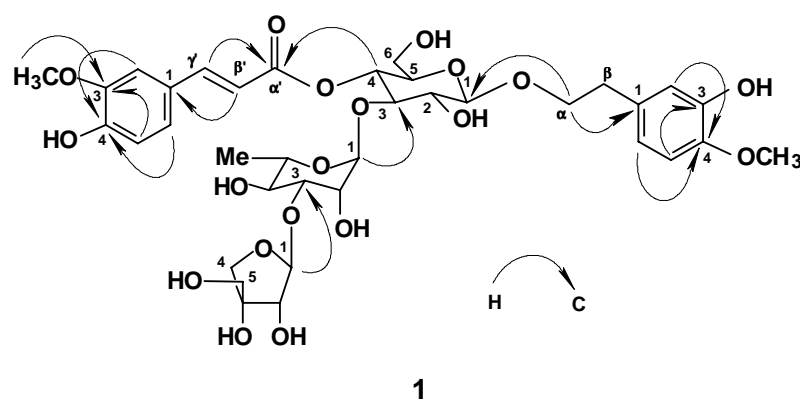
Table 1 ^1H NMR spectral data of compounds **1** and **2** in pyridine- d_5 (500MHz, δ in ppm from TMS and J in Hz)

Proton	1	2
Aglycone		
2	6.83 d (1.8)	6.62 d (1.7)
5	6.71 d (8.1)	6.63 d (7.6)
6	6.67 dd (8.1, 1.9)	6.50 dd (8.0, 1.8)
αa	3.89 m	3.84 m
αb	3.67 m	3.60 m
β	2.76 m	2.70 m
OMe	3.75 s	
Acyl moiety		
2	7.32 d (1.8)	7.04 d (1.8)
5	6.84 d (8.1)	6.76 d (8.0)
6	7.12 dd (8.0, 1.8)	6.99 dd (8.2, 1.8)
β'	6.44 d (15.8)	6.22 d (15.8)
γ'	7.58 d (15.8)	7.46 d (15.8)
OMe	3.83 s	
Glucosyl moiety		
1	4.41 d (7.6)	4.37 d (7.8)
2	3.20 m	3.17 m
3, 5	3.72 m	3.69 m
4	4.71 t (9.7)	4.68 t (9.7)
6		3.43 m
6a	3.45 m	
6b	3.40 m	
Rhamnosyl moiety		
1	5.06 br.s	5.02 br.s
2, 3	3.72 m	3.69 m
4	3.14 m	3.11 m
5	3.37 m	3.33 m
6	1.01 d (6.1)	0.96 d (6.1)
Apiosyl moiety		
1	4.81 d (2.7)	4.77 d (2.9)
2	3.72 m	3.69 m
4a	3.80 d (9.2)	3.78 d (9.4)
4b	3.58 d (9.2)	3.55 d (9.4)
5	3.32 m	3.27 m

Acidic hydrolysis of **1** gave glucose, rhamnose and apiose (identified by TLC

comparing with authentic samples). Additionally, its ^1H and ^{13}C NMR spectral data showed three anomeric signals at δ_{H} 4.41 (1H, d, $J = 7.6$ Hz) and δ_{C} 102.3 (d); δ_{H} 5.06 (1H, br. s) and δ_{C} 101.3 (d) and δ_{H} 4.81 (1H, d, $J = 2.7$ Hz) and δ_{C} 109.2 (d), which were attributed to glucose, rhamnose and apiose by the ^1H - ^1H COSY, HMQC and HMBC spectra of **1**. Further analysis of HMBC, HMQC-TOCSY spectra and comparison of the NMR spectral data of **1** with those of **2**, the ^1H and ^{13}C NMR spectral signals of sugar moieties could be assigned. The negative FABMS produced the fragment (m/z 651 [M-Api-1]), which revealed the apiose as a terminal sugar. From the HMBC spectrum of **1**,

Figure 2 The key ^1H - ^{13}C long-range correlations observed in the HMBC spectrum of **1**



some principal ^1H - ^{13}C long range correlations (**Figure 2**) were clearly observed between the protons (δ 3.75) of a methoxy group and C-3 (δ 148.0) of acyl moiety; the protons (δ 3.83) of a methoxy group and C-4 (δ 145.9) of aglycone moiety; H-5 (δ 6.84) of acyl moiety and C-3 (δ 148.0) of acyl moiety; H-2 (δ 7.32) and H-6 (δ 7.12) of acyl moiety and C-4 (δ 149.5) of acyl moiety, respectively; H-5 (δ 6.71) of aglycone and C-3 (δ 146.2) of aglycone; H-2 (δ 6.83), H-6 (δ 6.67) of aglycone and C-4 (δ 145.9) of aglycone, respectively; H-1 (δ 4.41) of glucose and C- α (δ 70.2) of aglycone; H-4 (δ 4.71) of glucose and C- α' (δ 166.0) of acyl moiety; H-1 (δ 5.06) of rhamnose and C-3 (δ 78.9) of glucose; and H-1 (δ 4.81) of apiose and C-3 (δ 76.1) of rhamnose. These facts indicated that the linkages among aglycone, acyl moiety, glucose, rhamnose and apiose of **1** were consistent with those of **2** and showed the two additional methoxy groups were attached to the C-3 position of acyl moiety and the C-4 position of aglycone, respectively. Therefore, serratumoside A was deduced to be 3-hydroxy-4-methoxy- β -phenethyl-O- β -D-apiofuranosyl-(1 \rightarrow 3)- α -L-rhamnopyranosyl-(1 \rightarrow 3)-4-O-feruloyl- β -D-glucopyranoside. Its structure was shown in **Figure 1**.

Table 2 ^{13}C NMR spectral data of compounds **1** and **2** in pyridine- d_5 (125.8 MHz, δ in ppm from TMS)

C	1	2	C	1	2
Aglycone			Glc moiety		
1	131.1 s	129.2 s	1	102.3 d	102.2 d
2	112.4 d	115.5 d	2	74.5 d	74.4 d
3	146.2 s	144.9 s	3	78.9 d	78.9 d
4	145.9 s	143.5 s	4	69.5 d	69.4 d
5	116.4 d	116.4 d	5	72.9 d	72.8 d
6	119.6 d	119.6 d	6	67.2 t	67.1 t
α	70.2 t	70.4 t	Rha moiety		
β	35.0 t	35.0 t	1	101.3 d	101.3 d
OMe	55.8 q		2	70.6 d	70.5 d
Acyl moiety			3	76.1 d	75.9 d
1	125.8 s	125.5 s	4	71.7 d	71.7 d
2	111.2 d	114.8 d	5	68.9 d	68.8 d
3	148.0 s	145.9 s	6	18.2 q	18.2 q
4	149.5 s	148.5 s	Api moiety		
5	115.6 d	115.8 d	1	109.2 d	109.2 d
6	123.3 d	121.6 d	2	76.1 d	75.9 d
α'	165.9 s	165.8 s	3	78.9 s	78.9 s
β'	114.0 d	113.4 d	4	73.5 t	73.4 t
γ'	146.3 d	145.6 d	5	63.2 t	63.2 t
OMe	55.8 q				

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