

Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *TEXSAN LS*. Molecular graphics: *TEXSAN ORTEP* (Johnson, 1965). Software used to prepare material for publication: *TEXSAN FINISH*.

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (\AA^2)

$$U_{\text{eq}} = \frac{1}{3} \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
O1	0.7028 (3)	0.1345	0.2978 (2)	0.0505 (1)
C2	0.8586 (4)	0.0572 (4)	0.3971 (3)	0.0473 (1)
N3	0.7758 (3)	-0.0860 (3)	0.4239 (2)	0.0540 (1)
C4	0.5611 (4)	-0.1265 (4)	0.3273 (2)	0.0457 (1)
C5	0.4877 (4)	0.0418 (4)	0.2706 (2)	0.0448 (1)
C6	0.3890 (4)	-0.2107 (5)	0.3927 (3)	0.0660 (1)
C7	0.3696 (4)	0.0488 (4)	0.1208 (2)	0.0461 (1)
C8	0.4690 (5)	-0.0169 (4)	0.0202 (3)	0.0613 (1)
C9	0.3508 (6)	-0.0141 (5)	-0.1163 (3)	0.0744 (1)
C10	0.1370 (7)	0.0537 (6)	-0.1525 (3)	0.0804 (1)
C11	0.0393 (5)	0.1204 (6)	-0.0538 (4)	0.0858 (1)
C12	0.1537 (4)	0.1188 (5)	0.0825 (3)	0.0645 (1)
O13	1.0419 (3)	0.1189 (3)	0.4493 (2)	0.0645 (1)

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Table 2. Geometric parameters (\AA , $^\circ$)

O1—C2	1.355 (3)	C5—C7	1.502 (3)
O1—C5	1.463 (3)	C7—C8	1.381 (4)
C2—N3	1.328 (4)	C7—C12	1.383 (4)
C2—O13	1.211 (3)	C8—C9	1.385 (4)
N3—C4	1.457 (3)	C9—C10	1.362 (5)
C4—C5	1.524 (4)	C10—C11	1.363 (5)
C4—C6	1.499 (4)	C11—C12	1.376 (4)
C2—O1—C5	108.6 (2)	C4—C5—C7	115.8 (2)
O1—C2—N3	109.9 (2)	C5—C7—C8	121.6 (2)
O1—C2—O13	121.4 (2)	C5—C7—C12	119.4 (2)
N3—C2—O13	128.7 (3)	C8—C7—C12	119.0 (2)
C2—N3—C4	112.4 (2)	C7—C8—C9	119.9 (3)
N3—C4—C5	100.0 (2)	C8—C9—C10	120.5 (3)
N3—C4—C6	113.8 (2)	C9—C10—C11	119.8 (3)
C5—C4—C6	114.5 (2)	C10—C11—C12	120.6 (3)
O1—C5—C4	103.9 (2)	C7—C12—C11	120.1 (3)
O1—C5—C7	111.2 (2)		

The scan rate was 3° min^{-1} (in ω). Weak reflections [$I < 10\sigma(I)$] were rescanned (maximum of two rescans) and the counts accumulated to improve accuracy. H atoms were placed at assumed positions ($\text{C}_n\text{H} = 0.95 \text{ \AA}$, $U = 1.2 \times U_{\text{eq}}$ of the associated atom) and fixed. The correct enantiomer was chosen on the basis of the known configuration of the starting material [(-)-(1*R*,2*S*)-norephedrine].

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Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55982 (10 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: CR1047]

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Structure of Vicogenin

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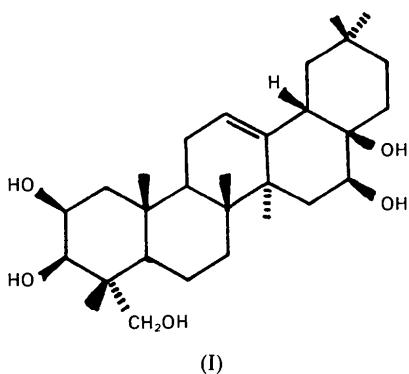
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Abstract

Vicogenin is a new 28-nortriterpenoid. Its structure has been elucidated as $2\alpha,3\beta,16\alpha,17\beta,23$ -penta-hydroxy-28-norolean-12-ene. In the crystal there are two water molecules per vicogenin molecule. Hydrogen bonds link the vicogenin molecules to each other and to the water molecules.

Comment

The title compound (I) has been isolated for the first time from the chloroform extract of the plant *Vicoa indica* DC(composital). The plant is said to exhibit antifertility activity when administered during the post-partum period in albino rats (Gandhi,



Sankaranarayanan, Ramesh Lal & Mathur, 1983) and anti-implantation activity (Mohana, Susan & Purushothaman, 1984). The configuration at the *D/E* ring junction was tentatively fixed as *cis* on the basis of biogenetic considerations and ^{13}C NMR spectra and the molecular formula was established by high-resolution mass spectrometry. In order to confirm the structure, particularly the nature of the configuration at the *D/E* ring junction, an X-ray crystallographic investigation was undertaken. 28-Noroleananes are relatively rare in nature. There is only one report of the extraction of 17-hydroxy-28-norolean-12-ene, a natural product (Linde, 1979).

The C—C bond lengths range from 1.507 (11) to 1.600 (11) Å with a mean value of 1.54 (2) Å. The endocyclic bond angles for tetrahedral atoms in rings *A*, *B*, *C*, *D* and *E* vary from 104.9 (6) to 125.4 (7) $^{\circ}$ with a mean value of 111.81 (7) $^{\circ}$, which compares well with that of related compounds (Hoge & Nordman, 1974; Subramanian, Selladurai, Sivakumar, Ponnuswamy & Sukumar, 1989). The rings *A*, *B*, *D* and *E* possess chair conformations and the conformation of ring *C* is intermediate between half-chair and sofa (Gzella, Zaprutko, Wrzeciono & Jaskólski, 1987) with the best rotational axis bisecting the C(8)—C(9) and C(13)—C(12) bonds.

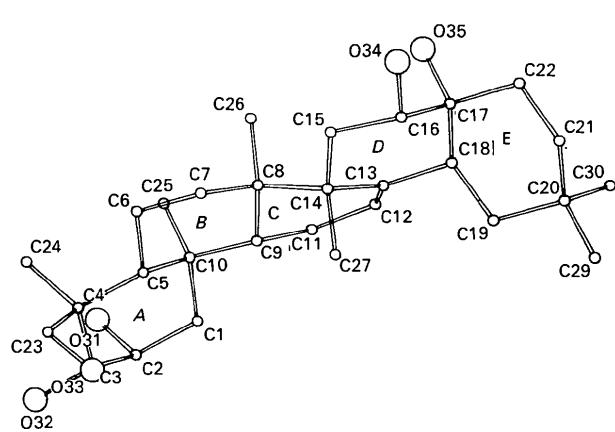


Fig. 1. The vicogenin molecule (with H atoms omitted for clarity) showing the atomic numbering scheme and ring identifiers.

$\{\Delta\text{C}_2[\text{C}(8)—\text{C}(9)] = 16.5^{\circ}\}$. The ring junctions *A/B*, *B/C* and *C/D* are *trans*, but the *D/E* ring junction is *cis* with both the hydroxyl group and the H atom pointing upwards, thus showing a *cis* di- β -configuration. The angles between the mean planes of the rings are *A/B* 15.20, *B/C* 165.49, *C/D* 165.11 and *D/E* 121.46 $^{\circ}$ (Maurin & Krygowski, 1987), making a ‘zigzag’ along the length of the molecule. The molecules are held together by a network of hydrogen bonds (Table 3) involving the hydroxyl groups and the water molecules.

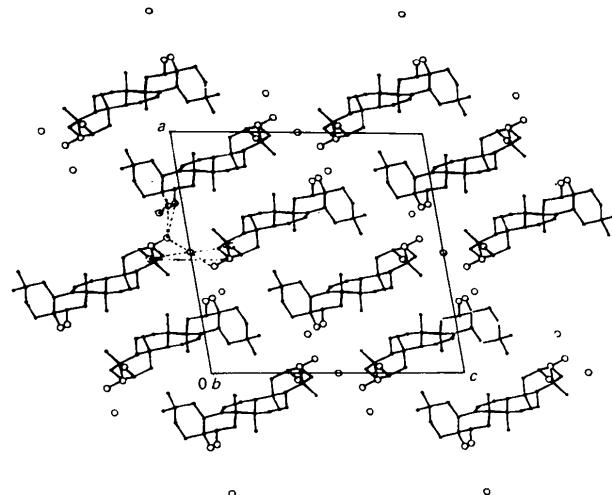


Fig. 2. Molecular packing viewed along the *b* axis.

Experimental

Crystal data



$$M_r = 512$$

Monoclinic

*I*2

$$a = 19.14 (3) \text{ \AA}$$

$$b = 7.635 (3) \text{ \AA}$$

$$c = 19.838 (1) \text{ \AA}$$

$$\beta = 99.84 (4)^{\circ}$$

$$V = 2856.36 \text{ \AA}^3$$

$$Z = 4$$

$$D_x = 1.192 \text{ Mg m}^{-3}$$

Cu $K\alpha$ radiation

$$\lambda = 1.5418 \text{ \AA}$$

Cell parameters from 20 reflections

$$\theta = 20\text{--}30^{\circ}$$

$$\mu = 0.634 \text{ mm}^{-1}$$

$$T = 293 \text{ K}$$

Needle

$$0.5 \times 0.4 \times 0.3 \text{ mm}$$

Colourless

Data collection

Enraf-Nonius CAD-4 diffractometer

$\omega/2\theta$ scans

Absorption correction: empirical

$$T_{\min} = 0.960, T_{\max} = 1.021$$

2220 measured reflections

2220 independent reflections

2025 observed reflections

$$[I > 3\sigma(I)]$$

$$R_{\text{int}} = 0.03$$

$$\theta_{\max} = 60^{\circ}$$

$$h = -21 \rightarrow 21$$

$$k = 0 \rightarrow 7$$

$$l = 0 \rightarrow 22$$

2 standard reflections monitored every 100

reflections

intensity variation: <2%

Refinement

Refinement on F by full-matrix least squares

Final $R = 0.056$

$wR = 0.060$

$S = 2.3$

2025 reflections

526 parameters

All H-atom parameters refined; positions found from difference Fourier syntheses

Cell refinement: *SDP* (Frenz, 1978). Data reduction: *SDP*. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1986). Program(s) used to refine structure: *SHELX76* (Sheldrick, 1976). Software used to prepare material for publication: *PARST* (Nardelli, 1983).

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (\AA^2)

$$U_{\text{eq}} = (U_{11} + U_{22} + U_{33})/3.$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
C(1)	0.5236 (4)	-0.1923 (10)	0.2686 (3)	0.39 (3)
C(2)	0.4881 (4)	-0.2416 (11)	0.1971 (4)	0.43 (2)
C(3)	0.4760 (4)	-0.0811 (11)	0.1509 (4)	4.15 (2)
C(4)	0.5417 (4)	0.0321 (11)	0.1491 (3)	3.65 (2)

$$w = 7.4521/[\sigma^2(F) + 0.000092F^2]$$

$$(\Delta/\sigma)_{\text{max}} = -0.156$$

$$\Delta\rho_{\text{max}} = 0.4765 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\text{min}} = -0.398 \text{ e \AA}^{-3}$$

Atomic scattering factors from *International Tables for X-ray Crystallography* (1974, Vol. IV, Table 2.2B)

Ring *B*

C(8)—C(9)—C(10)—C(5)	-54.9 (7)
C(7)—C(8)—C(9)—C(10)	45.1 (8)
C(6)—C(7)—C(8)—C(9)	-42.4 (9)

C(5)—C(6)—C(7)—C(8)	53.7 (9)
C(10)—C(5)—C(6)—C(7)	-64.3 (8)

Ring *C*

C(8)—C(9)—C(11)—C(12)	38.9 (8)
C(9)—C(11)—C(12)—C(13)	-7.6 (12)
C(11)—C(12)—C(13)—C(14)	0.8 (13)

C(12)—C(13)—C(14)—C(8)	-25.0 (10)
C(9)—C(8)—C(14)—C(13)	54.7 (7)

Ring *D*

C(14)—C(13)—C(18)—C(17)	-43.4 (9)
C(16)—C(17)—C(18)—C(13)	52.7 (9)
C(15)—C(16)—C(17)—C(18)	-58.1 (8)

C(14)—C(15)—C(16)—C(17)	54.3 (9)
C(13)—C(14)—C(15)—C(16)	-42.0 (9)

Ring *E*

C(17)—C(18)—C(19)—C(20)	-54.8 (9)
C(18)—C(19)—C(20)—C(21)	55.3 (9)
C(19)—C(20)—C(21)—C(22)	-54.4 (10)

C(20)—C(21)—C(22)—C(17)	54.3 (11)
C(18)—C(17)—C(22)—C(21)	-51.4 (10)

Table 3. Hydrogen-bond distances (\AA)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	Symmetry location of acceptor
O(31)—H(31) \cdots O(33)*	2.688 (8)	<i>x, y - 1, z</i>
O(31)—H(31) \cdots O(36)*	3.246 (6)	<i>x + 1/2, y - 1/2, z - 1/2</i>
O(32)—H(32) \cdots O(34)	3.142 (6)	<i>x - 1/2, y - 1/2, z - 1/2</i>
O(34)—H(34) \cdots O(32)	3.142 (6)	<i>x + 1/2, y + 1/2, z + 1/2</i>
O(35)—H(35) \cdots O(32)	2.834 (5)	<i>x + 1/2, y + 1/2, z + 1/2</i>
O(36)—H2(36) \cdots O(32)	3.021 (3)	<i>x - 1/2, y + 1/2, z + 1/2</i>
O(37)—H1(37) \cdots O(34)	2.678 (13)	<i>x - 1/2, y - 1/2, z - 1/2</i>
O(37)—H2(37) \cdots O(35)	2.714 (13)	<i>x - 1/2, y + 1/2, z - 1/2</i>

* Angle O(33) \cdots H(31) \cdots O(36) = 80^\circ; sum of angles at H(31) = 354°.