riding on the bonded atom with their $U_{\rm iso}$ fixed at $1.2U_{\rm eq}$ of the attached atom. All other H atoms were calculated to fit the geometry and their isotropic displacement parameters constrained to be the same and refined to 0.028 Å^2 . The absolute configuration of the molecule was not determined since it was known from the synthesis and was assigned according to the configurational angles given by Cano *et al.* (1985). Calculation of the Flack absolute structure parameter χ and its e.s.d. [-0.2 (4); Flack, 1983] further suggests that the refined absolute structure is correct.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: CAD-4 Software. Program(s) used to solve structure: SHELXS86 (Sheldrick, 1985). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: PLUTON (Spek, 1991) and PLATON (Spek, 1990). Software used to prepare material for publication: SHELXL93.

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Lists of atomic coordinates, displacement parameters, structure factors and complete geometry have been deposited with the IUCr (Reference: SK1000). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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A Clerodane Diterpene with Antibacterial Activity from Ajuga lupulina

Hao Chen,^a Ren-Xiang Tan,^a Zhi-Li Liu,^a Cong-Yuan Zhao^b and Jie Sun^c

^aDepartment of Biological Science and Technology, Nanjing University, Nanjing 210093, People's Republic of China, ^bDepartment of Chemistry, Nanjing University, Nanjing 210093, People's Republic of China, and ^cShanghai Institute of Organic Chemistry, Chinese Academy of Sciences, Shanghai 210032, People's Republic of China. E-mail: zh@netra.nju.edu.cn

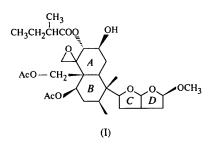
(Received 25 April 1996; accepted 4 November 1996)

Abstract

The structure of a new diterpene, $C_{30}H_{46}O_{11}$, with antibacterial activity against *Pseudomonas aeruginose* and *Escherichia coli*, isolated from the fresh whole plants of *Ajuga lupulina* (Labiatae) was established to be 2β -hydroxy-2-methylbutanoyl- 3α lupulin (3-deoxy-14,15-dihydro-2-hydroxy-15-methoxycaryoptinol 2-methylbutanoate), by means of X-ray crystallographic analysis. The present study confirms that the two six-membered rings are in ideal chair conformations.

Comment

As part of our ongoing research on diterpenoid compounds with antibacterial activity from Ajuga lupulina (Labiatae), a new clerodane diterpene, (I), with potential antibacterial activity against *Pseudomonas aeruginose* and *Escherichia coli*, was isolated from the fresh whole plants of Ajuga lupulina. In order to establish its structure, the X-ray analysis was undertaken. The results of the present study are in agreement with the structural elucidation based on spectroscopic data.



In rings A and B, the mean values of the valence angles (109.8 and 111.6°) are close to that of 111.6° for the chair form of cyclohexane obtained by Hendrickson (1967) from strain-energy minimization calculations. Rings C and D of the tetrahydrofurofuran

Acta Crystallographica Section C ISSN 0108-2701 © 1997 ring exhibited envelope and planar conformations, respectively. The bond lengths of rings C and D vary from 1.397 (2) to 1.546 (2) Å. The mean values of bond lengths and valence angles are close to those of the tetrahydrofuran ring (Shiro & Koyama, 1970; Chen, Tan, Liu, Lie & Chen, 1996). The stereochemistry of the title compound was established by comparison with the absolute configuration of ajugareptansin (Solans, Miravitlles, Declercq & Germain, 1979), clerodin (Kato, Munakata & Katayama, 1973; Rogers et al., 1979) and ajugarins IV and V (Kubo, Kido & Fukuyama, 1980; Kubo, Klocke, Miura & Fukuyama, 1982; Chen et al., 1996) (Fig. 1). Owing to the unfavorable value of the reflection-to-parameter ratio (2310/380 = 6.08), the numerical values of the bond distances and angles must be taken with caution.

The structure is stabilized by intermolecular hydrogen bonds between the O(4) and O(9ⁱ) atoms [O(4)-H(4A)]0.85, $H(4A) \cdots O(9^i)$ 2.04 Å and $O(4) - H(4A) \cdots O(9^i)$ 178.6°; symmetry code: (i) $-\frac{1}{2} + x$, $\frac{1}{2} - y$, 2 - z]. Due

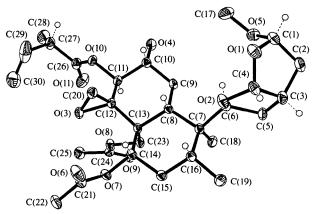


Fig. 1. View of the title molecule with the atom-numbering scheme. H atoms have been omitted for clarity. Displacement ellipsoids are at the 30% probability level.

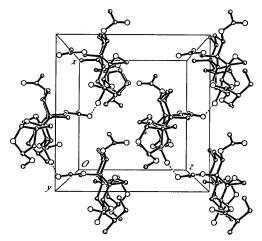


Fig. 2. Crystal-packing diagram of the title compound.

Experimental

The title compound (m.p. 445-447 K) was isolated from the fresh whole plant of Ajuga lupulina. After purification by preparative TLC, a single crystal was obtained from n-hexane-EtOAc (2:1).

Crystal data

| $C_{30}H_{46}O_{11}$ | Mo $K\alpha$ radiation |
|---|-----------------------------------|
| $M_r = 582.7$ | $\lambda = 0.7107 \text{ Å}$ |
| Orthorhombic | Cell parameters from 20 |
| P2 ₁ 2 ₁ 2 ₁ | reflections |
| a = 10.750(2) Å | $\theta = 18.71 - 21.39^{\circ}$ |
| b = 26.735 (6) Å | $\mu = 0.097 \text{ mm}^{-1}$ |
| c = 10.533(1) Å | T = 293 K |
| $V = 3027.2 (15) \text{ Å}^3$ | Prism |
| Z = 4 | $0.30 \times 0.20 \times 0.20$ mm |
| $D_x = 1.278 \text{ Mg m}^{-3}$ | Colorless |
| D_m not measured | |

 $R_{\rm int} = 0.004$

 $\theta_{\rm max} = 25^{\circ}$

 $h = 0 \rightarrow 11$

 $k = 0 \rightarrow 31$

 $l = 0 \rightarrow 12$

3 standard reflections every 200 reflections intensity decay: none

Data collection

| Rigaku AFC-7R diffractom- |
|------------------------------|
| eter |
| $\theta/2\theta$ scans |
| Absorption correction: none |
| 2995 measured reflections |
| 2945 independent reflections |
| 2310 reflections with |
| $F > 4\sigma(F)$ |

Refinement

| Refinement on F | $\Delta \rho_{\rm max} = 0.35 \ {\rm e} \ {\rm \AA}^{-3}$ |
|-------------------------------------|--|
| R = 0.0465 | $\Delta \rho_{\rm min} = -0.41 \ {\rm e} \ {\rm \AA}^{-3}$ |
| wR = 0.0614 | Extinction correction: |
| S = 2.38 | Zachariasen (1967) type |
| 2310 reflections | 2, Gaussian isotropic |
| 380 parameters | Extinction coefficient: |
| H atoms not refined | $4(4) \times 10^{-4}$ |
| $w = 1/[\sigma^2(F) + 0.0003F^2]$ | Scattering factors from Inter- |
| $(\Delta/\sigma)_{\rm max} = 0.020$ | national Tables for X-ray |
| | Crystallography (Vol. IV) |
| | |

Table 1. Selected geometric parameters (Å, °)

| O(1)—C(1) | 1.415 (1) | O(6)—C(21) | 1.188 (1) |
|--|---|---|---|
| O(3)—C(12) | 1.429 (2) | C(8)—C(13) | 1.593 (2) |
| O(3)—C(20) | 1.453 (2) | C(12)—C(20) | 1.480 (1) |
| O(4)—C(10) | 1.426 (1) | C(13)—C(14) | 1.531 (1) |
| O(5)—C(17) | 1.427 (2) | C(13)—C(23) | 1.538 (1) |
| C(12) - O(3) - C(20) $C(1) - O(5) - C(17)$ $O(5) - C(1) - C(2)$ $C(2) - C(3) - C(4)$ $C(2) - C(3) - C(5)$ $O(1) - C(4) - O(2)$ $O(2) - C(4) - C(3)$ $O(2) - C(6) - C(7)$ | 61.8 (1) 114.7 (1) 108.8 (1) 103.7 (1) 116.6 (1) 112.9 (1) 108.4 (1) 110.5 (1) | $\begin{array}{c} C(6) - C(7) - C(8) \\ O(4) - C(10) - C(9) \\ O(3) - C(12) - C(11) \\ O(3) - C(12) - C(20) \\ C(13) - C(12) - C(20) \\ C(8) - C(13) - C(12) \\ C(12) - C(13) - C(23) \\ O(6) - C(21) - O(7) \end{array}$ | 107.1 (1) 112.0 (1) 115.6 (1) 59.9 (1) 122.6 (1) 103.8 (1) 109.2 (1) 124.2 (2) |

For data collection, a scan width of $(1.00 + 0.30 \tan \theta)^{\circ}$ (in ω) and a scan speed of 16.0° min⁻¹ were used. A colorless prism crystal was mounted on a glass fiber. Cell constants and an orientation matrix for data collection were obtained from a least-squares refinement. The structure was solved by direct methods (SHELXS86; Sheldrick, 1985) and expanded using Fourier techniques (DIRDIF; Beurskens et al., 1992). The non-H atoms were refined anisotropically by full-matrix least squares. H atoms were placed in calculated positions (C-H 0.96 Å), assigned fixed isotropic displacement parameters and allowed to ride on their respective parent atoms. The contributions of these H atoms were included in the structurefactor calculations. The C(29) H atoms [s.o.f.'s for H(29B) and H(29C) were fixed at 0.6 and 0.4, respectively], and the C(30) and C(30') H atoms [s.o.f.'s for atoms H(30A)-H(30C) and H(30D)-H(30F) were fixed at 0.6 and 0.4, respectively] were found to be highly disordered. Neutral atom scattering factors were taken from Cromer & Waber (1974). Anomalous dispersion effects were included in F_{calc} (Ibers & Hamilton, 1964). The values for the mass attenuation coefficients are those of Creagh & Hubbel (1992). All calculations were performed on a PC586 using the SHELXTL-Plus (Sheldrick, 1991) program package.

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1988). Cell refinement: MSC/AFC Diffractometer Control Software. Data reduction: TEXSAN (Molecular Structure Corporation, 1985, 1992). Program(s) used to refine structure: SHELXTL-Plus. Software used to prepare material for publication: SHELXL93 (Sheldrick, 1993).

Lists of atomic coordinates, displacement parameters, structure factors and complete geometry have been deposited with the IUCr (Reference: NA1249). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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4,6,10,12-Tetraphenyl-7,9-dioxa-1,3-dithiacyclododecane

JORDI LLORCA,^a † ELIES MOLINS,^a ABELARDO FLORES-VELA,^b SAMUEL CRUZ-SÁNCHEZ^b AND EUSEBIO JUARISTI^b

^aInstitut de Ciència de Materials de Barcelona (CSIC), Campus Universitat Autònoma de Barcelona, 08193 Cerdanyola, Spain, and ^bDepartamento de Química, Centro de Investigación y de Estudios Avanzados del Instituto Politécnico Nacional, Ap. 14-740, 07000 Mexico DF, Mexico. E-mail: elies.molins@icmab.es

(Received 13 March 1996; accepted 20 January 1997)

Abstract

The title compound, $C_{32}H_{32}O_2S_2$, is a mixed oxa-thia crown macrocycle. The molecule has all heteroatoms at side positions of a quadrangular skeleton. Both sulfur and oxygen conformational arrangements are virtually the same.

Comment

Sulfur coronands possess solid-state conformations that differ significantly from those of their oxygen counterparts and yield metal complexes with unexpected reactivity and redox properties (Cooper, 1988). In this context, we have isolated the title compound, (I), which, in addition to its potential as a transition metal complexing agent, is an interesting candidate for conformational studies since it contains both S and O heteroatoms in a novel arrangement.

[†] Present address: Departament de Química Inorgànica, Universitat de Barcelona, Diagonal 647, 08028 Barcelona, Spain.