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1 **Triterpenoid saponins from *Anagallis monelli* ssp. *linifolia* (L.) Maire and their**
2 **chemotaxonomic significance**

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9

10 **Abstract**

11 Thirteen undescribed triterpenoid saponins named monellosides A-M, were isolated from the
12 aerial parts of *Anagallis monelli* ssp. *linifolia* (L.) Maire, together with ten known oleanane-type
13 glycosides. Their structures were elucidated by 1D and 2D-NMR spectroscopy (COSY,
14 TOCSY, HSQC, HMBC and ROESY) as well as high resolution mass spectrometry (HR-ESI-
15 MS) and acid hydrolysis. Monellosides A-M have a carbohydrate chain linked on the C-3 of
16 the aglycone with a common β -D-glucopyranosyl-(1 \rightarrow 4)- α -L-arabinopyranosyl sequence
17 which was further glycosylated by a glucose and/or a xylose. The sequence β -D-
18 xylopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-
19 arabinopyranosyl was common to all the 13,28-epoxy-oleanane core skeleton except one
20 compound. In order to discuss the reclassification of *Anagallis* in Primulaceae, we compared
21 saponins from species of Myrsinaceae and Primulaceae families and showed that these
22 species were characterized by a pentacyclic triterpenoid saponin with a 13,28-epoxy bridge
23 skeleton. Our phytochemical results increase the knowledge of saponins of the genus
24 *Anagallis*, their chemotaxonomy and stimulate the evaluation of the biological activities of
25 these saponins.

26

27 **Keywords:** *Anagallis monelli*, Primulaceae, Triterpenoid saponins, Chemotaxonomy

28

29 **1. Introduction**

30 The genus *Anagallis* currently belongs to the Primulaceae family, although recent studies
31 based on DNA sequences of three chloroplast genes and morphology have suggested its
32 placement in the Myrsinaceae as *Lysimachia* (Källersjö et al., 2000; Manns and Anderberg,
33 2009 and 2011). *Anagallis* contained about 28 species growing as mainly annual herbs,
34 distributed in Africa, Madagascar, Europe and South America. This genus is represented in
35 Algeria by four species *A. arvensis* L., *A. monelli* L., *A. repens* Pomel. and *A. enella* L. (Quezel
36 and Santa, 1963). The plant *Anagallis monelli* is an endemic herb from North Africa and is
37 represented in Algeria by two different subspecies, *Anagallis monelli* ssp. *collina* (Schousb.)
38 Maire and *Anagallis monelli* ssp. *linifolia* (L.) Maire (Quezel and Santa, 1963). Among the

39 studied species of the genus *Anagallis*, *A. arvensis* and *A. foemina* have been used in
40 traditional medicine in Navarre (Spain) against skin injuries like burns and wounds (López et
41 al., 2011). The whole plant of *A. arvensis* has been used for liver complications in Taiwan, for
42 skin diseases in Italy and for fish poisoning in rural areas of Nepal (Yasmeen et al., 2020).
43 Chemical investigations on *Anagallis* spp. have been mainly characterized by the presence of
44 saponins (Aliotta et al., 1992 ; Amoros and Girre, 1987; Glombitza and Kurth, 1987a and b;
45 Shoji et al., 1994a and b; Soberón et al., 2017), pentacyclic triterpenes (Aliotta et al., 1992;
46 De Napoli et al., 1992; Heitz et al., 1971), flavonoids and polyphenols (Ammar et al., 2008;
47 Ishikura, 1981; Kawashty et al., 1998; Rastogi and Norula, 1980), sterols (Rastogi and
48 Norula, 1980), in addition to alkaloids and quinones (Saxena and Rao, 2021). Triterpenoid
49 saponins were found in a wide variety of higher plants and display a wide range of
50 pharmacological activities, including haemolytic, expectorant, anti-inflammatory,
51 hypolipidemic, gastroprotective, immunomodulatory and antimicrobial properties (Netala et al.,
52 2015; Podolak et al., 2010). The potential anticancer activity of saponins has been suggested
53 by their cytotoxic, cytostatic, pro-apoptotic and anti-invasive effects (Koczurkiewicz et al.,
54 2015). The 13,28-epoxy-oleanane type saponins from the plant families Myrsinaceae and
55 Primulaceae show also a wide range of biological activities such as cytotoxic activities (Foubert
56 al., 2008; Podolak et al., 2013a).

57 *Anagallis monelli* ssp. *linifolia* (L.) Maire [Synonym of *Anagallis monelli* L.], also known under
58 the synonym *Lysimachia monelli* (L.) U. Mann and Anderb, is an herbaceous, perennial herb.
59 The 8 to 60 cm long stems are woody at the base. The leaves are opposite. The flowers in the
60 axils of the upper leaves are carried by pedicels of 12 to 40 mm, opposite or in 3 veinlets,
61 longer than the leaves. The lobes of the calyx of 3.6 to 7 mm, are lanceolate, with a scarious
62 margin, sometimes finely serrated (Valdes et al., 1987). In this work, we have studied the
63 chemical profile of *Anagallis monelli* ssp. *linifolia* and isolated 13 undescribed triterpenoid
64 saponins, namely monellosides A-M (1-13) and ten known triterpenoid saponins (14-23). The
65 chemophenetic significance of the isolated saponins was discussed by comparing saponins
66 from other Primulaceae species.

67

68 2. Results and discussion

69 2.1 Isolation and structural elucidation

70 The 70% EtOH extract from the aerial parts of *A. monelli* ssp. *linifolia* was sequentially
71 partitioned with EtOAc and *n*-BuOH, respectively. The *n*-BuOH soluble fraction was subjected
72 to a Diaion HP-20 resin chromatography to give three fractions (A-C). The saponin-containing
73 fraction (C), was subjected to further column chromatography to give ten known compounds
74 and thirteen undescribed saponins named monellosides A-M (1-13) (Fig. 1). Their structures
75 were elucidated by NMR techniques (¹H, ¹³C, COSY, TOCSY, ROESY, HSQC, and HMBC)
76 and mass spectrometry (HR-ESI-MS) and by comparison with literature data. The

77 monosaccharides of monellosides A-M (**1-13**) obtained by acid hydrolysis of an aliquot of the
78 saponin-containing fraction (C) were identified as L-arabinose, D-glucose and D-xylose by
79 comparison on TLC with authentic samples followed by measurement of their optical rotation
80 values after purification on TLC.

81 **Compound 1** was obtained as an amorphous white powder. Its molecular formula was
82 determined as C₅₂H₈₆O₂₂ based on the negative-ion HR-ESI-MS (1061.5537 [M-H]⁻, calcd
83 1061.5532). The ¹H NMR data (Table 1) showed the presence of seven signals corresponding
84 to the tertiary methyls at δ_H 1.28, 1.17, 1.08, 1.00, 0.98, 0.92 and 0.87 giving correlations with
85 seven carbons signals in the HSQC spectrum at δ_C 18.7 (C-27), 17.4 (C-26), 27.0 (C-23), 32.3
86 (C-29), 24.6 (C-30), 15.3 (C-25) and 15.3 (C-24), respectively. In addition, the HSQC spectrum
87 showed correlations at δ_H 4.33 (1H, d, *J* = 5.0 Hz, H-16) / δ_C 69.8 (C-16), δ_H 3.16 (1H, dd, *J* =
88 11.7, 4.3 Hz, H-3) / 90.0 (C-3) and δ_H 3.74 (1H, dd, *J* = 11.6, 4.9 Hz, H-22) / δ_C 74.2 (C-22).
89 Furthermore, a quaternary carbon signal at δ_C 86.9 due to C-13 was linked to an oxygenated
90 methylene at δ_H 3.67 (m, H₂-28) in the HMBC spectrum (Aliotta et al., 1992). HMBC spectrum
91 showed also correlations from H₂-28 and H-18 [δ_H 1.47 (1H, dd, *J* = 14.0, 2.5 Hz)] to two
92 oxygenated methines at δ_C 69.8 (C-16) and δ_C 74.2 (C-22) and from H-3 (δ_H 3.16) to carbons
93 at δ_C 27.0 (C-23), 15.3 (C-24) and a quaternary carbon at δ_C 39.2 (C-4) (Fig. 2). Taken
94 together, these data were indicative of a 13,28-epoxy-16,22-dihydroxyoleanan skeleton
95 (priverogenin B) (Kitagawa et al., 1972 ; Yosioka et al., 1967). This assumption was confirmed
96 by detailed analysis of the COSY, ROESY, HSQC and HMBC spectra which allowed the full
97 assignment of the proton and carbon resonances of the aglycone (Table 1). Correlations
98 observed between H-3/H-5 and H-5/H-9 in the ROESY spectrum indicated their α-axial
99 orientation and thus the β-orientation of the oxygen at C-3 (Aliotta et al., 1992). The 16α-
100 configuration of hydroxyl group was evident from the small coupling constant ³J_{H-16/H-15} value (*J*
101 = 5 Hz), characteristic of an equatorial H-16 proton (Lehbili et al., 2018), which was confirmed
102 by the correlations from H-16/ H-15_{ax} and H-16/ H₃-26 β-oriented in the ROESY spectrum. In
103 the same fashion, the coupling constants of H-22 at δ_H 3.74 (1H, dd, *J* = 11.6, 4.9 Hz) indicated
104 its axial orientation which was confirmed by the correlations H-22/ H-30 and H-22/ H-28 in the
105 ROESY spectrum; leading to the α-orientation of the oxygen at C-22 (Aliotta et al., 1992).
106 In addition, the HMBC correlation between the H-3 proton (δ_H 3.16) and an anomeric carbon
107 at δ_C 104.2 (C-1') indicated that a glycosidic moiety was linked to C-3. After acid hydrolysis,
108 the sugar units were identified as L-arabinose, D-glucose and D-xylose by co-TLC with
109 authentic sugar followed by measurement of the optical rotation values of each purified
110 monosaccharide. The 1D and 2D NMR spectra of compound **1** confirmed the presence of one
111 α-L-arabinopyranosyl unit [δ_H 4.41 (1H, d, *J* = 6.6 Hz, H-1'), δ_C 104.2, C-1'], two β-D-
112 glucopyranosyl units [δ_H 4.71 (1H, d, *J* = 7.7 Hz, H-1''), δ_C 103.0, C-1''; δ_H 4.54 (1H, d, *J* = 7.8
113 Hz, H-1'''), δ_C 103.4, C-1'''] and one β-D-xylopyranosyl unit [δ_H 4.53 (1H, d, *J* = 7.6 Hz, H-1''''),
114 δ_C 105.9, C-1'''''] (Table 2). The coupling constant of H-1' (³J_{H-1'/H-2'} = 6.6 Hz) and the axial

115 correlations observed between H-1'/H-3' and H-1'/H-5' in the ROESY spectrum of **1**, indicated
116 the α -anomer configuration of the arabinose unit. The large coupling constants from anomeric
117 protons (>7 Hz) of the xylose and glucoses units, indicated their β -configurations (Liang et al.,
118 2011). Extensive 2D-NMR analysis (COSY, TOCSY, ROESY, HSQC and HMBC) enabled the
119 full assignments of all proton and carbon resonances of each monosaccharide (Table 2).
120 HMBC correlation between H-1' and C-3 indicated that the arabinose unit was linked to C-3 of
121 the aglycone. The glycoside sequence of compound **1** was determined by analysis of the
122 HMBC and ROESY spectra. Thus, HMBC correlations were observed between H-1'' / C-2' (δ_C
123 78.0), H-1''' / C-4' (δ_C 78.8) and H-1'''' / C-2''' (δ_C 83.7) (Fig. 2). In addition, ROESY correlations
124 confirming the interglycosidic linkage and the point of attachment of the tetra-saccharide at the
125 C-3 of the aglycone were observed between H-1' / H-3, H-1'' / H-2', H-1''' / H-4' and H-1'''' / H-
126 2''' (Fig. 2). According to the above results, the structure of compound **1** was elucidated as 3-
127 O- β -D-xylopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-
128 arabinopyranosyl-priverogenin B, named monelloside A.

129 **Compound 2**, isolated as white amorphous powder, had a molecular formula of C₅₈H₉₆O₂₇
130 determined by the negative-ion HR-ESI-MS (1223.6061 [M-H]⁻, calcd 1223.6071), and differed
131 from **1** by 162 amu corresponding to a supplementary hexosyl group. Comparison of the NMR
132 data of **2** with those of **1** (Tables 1 and 2) showed that they shared the same aglycone but
133 differed by the presence in **2** of an additional hexose unit identified as β -D-glucopyranose
134 (GlcIII, $\delta_{H-1''''}$ 4.71). The HMBC correlation between H-1'''' and C-4''' (δ_C 78.5) indicated the
135 linkage of GlcIII to GlcII on C-4''' (Shoji et al., 1994a). Therefore, the structure of compound **2**
136 was established as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-xylopyranosyl-(1 \rightarrow 2)-] β -D-
137 glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-arabinopyranosyl-priverogenin B,
138 named monelloside B.

139 **Compound 3** was obtained as a white amorphous powder. Its molecular formula was
140 determined as C₅₄H₈₈O₂₃ on the basis of its negative-ion HR-ES-IMS (1103.5629 [M-H]⁻, calcd
141 1103.5638) and it differed from **1** by 42 amu corresponding to an additional acetyl group.
142 Extensive 1D and 2D-NMR analysis (Tables 1 and 2) showed that compound **3** differ from **2**
143 by the absence of a hydroxyl group at C-22 and the presence of a hydroxyl group at C-23 of
144 the aglycone, as in anagalligenin B (Mahato et al., 1991, Shoji et al., 1994b). Additional signals
145 assigned to an acetyl group linked to C-6'' of GlcI, was evidenced by the HMBC correlations
146 between a carbonyl carbon signal at δ_C 171.5 with methyl protons at δ_H 2.08 and the same
147 carbonyl carbon signal with H₂-6'' of GlcI ($\delta_{H-6''}$ 4.20 (1H, dd, $J = 11.5, 2.5$ Hz); indicating that
148 **3** was an acetylated derivative of desglucoanagalosine B (**20**) (Shoji et al., 1994b). According
149 to the above results, the structure of compound **3** was elucidated as 3-O- β -D-xylopyranosyl-
150 (1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)-[6-O-acetyl- β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-
151 arabinopyranosyl-anagalligenin B, named monelloside C

152 **Compound 4** had a molecular formula of $C_{59}H_{98}O_{28}$, determined on the basis of its negative-
153 ion HR-ESI-MS (1253.6169 $[M-H]^-$, calcd 1253.6166). Extensive 1D and 2D NMR analysis
154 showed that compounds **4** and **2** differed only by the aglycone part (Tables 1 and 2).
155 Compound **4** revealed six signals corresponding to tertiary methyls at δ_H 1.25 (H₃-27), 1.20
156 (H₃-26), 0.96 (H₃-25), 0.94 (H₃-29), 0.90 (H₃-30) and 0.73 (H₃-24) and two oxygenated methine
157 protons at δ_H 3.62 (H-3) and 3.83 (1H, d, $J=5.2$ Hz, H-16) correlating with six methyl carbons
158 signals at δ_C 18.4, 17.5, 15.8, 32.5, 23.5, 11.7 and two oxygenated methine carbons signals at
159 δ_C 82.5, 75.8, respectively in the HSQC spectrum. The disappearance of the hydroxyl group at
160 C-22 led to the deshielding of C-16. In addition, two oxygenated methylene protons signal at
161 δ_H 3.30 (1H, d, $J=11.5$ Hz, H-23a) and δ_H 3.73 (1H, d, $J=11.5$ Hz, H-23b) correlated with the
162 carbon signal at δ_C 63.2 (C-23), in the HSQC spectrum. HMBC cross-peaks from δ_{H-24} 0.72 to
163 δ_{C-23} 63.3 suggested the location of a hydroxyl function at C-23 (Bechkri et al., 2020). In
164 addition, a quaternary carbon signal at δ_C 87.6 (C-13) and a singlet resonance at δ_H 4.19 (1H)
165 corresponding to H-28 indicated the presence of 13,28-epoxy-oleanane skeleton. The H-28
166 proton correlated in the HSQC spectrum with C-28 (δ_C 105.5) indicating the presence of
167 another alkoxy unit. The resulting acetal was confirmed by the HMBC correlation between the
168 C-28 and the methyl of a methoxy group at δ_H 3.33 (s, CH₃). Assignments of other proton and
169 carbon signals of the aglycone were accomplished by extensive 2D-NMR analyses which led
170 to the elucidation of the aglycone part of **4** as 13,28-epoxy-(3 β ,16 α ,23)-trihydroxy-28-methoxy-
171 oleanane, which differ from anagalligenin B (Mahato et al., 1991) by the presence of a methoxy
172 group at C-28. Thus, compound **4** was identified as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-
173 xylopyranosyl-(1 \rightarrow 2)-] β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl(1 \rightarrow 2)-] α -L-
174 arabinopyranosyl-13,28-epoxy-3 β ,16 α ,23-trihydroxy-28-methoxy-oleanane or 3-O- β -D-
175 glucopyranosyl-(1 \rightarrow 4)-[β -D-xylopyranosyl-(1 \rightarrow 2)-] β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-
176 glucopyranosyl(1 \rightarrow 2)-] α -L-arabinopyranosyl-28-methoxy-anagalligenin B, named monelloside
177 D.

178 **Compound 5** was obtained as a white amorphous powder. Its molecular formula was
179 determined as $C_{53}H_{88}O_{23}$ based on its negative-ion mode HR-ESI-MS (1091.5646 $[M-H]^-$, calcd
180 1091.5638). Comparison of the NMR data of **5** with those of **4** (Tables 1 and 2) and analysis
181 of the NMR spectra showed that compounds **4** and **5** had the same aglycone moiety (28-
182 methoxy-anagalligenin B), while comparison of the ¹H and ¹³C NMR values of the
183 oligosaccharide part of **5** with those of **1** indicated that **5** had the same tetrasaccharide moiety
184 as in **1**, linked to C-3. Location of all proton and carbon signals was achieved by extensive 2D-
185 NMR analyses, which elucidated compound **5** as 3-O- β -D-xylopyranosyl-(1 \rightarrow 2)- β -D-
186 glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-arabinopyranosyl-13,28-epoxy-
187 3 β ,16 α ,23-trihydroxy-28-methoxy-oleanane, or 3-O- β -D-xylopyranosyl-(1 \rightarrow 2)- β -D-
188 glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-arabinopyranosyl-28-methoxy-
189 anagalligenin B, named monelloside E.

190 **Compound 6**, isolated as a white amorphous powder, had a molecular formula of $C_{62}H_{104}O_{28}$,
191 determined on the basis of its HR-ESI-MS negative-ion (1295.6633 [M-H]⁻, calcd 1295.6636).
192 Comparison of the ¹H and ¹³C NMR data of **6** to those of **4** and analysis of the 2D NMR spectra
193 of **6** showed that both possessed the same penta-saccharide chain (Tables 2 and 4), while
194 slight differences were observed in the aglycone part, notably those due to the D and E rings
195 (Tables 1 and 3). Compound **6** did not show the signals of a methoxy group bound to C-28 in
196 its NMR spectra, but proton and carbon signals for an *n*-butyloxy group [δ_C 12.8, 19.1, 31.7
197 and 66.5; δ_H 0.95, 1.42, 1.56, 3.33 and 3.70]. The COSY spectrum supported the presence of
198 the *n*-butyloxy group by correlations observed between CH₂-a (δ_H 3.33 and 3.70)/CH₂-b (δ_H
199 1.56), CH₂-b/CH₂-c (δ_H 1.42), and CH₂-c/CH₃-d (δ_H 0.95). The linkage of *n*-butyloxy group at
200 the C-28 position of the aglycone was evidenced by the HMBC correlation between δ_H 4.28
201 (1H, brs, H-28)/ δ_C 66.5 (C-a) and δ_H 1.56 (2H, t, *J* = 6.5Hz, CH₂-b)/ δ_C 12.8 (C-d). These
202 evidences led to the assignment of **6** as 3-*O*- β -D-glucopyranosyl-(1→4)-[β -D-xylopyranosyl-
203 (1→2)-] β -D-glucopyranosyl-(1→4)-[β -D-glucopyranosyl-(1→2)-] α -L-arabinopyranosyl-13,28-
204 epoxy-3 β ,16 α ,23-trihydroxy-28-*n*-butyloxy-oleanane, or 3-*O*- β -D-glucopyranosyl-(1→4)-[β -D-
205 xylopyranosyl-(1→2)-] β -D-glucopyranosyl-(1→4)-[β -D-glucopyranosyl-(1→2)-] α -L-
206 arabinopyranosyl-28-*n*-butyloxy-anagalligenin B, named monelloside F.

207 **Compound 7** was obtained as an amorphous white powder. Its molecular formula was
208 determined as $C_{56}H_{94}O_{23}$ on the basis of its negative-ion in HR-ESI-MS (1133.6107 [M-H]⁻,
209 calcd 1133.6108), and corresponds to the loss of 162 amu compared to **6**. Comparison of the
210 ¹H- and ¹³C-NMR data of **7** with those of **6** and **5** showed that the NMR data of **7** exhibited
211 many similarities with those of **6**, particularly for resonances assigned to *n*-butyloxy-
212 anagalligenin B (Table 3), whereas the ¹H and ¹³C NMR signals due to the saccharide moieties
213 showed that **7** and **5** shared the same tetra-saccharide chain (Tables 2 and 4). These data
214 were confirmed by extensive 2D-NMR analyses and the assignments of all proton and carbon
215 signals of **7**, leading to the elucidation of its structure as 3-*O*- β -D-xylopyranosyl-(1→2)- β -D-
216 glucopyranosyl-(1→4)-[β -D-glucopyranosyl-(1→2)-] α -L-arabinopyranosyl-13,28-epoxy-
217 3 β ,16 α ,23-trihydroxy-28-*n*-butyloxy-oleanane, named monelloside G.

218 **Compound 8**, isolated as an amorphous white powder, had a molecular formula of $C_{58}H_{94}O_{27}$
219 (negative-ion HR-ESI-MS (1221.5913 [M-H]⁻, calcd 1221.5904)). Comparison of the NMR data
220 of **8** with those of **4** (Tables 2 and 4) and analysis of the NMR spectra showed that **8** had the
221 same penta-saccharide moiety as **4**. Most of the aglycone NMR signals were directly attributed
222 by comparison with the corresponding signals of **4** (Tables 1 and 3), but detailed analysis of
223 the NMR spectra showed dissimilarities in the ring D and different chemical shifts for C-14, C-
224 15, C-16, C-27, and C-28, due to the replacement of the hydroxyl at C-16 by a carbonyl group
225 and the absence of signals due to the methoxy group at C-28 in **8** (Tables 1 and 3). The
226 position of the carbonyl group was indicated by HMBC correlations of H-28 [δ_H 3.93 and 3.48;
227 1H each, d, *J* = 8.4 Hz] and H-15 [δ_H 2.80 and 1.81; 1H each, d, *J* = 15.8 Hz, H-15] with the

228 ketocarbonyl at δ_c 214.3 (C-16) (Liang et al., 2006 ; Liang et al., 2011). The aglycone part was
229 identified as anagalligenone B (Mahato et al., 1991). According to the above results, the
230 structure of compound **8** was elucidated 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-xylopyranosyl-
231 (1 \rightarrow 2)-] β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)]- α -L-arabinopyranosyl-
232 anagalligenone B, named monelloside H.

233 **Compound 9**, isolated as an amorphous white powder, had a molecular formula of C₅₂H₈₄O₂₂,
234 determined on the basis of its HR-ESI-MS negative-ion (1059.5370 [M-H]⁻, calcd 1059.5376),
235 indicating the loss of 162 amu compared to **8**. The complete assignments of each ¹H- and ¹³C-
236 NMR signals were achieved by analysis of the 2D-NMR experiments. The ¹H and ¹³C NMR
237 data (Tables 3 and 4), were identical to those of compound **8**, except for signals due to the β -
238 D-glucopyranosyl (Glcl) unit linked in C-2' position of the arabinosyl unit which disappeared in
239 compound **9**. Accordingly, the structure of **9** was elucidated as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)-
240 [β -D-xylopyranosyl-(1 \rightarrow 2)-] β -D-glucopyranosyl-(1 \rightarrow 4)- α -L-arabinopyranosyl- anagalligenone
241 B, named monelloside I.

242 **Compound 10** was obtained as an amorphous white powder. Its molecular formula was
243 determined as C₅₈H₉₄O₂₇ on the basis of its HR-ESI-MS negative-ion (1221.5911 [M-H]⁻, calcd
244 1221.5904). The ¹H NMR and ¹³CNMR spectra of **10** (Tables 5 and 6) showed signals
245 assignable to six angular methyl groups at δ_H 0.75/ δ_{C-24} 11.9, 1.00/ δ_{C-25} 15.1, 0.76/ δ_{C-26} 16.5,
246 1.41/ δ_{C-27} 25.9, 0.93/ δ_{C-29} 32.0 and 0.98/ δ_{C-30} 23.1, one olefinic proton at δ_{H-12} 5.42 (1H, t, *J* =
247 3.6 Hz; δ_{C-12} 123.0), two oxygenated methine protons at δ_{H-3} 3.63 (1H, m; δ_{C-3} 82.5), and δ_{H-16}
248 4.29 (1H, t, *J* = 3.6 Hz; δ_{C-16} 72.5) and one oxygenated methylene (δ_{H-23} 3.30 and 3.73; δ_{C23}
249 63.3), corresponding to a 3 β ,16 α ,23-trihydroxyolean-12-en skeleton (Bechkri et al., 2020).
250 Moreover, an HSQC correlation between a proton singlet at δ_H 9.23 (1H, s, H₁-28) and its
251 carbon at δ_c 205.6 (C-28) indicated the presence of an aldehyde group. This was confirmed
252 by the HMBC correlations observed between H-18, H-16 with C-28 (Fig. 3) (Lakhal et al., 2014,
253 Zhang et al., 2002). The aglycone of compound **10** was identified as 3 β ,16 α ,23-
254 trihydroxyolean-12-en-28-al. Detailed analysis of the 2D-NMR spectra of **10** and comparison
255 of its NMR data with those of **2** (Tables 2 and 6) identified the same penta-saccharide as in **2**.
256 The HMBC correlations observed between the signal at δ_H 3.63 (H-3) and the anomeric carbon
257 of L-arabinose at $\delta_{C-1'}$ 103.3 1 in the HMBC spectrum (Fig. 3) confirmed that the glycosidic
258 moiety was linked to C-3 of the aglycone (Zhang et al., 2002). According to the above results,
259 the structure of compound **10** was elucidated as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-
260 xylopyranosyl-(1 \rightarrow 2)-] β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)]- α -L-
261 arabinopyranosyl-3 β ,16 α ,23-trihydroxyolean-12-en-28-al, named monelloside J.

262 **Compound 11** was obtained almost as an amorphous white powder. Its molecular formula
263 was determined as C₅₂H₈₄O₂₂ on the basis of its HR-ESI-MS negative-ion (1059.5367 [M-H]⁻,
264 calcd 1059.5376). Comparison of the NMR data of **11** with those of **10** (Tables 5 and 6) showed
265 that it had the same aglycone as **10** but differed in its saccharide units. The NMR spectroscopic

266 data of the saccharide part of **11** were identical to those of **1** (Tables 2 and 6). Extensive 2D-
267 NMR analysis enabled the full assignments of the same tetra-saccharide as in **1**. The
268 correlation between H-3 of aglycone (δ_{H} 3.64, 1H, dd, $J = 11.7, 4.3$ Hz,) and the anomeric
269 carbon of the arabinose unit ($\delta_{\text{C}-1}$ 103.3) on the HMBC spectrum confirmed that the glycosidic
270 moiety was linked to C-3. According to the above evidences, the structure of **11** was elucidated
271 as 3-O- β -D-xylopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)-] α -L-
272 arabinopyranosyl-3 β ,16 α ,23-trihydroxyolean-12-en-28-al, named monelloside K.

273 **Compound 12** was obtained as an amorphous white powder. Its molecular formula was
274 determined as C₅₃H₈₆O₂₃ on the basis of its HR-ESI-MS negative-ion (1089.5490 [M-H]⁻; calcd
275 1089.5482). Comparison of the ¹H- and ¹³C-NMR data of **12** with those of **10** (Tables 5 and 6)
276 showed that it had the same aglycone but differed in its saccharide units. The NMR
277 spectroscopic data of **12** were almost identical to those of **10**, except for the absence of the
278 signals of β -D-xylopyranosyl. Detailed analysis of the 2D-NMR spectra of **12** led to the
279 identification of its structure as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -
280 D-glucopyranosyl-(1 \rightarrow 2)-] α -L-arabinopyranosyl-3 β ,16 α ,23-trihydroxyolean-12-en-28-al,
281 named monelloside L.

282 **Compound 13** was obtained as an amorphous white powder. Its molecular formula was
283 determined as C₄₇H₇₆O₁₈ on the basis of its HR-ESI-MS negative-ion (927.4966 [M-H]⁻; calcd
284 927.4953). Comparison of ¹H- and ¹³C-NMR spectra of **13** (Tables 5 and 6) and **10** indicated
285 a similarity of these two compounds, with the exception of the disappearance of the glucose
286 (GlcIII) and xylose units in the glycosidic part of **13**. Extensive analysis of the 1D- and 2D-NMR
287 spectra of **13** established that the glycosidic sequence was 3-O-glc-(1 \rightarrow 4)-[glc-(1 \rightarrow 2)-]ara.
288 Thus, the structure of compound **13** was elucidated as 3-O- β -D-glucopyranosyl-(1 \rightarrow 4)- β -D-
289 glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)] α -L-arabinopyranosyl-3 β ,16 α ,23-
290 trihydroxyolean-12-en-28-al, named monelloside M.

291 The known compounds were identified as repandoside (**14**), lysikoianoside (**15**) (Dall'acqua et
292 al., 2010), capilliposide A (**16**) (Tian et al., 2006), anagallosaponin I (**17**), anagalloside C (**18**),
293 anagalloside B (**19**), desgluconagalloside B (**20**), (Shoji et al., 1994a), anagallosaponin IX (**21**)
294 (Hifnawy et al., 2020), anagallisin D (**22**) (Mahato et al., 1991), heterogenoside D (**23**) (Huang
295 et al., 2009) (Fig. S103 in supporting material). Their spectroscopic data were in perfect
296 agreement with those reported in the literature.

297

298 2.1 Chemophenetic significance

299 Phytochemical studies of the genus *Anagallis* revealed its richness of triterpenoids
300 saponins with an oleanane skeleton. Until now, 38 compounds were isolated from only one
301 species *Anagallis arvensis* (Aliotta et al., 1992; Amoros et al., 1987; De Napoli et al., 1992;
302 Glombitza and Kurth, 1987a; Kitagawa et al., 1976; Mahato et al., 1991; Shoji et al., 1994).
303 The present study, which was carried out on the Algerian subspecies *Anagallis monelli* ssp.

304 *linifolia* (L). Maire, allowed the isolation and identification of 13 undescribed
305 monodesmoside saponins (Fig. 1), as well as 10 known monodesmoside saponins (Fig.
306 S103 in supporting material). The aglycones of these triterpenoid glycosides were found
307 to be priverogenin B (**1-2**), anagalligenin B (**3, 19, 20**), 28-alkoxy-anagalligenin B (**4-7**),
308 anagalligenone B (**8, 9, 22**), 16,23-dihydroxy-oleanolicaldehyde (**10-13**), protoprimumagenin
309 A (**14-15**), 28-hydroxy-protoprimulagenin A (**16-17**), 22-acetyl-priverogenin B (**18**), 22-
310 acetoxo-anagalligenin B (**21**), and longispinogenin (**23**). All the 23 saponins isolated from *A.*
311 *monelli* ssp. *linifolia* in the present study shared a carbohydrate chain linked to C-3 of the
312 aglycone with a common sequence glc-(1→4)-ara- which was further glycosylated by glc
313 and/or ara. In 22 of the 23 isolated saponins, arabinose unit was substituted at its C-2' by
314 another glucose unit (glcII) to give the motif **S6**: glc-(1→4)-[glc-(1→2)]-ara-. In all the
315 bridged oleanane skeleton, except for compound **9** (Fig. 1, **S1, S2** and **S3**), a xylose unit
316 was linked to C-2 of glcI making the sequence **S2**: xyl-(1→2)-glc-(1→4)-[glc-(1→2)]-ara-,
317 and this sequence was also found in three (**10, 11, 23**) of the five compounds exhibiting
318 ring-opening of the 13,28-epoxy-oleanane skeleton (Fig. 1, F.S103). About half of the
319 compounds isolated from *A. monelli* ssp. *linifolia* were substituted at C-4 of glcII by a third
320 glucose unit (Fig. 1) to give the sequence **S1**: glc-(1→4)-[xyl-(1→2)]-glc-(1→4)-[glc-
321 (1→2)]-ara- which has been encountered in saponins isolated from *A. arvensis* (Glombitza
322 and Kurth 1987b). Similarly, the sequences **S4**: glc-(1→4)-[xyl-(1→2)]-glc-(1→4)-ara-, **S5**:
323 glc-(1→4)-glc-(1→4)-[glc-(1→2)]-ara- and **S6**: glc-(1→4)-[glc-(1→2)]-ara- at C-3 of
324 compounds **9, 12** and **13**, respectively, were previously encountered in saponins isolated
325 from of *A. arvensis* (Shoji et al., 1994a; Mahato et al., 1991). In addition, all aglycone parts
326 of *A. monelli* ssp. *linifolia* with 13 β ,28-epoxy bridge were previously identified in saponins
327 isolated from *A. arvensis* (Shoji et al., 1994b), except for compounds **4-7**, which showed
328 novel features including the C-28 alkoxy group. The known anagallosaponin I (**17**),
329 anagalloside C (**18**), anagalloside B (**19**), desgluconagalloside B (**20**), anagallosaponin IX (**21**)
330 anagallisin D (**22**) were previously isolated from *A. arvensis* (Shoji et al., 1994a; Mahato et al.,
331 1991). It worth to note that the aglycones of compounds **10-13** were formerly identified in
332 saponins from *Lysimachia candida* (Zhang et al., 2002), while repandoside (**14**),
333 lysikoianoside (**15**), capilliposide A (**16**) and heterogenoside D (**23**) were previously isolated
334 from *Cyclamen repandum* (Dall'acqua et al., 2010), *Lysimachia sikokiana* (Dall'acqua et al.,
335 2010; Kohda et al., 1989) (**14, 15**), *Lysimachia capillipes* (**16**) (Tian et al., 2006) and
336 *Lysimachia heterogenea* (**23**) (Huang et al., 2009).

337 This genus *Anagallis* has traditionally been assigned to the family Primulaceae (Quezel and
338 Santa, 1963). However, data from phylogenetic analyses suggest that *Anagallis*, along with
339 eight other genera, *Lysimachia*, *Trientalis*, *Glaux*, *Asterolinon*, *Pelletiera*, *Coris*, *Ardisiandra*,
340 and *Cyclamen*, be re-located to the family Myrsinaceae (Källersjö et al., 2000; Hao et al.,
341 2004). Moreover, Manns and Anderberg (2009 and 2011) have suggested to re-located

342 *Anagallis* species in *Lysimachia* genus. In order to discuss the reclassification of *Anagallis*
343 species in the *Lysimachia* genus, both transferred from the Primulaceae to the Myrsinaceae,
344 we analysed some saponin characteristics of species from the Myrsinaceae and Primulaceae
345 families which showed that these species were characterized by a pentacyclic triterpenoid
346 saponin with 16- α -hydroxy and 13 β ,28-epoxy bridge skeleton (Foubert et al., 2008). Actually, the
347 Primulaceae family is divided into four subfamilies: Maesoideae, Theophrastoideae,
348 Primuloideae and Myrsinoideae. *Anagallis* was in the Myrsinoideae subfamily, which is
349 consistent of 41 genera and 1435 species including the genus *Ardisia*, *Cyclamen*, *Lysimachia*,
350 and *Myrsine* (Stevens, 2017). So the species previously included in Myrsinaceae family are in
351 the subfamily Myrsinoideae in the Primulaceae family.

352 In the present species, we have identified a four-unit branched sugar chain linked at C-3 of the
353 aglycone **S2** { xyl-(1 \rightarrow 2)-glc-(1 \rightarrow 4)-[glc-(1 \rightarrow 2)]ara-} (Fig. 4). This chain was found in
354 saponins of *Anagallis* (Glombitza et al., 1987b, Shoji et al., 1994a and b), *Lysimachia* (Kohda
355 et al., 1989, Podolack et al., 2013), *Cyclamen* (Altunkeyik et al., 2012; Bencharif-Betina et al.,
356 2012, Dall'acqua et al., 2010, El Hosry et al., 2014), *Ardisia* (Jia et al., 1994), *Myrsine* (Bloor
357 and Qi, 1994), and *Androsace* genera (Waltho et al., 1986) (Table 7). This tetrasaccharide
358 sequence **S2** can be substituted with glucose at C-4 of glcII or glcI in the case of *Anagallis*
359 saponins (Shoji et al., 1994a and b; Soberón et al., 2017), at C-3 or C-6 of glcII in the case of
360 *Cyclamen* saponins (Çalis et al., 1997), or at C-4 of the terminal glucose (glcI) in the case of
361 *Androsace saxifragaefolia* saponins (Waltho et al., 1986); however, it was the only study found
362 for the genus *Androsace*. In addition, this common chain was substituted with xylose at C-4 of
363 glcII in the case of *Lysimachia* saponins (Podolak et al., 2013b), or with rhamnose at C-3 of
364 glcII in saponins of *Ardisia gigantifolia* (Mu et al., 2010). Many saponins contain rhamnose in
365 the sequence of the sugar moiety in the genera *Ardisia* and *Myrsine* (Foubert et al., 2008). For
366 *Primula denticulata*, it was the only species that had similarity in saponin content with the genus
367 *Anagallis*, the other species had completely different osidic chains (Foubert et al., 2008). We can
368 conclude that the genera *Anagallis*, *Lysimachia*, and *Cyclamen* are very similar with respect
369 to chemotaxonomic markers and thus can confirm their place in the Myrsinoideae subfamily.
370 Triterpene saponins identified so far from the genus *Lysimachia*, generally have oleanane-
371 derived sapogenols of two structural types: I. 13 β ,28-epoxy and II. Δ^{12} -17-CH₂OH.
372 Compounds of type I are considered very rare and are found almost exclusively in the
373 Myrsinaceae and Primulaceae families (Foubert et al., 2008; Podolack et al., 2013). In the
374 present study, eighteen type I and one type II triterpene saponins were obtained from *Anagallis*
375 *monelli* ssp. *linifolia*. It is interesting to observe that the known compounds **14-23** were reported
376 previously in various species belonging exclusively to genera traditionally assigned to the
377 family Myrsinaceae or genera which were re-located to this family from the Primulaceae, i.e.,
378 *Anagallis*, *Lysimachia* and *Cyclamen*. Four of the ten known compounds isolated from *A.*
379 *monelli* ssp. *linifolia* (**14-16**, and **23**), having the sequence **S2** at C-3 of the aglycone were

380 previously isolated from *Lysimachia* species. They may therefore be considered as
381 chemotaxonomic markers for this family, and provide chemical support for phylogenetic
382 analyses, which suggest the transfer of the genus *Lysimachia* to the family Myrsinaceae. It
383 worth to note that *Lysimachia* saponins possess a second tetrasaccharide C-3 linked chain, in
384 which a rhamnose can replace the xylose at C-2 of glcII (Dai et al., 2017, Podolak et al., 2013b).
385 The present study reinforces previous reports which indicated that sapogenols with 13 β ,28-
386 epoxy-bridge are the predominant triterpenoid skeleton in species of the Myrsinaceae,
387 including the genus *Anagallis* and can be considered as a chemotaxonomic marker for this
388 plant family (Foubert et al., 2008). A branched four-unit sugar chain **S2**, with arabinose
389 substituted at C-2 with glucose and at C-4 with glucose and terminal xylose, seems to be
390 typical for these Primulaceae saponins.

391 In conclusion, in our phytochemical research on *A. monelli* ssp. *inifolia*, the chemotaxonomic
392 significance associated with *Anagallis* was discussed. Our results confirmed the richness of
393 *Anagallis* species in oleanan-type glycosides and showed that the sequence **S1** (glc-(1 \rightarrow 4)-
394 [xyl-(1 \rightarrow 2)-]glc-(1 \rightarrow 4)-[glc-(1 \rightarrow 2)-]ara-) can be suggested as a chemotaxonomic marker
395 for the genus *Anagallis*, but that proposal needs to be confirmed since only the species
396 *Anagallis arvensis* had been studied before this work. Finally, our phytochemical results
397 increase the knowledge on saponins of the genus *Anagallis* and its family Primulaceae and
398 stimulate to evaluate the biological activities of these saponins.

399

400 **3. Experimental**

401 *3.1 General experimental procedures*

402 The values of the optical rotations were measured by an Anton Paar MCP 5100 polarimeter
403 (Graz, Austria). 1D- and 2D-NMR spectra (^1H - and ^{13}C -NMR, ^1H - ^1H COSY, ROESY, HSQC
404 and HMBC) were recorded on a Bruker Avance III-600 spectrometer (Karlsruhe, Germany)
405 equipped with a 5 mm TCI cryoprobe. 2D-NMR experiments were performed using a standard
406 Bruker microprograms (TopSpin 4.0.6 software). HR-ESI-MS experiments were performed on
407 a Waters SYNAPT G2-Si High Resolution Q-TOF Mass Spectrometer equipped with
408 electrospray ionization (ESI) source (Waters Corp., Manchester, UK). Flash chromatography
409 was performed on a GRACE Reveleris X1 system (Flawil, Switzerland) equipped with a
410 Reveleris Navigator and dual UV and ELSD detection using Grace® cartridges (silica gel or
411 RP-C₁₈). Preparative high-performance liquid chromatography (HPLC) was performed on a
412 Gilson PLC 2050 (Saint-Avé, France) equipped with Gilson Glider software, Armen pump and
413 Ecom UV detector, using a RP-C₁₈ column (Interchim uptisphere strategy C18-HQ, 5 μm ,
414 250x21.2 mm). The mobile phase was composed of H₂O with TFA (0.0025%)/CH₃CN with a
415 flow rate 250 mL/min. The chromatograms were monitored at 205, 215 and 360 nm. Semi-
416 preparative HPLC was performed on an Agilent LC Series instrument (1200 Infinity Series –
417 1220, Les Ulis, France) equipped with an agilent G1329A sample injector, Jasco CO-4060

418 column oven (Lisses, France), agilent G1311A pump, Ultimate DAD3000 thermofisher detector
419 (Villebon sur Yvette, France) and chromeleon® 7.2 software. An RP-C₁₈ prep column
420 (Interchim uptisphere strategy C18(2), 5µm, 250x10 mm, Montlucon, France). The mobile
421 phase for semi-preparative HPLC was a mixture of H₂O with TFA (0.0025%) and CH₃CN with
422 a flow rate of 5 mL/min. The chromatograms were monitored at 205, 215 and 360 nm.
423 Analytical HPLC experiments were performed using a ThermoFisher Ultimate 3000 (Thermo
424 Fischer Scientific, Villebon sur Yvette, France), equipped with an LPG 3400 SD pump, a WPS
425 3000 SL injector and a UV-DAD-3000 detector with Chromeleon® software version 6.8 and an
426 Interchim uptisphere strategy C18(2) column, 5µm, 250x10 mm, using the same eluent as
427 semi-preparative HPLC with a flow rate of 1 mL/min. The chromatograms were monitored at
428 205, 215 and 360 nm. Thin layer chromatography (TLC) was carried out on silica gel plates
429 (Merck 60 F₂₅₄, Darmstadt, Germany) and visualized under UV lamps at 254 and 366 nm, then
430 by spraying with 50% H₂SO₄, followed by heating. All solvents used for Flash chromatography
431 were of analytical grade (Carlo Erba Reactifs SDS, Val de Reuil, France), and solvents used
432 for analytical, semi-preparative and preparative HPLC were of HPLC grade (Carlo Erba
433 Reactifs SDS, Val de Reuil, France). Trifluoroacetic acid (TFA) was purchased from Carlo Erba
434 Reactifs SDS (Val de Reuil, France).

435 3.2 Plant material

436 The aerial parts of *Anagallis monelli* ssp. *linifolia* (L.) Maire were collected at Bouililhet, in the
437 province of Oum el Bouaghi, northeast Algeria (latitude 35°43'53.1"N, longitude 6°41'34.0"E
438 and altitude of 970 m) in May 2019. The plant was authenticated by Mr. Kamel Kabouche. A
439 voucher specimen (LOST.Am.05.9.19) was preserved in the LOST laboratory of Université
440 des Freres Mentouri-Constantine 1, Algeria.

441 3.3. Extraction and isolation

442 The dried powder of the aerial parts of *Aanagallis monelli* ssp. *linifolia* (L.) Maire (1 kg) was
443 macerated in 70% EtOH (4 × 5 L, 24 h) at room temperature. After filtration and removal of the
444 solvent by evaporation under reduced pressure, the dried 70% EtOH extract (300g, 30% yield)
445 was dissolved in H₂O and then partitioned successively with EtOAc and *n*-BuOH. The *n*-BuOH
446 extract (70 g) was fractionated by Diaion HP-20 resin column (4.3 40 cm), which was eluted
447 with H₂O-MeOH (25, 50 and 100%, each 2 L), to obtain fractions A (10g), B (6g) and C (40g),
448 respectively. Fraction C (40 g) (the saponin-containing fraction) was applied to vacuum liquid
449 chromatography (VLC) over silica gel, using as eluent a mixture of CHCl₃-MeOH-H₂O (9:1:0,
450 8:2:0, 7:3:0, 7:3:0.1, 7:3:0.2, 7:3:0.5 and 100% MeOH) to give 7 fractions (C1-C7). The C4, C6
451 and C7 fractions (5.4 g), (3.9 g) and (5.1 g) were subjected to flash chromatography on RP-
452 C₁₈, eluted by a gradient of (20→60% CH₃CN, in 35 min), to afford the fractions C4₁-C4₃₂, C6₁-
453 C6₃₂ and C7₁-C7₃₁, respectively. Fractions C4₂₁ (1.1 g), C4₂₂₋₂₄ (600 mg) and C4₂₆₋₂₇ (500 mg)
454 were purified by silica gel flash chromatography, using a gradient system of CHCl₃-MeOH-H₂O
455 (8:2:0 → 7:3:2, in 45 min) to give 15 fractions C4_{21F1}-C4_{21F8}, C4_{22-24F1}-C4_{22-24F14} and C4_{26-27F1}-

456 C₄_{26-27F10}, respectively. Fraction C₄_{21F7-8} (130 mg) was purified by flash chromatography over
457 RP-C₁₈, eluted by a gradient of (25→60% CH₃CN, in 35 min), to obtain 4 fractions C₄_{21F7-8f1-4},
458 including compound **1** (32.4mg) as a pure compound in fraction C₄_{21F7-8f3}. The purification of
459 fraction C₄_{21F7-8f4} (78.5 mg) was realised by semi-prep. HPLC with a gradient of (30→70%
460 CH₃CN, in 30 min) to give compounds **11** (2.5mg, *t_R* 15.8 min), **5** (5.3mg, *t_R* 17.1 min) and **7**
461 (1.1mg, *t_R* 26.1 min). Fractions C₄_{21F10-11} and C₄_{22-24F14f5-6} were purified separately by
462 preparative HPLC using the same gradient (30→50% CH₃CN, in 60 min) to afford compounds
463 **6** (1.5mg), **2** (2.7mg) and **18** (1.3mg) and compounds **17** (1.5mg), **19** (2.4mg), **14** (10.2mg), **20**
464 (1.4mg) and **8** (2.8mg), respectively. Fractions C₄_{26-27F4} and C₆₁₆ were purified separately by
465 semi-prep HPLC with a gradient of (20→80% CH₃CN, in 45 min) to yield compounds **15**
466 (2.8mg, *t_R* 8.1 min), **9** (1.5mg, *t_R* 11.8 min), and **3** (2.2mg, *t_R* 12.8 min) from C₄_{26-27F4} and
467 compounds **12** (1.0mg, *t_R* 18.1 min), **10** (1.4mg, *t_R* 17.6 min), and **13** (1.5mg, *t_R* 19.4 min) from
468 C₆₁₆. The fractions C₆₁₄ and C₇₂₂ were purified by preparative HPLC, eluted by the gradient
469 (20→30% CH₃CN, in 45 min) to give compounds **23** (1.4mg), **22** (1.2mg), and **4** (3.0mg) from
470 C₆₁₄, whereas the gradient (30→60% CH₃CN, in 45 min) was used for C₇₂₂ to give compounds
471 **16** (1.4mg) and **21** (1.8mg).

472 3.4. Monelloside A (**1**)

473 Amorphous white powder, [α]_D²⁰ -9.5 (c 0.10, MeOH) ; ¹H (600 MHz, CD₃OD) and ¹³C (150
474 MHz, CD₃OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z*: 1061.5537 [M-H]⁻ (calcd for
475 C₅₂H₈₅O₂₂, 1061.5532).

476 3.5. Monelloside B (**2**)

477 Amorphous white powder, [α]_D²⁰ -3.0 (c 0.10, MeOH) ; ¹H (600 MHz, CD₃OD) and ¹³C (150
478 MHz, CD₃OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z*: 1223.6071 [M-H]⁻ (calcd for
479 C₅₈H₉₅O₂₇, 1223.6061).

480 3.6. Monelloside C (**3**)

481 Amorphous white powder, [α]_D²⁰ -0.9 (c 0.20, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
482 MHz, CD₃OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z* : 1103.5629 [M-H]⁻ (calcd for
483 C₅₄H₈₇O₂₃, 1103.5638).

484 3.7. Monelloside D (**4**)

485 Amorphous white powder; [α]_D²⁰ + 3.2 (c 0.28, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
486 MHz, CD₃OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z* : (1253.6169 [M-H]⁻ (calcd for
487 C₅₉H₉₇O₂₈, 1253.6166).

488 3.8. Monelloside E (**5**)

489 Amorphous white powder; [α]_D²⁰ + 12.8 (c 0.25, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
490 MHz, CD₃OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z* :1091.5646 [M-H]⁻ (calcd for
491 C₅₃H₈₇O₂₃, 1091. 5638).

492 3.9. Monelloside F (**6**)

493 Amorphous white powder; $[\alpha]^{20}_D + 1.3$ (c 0.15, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
494 MHz, CD_3OD) data: see Tables 3 and 4 ; HR-ESI-MS m/z : 1295.6633 $[\text{M-H}]^-$ (calcd for
495 $\text{C}_{62}\text{H}_{103}\text{O}_{28}$, 1295.6636).

496 3.10. Monelloside G (7)

497 Amorphous white powder; $[\alpha]^{20}_D + 10.0$ (c 0.11, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
498 MHz, CD_3OD) data: see Tables 3 and 4 ; HR-ESI-MS m/z : 1133.6107 $[\text{M-H}]^-$ (calcd for
499 $\text{C}_{56}\text{H}_{93}\text{O}_{23}$, 1133.6108).

500 3.11. Monelloside H (8)

501 Amorphous white powder, $[\alpha]^{20}_D -9.3$ (c 0.14, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
502 MHz, CD_3OD) data: see Tables 3 and 4; HR-ESI-MS m/z : 1221.5913 $[\text{M-H}]^-$ (calcd for
503 $\text{C}_{58}\text{H}_{93}\text{O}_{27}$, 1221.5904).

504 3.12. Monelloside I (9)

505 Amorphous white powder; $[\alpha]^{20}_D -5.5$ (c 0.11, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
506 MHz, CD_3OD) data: see Tables 3 and 4; HR-ESI-MS m/z : 1059.5370 $[\text{M-H}]^-$ (calcd for
507 $\text{C}_{52}\text{H}_{83}\text{O}_{22}$, 1059.5376).

508 3.13. Monelloside J (10)

509 Amorphous white powder; $[\alpha]^{20}_D + 15.7$ (c 0.14, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
510 MHz, CD_3OD) data: see Tables 5 and 6 ; HR-ESI-MS m/z : 1221.5911 $[\text{M-H}]^-$ (calcd $\text{C}_{58}\text{H}_{93}\text{O}_{27}$,
511 1221.5904).

512 3.14. Monelloside K (11)

513 Amorphous white powder; $[\alpha]^{20}_D -1.2$ (c 0.25, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
514 MHz, CD_3OD) data: see Tables 5 and 6 ; HR-ESI-MS m/z : 1059.5367 $[\text{M-H}]^-$ (calcd for
515 $\text{C}_{52}\text{H}_{83}\text{O}_{22}$, 1059.5376).

516 3.15. Monelloside L (12)

517 Amorphous white powder; $[\alpha]^{20}_D -1.7$ (c 0.12, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
518 MHz, CD_3OD) data: see Tables 5 and 6 ; HR-ESI-MS m/z : 1089.5490 $[\text{M-H}]^-$ (calcd for
519 $\text{C}_{53}\text{H}_{85}\text{O}_{23}$, 1089.5482).

520 3.16. Monelloside M (13)

521 Amorphous white powder; $[\alpha]^{20}_D + 2.5$ (c 0.12, MeOH); ^1H (600 MHz, CD_3OD) and ^{13}C (150
522 MHz, CD_3OD) data: see Tables 5 and 6 ; HR-ESI-MS m/z : 927.4966 $[\text{M-H}]^-$ (calcd for
523 $\text{C}_{47}\text{H}_{75}\text{O}_{18}$, 927.4953).

524 3.17. Acid hydrolysis :

525 The acid hydrolysis of fraction C (200 mg) rich in saponins was realized with 35 mL of 2 N TFA
526 (trifluoroacetic acid, aqueous solution) at 90 °C for 4h. After extraction with CH_2Cl_2 (10 mL \times
527 3), the aqueous phase was concentrated under vacuum to obtain the sugar residues (100 mg).
528 Three sugars was confirmed by comparison on TLC with pure samples of glucose, xylose and
529 arabinose, using (MeCOEt:iso-PrOH:Me₂CO:H₂O, 20:10:7 :2). The purification of sugars by
530 preparative TLC using the same solvent system afford L-arabinose [5.9 mg, $R_f = 0.52$, $[\alpha]^{20}_D$

531 +31.7 (c 0.5, H₂O)]; D-glucose [10 mg, R_f = 0.46, [α]²⁰_D +56 (c 0.9, H₂O)] and D-xylose [3.2 mg,
532 R_f = 0.63, [α]²⁰_D +15.3 (c 0.3, H₂O)].

533

534 **Declaration of competing interest**

535 The authors declare that they have no known competing financial interests or personal
536 relationships that could have appeared to influence the work reported in this paper.

537

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544

545 **Appendix A. Supplementary data**

546 The original spectra of monellosides A-M, including ¹H and ¹³C NMR, 2D NMR, and HR-ESI-
547 MS are given as supplementary data

548

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727 *Lysimachia candida*. Zhongcaoyao 33(6), 481-483.
728
729

730 **List of Figure caption**

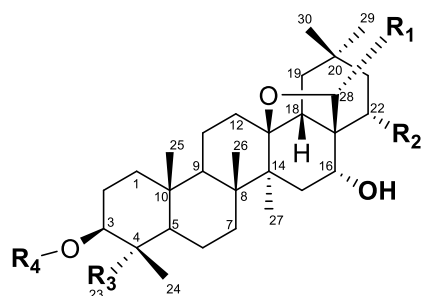
731 **Fig. 1.** Structures of compounds **1–13** isolated from *Anagallis monelli* ssp. *linifolia*

732 **Fig. 2.** Key HMBC and ROESY correlations for compound **1**.

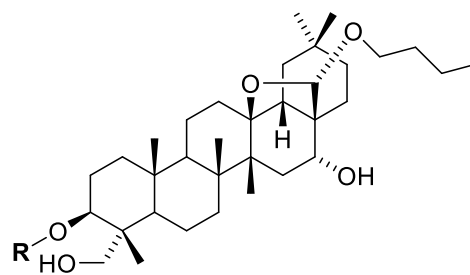
733 **Fig. 3.** Key HMBC correlations for compound **10**.

734 **Fig. 4.** The common carbohydrate chain linked on the C-3 of 13,28-epoxy-3,16-oleananediol
735 derivatives skeleton structure of species in the Myrsinaceae and Primulaceae

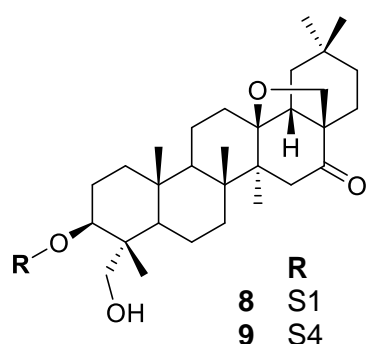
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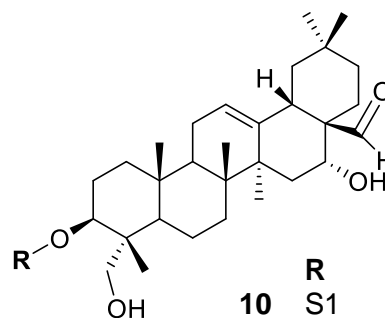
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1	H	OH	CH ₃	S2
2	H	OH	CH ₃	S1
3	H	H	CH ₂ OH	S3
4	OCH ₃	H	CH ₂ OH	S1
5	OCH ₃	H	CH ₂ OH	S2



	R
6	S1
7	S2



	R
8	S1
9	S4



	R
10	S1
11	S2
12	S5
13	S6

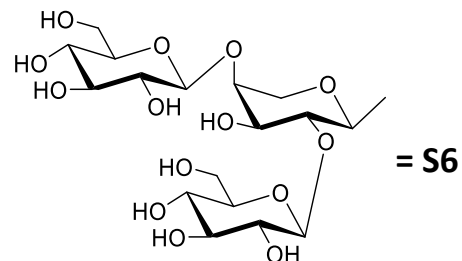
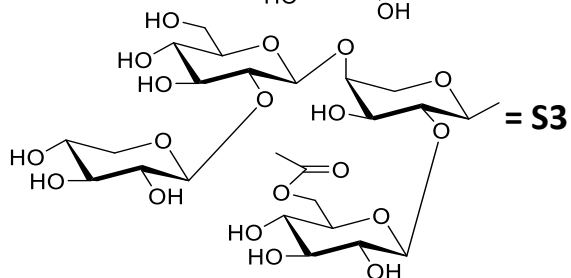
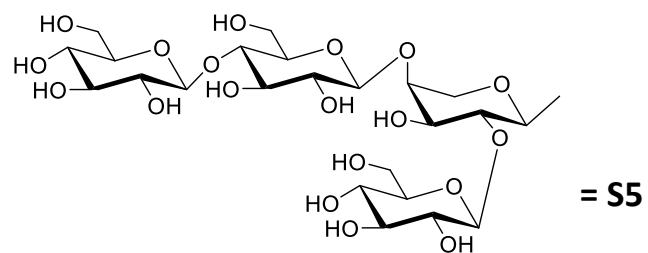
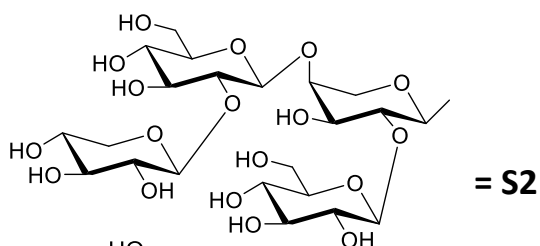
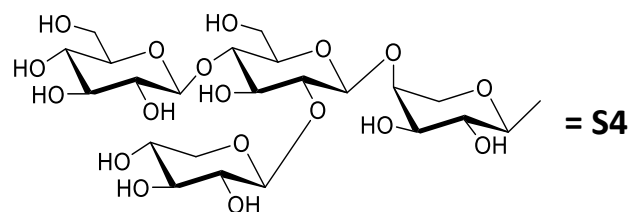
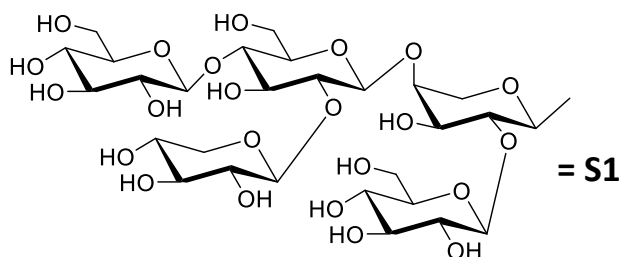


Fig. 1. Structures of compounds 1–13 isolated from *Anagallis monelli* ssp. *Linifolia*

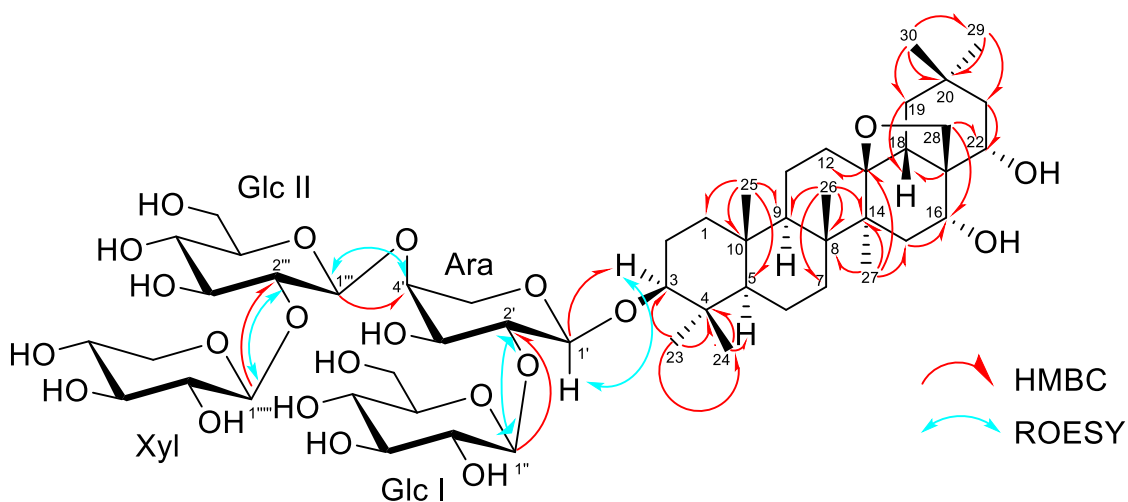


Fig. 2. Key HMBC and ROESY correlations for compound **1**.

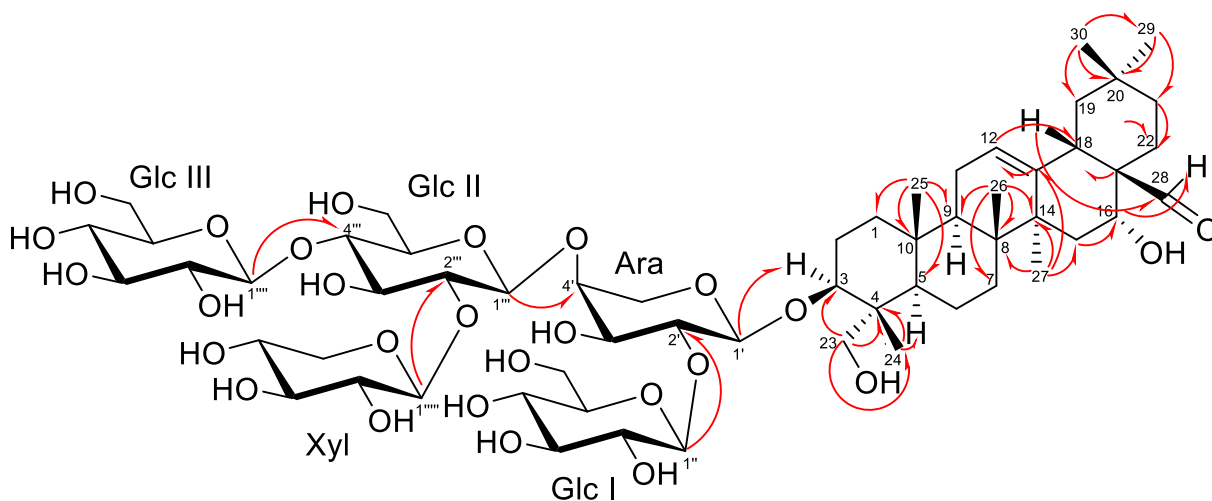


Fig. 3. Key HMBC correlations for compound **10**.

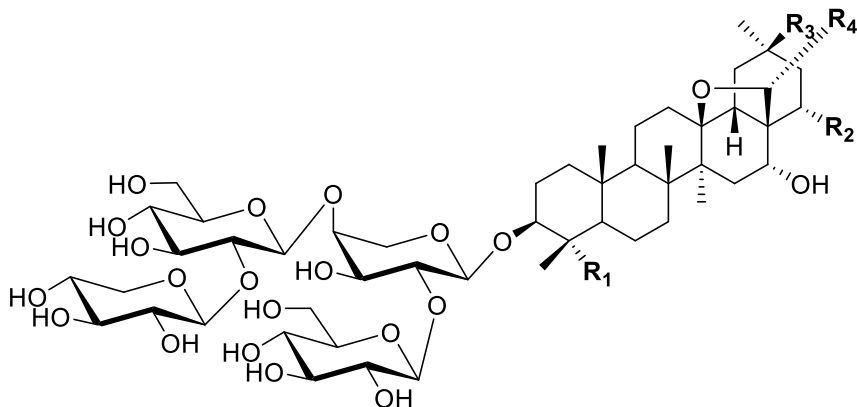


Fig. 4. The common carbohydrate chain linked on the C-3 of 13,28-epoxy-3,16-oleananediol derivatives skeleton structure of species in the Myrsinaceae and Primulaceae

Table 1

¹³C NMR and ¹H NMR spectroscopic data of aglycone moieties for compounds 1–5 in CD₃OD.^{a, b}

Position	1		2		3		4		5	
	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H
1	38.8	0.98 m 1.77 m	38.8	0.97 m 1.77 m	38.5	0.95 m 1.75 m	38.6	0.96 m 1.77 m	38.6	0.96 m 1.77 m
2	25.8	1.76 m 1.88 m	25.8	1.76 m 1.87 m	25.3	1.77 t (12.5) 1.88 m	25.2	1.80 m 1.89 m	25.2	1.80 m 1.89 m
3	90.0	3.16 dd (11.7, 4.3)	90.0	3.16 dd (11.7, 4.3)	81.9	3.58 dd (11.9, 4.7)	82.5	3.62 m	82.5	3.62 m
4	39.2	–	39.2	–	42.9	–	42.9	–	42.9	–
5	55.4	0.74 dd (9.4, 4.5)	55.4	0.74 dd (11.3, 2.8)	46.4	1.19 m	46.8	1.17 dd (9.9, 2.7)	46.8	1.17 dd (9.9, 2.7)
6	17.4	1.47 m 1.54 m	17.3	1.46 m 1.54 m	16.8	1.41 m 1.45 m	16.9	1.42 m 1.46 m	16.9	1.42 m 1.46 m
7	33.7	1.26 t (13.4) 1.56 m	33.7	1.25 m 1.56 m	33.2	1.19 m 1.67 td (13.2, 3.8)	33.2	1.19 m 1.69 m	33.2	1.19 m 1.69 m
8	42.1	–	42.1	–	41.8	–	41.9	–	41.9	–
9	50.0	1.28 m	49.9	1.27 m	49.9	1.32 m	50.0	1.29 m	50.0	1.29 m
10	36.4	–	36.4	–	36.2	–	36.2	–	36.2	–
11	18.5	1.49 m 1.64 dd (12.5, 4.4)	18.5	1.49 m 1.64 dd (12.7, 3.4)	18.5	1.49 m 1.64 m	18.6	1.47 m 1.69 m	18.6	1.47 m 1.69 m
12	32.1	1.31 m 2.06 m	32.1	1.31 m 2.06 m	31.9	1.29 m 2.07 m	32.3	1.32 m 2.01 m	32.3	1.32 m 2.01 m
13	86.9	–	86.9	–	87.0	–	87.6	–	87.6	–
14	44.5	–	44.5	–	44.0	–	42.7	–	42.7	–
15	35.5	1.28 m 2.09 m	35.5	1.27 m 2.08 m	35.6	1.23 d (15.7) 2.13 m	35.7	1.22 d (15.2) 2.01 m	35.7	1.22 d (15.2) 2.01 m
16	69.8	4.33 d (5.0)	69.8	4.32 br s	76.7	3.90 d (4.7)	75.8	3.83 d (5.2)	75.8	3.83 d (5.2)
17	49.1	–	49.1	–	43.9	–	47.9	–	47.9	–
18	50.7	1.47 dd (14.0, 2.5)	50.7	1.46 dd (12.5, 2.1)	51.0	1.52 dd (14.0, 4.7)	46.7	1.71 m	46.7	1.71 m
19	37.6	1.15 m 2.46 dd (14.1, 12.3)	37.6	1.15 m 2.45 dd (13.9, 12.5)	38.4	2.40 dd (14.0, 12.2) 1.21 d (12.6)	38.4	1.17 m 2.30 dd (14.5, 12.0)	38.4	1.17 m 2.30 dd (14.5, 12.0)
20	32.5	–	32.5	–	31.0	–	31.0	–	31.0	–
21	44.7	1.40 dd (12.3, 4.9) 2.20 t (12.3)	44.7	1.39 m 2.19 t (11.7)	36.0	1.15 m 2.10 m	36.0	1.15 m 2.06 t (13.3)	36.0	1.15 m 2.06 t (13.3)
22	74.2	3.74 dd (11.6, 4.9)	74.2	3.72 m	30.8	1.52 td (13.2, 4.4) 1.79 d (13.2)	25.1	1.98 m 1.42 dm (12.4)	25.1	1.98 m 1.42 dm (12.4)
23	27.0	1.08 s	27.0	1.07 s	62.9	3.21 d (11.2) 3.79 d (11.2)	63.2	3.30 d (11.5) 3.73 d (11.5)	63.2	3.30 d (11.5) 3.73 d (11.5)
24	15.3	0.87 s	15.3	0.86 s	11.9	0.68 s	11.7	0.73 s	11.7	0.73 s
25	15.3	0.92 s	15.3	0.92 s	15.8	0.93 s	15.8	0.96 s	15.8	0.96 s
26	17.4	1.17 s	17.4	1.17 s	17.4	1.17 s	17.5	1.20 s	17.5	1.20 s
27	18.7	1.28 s	18.7	1.28 s	18.5	1.27 s	18.4	1.25 s	18.4	1.25 s
28	76.7	3.67 m	76.7	3.67 s	77.3	3.14 d (7.6) 3.52 d (7.6)	105.5	4.19 s	105.5	4.19 s
29	32.3	1.00 s	32.3	1.00 s	32.5	0.97 s	32.5	0.94 s	32.5	0.94 s
30	24.6	0.98 s	24.6	0.96 s	23.5	0.93 s	23.5	0.90 s	23.5	0.90 s
OAc					171.5	–	–	–	–	–
CH ₃					19.6	2.08 s	53.9	3.33 s	53.9	3.33 s

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 500 or 600 MHz (¹H) and at 125 or 150 MHz (¹³C).

Table 2

¹³C NMR and ¹H NMR spectroscopic data of the sugar moieties for compounds 1–5 in CD₃OD.^{a, b}

Position	1		2		3		4		5	
	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H
C3	Ara		Ara		Ara		Ara		Ara	
1'	104.2	4.41 d (6.6)	104.2	4.42 d (6.3)	103.4	4.43 d (6.8)	103.3	4.46 d (6.3)	103.3	4.46 d (5.8)
2'	78.0	3.82 m	78.0	3.82 m	79.9	3.73 dd (8.2, 6.8)	78.3	3.79 m	78.3	3.79 dd (8.7, 5.8)
3'	72.9	3.81 m	72.9	3.81 m	73.3	3.79 d (9.0)	73.1	3.80 m	73.0	3.80 dd (5.8, 3.3)
4'	78.8	3.90 m	78.8	3.90 m	78.7	3.92 m	78.8	3.91 m	78.7	3.91 m
5'	64.4	3.55 dd (12.5, 3.8) 4.23 dd (12.5, 2.8)	64.3	3.55 dm (12.5) 4.22 d (12.5)	64.5	3.53 d (12.5) 4.19 dd (11.9, 5.4)	64.4	3.55 dm (12.6) 4.22 dd (12.6, 2.9)	64.4	3.55 dd (12.7, 1.2) 4.24 dd (12.7, 2.5)
	Glc I		Glc I		Glc I		Glc I		Glc I	
1''	103.0	4.71 d (7.7)	103.1	4.43 d (8.3)	103.7	4.71 d (7.6)	102.9	4.73 d (7.7)	102.9	4.73 d (7.8)
2''	74.6	3.21 dd (8.9, 7.7)	73.5	3.24 t (8.9)	74.6	3.25 t (7.6)	74.6	3.24 t (9.4, 7.7)	74.6	3.22 dd (9.1, 7.8)
3''	76.4	3.39 t (9.3)	76.6	3.30 m	76.4	3.41 t (9.0)	76.4	3.39 t (9.2)	76.4	3.40 t (9.1)
4''	70.6	3.19 t (9.2)	69.9	3.32 m	69.8	3.35 t (9.5)	70.4	3.20 t (9.3)	70.4	3.21 t (9.1)
5''	76.7	3.29 m	76.7	3.30 m	76.8	3.46 m	76.7	3.29 m	76.7	3.29 m
6''	61.9	3.62 dd (12.0, 6.7) 3.86 dd (12.0, 2.0)	61.0	3.68 dd (10.4, 2.6) 3.89 m	63.5	4.20 dd (11.5, 2.5) 4.34 dd (11.5, 2.0)	61.7	3.62 dd (12.1, 6.7) 3.87 dd (12.1, 2.1)	61.7	3.62 dd (11.9, 6.6) 3.86 dd (11.9, 4.3)
	Glc II		Glc II		Glc II		Glc II		Glc II	
1'''	103.4	4.54 d (7.8)	103.3	4.57 d (8.8)	103.2	4.56 d (7.2)	103.2	4.58 d (7.8)	103.3	4.54 d (7.8)
2'''	83.7	3.41 dd (8.9, 7.8)	83.1	3.48 t (9.0)	83.6	3.44 dd (9.0, 7.2)	83.1	3.48 dd (9.0, 7.8)	83.7	3.40 t (8.5)
3'''	76.2	3.57 t (8.9)	74.7	3.75 t (9.0)	76.2	3.56 t (9.0)	74.6	3.75 t (9.1)	76.2	3.55 t (9.3)
4'''	69.7	3.35 m	78.5	3.63 t (8.9)	69.6	3.37 t (9.0)	78.5	3.63 t (9.2)	69.7	3.35 t (9.4)
5'''	76.7	3.34 m	75.1	4.30 m	76.5	3.28 m	75.1	3.44 dt (9.6, 3.6)	76.5	3.30 m
6'''	61.2	3.69 dd (12.4, 6.2) 3.88 d (12.4)	60.3	3.89 m 3.88 m	61.1	3.68 dd (12.2, 5.4) 3.87 d (12.2, 2.2)	60.3	3.88 m 3.88 m	61.1	3.68 dd (11.9, 5.6) 3.87 dd (11.9, 4.1)
			Glc III				Glc III			
1''''			102.9	4.71 d (7.3)			103.1	4.43 d (7.9)		
2''''			74.6	3.10 dd (8.9, 7.3)			73.5	3.23 dd (9.0, 7.9)		
3''''			76.4	3.39 m			76.4	3.37 t (9.0)		
4''''			70.6	3.18 t (8.9)			69.9	3.32 m		
5''''			76.7	3.30 m			76.7	3.35 m		
6''''			61.9	3.60 dd (11.0, 4.7) 3.86 d (11.0)			61.0	3.68 dd (11.7, 5.2) 3.89 dd (11.7, 1.9)		
	Xyl		Xyl		Xyl		Xyl		Xyl	
1'''''	105.9	4.53 d (7.6)	105.9	4.56 d (7.8)	105.9	4.55 d (6.8)	106.0	4.56 d (7.5)	105.9	4.53 d (7.8)
2'''''	74.5	3.27 dd (8.7, 7.6)	74.6	3.26 t (7.8)	74.5	3.27 dd (9.0, 6.8)	74.6	3.26 dd (8.2, 7.5)	74.5	3.27 dd (9.2, 7.8)
3'''''	76.5	3.39 m	76.1	3.37 m	76.3	3.36 t (9.0)	76.1	3.37 m	76.2	3.37 t (9.2)
4'''''	69.6	3.54 m	69.6	3.52 m	69.5	3.52 m	69.6	3.52 m	69.6	3.53 m
5'''''	66.0	3.32 m 4.01 dd (11.3, 5.2)	66.0	3.32 m 4.01 dd (11.5, 5.7)	66.0	3.30 m 3.99 dd (11.5, 5.4)	66.0	3.32 m 4.00 dd (11.4, 5.2)	66.0	3.31 m 4.00 dd (11.4, 5.4)
OAc					171.5	-				
CH ₃					19.6	2.08 S				

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 500 or 600 MHz (¹H) and at 125 or 150 MHz (¹³C).

Table 3¹³C NMR and ¹H NMR spectroscopic data of aglycone moieties for compounds **6–9** in CD₃OD.^{a, b}

Position	6		7		8		9	
	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H
1	38.6	0.95 m 1.77 m	38.6	0.95 m 1.77 m	38.5	0.97 m 1.78 m	38.5	0.95 m 1.76 m
2	25.2	1.81 m 1.89 m	25.2	1.82 t (12.9) 1.90 m	25.2	1.81 m 1.90 m	25.1	1.77 m 1.90 m
3	82.5	3.62 m	82.5	3.63 m	82.3	3.63 d (8.9)	81.7	3.63 dd (12.0, 4.8)
4	42.9	–	42.9	–	42.9	–	42.7	–
5	46.8	1.17 m	46.8	1.18 m	46.6	1.18 dd (10.2, 2.9)	46.7	1.19 dd (9.5, 5.3)
6	16.9	1.42 m 1.46 m	16.9	1.42 m 1.43 m	16.7	1.47 m 1.49 m	16.7	1.47 m 1.49 m
7	33.2	1.18 m 1.68 m	33.2	1.19 m 1.68 m	32.7	1.11 dt (12.7, 3.2) 1.62 m	32.8	1.11 dt (12.7, 3.2) 1.60 m
8	41.9	–	41.9	–	42.5	–	42.5	–
9	50.0	1.28 m	49.9	1.28 m	49.8	1.26 m	49.8	1.26 m
10	36.2	–	36.2	–	36.1	–	36.1	–
11	18.6	1.47 m 1.68 m	18.6	1.47 m 1.68 m	18.3	1.57 m 1.74 m	18.3	1.57 m 1.74 m
12	32.4	1.31 m 2.02 m	32.4	1.31 t (13.6) 2.01 m	31.1	1.51 m 2.05 m	31.1	1.51 m 2.05 m
13	87.4	–	87.4	–	86.2	–	86.2	–
14	42.7	–	42.7	–	49.7	–	49.7	–
15	35.7	1.22 m 2.01 m	35.7	1.22 m 2.01 m	45.0	1.81 d (15.8) 2.80 d (15.8)	45.0	1.81 d (15.8) 2.80 d (15.8)
16	75.9	3.82 d (4.6)	75.9	3.82 m	214.3	–	214.3	–
17	48.0	–	48.4	–	55.8	–	55.8	–
18	46.7	1.76 m	46.7	1.77 m	54.5	2.02 dd (12.1, 2.7)	54.5	2.01 dd (13.8, 3.2)
19	38.5	1.17 m 2.28 t (12.8)	38.5	1.18 m 2.29 dd (14.2, 12.6)	39.6	1.36 t (13.8) 1.48 m	39.6	1.36 t (13.8) 1.48 m
20	31.1	–	31.0	–	31.1	–	31.1	–
21	36.1	1.16 m 2.04 m	36.1	1.17 m 2.05 m	35.0	1.23 m 1.57 m	35.0	1.23 m 1.57 m
22	25.2	1.41 m 2.04 m	25.2	1.41 m 2.05 m	24.2	1.25 m 2.11 d (11.7)	24.2	1.25 m 2.11 dt (12.7, 2.1)
23	63.2	3.30 m 3.72 m	63.2	3.30 m 3.73 d (11.6)	63.1	3.29 m 3.73 d (11.3)	63.1	3.31 m 3.65 d (11.1)
24	11.7	0.73 s	11.7	0.73 s	11.7	0.74 s	11.7	0.75 s
25	15.9	0.95 s	15.9	0.96 s	15.5	0.96 s	15.6	0.96 s
26	17.5	1.20 s	17.5	1.20 s	17.8	1.28 s	17.8	1.28 s
27	18.3	1.24 s	18.3	1.25 s	20.8	1.08 s	20.8	1.08 s
28	104.1	4.28 brs	104.1	4.28 brs	74.7	3.48 d (8.4) 3.93 d (8.4)	74.7	3.48 d (8.5) 3.93 d (8.5)
29	32.5	0.95 s	32.3	0.94 s	32.4	0.95 s	32.4	0.95 s
30	23.4	0.91 s	23.5	0.91 s	22.4	0.91 s	22.4	0.91 s
a	66.5	3.33 m 3.70 m	66.5	3.34 m 3.71 m				
b	31.7	1.56 t (6.5)	31.7	1.56 t (6.7)				
c	19.1	1.42 m	19.1	1.42 m				
d	12.8	0.95 t (6.9)	12.8	0.95 t (6.9)				

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 600 and 150 MHz (¹H and ¹³C).

Table 4

¹³C NMR and ¹H NMR spectroscopic data of the sugar moieties for compounds **6–9** in CD₃OD.^{a, b}

Position	6		7		8		9	
	$\bar{\delta}_C$	$\bar{\delta}_H$	$\bar{\delta}_C$	$\bar{\delta}_H$	$\bar{\delta}_C$	$\bar{\delta}_H$	$\bar{\delta}_C$	$\bar{\delta}_H$
C3	Ara		Ara		Ara		Ara	
1'	103.3	4.46 d (6.3)	103.3	4.46 d (5.4)	103.3	4.46 d (6.1)	105.1	4.28 d (7.2)
2'	78.3	3.79 m	78.3	3.79 m	78.3	3.79 m	72.7	3.43 m
3'	73.1	3.80 m	73.1	3.80 m	73.1	3.80 m	73.2	3.47dd (9.2,4.0)
4'	78.8	3.91 m	78.7	3.90 m	78.7	3.91 m	80.0	3.85 dm (2.8)
5'	64.4	3.55 dd (11.9, 1.5) 4.22 d (11.9)	64.4	3.54 m 4.24 dd (12.7, 3.3)	64.4	3.55 m 4.22 dd (12.7, 2.8)	65.5	3.56 d (12.8) 4.20 dd (12.8, 2.0)
	Glc I		Glc I		Glc I			
1''	102.9	4.73 d (7.7)	102.9	4.73 d (7.6)	102.9	4.72 d (7.7)		
2''	74.6	3.21 dd (9.2, 7.7)	74.6	3.22 dd (8.8, 7.6)	74.6	3.21 dd (8.9, 7.7)		
3''	76.4	3.39 m	76.4	3.39 t (9.2)	76.4	3.39 t (9.4)		
4''	70.4	3.20 m	70.4	3.21 t (9.1)	70.4	3.20 t (9.4)		
5''	76.7	3.29 m	76.7	3.30 m	76.7	3.29 m		
6''	61.7	3.62 dd (11.9, 4.5) 3.86 dd (11.9, 2.3)	61.7	3.63 m 3.86 dd (11.9, 4.8)	61.7	3.61 dd (11.9, 6.6) 3.86 dd (11.9, 2.0)		
	Glc II		Glc II		Glc II		Glc II	
1'''	103.2	4.58 d (7.7)	103.3	4.54 d (7.8)	103.2	4.58 d (7.7)	103.7	4.56 d (8.0)
2'''	83.1	3.48 dd (8.9, 7.7)	83.8	3.40 t (8.7)	83.1	3.49 dd (8.9, 7.7)	84.1	3.50 dd (9.0, 8.0)
3'''	74.6	3.74 m	76.1	3.56 m	74.6	3.74 t (9.1)	74.5	3.74 t (9.0)
4'''	78.5	3.62 t (9.2)	69.7	3.35 m	78.5	3.63 t (9.4)	78.3	3.66 t (9.2)
5'''	75.1	3.44 m	76.5	3.34 m	75.1	3.44 m	75.0	4.43 m
6'''	60.3	3.87 m 3.89 m	60.1	3.69 dd (11.9, 5.7) 3.87 d (11.9, 4.8)	60.3	3.87 m 3.89 m	60.3	3.87 m 3.89 m
	Glc III				Glc III		Glc III	
1''''	103.1	4.43 d (8.6)			103.1	4.43 d (7.9)	103.1	4.44 d (7.6)
2''''	73.5	3.24 m			73.5	3.24 dd (8.8, 7.9)	73.5	3.24 dd (8.8, 7.6)
3''''	76.4	3.38 t (8.9)			76.1	3.37 m	76.4	3.37 t (9.0)
4''''	69.9	3.33 m			69.9	3.33 m	69.9	3.32 m
5''''	76.6	3.34 m			76.7	3.43 m	76.6	3.30 m
6''''	61.0	3.67 dd (11.6, 3.4) 3.88 m			60.9	3.68 dd (11.9, 5.6) 3.89 m	61.0	3.68 dd (12.0, 5.6) 3.89 dd (12.0, 2.0)
	Xyl		Xyl		Xyl		Xyl	
1'''''	106.0	4.56 d (7.9)	105.9	4.52 d (7.6)	105.9	4.56 d (7.6)	106.6	4.52 d (7.6)
2'''''	74.6	3.26 dd (8.9, 7.9)	74.5	3.27 t (7.9)	74.6	3.25 t (8.1)	74.8	3.25 dd (9.6, 7.6)
3'''''	76.1	3.37 m	76.2	3.35 m	76.4	3.37 m	76.2	3.37 t (9.6)
4'''''	69.6	3.53 m	69.5	3.52 m	69.5	3.52 m	69.4	3.52 m
5'''''	66.0	3.32 t (11.0) 4.00 dd (11.0, 5.5)	66.0	3.31 m 4.00 dd (11.3, 5.4)	66.0	3.32 m 4.00 dd (11.4, 5.4)	65.7	3.28 t (11.6) 4.00 dd (11.6, 5.6)

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 600 MHz and 150 (¹H and ¹³C).

Table 5¹³C NMR and ¹H NMR spectroscopic data of the aglycone moieties for compounds **10–13** in CD₃OD.^{a, b}

Position	10		11		12		13	
	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H
1	38.2	1.00 m 1.65 m	38.2	1.00 m 1.66 m	38.2	1.00 m 1.66 dt (13.3, 3.6)	38.2	1.00 m 1.65 m
2	25.0	1.80 t (14.3) 1.89 m	25.0	1.80 t (12.9) 1.89 m	24.9	1.79 m 1.89 m	24.9	1.80 t (12.6) 1.89 m
3	82.5	3.63 m	82.4	3.64 dd (11.7, 4.3)	82.4	3.63 m	82.4	3.64 m
4	42.8	–	42.7	–	42.7	–	42.7	–
5	46.9	1.24 d (11.7)	46.8	1.24 d (11.7)	46.8	1.24 d (11.7)	46.8	1.24 d (10.9)
6	17.4	1.37 m 1.52 m	17.4	1.38 m 1.52 m	17.3	1.37 m 1.51 m	17.4	1.37 m 1.52 m
7	32.3	1.31 m 1.69 m	32.2	1.31 m 1.70 m	32.2	1.31 m 1.69 m	32.2	1.31 m 1.70 m
8	39.4	–	39.4	–	39.4	–	39.4	–
9	46.7	1.69 t (8.5)	46.6	1.69 m	46.7	1.68 t (8.8)	46.7	1.69 m
10	36.2	–	36.2	–	36.2	–	36.2	–
11	23.1	1.93 m 1.94 m	23.1	1.93 m 1.94 m	23.1	1.92 m 1.94 m	23.1	1.92 m 1.93 m
12	123.0	5.42 t (3.6)	123.0	5.42 brs	123.0	5.42 t (3.5)	123.0	5.42 t (3.7)
13	142.9	–	142.9	–	142.9	–	142.9	–
14	41.4	–	41.4	–	41.4	–	41.4	–
15	34.6	1.40 m 1.82 dd (14.7, 4.0)	34.5	1.40 m 1.82 d (12.9)	34.5	1.40 dd (14.2, 2.7) 1.82 dd (14.2, 3.6)	34.5	1.40 m 1.82 dd (14.5, 3.7)
16	72.5	4.29 t (3.6)	72.4	4.29 brs	72.4	4.29 t (3.5)	72.4	4.28 t (3.7)
17	50.8	–	50.8	–	50.9	–	50.8	–
18	40.4	2.68 dd (14.5, 3.9)	40.4	2.68 dd (13.8, 3.3)	40.4	2.68 dd (14.2, 4.4)	40.4	2.68 dd (13.5, 4.3)
19	46.1	1.14 dd (13.3, 4.9) 2.27 t (13.3)	46.0	1.13 d (13.8) 2.27 t (13.8)	46.0	1.14 dd (12.7, 3.6) 2.27 t (14.2)	46.0	1.13 dd (13.5, 4.3) 2.27 t (13.5)
20	29.9	–	29.9	–	29.9	–	29.9	–
21	34.2	1.20 m 1.94 m	34.2	1.21 m 1.94 m	34.2	1.20 m 1.93 m	34.2	1.21 m 1.94 m
22	25.9	1.56 m	25.9	1.55 m	25.9	1.54 m	25.9	1.54 m
23	63.3	3.30 m 3.73 d (11.6)	63.2	3.30 m 3.73 d (11.4)	63.2	3.30 m 3.70 d (11.4)	63.3	3.31 m 3.68 d (11.6)
24	11.9	0.75 s	11.9	0.74 s	11.9	0.73 s	11.9	0.73 s
25	15.1	1.00 s	15.1	1.00 s	15.1	1.00 s	15.1	1.01 s
26	16.5	0.76 s	16.5	0.75 s	16.5	0.76 s	16.5	0.75 s
27	25.9	1.41 s	25.9	1.41 s	25.9	1.41 s	25.9	1.41 s
28	205.6	9.23 s	205.6	9.23 s	205.6	9.23 s	205.6	9.23 s
29	32.0	0.93 s	32.0	0.92 s	32.0	0.92 s	32.0	0.92 s
30	23.1	0.98 s	23.1	0.98 s	23.1	0.98 s	23.1	0.98 s

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 600 and 150 MHz (¹H and ¹³C).

Table 6

¹³C NMR and ¹H NMR spectroscopic data of the sugar moieties for compounds **10–13** in CD₃OD.^{a, b}

Position	10		11		12		13	
	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H	δ_C	δ_H
C3	Ara		Ara		Ara		Ara	
1'	103.3	4.46 d (6.4)	103.3	4.46 d (5.1)	102.9	4.56 d (6.2)	102.9	4.55 d (5.3)
2'	78.3	3.79 m	78.2	3.79 m	77.5	3.89 t (7.7)	77.6	3.89 m
3'	73.1	3.80 m	73.1	3.80 m	72.2	3.87 dd (8.2, 2.3)	72.3	3.87 m
4'	78.8	3.91 m	78.8	3.90 m	77.3	3.98 dd (6.6, 2.7)	77.2	3.99 m
5'	64.4	3.55 d (12.6) 4.21 dd (12.6, 2.6)	64.5	3.55 m 4.24 dd (12.8, 3.3)	63.4	3.57 dd (12.4, 1.9) 4.15 dd (12.4, 4.3)	63.5	3.57 dd (12.4, 1.9) 4.15 dd (12.4, 4.2)
	Glc I		Glc I		Glc I		Glc I	
1''	103.0	4.73 d (7.7)	102.9	4.73 d (7.6)	103.1		103.0	
2''	74.6	3.21 dd (9.2, 7.7)	74.5	3.21 dd (8.9, 7.6)	74.4	4.66 d (7.8)	74.4	4.65 d (7.7)
3''	76.5	3.40 t (9.2)	76.4	3.39 t (8.9)	76.6	3.22 dd (8.9, 7.8)	76.6	3.22 dd (9.1, 7.7)
4''	70.4	3.20 t (9.2)	70.4	3.19 t (9.5)	70.4	3.37 t (8.9)	70.4	3.37 t (9.0)
5''	76.7	3.29 m	76.7	3.29 m	76.8	3.22 t (9.1)	76.9	3.22 t (9.5)
6''	61.8	3.62 dd (12.0, 6.6) 3.87 dd (12.0, 1.6)	61.8	3.60 dd (12.0, 4.9) 3.87 dd (12.0, 4.6)	61.6	3.28 m 3.63 dd (12.0, 6.7)	61.6	3.29 m 3.63 dd (11.8, 6.6)
	Glc II		Glc II		Glc II		Glc II	
1'''	103.2	4.59 d (7.7)	103.4	4.54 d (7.8)	104.1	3.87 dd (12.0, 2.3)	104.3	3.87 dd (11.8, 2.4)
2'''	83.1	3.42 dd (9.2, 7.7)	83.8	3.40 m	73.8		74.0	
3'''	74.6	3.75 t (9.2)	76.2	3.56 t (9.2)	74.8		76.5	
4'''	78.5	3.64 t (9.5)	69.6	3.34 m	79.0	4.53 d (7.8)	70.1	4.53 d (7.7)
5'''	75.1	3.43 m	76.5	3.29 m	75.1	3.33 m	76.7	3.27 dd (8.9, 7.7)
6'''	60.3	3.87 d (12.4) 3.89 m	61.1	3.68 dd (11.7, 5.2) 3.88 m	60.4	3.51 t (8.9) 3.59 t (8.9) 3.42 m	61.3	3.35 m 3.29 m 3.29 m
	Glc III				Glc III			
1''''	103.1	4.43 d (7.9)			103.2	3.87 dm (12.5)		3.67 d (11.8, 5.5)
2''''	73.5	3.24 t (8.2)			73.5	3.91 d (12.5, 2.6)		3.88 d (11.8, 2.4)
3''''	76.5	3.39 t (8.9)			76.4			
4''''	69.9	3.33 m			69.9	4.42 d (7.8)		
5''''	76.7	3.36 m			76.6	3.23 dd (8.6, 7.7)		
6''''	61.0	3.68 dd (11.8, 5.5) 3.89 m			61.0	3.38 t (8.6) 3.32 m 3.35 m		
	Xyl		Xyl					
1'''''	106.0	4.56 d (7.9)	105.9	4.52 d (8.2)				3.68 dd (11.8, 5.5)
2'''''	74.6	3.27 dd (9.2, 7.7)	74.5	3.26 dd (9.5, 8.2)				3.89 dd (11.8, 2.3)
3'''''	76.1	3.37 t (9.2)	76.2	3.37 t (9.5)				
4'''''	69.6	3.53 m	69.5	3.52 m				
5'''''	66.0	3.32 m 4.01 dd (11.3, 5.3)	66.0	3.32 m 4.00 dd (11.4, 4.9)				

^a in ppm, *J* in parentheses in Hz.^b NMR spectra recorded at 600 and 150 MHz (¹H and ¹³C).

Table 7: 3-O- β -D-xyl-(1 \rightarrow 2)-glc-(1 \rightarrow 4)-[glc-(1 \rightarrow 2)-]-arabinopyranosyl-13,28-epoxy-3,16-oleananediol derivatives skeleton of species in the Myrsinaceae and Primulaceae

R1	R2	R3	R4	Name of plant	Reference
Me	H	Me	H	<i>Cyclamen repandum</i> <i>Lysimachia sikokiana</i> <i>Lysimachia vulgaris</i> <i>Lysimachia clethroides</i> <i>Myrsine australis</i>	Dall'acqua et al., 2010 Kohda et al., 1989 Podolack et al., 1998 Podolack et al., 2013 Bloor and Qi, 1994
CH ₂ OH	H	Me	H	<i>Anagallis arvensis</i> <i>Cyclamen africanum</i> <i>Cyclamen repandum</i> <i>Lysimachia ciliata</i> <i>Lysimachia ephemerum</i> <i>Lysimachia heterogenea</i>	Glombitza et al., 1987b Bencharif-Betina et al., 2012 Dall'acqua et al., 2010 Podolack et al., 2013 Podolack et al., 2013 Huang et al., 2011
Me	H	CH ₂ OH	H	<i>Ardisia crenata</i> <i>Lysimachia thyrsoiflora</i>	Jia et al., 1994 Podolack et al., 2013
Me	H	CHO	H	<i>Androsace saxifragaefolia</i> <i>Ardisia crispa</i> <i>Cyclamen africanum</i> <i>Cyclamen libanoticum</i> <i>Cyclamen persicum</i> <i>Cyclamen repandum</i> <i>Cyclamen spp.</i> <i>Cyclamen trocopteranthum</i> <i>Lysimachia nummularia</i> <i>Lysimachia punctata</i> <i>Myrsine australis</i> <i>Myrsine pellucida</i> <i>Myrsine salicina</i> <i>Primula denticulata</i>	Waltho et al., 1986 Jansakul et al., 1987 Bencharif-Betina et al., 2012 El Hosry et al., 2014 El Hosry et al., 2014 Dall'acqua et al., 2010 Rezniek et al., 1989 Mihci-Gaidi et al., 2010 Podolack et al., 2013 Podolack et al., 2013 Bloor and Qi, 1994 Lavaud et al., 1994 Bloor and Qi, 1994 Ahmad et al., 1988
CH ₂ OH	H	CHO	H	<i>Cyclamen coum var. coum</i> <i>Cyclamen mirabilis</i>	Calis et al., 1997b Calis et al., 1997a
Me	H	COOH	H	<i>Cyclamen hederifolium</i>	Altunkeyik et al., 2012
Me	H	Me	OH	<i>Myrsine australis</i>	Bloor and Qi, 1994
Me	OH	Me	H	<i>Anagallis arvensis</i>	Shoji et al., 1994b
Me	OAc	Me	H	<i>Anagallis arvensis</i> <i>Lysimachia ciliata</i> <i>Lysimachia ephemerum</i>	Shoji et al., 1994a Podolack et al., 2013 Podolack et al., 2013
Me	OH	Me	OH	<i>Lysimachia capillipes</i>	Tian et al., 2006