



Acylated oleanane-type saponins from *Ganophyllum giganteum*



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ABSTRACT

Five oleanane-type saponins, 3-O-β-D-glucuronopyranosylzanhic acid 28-O-β-D-xylopyranosyl-(1→3)-[α-L-rhamnopyranosyl-(1→2)]-(4-O-acetyl)-β-D-fucopyranosyl ester (**1**), 3-O-β-D-glucopyranosylzanhic acid 28-O-β-D-xylopyranosyl-(1→3)-[α-L-rhamnopyranosyl-(1→2)]-(4-O-acetyl)-β-D-fucopyranosyl ester (**2**), zanhic acid 28-O-β-D-xylopyranosyl-(1→3)-[α-L-rhamnopyranosyl-(1→2)]-(4-O-acetyl)-β-D-fucopyranosyl ester (**3**), zanhic acid 28-O-α-L-rhamnopyranosyl-(1→2)-4-O-[(3'-hydroxy-2'-methyl-butyroyloxy)-3-hydroxy-2-methyl-butyroyloxy]-β-D-fucopyranosyl ester (**4**), medicagenic acid 28-O-α-L-rhamnopyranosyl-(1→2)-4-O-[(3'-hydroxy-2'-methyl-butyroyloxy)-3-hydroxy-2-methyl-butyroyloxy]-β-D-fucopyranosyl ester (**5**), were isolated from the root barks of *Ganophyllum giganteum*. Compounds **4** and **5** possessed an unusual substitution of the C-4 position of the β-D-fucopyranosyl moiety by a C10 ester group formed by two symmetrical C5 nilic acid. From a chemotaxonomic point of view, their structures are in accordance with the previous saponins isolated from the Doratoxyleae tribe of the Sapindaceae family. Their cytotoxicity and anti-inflammatory activity were also evaluated.

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Introduction

In the order of the Sapindales, the Sapindaceae, known as the soapnut family, is an important family represented by some 150 genera with almost 2000 species (Delaude, 1993). According to the APG III classification, the Sapindaceae is divided into four subfamilies: Dodonaeoideae, Hippocastanoideae, Sapindoideae, and Xanthoceroideae for certain authors only (Buerki et al., 2010). Plants of this family are various, from trees to herbaceous plants, and many species possess an economical importance: in food industry with the lychee or rambutan, in phytotherapy with the genus *Paullinia* (Guarana), for their wood (*Acer*), as ornamentals (*Koelroetia*), and as detergents (*Sapindus*). The latter genera are rich in triterpene saponins, and their fruits were traditionally used in tropical countries to clean fabrics and to kill fish (Delaude, 1993). In the field of our researches on natural saponins, we have investigated the structure and biological activity of some saponins isolated from the roots of a Sapindaceae species named *Ganophyllum giganteum* (A. Chev.) Hauman. This tree, found in

the Democratic Republic of Congo, belongs to the Dodonaeoideae subfamily and the Doratoxyleae tribe, and is used in wooden factory (Delaude, 1993). *Ganophyllum giganteum* is also described in Gabonese traditional medicine and previous phytochemical works showed a good cytotoxic activity of an hydromethanolic extract of the leaves against human monocytes (Lamidi et al., 2005). Moreover, investigations of the crude saponin mixture resulted in the characterization of two aglycons: medicagenic acid and zanhic acid, which appeared as chemotaxonomic markers of the Doratoxyleae tribe (Delaude, 1993; Dimbi et al., 1984). In this paper, we described the isolation by successive chromatographic steps of five new triterpene saponins (**1–5**) from the aqueous methanolic extract of the root barks (Fig. 1). Their structural determination was performed by a detailed analysis of their spectral data including 600 MHz 2D-NMR (COSY, TOCSY, ROESY, HSQC, HMBC), and mass spectrometry. Cytotoxicity and anti-inflammatory activity of the compounds **2–5** were also evaluated.

Results and discussion

After extraction by MeOH 70% of the root bark of *G. giganteum*, the resulting extract was subjected to multiple chromatographic

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steps over silica gel to yield compounds **1–5**. The elucidation of their structures was performed mainly by 600 MHz 2D NMR analysis (^1H – ^1H COSY, TOCSY, ROESY, HSQC, HMBC), in combination with mass spectrometry (HRESIMS and FAB- and ESIMS). All saponins were isolated as white amorphous powders.

The HRESIMS of compound **1** established its molecular formula as $\text{C}_{55}\text{H}_{84}\text{O}_{26}$ and **1** showed in ESIMS (negative-ion mode) a quasi-molecular ion peak $[\text{M}-\text{H}]^-$ at m/z 1159 indicating a molecular weight of 1160.

The ^1H NMR spectrum of the aglycon part of **1** showed signals of six angular methyl groups as singlets (Table 1), one olefinic proton at δ_{H} 5.25 (*t*-like), three oxygens bearing methine protons at δ_{H} 4.00 (*d*, $J = 3.6$ Hz), 4.10 (*m*), and δ_{H} 4.30 (*m*). The HMBC correlation between one angular methyl group at δ_{H} 1.15 (*s*, H_3 -24) and δ_{C} 84.1 (C-3) confirmed the location of the first secondary alcoholic function at C-3. The COSY correlation between δ_{H} 4.00 (*d*, $J = 3.6$ Hz, H-3) and δ_{H} 4.10 (*m*) allowed the location of the second secondary alcoholic function at C-2. The C-16 position of the third one was deduced by the HMBC correlation between δ_{H} 4.30 (*m*) and δ_{C} 40.5 (C-18) and the COSY correlation between δ_{H} 4.30 (*m*) and δ_{H} 1.60 (H-15). A cross-peak in the ROESY spectrum between H-3 at δ_{H} 4.00 and the α -axial H-5 at δ_{H} 1.39 suggested a α -axial orientation of the H-3. The α -equatorial orientation of the H-2 was determined by the multiplicity and the coupling constant of H-3 as a doublet ($J = 3.6$ Hz) and from the connectivity observed in the ROESY spectrum between H-3 at δ_{H} 4.00 and H-2 at δ_{H} 4.10 ppm. For C-16, the ROESY correlations between δ_{H} 0.69 (*s*, H_3 -26) and δ_{H} 1.60 ($\text{H}_{\beta\text{ax}}$ -15), and between δ_{H} 1.60 ($\text{H}_{\beta\text{ax}}$ -15) and δ_{H} 4.30 (*m*, H-16), suggested a β -equatorial orientation of the H-16 and thus a α -axial orientation of the OH group. The structure of the aglycon of **1** was thus recognized as the triterpene zanhic acid, and was in

full agreement with literature data (Lavaud et al., 1998; Tava et al., 2005).

The monosaccharides obtained by acid hydrolysis of **1** were identified by comparison on TLC with authentic samples as glucuronic acid (GlcA), fucose (Fuc), xylose (Xyl) and rhamnose (Rha). The absolute configurations of the sugars were determined to be D for GlcA, Fuc and Xyl, and L for Rha, by GC analysis by a method previously described (Hara et al., 1987). The relatively large $^3J_{\text{H-1,H-2}}$ values of the GlcA, Fuc, and Xyl (7.3–8.1 Hz) in their pyranose form indicated a β anomeric orientation for GlcA, Fuc, and Xyl (Mimaki et al., 2004). The large $^1J_{\text{H-1,C-1}}$ value of the Rha (166 Hz) confirmed that the anomeric proton was equatorial (α -pyranoid anomeric form).

The ^1H NMR spectrum of **1** displayed in the sugar region signals of four anomeric protons at δ_{H} 5.37 (*d*, $J = 8.1$ Hz), 5.11 (*br s*), 4.31 (*d*, $J = 7.3$ Hz), and 4.26 (*d*, $J = 7.7$ Hz), which gave correlations, in the HSQC spectrum, with four anomeric carbon signals at δ_{C} 93.1, 100.5, 104.1 and 101.5, respectively. Complete assignments of the resonances of each sugar were achieved by extensive 2D NMR analyses (COSY, TOCY, HSQC, HMBC) (Table 2). Units of one β -D-fucopyranosyl, one α -L-rhamnopyranosyl, one β -D-xylopyranosyl, and one β -D-glucuronopyranosyl were thus identified. A cross-peak observed in the ROESY spectrum at δ_{H} 4.26 (*d*, $J = 7.7$ Hz, GlcA-1)/ δ_{H} 4.00 (*d*, $J = 3.6$, H-3), and a HMBC correlation at δ_{H} 4.26 (*d*, $J = 7.7$ Hz, GlcA-1)/ δ_{C} 84.1 (C-3), revealed a substitution at C-3 of the aglycon by a glucuronopyranosyl moiety. Moreover, the C-28 position at δ_{C} 174.9 was esterified by an oligosaccharidic chain which was analysed mainly on the basis of ROESY and HMBC spectra: The ROESY cross-peaks at δ_{H} 5.11 (*br s*, Rha-1)/ δ_{H} 3.68 (*dd*, $J = 8.9$, 8.1 Hz, Fuc-2), and δ_{H} 4.31 (*d*, $J = 7.3$ Hz, Xyl-1)/ δ_{H} 3.99 (*dd*, $J = 8.9$, 3.3 Hz, Fuc-3) revealed the (1 \rightarrow 2) linkage

Table 1

^{13}C , ^1H NMR spectroscopic data of the aglycon moieties of **1–5** in $\text{DMSO}-d_6$ (δ_{H} ppm, J in Hz).^a

	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	44.0	1.08, 1.95 <i>br d</i> (13.8)	43.7	1.94, nd	43.6	1.94, nd	43.6	1.10, 1.94	43.7	1.10, 2.70
2	68.1	4.10 <i>m</i>	68.4	4.02 <i>m</i>	68.2	4.02 <i>m</i>	68.1	4.04 <i>m</i>	68.3	4.04 <i>m</i>
3	84.1	4.00 <i>d</i> (3.6)	84.1	4.00	75.9	3.15	75.9	3.16	76.0	3.16
4	51.1	–	50.7	–	50.8	–	50.7	–	50.7	–
5	51.0	1.39	50.7	nd	50.7	nd	50.7	1.50	50.8	nd
6	19.7	1.32, nd	19.7	1.35, nd	19.7	nd	19.6	1.25, 1.36	19.7	1.34, nd
7	32.3	1.24, 1.34	32.3	1.22, 1.35	32.3	1.22, 1.35	32.3	1.22, 1.36	32.2	1.22, 1.37
8	40.0	–	39.4	–	39.6	–	39.5	–	39.3	–
9	46.4	1.46	46.5	1.47	46.4	1.48	46.3	1.48	47.3	1.43
10	36.0	–	35.6	–	35.6	–	35.7	–	35.7	–
11	23.0	1.85, 1.87	23.0	1.85, 1.87	23.0	1.85, 1.87	23.0	1.84, 1.87	23.0	1.84, 1.91
12	121.4	5.25 <i>t</i> -like	123.0	5.24 <i>t</i> -like	121.4	5.24 <i>t</i> -like	121.5	5.24 <i>t</i> -like	121.7	5.20 <i>t</i> -like
13	143.3	–	143.3	–	143.3	–	143.2	–	143.1	–
14	41.2	–	41.2	–	41.2	–	41.1	–	41.4	–
15	34.5	1.29, 1.60	34.7	1.27, 1.58	34.6	1.26, 1.60	34.6	1.26, 1.64	27.2	1.04, 1.53
16	72.5	4.30 <i>m</i>	72.5	4.31 <i>m</i>	72.3	4.31 <i>m</i>	72.4	4.32 <i>m</i>	22.6	1.54, 1.96
17	48.1	–	48.1	–	48.1	–	48.1	–	46.0	–
18	40.5	2.83 <i>dd</i> (12.6, 3.0)	40.4	2.83 <i>dd</i> (13.5, 3.0)	40.5	2.83 <i>dd</i> (14.2, 3.2)	40.5	2.85	41.1	2.73
19	46.6	0.98, 2.22	46.5	0.98, 2.23	46.6	0.98, 2.23	46.6	0.98, 2.22	45.6	1.07, 1.66
20	30.0	–	30.0	–	30.0	–	30.0	–	30.2	–
21	34.9	1.08, 1.88	34.9	1.09, 1.88	34.9	1.09, 1.86	34.9	1.10, 1.89	33.2	1.17, 1.35
22	30.6	1.69, 1.81	30.5	1.68, 1.80	30.2	1.69, 1.81	30.2	1.69, 1.80	31.4	1.48, 1.65
23	183.0	–	183.5	–	183.0	–	183.5	–	183.0	–
24	14.5	1.15 <i>s</i>	13.9	1.17 <i>s</i>	13.9	1.18 <i>s</i>	13.9	1.18 <i>s</i>	13.9	1.19 <i>s</i>
25	16.6	1.18 <i>s</i>	16.3	1.17 <i>s</i>	16.3	1.19 <i>s</i>	16.3	1.19 <i>s</i>	16.2	1.18 <i>s</i>
26	16.7	0.69 <i>s</i>	16.6	0.68 <i>s</i>	16.7	0.69 <i>s</i>	16.7	0.68 <i>s</i>	16.7	0.70 <i>s</i>
27	26.3	1.30 <i>s</i>	26.4	1.29 <i>s</i>	26.4	1.30 <i>s</i>	26.4	1.30 <i>s</i>	25.5	1.08 <i>s</i>
28	174.9	–	174.9	–	175.0	–	175.0	–	175.4	–
29	32.7	0.83 <i>s</i>	32.7	0.83 <i>s</i>	32.7	0.83 <i>s</i>	32.7	0.83 <i>s</i>	32.7	0.84 <i>s</i>
30	24.2	0.90 <i>s</i>	24.2	0.90 <i>s</i>	24.2	0.90 <i>s</i>	24.2	0.91 <i>s</i>	23.4	0.88 <i>s</i>

^a Overlapped proton signals are reported without designated multiplicity. Nd: not determined.

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