173.03 (39)°], although it is present with the bulky Phe side chain in close proximity to the Boc group (Benedetti *et al.*, 1980). On the other hand, the value of the torsion angle N(8)-C(9)-C(17)-O(18) between the urethane moiety and the next peptide group is 67.86 (31)°, showing that the Boc derivative occurs in the crystal in a bent conformation.

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Structure of Spirosta-5,25(27)-diene- 3β ,11 α -diol

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 $C_{27}H_{40}O_4$, $M_r = 428.61$, orthorhombic. Abstract. a = 6.369 (1), $b = 12 \cdot 240$ (3), $P2_{1}2_{1}2_{1}$, *c* = $V = 2381 (1) \text{ Å}^3$, Z=4, $D_r =$ 30.547 (5) Å, 1.195 g cm^{-3} , $\lambda(\text{Mo } K\alpha) = 0.71073 \text{ Å}$, $\mu = 0.73 \text{ cm}^{-1}$, F(000) = 936, T = 293 (2) K, R = 0.045 for 1615 unique observed reflections. The compound was isolated from Helleborus serbicus Adam 1906 (Ranunculaceae). The rings have the following conformations: A, C and F chair; B and E intermediate between half chair and envelope; D half chair. The molecules are linked by $O-H\cdots O$ hydrogen bonds into chains along the b axis. The molecule is very similar to molecule (I) spirosta-5,25(27)-diene- 1β , 3β , 11α -triol of monohydrate [Kálmán, Argay, Ribár, Živanov-Stakić & Vladimirov (1985). Acta Cryst. C41, 1645-1647].

Introduction. This new steroid sapogenin was isolated from the subterranean organs of the plant *Helleborus* serbicus Adam 1906 (Ranunculaceae) by column chromatography and preparative thin-layer chromatography on silica gel using mobile-phase isopropyl ether-methanol 95:5 (v/v) and chloroform-

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methanol 95:5. The crystals reached a melt at 496–498 K. The chemical structure (1) proposed from chemical and spectroscopic studies was substantiated by X-ray diffraction.



Experimental. Colourless crystal ca $0.4 \times 0.6 \times 0.9$ mm. Enraf-Nonius CAD-4 diffractometer, graphite-monochromated Mo K α radiation. $0.0327 \leq \sin\theta/\lambda \leq 0.6606$ Å⁻¹, $\omega-2\theta$ scan, h 0-8, k 0-16, l 0-40. Cell parameters by least-squares fit for 18 centred reflections. Systematic absences: h = 2n + 1 in h00, k = 2n + 1 in 0k0, l = 2n + 1 in 00l. Of 2779 unique reflections, 1615 taken as observed with $I > 2.5\sigma(I)$. No absorption correction performed. Three standard

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reflections, intensity variation < 2%. Structure solved by SHELX76 (Sheldrick, 1976) using 458 reflections with E > 1.20 and successive structure-factor, Fourier and difference-Fourier calculations. Full-matrix refinement. $\sum w(\Delta F)^2$ minimized for 31 heavy atoms (336) parameters). Final R = 0.045, wR = 0.065, w = $[\sigma^2(F) + 61.87 \times 10^{-4}(F)^2]^{-1}$. Residual electron density within $+0.2 e \text{ Å}^{-3}$. Positions of H atoms generated from assumed geometries, remaining H atoms bound to O atoms and C27 found in a difference Fourier map; all H atoms refined isotropically. Max. Δ/σ for H atoms 0.83. Scattering factors as in SHELX76. Calculation on an IBM 43/41 computer.

Table 1. Final fractional coordinates for non-H atoms and equivalent isotropic temperature parameters (with e.s.d.'s in parentheses)

$$U_{\rm eq} = \frac{1}{3} (U_{11} + U_{22} + U_{33})$$

	x	у	Ζ	$U_{eq}(\dot{A}^2)$
21	0.5555 (8)	0.3632 (4)	0.1191 (2)	0.060 (3)
22	0.5974 (10)	0.3523 (5)	0.0704 (2)	0.073 (4)
73	0.6405 (11)	0.2339 (5)	0.0584 (2)	0.081 (4)
24	0.4668 (10)	0.1608 (4)	0.0742 (2)	0.068 (3)
25	0.4220 (7)	0.1778 (4)	0.1220(1)	0.052 (3)
76	0.4360 (8)	0.0937(4)	0.1495 (2)	0.060 (3)
27	0.3973 (10)	0.1009 (4)	0.1973 (2)	0.062 (3)
28	0.2642 (7)	0.2005 (3)	0.2086(1)	0.045 (2)
29	0.3578 (7)	0.3024 (3)	0.1867 (1)	0.043 (2)
210	0.3725 (7)	0.2926 (3)	0.1364 (1)	0.045 (2)
211	0.2451 (7)	0.4077 (3)	0.2030 (2)	0.046 (3)
C12	0.2332 (7)	0.4134 (3)	0-2528 (1)	0.048 (3)
C 13	0-1285 (7)	0.3132 (3)	0.2726 (1)	0.043 (2)
C14	0.2528 (7)	0.2142 (3)	0.2577 (1)	0.045 (2)
C15	0.1694 (8)	0.1222 (4)	0.2861 (1)	0.051 (3)
C16	0-1496 (7)	0.1791 (4)	0.3304 (1)	0.048 (2)
C17	0.1525 (7)	0.3011 (3)	0.3226 (1)	0.046 (2)
C18	-0.1023 (6)	0.3060 (4)	0.2586(1)	0.052 (3)
C 19	0-1653 (8)	0.3283 (5)	0.1149 (2)	0.062 (3)
C20	-0.0183 (8)	0-3462 (4)	0.3531 (1)	0.053 (3)
C21	0.0490 (11)	0.4476 (4)	0.3802 (2)	0.075 (4)
C22	-0.0823 (7)	0.2471 (4)	0.3799 (1)	0.052 (3)
C23	-0.3085 (8)	0.2470 (5)	0.3946 (2)	0.067 (3)
C24	-0.3569 (10)	0.1534 (5)	0.4257 (2)	0.083 (4)
C25	-0.1982 (12)	0.1474 (6)	0-4615 (2)	0.094 (5)
C26	0.0185 (10)	0.1484 (5)	0.4432 (2)	0.074 (4)
C27	-0.2340 (21)	0.0900 (9)	0-4986 (3)	0.140 (8)
03	0.6585 (11)	0.2248 (5)	0.0118 (1)	0.130 (4)
011	0.3495 (6)	0.5022 (3)	0.1878 (1)	0.061 (2)
016	-0.0495 (5)	0.1580 (2)	0.3516(1)	0.054 (2)
022	0.0554(5)	0.2407(3)	0.4157(1)	0.062 (2)



Fig. 1. Perspective view of the molecule with atomic numbering. Numbers are for C unless indicated otherwise. The H atoms are shown but not labelled.

Discussion. Atomic coordinates of non-H atoms are in Table 1. The molecular geometry with atomic numbering is shown in Fig. 1. The bond lengths and angles for non-H atoms are listed in Table 2.*

Rings A and C have normal chair conformations while ring D assumes a half-chair shape [puckering parameters of Cremer & Pople (1975): Q =0.463(5) Å, $\varphi = 202.2(6)^{\circ}$; lowest asymmetry factor $fC_{2}(C16) = 0.026 (5) \text{ Å}$ (Kálmán, Czugler & Simon, 1982)] with a pseudo C_2 symmetry axis bisecting C13-C14. The flexible B ring possesses an intermediate form between envelope and half chair characterized by the puckering parameters Q = 0.482 (5) Å, $\varphi = 194.0$ (6)° and $\theta = 54.1$ (6)°. Ring E fused to ring D with a cis junction also exhibits an intermediate form

* Lists of structure factors, anisotropic thermal parameters and H-atom parameters have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 43183 (12 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

Table 2. Bond lengths (Å) and angles (°) with e.s.d.'s in parentheses

Cl-C2	1.517 (7)	C13-C14	1.518 (5)
Cl-C10	1.545 (6)	C13-C17	1.542 (6)
C2-C3	1.520 (8)	C13-C18	1.534 (6)
C3-C4	1.502 (8)	C14-C15	1.517 (6)
C3O3	1.433 (6)	C15-C16	1.528 (6)
C4C5	1.503 (7)	C16-C17	1.512 (6)
C5C6	1.333 (6)	C16-O16	1.448 (5)
C5C10	1.506 (6)	O16-C22	1.407 (5)
C6-C7 C7-C8 C8-C9 C8-C14	1.483 (7) 1.525 (6) 1.535 (6) 1.512 (6)	C17–C20 C20–C21 C20–C22 C22–C23 C22–C23	1-534 (6) 1-552 (6) 1-520 (7) 1-509 (7)
C9-C10	1.544 (6)	C22-C22	1.404 (3)
C9-C11	1.557 (6)	O22-C26	1.428 (6)
C10-C19	1.538 (6)	C23-C24	1.518 (7)
C11-C12	1.524 (6)	C24-C25	1.492 (9)
C12–C13	1.522 (6)	C25-C27	1.489 (9)
C2-C1-C10 C1-C2-C3	114-8 (4) 110-6 (4)	C17-C13-C18 C17-C13-C14 C18 C13 C14	111.5 (4) 99.7 (3)
C2-C3-C4 C2-C3-O3 O3-C3-C4	109-1 (5) 109-4 (5)	C13-C14-C15 C13-C14-C15 C13-C14-C8	103.8 (3) 114.3 (3)
C3-C4-C5 C4-C5-C10 C4-C5-C6	111.7 (4) 116.9 (4) 119.3 (4)	C14-C15-C16 C15-C16-C17	101.3 (3) 108.1 (3)
C6-C5-C10	123·7 (4)	C15-C16-O16	112-9 (4)
C5-C6-C7	123·9 (5)	O16-C16-C17	104-9 (4)
C6-C7-C8	111·6 (4)	C16-O16-C22	105-5 (3)
C7–C8–C9	109·6 (3)	C16-C17-C20	104-6 (4)
C7–C8–C14	109·8 (3)	C16-C17-C13	104-4 (3)
C14–C8–C9	111·1 (3)	C20-C17-C13	119-7 (4)
C8-C9-C10	113·2 (3)	C17–C20–C21	114-6 (4)
C8-C9-C11	110·7 (3)	C17–C20–C22	103-3 (4)
C11-C9-C10	114·2 (3)	C21–C20–C22	115-0 (4)
C9-C10-C1	110-1 (4)	C20-C22-C23	114-7 (4)
C9-C10-C5	112-1 (3)	C20-C22-O22	107-3 (4)
C9-C10-C19	110-6 (4)	C20-C22-O16	104-3 (3)
C5C10C19 C1C10C19	108.6 (4) 110.0 (4) 105.3 (4)	023-C22-016 022-C22-C23 022-C22-016	108-9 (4) 111-3 (4) 110-0 (4)
C9-C11-O11	110-8 (4)	C26O22C22	113-5 (4)
C9-C11-C12	112-3 (3)	C22C23C24	112-4 (5)
C11-C12-C13 C12-C13-C14	108-5 (2) 112-5 (3) 107-2 (3)	C23-C24-C25 C24-C25-C26 C24-C25-C27	110.7 (5) 121.1 (8)
C12-C13-C17	115-4 (3)	C27–C25–C26	118-3 (8)
C12-C13-C18	110-8 (4)	C25–C26–O22	112-3 (5)

between envelope and half chair with the puckering 2.18(5) Å, $\angle O(11)$ -H...O(16) 173(4)°], which forms parameters $Q = 0.382 (5) \text{ Å}, \quad \varphi = 263.2 (6)^{\circ}$ and lowest asymmetry factors $fC_2(C17) = 0.035(5)$ and $fC_{\rm s}(016) = 0.074$ (6) Å. The general features of the molecule resemble those of molecule (I) of spirosta-5.25(27)-diene-1β.3β,11α-triol monohydrate (Kálmán, Argay, Ribár, Živanov-Stakić & Vladimirov, 1985). However, there is one difference, namely the B ring of molecule (I) has a pronounced envelope conformation.

There is one hydrogen bond between symmetrically related molecules $[O(11)-H\cdots O(16)(-x, 0.5+y)]$ 0.5 - z): $O(11)\cdots O(16)$ 2.956 (6), $H\cdots O(16)$ a helix about the twofold screw b axis.

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The Structure of 9α -Acetoxy- 8β -[2'-(N-morpholino)ethyl]-10 β -methyl-trans-2-decalone*

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Abstract. $C_{19}H_{31}NO_4$, $M_r = 337.5$, monoclinic, $P2_1/c$, a = 8.289 (8), b = 10.631 (1), c = 21.500 (2) Å, $\beta =$ 94.25 (1)°, $U = 1889 \cdot 1$ (4) Å³, Z=4, $D_r =$ $\lambda(\operatorname{Cu} K\alpha) = 1.5418 \text{ Å}, \quad \mu(\operatorname{Cu} K\alpha) =$ 1.186 g cm^{-3} , $6 \cdot 6 \text{ cm}^{-1}$, F(000) = 736, room-temperature data collection, final R = 0.062 for 1930 reflections. The decalone skeleton adopts a chair-chair conformation which is essentially undistorted even in the presence of the 1,3-diaxial substituents. The mean torsion angle in ring A, the cyclohexanone ring, is 51.4° ; the mean in ring B is 53.4° . The N-morpholinoethyl substituent is extended with an all-trans orientation of bonds C(9)-C(8), C(8)-C(17), C(17)-C(18), C(18)-N(19), N(19)-C(24). The mean distances in the morpholine ring are C-N 1.458, C-C 1.499, and C-O 1.410 Å.

Introduction. Reactions in which a proton alpha‡ to a carbonyl group is abstracted by a general base are important in the function of many enzymes (Spencer, 1979; Westheimer, 1970; Wood, 1971). In our earlier research (Roberts, Ferran, Gula & Spencer, 1980; Hupe, Kendall & Spencer, 1973) we focused on elucidating why these biochemical alpha proton transfers so often occur via iminium ion formation, and concluded that this pathway has an inherent kinetic advantage, at physiological pH, of ca 10⁵ compared with direct deprotonation of the carbonyl compound.



The current emphasis in our research concerns the geometry of alpha proton abstraction. Having the functional groups involved in a reaction localized in the correct positions, rather than free in solution, is generally considered to be an important component of enzymatic catalysis (Jencks, 1969). Accordingly, we have been trying to assess the further contribution to enzymatic rate acceleration of alpha proton transfer

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^{* 4}a-Methyl-8-[2'-(N-morpholino)ethyl]-2-oxo-trans-decahydro-8a-naphthyl acetate.

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[‡] The word 'alpha' is used to denote the position of carbons relative to the carbonyl group and the symbols α and β are used to denote substituents underneath and above, respectively, the plane of the decalin ring system.