TRITERPENE GLYCOSIDES OF Astragalus AND THEIR GENINS. LXXII. X-RAY CRYSTAL STRUCTURE OF CYCLOORBICOSIDE D

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The structure of the new cycloartane glycoside cycloorbicoside D (1), which was isolated from *Astragalus orbiculatus* Ledeb. (Leguminosae), was determined based on chemical transformations and spectral data as 23ξ , 24ξ -cycloartan- 3β , 6α , 16β , 23, 24, 25-hexaol 3-O- β -D-xylopyranoside [1]. The methods used previously [1] were not capable of resolving the issue of the stereochemistry of asymmetric C-23 and C-24 in the glycoside. The present communication addresses this problem.

Figure 1 shows the molecular structure of **1** that was established by an x-ray crystal structure (XCS). Glycoside **1** contains six-membered rings A and C with practically ideal chair conformations (four atoms of the ring lie in a plane within ± 0.015 Å). Ring B has the twist-boat conformation due to the presence of β -orientated 9,19-cyclopropane. Five-membered ring D takes the conformation of an envelope. As expected, the ring fusions are A/B *trans*, B/C *cis*, and C/D *trans*. The location and configuration of all substituents were confirmed. The conformation of the side chain favors the formation of intramolecular H-bonds (IHB) between the C-23 and C-25 hydroxyls (O23—O25 distance 2.60 Å).

Taking into account the (3S,5S,6S,8R,13R,14S,16S,17S,20R) stereochemistry for **1**, the molecular structure indicates that asymmetric C-23 and C-24 have the 23R,24R configurations.

Thus, **1** has the structure 23R,24R-cycloartan- $3\beta,6\alpha,16\beta,23,24,25$ -hexaol 3-O- β -D-xylopyranoside.

Single crystals of $\bf 1$ were the dihydrate ($C_{35}H_{60}O_{10}\cdot 2H_2O$). The packing and intermolecular contacts in the crystal revealed O...H–O H-bonds between the hydroxyls and the waters of crystallization that formed a three-dimensional framework of weak interactions.

Compound 1 was recrystallized from CHCl₃:CH₃OH:H₂O (70:23:4) at 19°C, mp 266-267°C. Single crystals of 1 were transparent elongated prisms: a = 13.366(3), b = 6.074(1), c = 22.663(5) Å, $\beta = 104.26(3)$ °, V = 1783.2(6) Å³, $\rho_{calc} = 1.261$ g/cm³, space group $P2_1$, Z = 2. Unit cell constants and intensities of reflections were measured on a STOE Stadi-4 four-circle diffractometer ($\theta/2\theta$ -scanning) using Mo K α -radiation (graphite monochromator). Absorption corrections were not applied.

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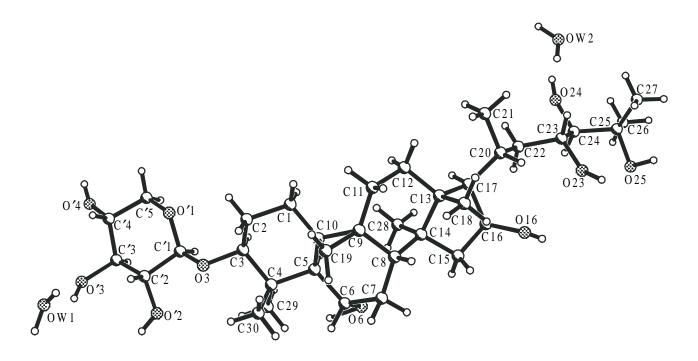


Fig. 1. Molecular structure and atomic numbering for cycloorbicoside D (water of crystallization also shown).

The structure was solved by direct methods using the SHELXS-97 programs. The structure was refined using the SHELXL-97 program. All nonhydrogen atoms were refined using anisotropic full-matrix least-squares methods (over F^2). The positions of H atoms were found geometrically and refined with fixed isotropic thermal parameters $U_{iso} = nU_{eq}$, where n = 1.5 for methyls and 1.2 for others and U_{eq} is the equivalent isotropic parameter of the corresponding C atom. Hydrogen atoms of hydroxyls and the water of crystallization were found in a difference electron-density synthesis. The final agreement factors (R) were 0.0860 for 2203 reflections [$I > 2\sigma(I)$] (wR2 = 0.1766) and 0.1159 for all 3031 reflections (wR2 = 0.1950).

The data from the XCS were deposited in the Cambridge Crystallographic Data Centre (CCDC 604459).

REFERENCES

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