DESANGELOYLSHAIRIDIN, X-RAY STRUCTURE DETERMINATION OF A SESQUITERPENE LACTONE FROM GUILLONEA SCABRA

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Key Word Index—Guillonea scabra; Umbelliferae; guaianolide; desangeloylshairidin; absolute configuration; conformational analysis.

Abstract—The structure and absolute configuration of desangeloylshairidin, a guaianolide isolated from *Guillonea scabra*, have been established by X-ray diffraction analysis. No conformational change was observed in its seven-membered ring between the crystal and deuterochloroform solution states.

INTRODUCTION

In previous communications [1, 2] we reported the isolation of two new sesquiterpene constituents of the roots of Guillonea scabra Cav. Cosson., an umbelliferous plant endemic in the Iberian Peninsula. These were desangeloylshairidin (1) and guillonein (2). Recently, the structure and absolute configuration of the latter was rigorously established by X-ray analysis [2]. The observation [2] that guillonein (2), in going from the crystal to deuterochloroform solution, undergoes a conformational change in its seven-membered ring, left open an alternative to the proposed structure of desangeloylshairidin (1). This prompted us to obtain the X-ray diffraction crystalline molecular structure of 1 to definitely establish its structure and absolute configuration, which is reported in this paper.

RESULTS AND DISCUSSION

Figure 1 shows the X-ray absolute molecular model of desangeloylshairidin (1). The seven-membered ring has a chair conformation, with C-7 at 0.55 Å out of the plane C-5–C-6–C-8–C-9 and C-1 = C-10 at 1.06 Å. This configuration and conformation are the same as that found in guillonein (2) [2]. The Cremer [3] parameters of this ring in 1 and 2 are: $\theta_2 = 38^\circ$, $\Phi_2 = 51^\circ$, $\Phi_3 = 79^\circ$, QT = 0.60 Å and $\theta_2 = 35^\circ$, $\Phi_2 = 56^\circ$, $\Phi_3 = 79^\circ$, QT = 0.77 Å, respectively. This calculation starts at C-1 in the sense of C-1

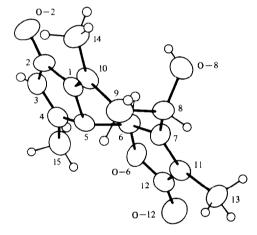


Fig. 1. Absolute X-ray molecular model of desangeloylshairidin (1).

 \rightarrow C-5. The double bond C-1=C-10 of 1.345 (6) Å in desangeloylshairidin (1) increases the planarity around, making the group C-1-C-2-C-3-C-4-C-5-C-9-C-10-C-14-C-15-O-2 planar. The crystal structure of 1 is built by one intermolecular hydrogen bond between O-8-H.... O-2 with O-8.... O-2 = 2.745 (6) Å and the angle at the hydrogen atom is 172°.

Comparison of the X-ray analyses of 1 and 2 shows a complete analogy between their intermolecular structures. As discussed in a previous work [2], the ${}^3J_{8,9}$ and ${}^3J_{8,9}$ coupling values, obtained from the 1H NMR spectrum of compound 2 in deuterochloroform solution, were not compatible with the corresponding torsion angles observed in the crystal, so that we concluded that a conformational change of the seven-membered ring from the chair C(7) to the twist-hinge TH'(7) form takes place in going from crystal to solution. This observation and the observed ${}^3J_{8,9} = 6.3$ Hz and ${}^3J_{8,9} = 1.0$ Hz values of desangeloylshairidin (1), opened the possibility of two alternatives for the structure of this compound in deutero-

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chloroform solution: (a) a C(7) conformation for the seven-membered ring with an α -configuration for the C-8 hydroxyl substituent; or (b) a TH'(7) conformation with a β -configuration for the hydroxyl grouping. The results obtained in this work definitely show that the first alternative prevails, as was put forward earlier [1, 2] on the basis of characteristic esterification shifts. It is noteworthy that, in contrast to guillonein (2) [2], desangeloyl-shairidin (1) maintains in deuterochloroform solution the seven-membered ring conformation shown in the crystal-line state.

EXPERIMENTAL

For the isolation of desangeloylshairidin from Guillonea scabra, see ref. [1].

X-ray structure determination of 1. Crystals of $C_{15}H_{16}O_4$ are monoclinic, space group $P2_1$, with unit cell parameters a = 10.7632(6), b = 6.7807(3), c = 9.1438(5) A, $\beta = 106.219(2)^\circ$, Z = 2 and $D_c = 1.349$ g/cm³.

The data were collected on a four circle diffractometer with graphite-monochromated CuK α radiation. The intensities of 1184 independent Friedel pairs up to $\theta = 65^{\circ}$ were measured in the $\omega/2\theta$ scan mode with a scan rate of 0.025° /sec. No absorption

correction was done ($\mu=7.63~{\rm cm}^{-1}$), and no crystal decomposition was observed throughout the expt. The structure was solved by MULTAN [4] and refined using the 1020 observed reflexions with $I>2\sigma(I)$. All H atoms were verified on a difference map and were included in the refinement as fixed isotropic contributors. A convenient weighting scheme was used to have no dependence of $\langle w\Delta^2F \rangle$ vs. $\langle F_0 \rangle$ and $\langle \sin\theta/\lambda \rangle$. The last cycles of weighted refinement, including both hkl and hkl reflexions, gave R=0.047 and $R_w=0.060$ [5].

The absolute configuration was determined comparing the 44 more relevant Bijvoet pairs with $F_0>10\sigma$ (F_0), $\Delta F_c>0.04$, 2.5 < $F_0<10$ and 0.37 < $\sin\theta/\lambda<0.60$, which include data with minimum exptal error. The averaged Bijvoet difference was 0.177 for the right enantiomer vs. 0.199 for the wrong one*.

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^{*}A list of atomic, positional and thermal parameters, bond distances and angles, torsional angles, conformational parameters and F_0 – F_c tables are deposited at the Cambridge Crystallographic Data Centre.