

CAULOSIDE D A NEW TRITERPENOID GLYCOSIDE FROM *CAULOPHYLLUM ROBUSTUM* MAXIM: IDENTIFICATION OF CAULOSIDE A*

L. I. STRIGINA, N. S. CHETYRINA, V. V. ISAKOV, YU. N. ELKIN, A. K. DZIZENKO and G. B. ELYAKOV
Pacific Institute of Bioorganic Chemistry, F. E. Science Centre, Academy of Science of USSR, Vladivostok 22, U.S.S.R.

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Abstract—The structure of cauloside D, one of the main saponins isolated from *Caulophyllum robustum* roots, was shown to be 3-*O*- α -L-arabinopyranosyl hederagenin-28-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 4)- β -D-glucopyranosyl(1 \rightarrow 6)- β -D-glucopyranoside with the aid of methylation and enzymatic hydrolysis by the digestive juice of the *Eulota maackii*. Cauloside A was shown to be identical with saponin A, isolated from *C. robustum* Maxim. previously. The composition of the digestive juice of *E. maackii* was shown to include enzymes that catalyse the cleavage of α -arabinosidic, β -1,6-glucosidic and acyl-*O*- β -glucosidic linkages.

INTRODUCTION

Murakami *et al.* [1] and ourselves [2-4] have reported studies on the triterpenoid glycosides of *Caulophyllum robustum* Maxim. (Berberidaceae). We now describe the results obtained in a study of caulosides D (1) and A (9).

RESULTS AND DISCUSSION

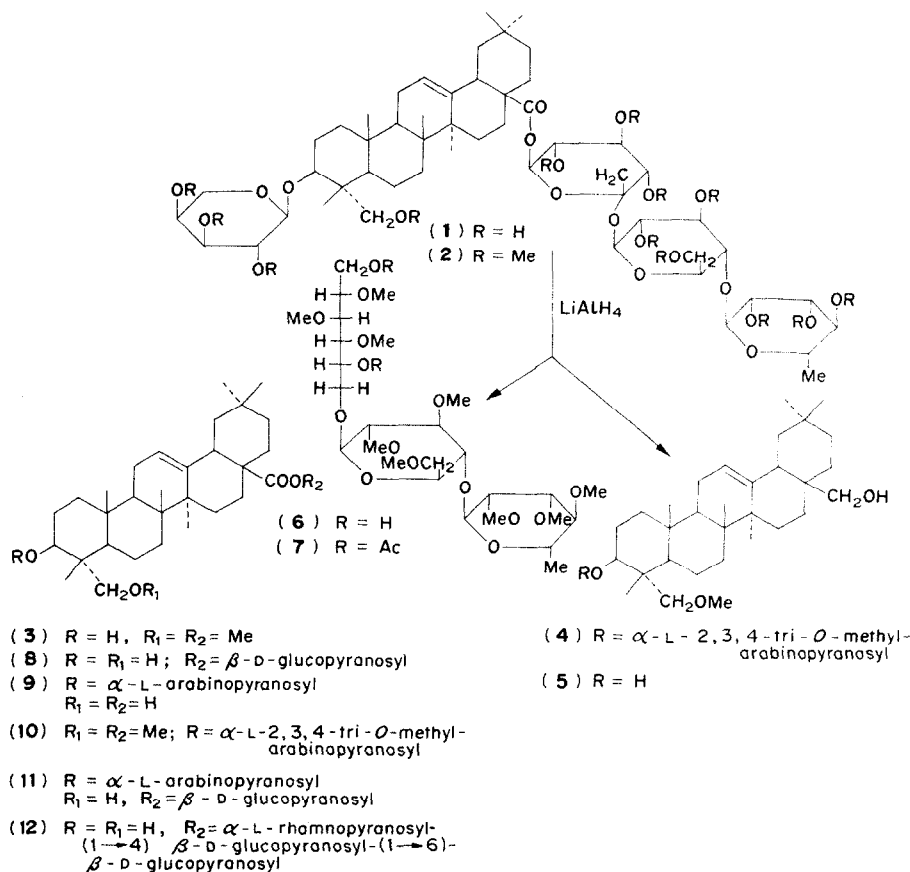
Acid hydrolysis of permethylated cauloside D (2), synthesized by the method of Purdie [5], yielded methyl-23-*O*-methyl-hederagenin (3) and a mixture of methyl glycosides. The latter mixture was acetylated and assayed with the aid of GLC-MS while the methyl ethers were examined by MS using the technique reported previously [6]. The compounds identified were 2,3,4-tri-*O*-methyl-rhamnose, 2,3,4-tri-*O*-methyl-arabinose, 2,3,6-tri-*O*-methyl-4-*O*-acetylglucose and 2,3,4-tri-*O*-methyl-6-*O*-acetyl-glucose. Reductive cleavage of (2) with lithium aluminum hydride resulted in a methylated progenin and an oligosaccharide.

Structure (4) corresponds to the progenin, since upon acid hydrolysis it yielded 23-methoxyerythrodil [7] and 2,3,4-tri-*O*-methyl-L-arabinopyranoside. TLC was used to identify 2,3,4-tri-*O*-methyl-L-rhamnose, 2,3,6-tri-*O*-methyl-D-glucose and 2,3,4-tri-*O*-methyl-D-sorbitol in the acid-hydrolysis products of the oligosaccharide. The reduction of 2,3,4-tri-*O*-methyl-D-glucose in the course of splitting (2) with lithium aluminum hydride indicated the direct bonding of the former with a hederagenin carboxyl group.

The structure (6), suggested for the oligosaccharide, was confirmed by the MS data for the oligosaccharide acetate (7). The detection in the MS of (7) of ion peaks at *m/e* 640 (M^+ -60) and 567 (M^+ -60-73) confirmed that (6) is a trisaccharide. Ion peaks at *m/e* 627, 539 and 117 were caused by the fragmentation of 2,3,4-tri-*O*-methyl-1,5-di-*O*-acetyl-D-sorbitol [8].

The configurations of the glycosidic bonds in cauloside D were established on examining the NMR spectra of the acetates of some progenins obtained from cauloside D by enzymatic hydrolysis. Earlier, we reported that incubation of caulosides D and E with the digestive juice or liver

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homogenate of *Eulota maackii* resulted in hederagenin and several progenins [9]. Progenins (8), (9), (11), and (12) were obtained after a chromatographic separation of the mixture obtained from the enzymatic hydrolysis. Acid hydrolysis of each of the progenins (8), (9), (11) and (12) and subsequent analysis of the reaction mixture on TLC resulted in the identification of glucose and rhamnose (12), arabinose and glucose (11), L-arabinose (9), and D-glucose (8). Accordingly each of the progenins (8) and (12) had one carbohydrate chain linked to a carboxyl group. Progenin (11) had two carbohydrate chains.

The integral intensity (18H) of the acetyl group signal in the NMR spectrum of the acetate of progenin (8) showed it to be a monoglucoside at C-28 of hederagenin. The anomeric proton signal of glucose (δ 5.58 ppm, J 8 Hz) indicated a β -configuration of the glycosidic bond. The β -configuration of the D-glucopyranosyl residue was confirmed by the difference in molecular rotation

of (8) and hederagenin: $\Delta [M]_D$ [(8)-hederagenin] -144.15° , $[M]_D$ methyl-D-glucosides: $\beta -62^\circ$; $\alpha +276^\circ$.

Progenin (9) is the 3-O- α -L-arabinopyranoside of hederagenin, since in the NMR spectrum of the corresponding acetate there was the anomeric proton signal of arabinose (δ 4.41 ppm, J 7 Hz), which indicated an α -configuration of the glycosidic bond. This was confirmed by the difference in molecular rotation of (9) and hederagenin: $\Delta [M]$ (9-hederagenin) -85.33° , $[M]_D$ methyl-L-arabinosides: $\alpha +28^\circ$; $\beta +404^\circ$ [10].

In the NMR spectrum of the acetate of progenin (11), the anomeric proton signals of D-glucose (δ 5.58 ppm, J 8 Hz) and arabinose (δ 4.41 ppm, J 7 Hz) indicated the β - and α -configurations respectively of the corresponding glycosidic bonds.

The integral intensity (33H) of the proton signal in the NMR spectrum of the acetate of progenin (12) showed it to be a triglycoside. The anomeric

proton signals of the two glucose molecules (δ 5.58 ppm, J 8 Hz and δ 4.60 ppm, J 7 Hz) indicated a β -configuration of the corresponding glycosidic bonds and a broad signal at δ 4.80 ppm, J 1 Hz, showed that rhamnose was linked by an α -glycosidic bond. An α -configuration of the L-rhamnopyranosyl residue in progenin (12) was confirmed by the difference in molecular rotation of cauloside D [2] and saponin D from *Akebia quinata* Decne [11]: $\Delta [\text{M}]_D$ (cauloside D and saponin D) -139° , $[\text{M}]_D$ methyl-L-rhamnosides: $\alpha -111^\circ$, $\beta +170^\circ$. Thus, the study of the progenins obtained by enzymatic hydrolysis allowed the determination of the configurations of all the glycosidic bonds in cauloside D, which is thus a novel glycoside of hederagenin with the structure of 3-*O*- α -L-arabinopyranosyl hederagenin 28-*O*- α -L-rhamnopyranosyl (1 \rightarrow 4)- β -D-glucopyranosyl (1 \rightarrow 6)- β -D-glucopyranoside (1). The structures of the products obtained by enzymatic hydrolysis shows that the digestive juice of *E. maackii* contains enzymes that catalyze the hydrolysis of the β -1,6-glucosidic, acyl-*O*- β -glucosidic and α -arabinosidic bonds.

Acid hydrolysis of permethylated cauloside A resulted in methyl-23-*O*-methyl-hederagenin and 2,3,4-tri-*O*-methyl-L-arabinose identical by TLC with authentic samples. The NMR spectrum of the acetate of cauloside A showed an α -configuration for the glycosidic bond (δ 4.37 ppm, J 10 Hz). Hence, cauloside A is a 3-*O*- α -L-arabinopyranoside of hederagenin (9) and is identical to saponin A, isolated previously from *C. robustum* [1].

EXPERIMENTAL

All m.p.s are uncorr. Column chromatography was on Si gel (150–200 mesh) with solvent systems: (1) C_6H_6 -EtOAc; (2) EtOAc-MeOH; and (3) CHCl_3 -EtOH. TLC was on Si gel or Silufol R with solvent systems: (4) toluene-EtOH (9:1); (5) CHCl_3 -MeOH- H_2O (2:1:0, 21); (6) CHCl_3 -MeOH (9:1); and (7) CHCl_3 -MeOH (2:1). GLC used glass columns (1.5 \times 6 mm) packed with 3% QF-1 on Chromosorb W (100–200 mesh), carrier gas (Ar): 50 ml/min, temp: 120–240°.

Permethylation of (1). (1) (1.5 g) was methylated twice by the Purdie method [5]. The reaction mixture (1.45 g) was chromatographed on a Si gel column in system 1 (3:1 \rightarrow 1:2). Fractions were analyzed by TLC with system 4. Yield: 0.49 g of 2, mp 85–87° (hexane), IR: ν_{max} 1730 cm^{-1} (COOR), no OH.

Methanolysis of (2). (2) (0.07 g) was refluxed for 5 hr with 3 ml 42% HClO_4 in MeOH (1:5). The reaction mixture was diluted with H_2O , and the aglycone filtered, washed with H_2O , dried and methylated with CH_2N_2 in Et_2O to give

(3) mp 190–194° (MeOH), $[\alpha]_D +70.4^\circ$ (CHCl_3). (Found, C, 76.29; H, 10.62; OCH_3 , 12.36. Calc. for $\text{C}_{32}\text{H}_{52}\text{O}_4$: C, 76.12; H, 10.60; OCH_3 , 12.09%). The aq soln was neutralized with Dowex 1 \times 2 (HCO_3^-), evaporated to a syrup, refluxed in abs. MeOH in the presence of Amberlite IR-120 (H^+), and acetylated with Ac_2O - $\text{C}_5\text{H}_5\text{N}$ at room temp. GLC was run in the presence of synthetic standards: 2,3,4-tri-*O*-methyl- β -D-xyloside ($T=0$) and 2,3,4,5,6-penta-*O*-acetyl-D-galactononitrile ($T=100$). The methyl pyranosides were identified as 2,3,4-tri-*O*-methyl-rhamnose ($T_R=2$, $T_x=7$), 2,3,4-tri-*O*-methyl-arabinose ($T_R=11$, $T_x=11.5$), 2,3,4-tri-*O*-methyl-6-*O*-acetyl-glucose ($T_R=35$, $T_x=42$), and 2,3,6-tri-*O*-methyl-4-*O*-acetyl-glucose ($T_R=39$, $T_x=44$).

LiAlH_4 Reduction of (2). (2) (0.47 g) in THF (5 ml) and LiAlH_4 (0.25 g) were refluxed for 3 hr. The LiAlH_4 residue was decomposed with H_2O , neutralized to pH 7 by 2N H_2SO_4 , evaporated and extracted, first with Et_2O and then with CHCl_3 . The Et_2O extracts were washed with H_2O , evaporated to dryness and chromatographed in system (1) (6:1 \rightarrow 2:1) to yield (4) mp 93.5°. The CHCl_3 extracts were washed with H_2O , evaporated to dryness and chromatographed in system 2 (50:1 \rightarrow 20:1) to yield (6) (0.27 g) syrup, $[\alpha]_D -32.3^\circ$ (CHCl_3). NMR: δ 1.31 (3H, s, J 7.71 Hz, C'-6Me rhamnose) 4.31 (1H, d, J 8 Hz, C'-1H glucose, 4.97 (1H, s, J 1.5 Hz, C'-1H rhamnose).

Methanolysis of (4). (4) (0.045 g) was hydrolyzed as described for (2). The residue was separated and chromatographed in system 1 (5:1 \rightarrow 2:1) to yield (5) (0.025 g), mp 188–190° (MeOH), $[\alpha]_D +54.2^\circ$ (CHCl_3). 2,3,4-Tri-*O*-methyl-L-arabinose, syrup, $[\alpha]_D +114^\circ$ (H_2O) was identified in the aq. portion following neutralization with Dowex 1 \times 2 (HCO_3^-) by TLC on Silufol in system 6.

Methanolysis of (6). (6) (0.05 g) was hydrolyzed as described for (4). 2,3,4-Tri-*O*-methyl-L-rhamnose, syrup, $[\alpha]_D +28.5^\circ$ (H_2O); 2,3,6-tri-*O*-methyl-D-glucose, syrup, $[\alpha]_D +58.4^\circ$ (MeOH); and 2,3,4-tri-*O*-methyl-D-sorbitol, syrup, were obtained after chromatography in system 3 (100:1 \rightarrow 100:15).

Enzymatic hydrolysis of (1). (1) (1.3 g) was dissolved in H_2O (200 ml) and incubated at 38° for 10 days with *E. maackii* digestive juice, periodically adding new portions of the juice. The reaction was checked by TLC in system (5). The hydrolyzate was extracted with *n*-BuOH, and the extracts were evaporated to dryness. After chromatography in system (3) (10:1 \rightarrow 1:2), the following products were obtained: (12) (0.05 g), mp 198–205° (MeCOMe + MeOH), $[\alpha]_D 0^\circ$ (MeOH); acetate of (12), amorphous substance, $[\alpha]_D 0^\circ$ (CHCl_3); (11) (0.062 g), mp 210–212° $[\alpha]_D +34^\circ$ (MeOH); acetate of (11), mp 118–120°, $[\alpha]_D +11.2^\circ$ (CHCl_3); (9) (0.1 g), mp 226–228° (MeOH), $[\alpha]_D +46.8^\circ$ (MeOH- CHCl_3); (8), mp 222–225°, $[\alpha]_D +35.3^\circ$ (CHCl_3); acetate of (8), mp 110–114° (EtOH), $[\alpha]_D +32.1^\circ$ (CHCl_3).

Permethylation of (9). (9) (0.27 g) was methylated by the Hakomori method [12]. The reaction product was extracted with CHCl_3 , evaporated to a foamy state, treated with CH_2N_2 in Et_2O , and chromatographed system (1) (50:1 \rightarrow 20:1) to give (10) (0.17 g). IR (CHCl_3): ν_{max} 1730 cm^{-1} , no OH.

Methanolysis of (10). (10) (0.049 g) was hydrolyzed as described for (2). The mixture was diluted with H_2O , and the aglycone filtered off to obtain (3), mp 190–192° (MeOH). The aq. residue was neutralized, evaporated to syrup and chromatographed on Silufol R in system 6 to identify 2,3,4-tri-*O*-methyl-L-arabinose, $[\alpha]_D +114^\circ$ (H_2O).

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