

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

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

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

1. Introduction

 The genus *Anagallis* currently belongs to the Primulaceae family, although recent studies based on DNA sequences of three chloroplast genes and morphology have suggested its placement in the Myrsinaceae as *Lysimachia* (Källersjö et al., 2000; Manns and Anderberg, 2009 and 2011). *Anagallis* contained about 28 species growing as mainly annual herbs, distributed in Africa, Madagascar, Europe and South America. This genus is represented in Algeria by four species *A. arvensis* L., *A. monelli* L*., A. repens* Pomel. and *A. enella* L. (Quezel and Santa, 1963). The plant *Anagallis monelli* is an endemic herb from North Africa and is represented in Algeria by two different subspecies, *Anagallis monelli* ssp. *collina* (Schousb.) Maire and *Anagallis monelli* ssp. *linifolia* (L.) Maire (Quezel and Santa, 1963). Among the studied species of the genus *Anagallis*, *A. arvensis* and *A. foemina* have been used in traditional medicine in Navarre (Spain) against skin injuries like burns and wounds (López et al., 2011). The whole plant of *A. arvensis* has been used for liver complications in Taiwan, for skin diseases in Italy and for fish poisoning in rural areas of Nepal (Yasmeen et al., 2020). Chemical investigations on *Anagallis* spp. have been mainly characterized by the presence of saponins (Aliotta et al., 1992 ; Amoros and Girre, 1987; Glombitza and Kurth, 1987a and b; Shoji et al., 1994a and b; Soberón et al., 2017), pentacyclic triterpenes (Aliotta et al., 1992; De Napoli et al., 1992; Heitz et al.,1971), flavonoids and polyphenols (Ammar et al., 2008; Ishikura, 1981; Kawashty et al., 1998; Rastogi and Norula, 1980), sterols (Rastogi and Norula, 1980), in addition to alkaloids and quinones (Saxena and Rao, 2021). Triterpenoid saponins were found in a wide variety of higher plants and display a wide range of pharmacological activities, including haemolytic, expectorant, anti-inflammatory, hypolipidemic, gastroprotective, immunomodulatory and antimicrobial properties (Netala et al., 2015; Podolak et al., 2010). The potential anticancer activity of saponins has been suggested by their cytotoxic, cytostatic, pro-apoptotic and anti-invasive effects (Koczurkiewicz et al., 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 al., 2008; Podolak et al., 2013a).

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

2.1 Isolation and structural elucidation

 The 70% EtOH extract from the aerial parts of *A. monelli* ssp. *linifolia* was sequentially partitioned with EtOAc and *n*-BuOH, respectively. The *n*-BuOH soluble fraction was subjected to a Diaion HP-20 resin chromatography to give three fractions (A-C). The saponin-containing fraction (C), was subjected to further column chromatography to give ten known compounds 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) and mass spectrometry (HR-ESI-MS) and by comparison with literature data. The monosaccharides of monellosides A-M (**1**-**13**) obtained by acid hydrolysis of an aliquot of the saponin-containing fraction (C) were identified as L-arabinose, D-glucose and D-xylose by comparison on TLC with authentic samples followed by measurement of their optical rotation values after purification on TLC.

81 **Compound 1** was obtained as an amorphous white powder. Its molecular formula was 82 determined as $C_{52}H_{86}O_{22}$ 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 $[\delta_H$ 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 16*α*-100 configuration of hydroxyl group was evident from the small coupling constant ${}^{3}J_{H-16/H-15}$ 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 $(\delta_H 3.16)$ 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''''), *i*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 correlations observed between H-1'/H-3' and H-1'/H-5' in the ROESY spectrum of **1**, indicated the *α*-anomer configuration of the arabinose unit. The large coupling constants from anomeric protons (>7 Hz) of the xylose and glucoses units, indicated their *β*-configurations (Liang et al., 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). HMBC correlation between H-1' and C-3 indicated that the arabinose unit was linked to C-3 of 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 confirming the interglycosidic linkage and the point of attachment of the tetra-saccharide at the C-3 of the aglycone were observed between H-1'/ H-3, H-1'' / H-2', H-1''' / H-4' and H-1''''' / H- 2''' (Fig. 2). According to the above results, the structure of compound **1** was elucidated as 3- *O*-*β*-D-xylopyranosyl-(1→2)-*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-priverogenin B, named monelloside A.

Compound 2, isolated as white amorphous powder, had a molecular formula of $C_{58}H_{96}O_{27}$ 130 determined by the negative-ion HR-ESI-MS (1223.6061 [M-H], calcd 1223.6071), and differed from **1** by 162 amu corresponding to a supplementary hexosyl group. Comparison of the NMR data of **2** with those of **1** (Tables 1 and 2) showed that they shared the same aglycone but differed by the presence in **2** of an additional hexose unit identified as *β*-D-glucopyranose 134 (GlcIII, δ_{H-1} ⁻⁻ 4.71). The HMBC correlation between H-1⁻¹¹ and C-4⁻¹ (δ_C 78.5) indicated the linkage of GlcIII to GlcII on C-4''' (Shoji et al., 1994a). Therefore, the structure of compound **2** was established as 3-*O*-*β*-D-glucopyranosyl-(1→4)-[*β*-D-xylopyranosyl-(1→2)-]*β*-D- glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-priverogenin B, named monelloside B.

 Compound 3 was obtained as a white amorphous powder. Its molecular formula was 140 determined as $C_{54}H_{88}O_{23}$ on the basis of its negative-ion HR-ES-IMS (1103.5629 [M-H], calcd 1103.5638) and it differed from **1** by 42 amu corresponding to an additional acetyl group. Extensive 1D and 2D-NMR analysis (Tables 1 and 2) showed that compound **3** differ from **2** by the absence of a hydroxyl group at C-22 and the presence of a hydroxyl group at C-23 of the aglycone, as in anagalligenin B (Mahato et al., 1991, Shoji et al., 1994b). Additional signals 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 (δ_{H-6"} 4.20 (1H, dd, J = 11.5, 2.5 Hz); indicating that **3** was an acetylated derivative of desglucoanagalosine B (**20**) (Shoji et al., 1994b). According to the above results, the structure of compound **3** was elucidated as 3-*O*-*β*-D-xylopyranosyl- (1→2)-*β*-D-glucopyranosyl-(1→4)-[6-*O*-acetyl-*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-anagalligenin B, named monelloside C

152 **Compound 4** had a molecular formula of C₅₉H₉₈O₂₈, 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 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 *δ*^C 82.5, 75.8, respectively in the HSQC spectrum. The disappearance of the hydroxyl group at 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 *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) 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 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 carbon signals of the aglycone were accomplished by extensive 2D-NMR analyses which led to the elucidation of the aglycone part of **4** as 13,28-epoxy-(3*β*,16*α*,23)-trihydroxy-28-methoxy- oleanane, which differ from anagalligenin B (Mahato et al., 1991) by the presence of a methoxy group at C-28. Thus, compound **4** was identified as 3-*O*-*β*-D-glucopyranosyl-(1→4)-[*β*-D-

xylopyranosyl-(1→2)-]*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl(1→2)-]*α*-L-

arabinopyranosyl-13,28-epoxy-3*β*,16*α*,23-trihydroxy-28-methoxy-oleanane or 3-*O*-*β*-D-

glucopyranosyl-(1→4)-[*β*-D-xylopyranosyl-(1→2)-]*β*-D-glucopyranosyl-(1→4)-[*β*-D-

 glucopyranosyl(1→2)-]*α*-L-arabinopyranosyl-28-methoxy-anagalligenin B, named monelloside D.

 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 1091. 5638). Comparison of the NMR data of **5** with those of **4** (Tables 1 and 2) and analysis of the NMR spectra showed that compounds **4** and **5** had the same aglycone moiety (28- 182 methoxy-anagalligenin B), while comparison of the ${}^{1}H$ and ${}^{13}C$ NMR values of the oligosaccharide part of **5** with those of **1** indicated that **5** had the same tetrasaccharide moiety as in **1**, linked to C-3. Location of all proton and carbon signals was achieved by extensive 2D- NMR analyses, which elucidated compound **5** as 3-*O*-*β*-D-xylopyranosyl-(1→2)-*β*-D- glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-13,28-epoxy-3*β*,16*α*,23-trihydroxy-28-methoxy-oleanane, or 3-*O*-*β*-D-xylopyranosyl-(1→2)-*β*-D-

glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-28-methoxy-

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). Comparison of the ¹H and ¹³C NMR data of **6** to those of **4** and analysis of the 2D NMR spectra of **6** showed that both possessed the same penta-saccharide chain (Tables 2 and 4), while slight differences were observed in the aglycone part, notably those due to the D and E rings (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 $\lceil \delta c \rceil$ 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 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 evidences led to the assignment of **6** as 3-*O*-*β*-D-glucopyranosyl-(1→4)-[*β*-D-xylopyranosyl- (1→2)-]-*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-13,28- epoxy-3*β*,16*α*,23-trihydroxy-28-*n*-butyloxy-oleanane, or 3-*O*-*β*-D-glucopyranosyl-(1→4)-[*β*-D-

xylopyranosyl-(1→2)-]-*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-

arabinopyranosyl-28-*n*-butyloxy-anagalligenin B, named monelloside F.

 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]. calcd 1133.6108), and corresponds to the loss of 162 amu compared to **6**. Comparison of the H- and ¹³C-NMR data of **7** with those of **6** and **5** showed that the NMR data of **7** exhibited 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 showed that **7** and **5** shared the same tetra-saccharide chain (Tables 2 and 4). These data were confirmed by extensive 2D-NMR analyses and the assignments of all proton and carbon signals of **7**, leading to the elucidation of its structure as 3-*O*-*β*-D-xylopyranosyl-(1→2)-*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-13,28-epoxy-

3*β*,16*α*,23-trihydroxy-28-*n*-butyoxy-oleanane, named monelloside G.

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 of **8** with those of **4** (Tables 2 and 4) and analysis of the NMR spectra showed that **8** had the same penta-saccharide moiety as **4**. Most of the aglycone NMR signals were directly attributed by comparison with the corresponding signals of **4** (Tables 1 and 3), but detailed analysis of the NMR spectra showed dissimilarities in the ring D and different chemical shifts for C-14, C- 15, C-16, C-27, and C-28, due to the replacement of the hydroxyl at C-16 by a carbonyl group 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→4)-[*β*-D-xylopyranosyl-
- 231 (1→2)-]*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)]-*α*-L-arabinopyranosyl-
- 232 anagalligenone B, named monelloside H.

Compound 9, isolated as an amorphous white powder, had a molecular formula of $C_{52}H_{84}O_{22}$, 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 ${}^{1}H$ and ${}^{13}C$ NMR data (Tables 3 and 4), were identical to those of compound **8**, except for signals due to the *β*- D-glucopyranosyl (GlcI) unit linked in C-2' position of the arabinosyl unit which disappeared in compound **9**. Accordingly, the structure of **9** was elucidated as 3-*O*-*β*-D-glucopyranosyl-(1→4)- [*β*-D-xylopyranosyl-(1→2)-]*β*-D-glucopyranosyl-(1→4)-*α*-L-arabinopyranosyl- anagalligenone B, named monelloside I.

- 242 **Compound 10** was obtained as an amorphous white powder. Its molecular formula was 243 determined as $C_{58}H_{94}O_{27}$ 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 δ_{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→4)-[*β*-D-260 xylopyranosyl-(1→2)-]*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→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 $C_{52}H_{84}O_{22}$ 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

 data of the saccharide part of **11** were identical to those of **1** (Tables 2 and 6). Extensive 2D- 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 (δ_{C-1'} 103.3) on the HMBC spectrum confirmed that the glycosidic moiety was linked to C-3. According to the above evidences, the structure of **11** was elucidated as 3-*O*-*β*-D-xylopyranosyl-(1→2)-*β*-D-glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-3*β*,16*α*,23-trihydroxyolean-12-en-28-al, named monelloside K.

 Compound 12 was obtained as an amorphous white powder. Its molecular formula was 274 determined as $C_{53}H_{86}O_{23}$ 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) showed that it had the same aglycone but differed in its saccharide units. The NMR spectroscopic data of **12** were almost identical to those of **10,** except for the absence of the signals of *β*-D-xylopyranosyl. Detailed analysis of the 2D-NMR spectra of **12** led to the identification of its structure as 3-*O*-*β*-D-glucopyranosyl-(1→4)-*β*-D-glucopyranosyl-(1→4)-[*β*-

D-glucopyranosyl-(1→2)-]*α*-L-arabinopyranosyl-3*β*,16*α*,23-trihydroxyolean-12-en-28-al,

named monelloside L.

 Compound 13 was obtained as an amorphous white powder. Its molecular formula was 283 determined as $C_{47}H_{76}O_{18}$ 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 a similarity of these two compounds, with the exception of the disappearance of the glucose (GlcIII) and xylose units in the glycosidic part of **13**. Extensive analysis of the 1D- and 2D-NMR spectra of **13** established that the glycosidic sequence was 3-*O*-glc-(1→4)-[glc-(1→2)-]ara. Thus, the structure of compound **13** was elucidated as 3-*O*-*β*-D-glucopyranosyl-(1→4)-*β*-D-

glucopyranosyl-(1→4)-[*β*-D-glucopyranosyl-(1→2)]*α*-L-arabinopyranosyl-3*β*,16*α*,23-

trihydroxyolean-12-en-28-al, named monelloside M.

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

2.1 Chemophenetic significance

Phytochemical studies of the genus *Anagallis* revealed its richness of triterpenoids

saponins with an oleanane skeleton. Until now, 38 compounds were isolated from only one

species *Anagallis arvensis* (Aliotta et al., 1992; Amoros et al., 1987; De Napoli et al., 1992;

Glombitza and Kurth, 1987a; Kitagawa et al., 1976; Mahato et al., 1991; Shoji et al., 1994).

The present study, which was carried out on the Algerian subspecies *Anagallis monelli* ssp*.*

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

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

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

 In the present species, we have identified a four-unit branched sugar chain linked at C-3 of the aglycone **S2** { xyl-(1→2)-glc-(1→4)-[glc-(1→2)-]ara-} (Fig. 4). This chain was found in saponins of *Anagallis* (Glombitza et al., 1987b, Shoji et al., 1994a and b), *Lysimachia* (Kohda et al., 1989, Podolack et al., 2013), *Cyclamen* (Altunkeyik et al., 2012; Bencharif-Betina et al., 2012, Dall'acqua et al., 2010, El Hosry et al., 2014), *Ardisia* (Jia et al., 1994), *Myrsine* (Bloor and QI, 1994)*,* and *Androsace* genera (Waltho et al., 1986) (Table 7). This tetrasaccharide sequence **S2** can be substituted with glucose at C-4 of glcII or glcI in the case of *Anagallis* saponins (Shoji et al., 1994a and b; Soberón et al., 2017), at C-3 or C-6 of glcII in the case of *Cyclamen* saponins (Çalis et al., 1997), or at C-4 of the terminal glucose (glcI) in the case of *Androsace saxifragaefolia* saponins (Waltho et al., 1986); however, it was the only study found for the genus *Androsace*. In addition, this common chain was substituted with xylose at C-4 of glcII in the case of *Lysimachia* saponins (Podolak et al., 2013b), or with rhamnose at C-3 of glcII in saponins of *Ardisia gigantifolia* (Mu et al., 2010). Many saponins contain rhamnose in the sequence of the sugar moiety in the genera *Ardisia* and *Myrsine* (Foubert al., 2008). For *Primula denticulata*, it was the only species that had similarity in saponin content with the genus *Anagallis*, the other species had completely different osidic chains (Foubert al., 2008). We can conclude that the genera *Anagallis*, *Lysimachia,* and *Cyclamen* are very similar with respect to chemotaxonomic markers and thus can confirm their place in the Myrsinoideae subfamily.

 Triterpene saponins identified so far from the genus *Lysimachia*, generally have oleanane-371 derived sapogenols of two structural types: I. 13β,28-epoxy and II. Δ¹²-17-CH₂OH. Compounds of type I are considered very rare and are found almost exclusively in the Myrsinaceae and Primulaceae families (Foubert et al., 2008; Podolack et al., 2013). In the present study, eighteen type I and one type II triterpene saponins were obtained from *Anagallis monelli* ssp*. linifolia*. It is interesting to observe that the known compounds **14**-**23** were reported previously in various species belonging exclusively to genera traditionally assigned to the family Myrsinaceae or genera which were re-located to this family from the Primulaceae, i.e., *Anagallis, Lysimachia* and *Cyclamen*. Four of the ten known compounds isolated form *A. monelli* ssp. *linifolia* (**14**-**16**, and **23**), having the sequence **S2** at C-3 of the aglycone were previously isolated from *Lysimachia* species. They may therefore be considered as chemotaxonomic markers for this family, and provide chemical support for phylogenetic analyses, which suggest the transfer of the genus *Lysimachia* to the family Myrsinaceae. It worth to note that *Lysimachia* saponins possess a second tetrasacharide C-3 linked chain, in which a rhamnose can replace the xylose at C-2 of glcII (Dai et al., 2017, Podolak et al., 2013b). The present study reinforces previous reports which indicated that sapogenols with 13*β*,28- epoxy-bridge are the predominant triterpenoid skeleton in species of the Myrsinaceae, including the genus *Anagallis* and can be considered as a chemotaxonomic marker for this plant family (Foubert et al., 2008). A branched four-unit sugar chain **S2**, with arabinose substituted at C-2 with glucose and at C-4 with glucose and terminal xylose, seems to be typical for these Primulaceae saponins.

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

3. Experimental

3.1 General experimental procedures

 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 ¹³C-NMR, ¹H-¹H COSY, ROESY, HSQC and HMBC) were recorded on a Bruker Avance III-600 spectrometer (Karlsruhe, Germany) equipped with a 5 mm TCI cryoprobe. 2D-NMR experiments were performed using a standard Bruker microprograms (TopSpin 4.0.6 software). HR-ESI-MS experiments were performed on a Waters SYNAPT G2-Si High Resolution Q-TOF Mass Spectrometer equipped with electrospray ionization (ESI) source (Waters Corp., Manchester, UK). Flash chromatography was performed on a GRACE Reveleris X1 system (Flawil, Switzerland) equipped with a 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 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, 5µm, 414 250x21.2 mm). The mobile phase was composed of H_2O with TFA (0.0025%)/CH₃CN with a flow rate 250 mL/min. The chromatograms were monitored at 205, 215 and 360 nm. Semi- preparative HPLC was performed on an Agilent LC Series instrument (1200 Infinity Series – 1220, Les Ulis, France) equipped with an agilent G1329A sample injector, Jasco CO-4060 column oven (Lisses, France), agilent G1311A pump, Ultimate DAD3000 thermofisher detector 419 (Villebon sur Yvette, France) and chromeleon® 7.2 software. An RP- C_{18} prep column (Interchim uptisphere strategy C18(2), 5µm, 250x10 mm, Montlucon, France). The mobile 421 phase for semi-preparative HPLC was a mixture of H_2O with TFA (0.0025%) and CH₃CN with a flow rate of 5 mL/min. The chromatograms were monitored at 205, 215 and 360 nm. Analytical HPLC experiments were performed using a Thermofisher Ultimate 3000 (Thermo Fischer Scientific, Villebon sur Yvette, France), equipped with an LPG 3400 SD pump, a WPS 3000 SL injector and a UV-DAD-3000 detector with Chromeleon® software version 6.8 and an 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 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 were of analytical grade (Carlo Erba Reactifs SDS, Val de Reuil, France), and solvents used for analytical, semi-preparative and preparative HPLC were of HPLC grade (Carlo Erba Reactifs SDS, Val de Reuil, France). Trifluoroacetic acid (TFA) was purchased from Carlo Erba Reactifs SDS (Val de Reuil, France).

3.2 Plant material

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

- *3.3. Extraction and isolation*
- The dried powder of the aerial parts of *Aanagallis monelli* ssp. *linifolia* (L.) Maire (1 kg) was 443 macerated in 70% EtOH (4 x 5 L, 24 h) at room temperature. After filtration and removal of the solvent by evaporation under reduced pressure, the dried 70% EtOH extract (300g, 30% yield) was dissolved in H2O and then partitioned successively with EtOAc and *n*-BuOH. The *n*-BuOH 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), 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, 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 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 \rightarrow 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 \rightarrow 7:3:2, in 45 min) to give 15 fractions C_{421F1} -C 4_{21F8} , $C_{422-24F1}$ -C $4_{22-24F14}$ and $C_{426-27F1}$ -

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 \rightarrow 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 C421F7-8f3. The purification of 459 fraction C4_{21F7-8f4} (78.5 mg) was realised by semi-prep. HPLC with a gradient of (30 \rightarrow 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 C421F10-11 and C422-24F14f5-6 were purified separately by 462 preparative HPLC using the same gradient $(30\rightarrow50\% \text{ CH}_3\text{CN}, \text{in } 60 \text{ min})$ 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 C426-27F4 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 C426-27F4 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 (2030% CH3CN, in 45 min) to give compounds **23** (1.4mg), **22** (1.2mg)**,** and **4** (3.0mg) from 470 C6₁₄, whereas the gradient (30 \rightarrow 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, $[\alpha]^{20}$ -9.5 (*c* 0.10, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- MHz, CD3OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z*: 1061.5537 [M-H]- 474 (calcd for
- 475 C₅₂H₈₅O₂₂, 1061.5532).
- 476 *3.5. Monelloside B (2)*
- 477 Amorphous white powder, $\text{[}\alpha \text{]}^{20}\text{D}$ -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, $[\alpha]^{20}$ -0.9 (*c* 0.20, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- MHz, CD3OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z* : 1103.5629 [M-H]- 482 (calcd for
- 483 C₅₄H₈₇O₂₃, 1103.5638).
- 484 *3.7. Monelloside D (4)*
- 485 Amorphous white powder; $[\alpha]^{20}$ _D + 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; $[\alpha]^{20}$ _D + 12.8 (*c* 0.25, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- MHz, CD3OD) data: see Tables 1 and 2 ; HR-ESI-MS *m/z* :1091.5646 [M-H]- 490 (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, CD₃OD) and ¹³C (150
- 494 MHz, CD₃OD) data: see Tables 3 and 4; HR-ESI-MS m/z : 1295.6633 [M-H]⁻ (calcd for
- 495 C₆₂H₁₀₃O₂₈, 1295.6636).
- 496 *3.10. Monelloside G (7)*
- 497 Amorphous white powder; $[\alpha]^{20}$ _D + 10.0 (*c* 0.11, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- MHz, CD3OD) data: see Tables 3 and 4 ; HR-ESI-MS *m/z* : 1133.6107 [M-H]- 498 (calcd for
- 499 C₅₆H₉₃O₂₃, 1133.6108).
- 500 *3.11. Monelloside H (8)*
- 501 Amorphous white powder, $\left[\alpha\right]^{20}$ -9.3 (c 0.14, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- 502 MHz, CD₃OD) data: see Tables 3 and 4; HR-ESI-MS m/z : 1221.5913 [M-H]⁻ (calcd for
- 503 C58H93O27, 1221.5904).
- 504 *3.12. Monelloside I (9)*
- 505 Amorphous white powder; $\left[\alpha\right]^{20}$ -5.5 (c 0.11, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- 506 MHz, CD₃OD) data: see Tables 3 and 4; HR-ESI-MS m/z : 1059.5370 [M-H] (calcd for
- 507 C₅₂H₈₃O₂₂, 1059.5376).
- 508 *3.13. Monelloside J (10)*
- 509 Amorphous white powder; $[\alpha]^{20}$ _D + 15.7 (*c* 0.14, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- 510 MHz, CD₃OD) data: see Tables 5 and 6; HR-ESI-MS m/z : 1221.5911 [M-H]⁻ (calcd C₅₈H₉₃O₂₇, 511 1221.5904).
-
- 512 *3.14. Monelloside K (11)*
- 513 Amorphous white powder; $\text{[}\alpha \text{]}^{20}\text{D}$ -1.2 (*c* 0.25, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- 514 MHz, CD₃OD) data: see Tables 5 and 6 : HR-ESI-MS m/z : 1059.5367 [M-H]⁻ (calcd for
- 515 $C_{52}H_{83}O_{22}$, 1059.5376).
- 516 *3.15. Monelloside L (12)*
- 517 Amorphous white powder; $\left[\alpha\right]^{20}$ -1.7 (c 0.12, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150
- 518 MHz, CD₃OD) data: see Tables 5 and 6; HR-ESI-MS m/z : 1089.5490 [M-H]⁻ (calcd for
- 519 C₅₃H₈₅O₂₃, 1089.5482).
- 520 *3.16. Monelloside M (13)*
- 521 Amorphous white powder; $[\alpha]^{20}$ _D + 2.5 (*c* 0.12, MeOH); ¹H (600 MHz, CD₃OD) and ¹³C (150 522 MHz, CD₃OD) data: see Tables 5 and 6; HR-ESI-MS m/z : 927.4966 [M-H]⁻ (calcd for
- 523 $C_{47}H_{75}O_{18}$, 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 CH₂Cl₂ (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, [α]²⁰_D

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- **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

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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

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

^ain ppm, *J* in parentheses in Hz.

 b NMR spectra recorded at 500 or 600 MHz (1H) and at 125 or 150 MHz (13C).

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^ain ppm, *J* in parentheses in Hz.

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¹³C NMR and ¹H NMR spectroscopic data of aglycone moieties for compounds **6**–**9** in CD3OD.a, b

^ain ppm, *J* in parentheses in Hz.

 b NMR spectra recorded at 600 and 150 MHz (1 H and 13 C).

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

^a in ppm, *J* in parentheses in Hz.

 b NMR spectra recorded at 600 MHz and 150 (1 H and 13 C).

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

^ain ppm, *J* in parentheses in Hz.

b NMR spectra recorded at 600 and 150 MHz (1H and 13C).

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

^ain 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→2)-glc-(1→4)-[glc-(1→2)-]-arabinopyranosyl-13,28-epoxy-3,16-oleananediol derivatives skeleton of species in the Myrsinaceae and Primulaceae

