

Structure of 8-(2,6-Dideoxy- β -ribo-hexopyranosyl)-5-hydroxy-2-(4-hydroxyphenyl)-7-methoxy-4H-1-benzopyran-4-one Sesquihydrate, Aciculatin

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Abstract. $C_{22}H_{22}O_8 \cdot 1.5H_2O$, $M_r = 441.4$, monoclinic, $I2$, $a = 21.037$ (7), $b = 7.371$ (2), $c = 27.512$ (4) Å, $\beta = 100.34$ (2)°, $V = 4196.8$ (8) Å³, $Z = 8$, $D_x = 1.397$ g cm⁻³, $\lambda(\text{Mo } K\alpha) = 0.71073$ Å, $\mu = 1.029$ cm⁻¹, $F(000) = 1864$, $T = 295$ K, final R (on F) = 0.058 for 2113 observed reflections with $I \geq 2\sigma(I)$. Aciculatin crystallizes as a sesquihydrate with two independent molecules per asymmetric unit. The 2,6-dideoxy- β -ribo-hexopyranosyl ring is linked to C8 of the flavone through a β -C-glycosidic bond. Intramolecular hydrogen bonding occurs between the hydroxyl and ketonic O atoms of the flavone ring system.

Introduction. Flavonoids are a naturally occurring class of plant pigments which frequently occur as glycosides containing either a glucose or rhamnose sugar moiety (Mayer & Cook, 1943). These pigments display a variety of therapeutic properties including anti-carcinogenic, anti-inflammatory, antioxidant and anti-allergenic activities (Glusker & Rossi, 1986).

The present crystallographic study was carried out to confirm the molecular structure of aciculatin, a flavonoid isolated from the plant *Chrysopogon aciculatis* (Carte, DeBrosse, Offen, Carr, Hemling, Berry, Mackenzie-LoCastro & Westley, 1991).

Experimental. Yellow plates were obtained from slow evaporation of an aqueous ethanol solution at room temperature. A suitable crystal of approximate dimensions 0.3 × 0.3 × 0.05 mm was mounted on the tip of a glass fiber with epoxy resin and used in the diffraction study.

Intensity data were collected at room temperature on an Enraf-Nonius CAD-4 diffractometer with graphite-monochromated Mo $K\alpha$ radiation. Lattice parameters obtained by least-squares refinement of angular settings from 25 reflections in 2θ range 24–30°. Intensity data (5603 reflections) collected using variable speed ω - 2θ scans with $2 \leq 2\theta \leq 56^\circ$ ($0 \leq h \leq 27$, $0 \leq k \leq 9$, $-36 \leq l \leq 36$). Three standard reflections ($\bar{3}, 3, \bar{10}$, $4, \bar{3}, 11$, $9\bar{3}6$) monitored every 3 h of X-ray exposure time showed intensity changes of

–5.6%; a correction for deterioration was made (correction: min. 1.000, max. 1.0290). Symmetry-equivalent data were averaged, $R_{\text{int}} = 4.7\%$ (on I); the 5460 unique reflections were corrected for Lorentz and polarization effects.

The structure was solved by direct methods with SHELXS86 (Sheldrick, 1985). The y coordinate of atom OW1 was held fixed to define the origin. Non-H atoms were refined with isotropic temperature factors, then with anisotropic displacement parameters. Most of the H-atom positions were located from difference Fourier maps; those not located were calculated at C—H = 1.00 Å. H-atom coordinates were held fixed at the located or calculated positions along with fixed isotropic temperature factors ($1.3 \times B_{\text{eq}}$ of the adjacent atom). The positions of HW42 and hydroxyl hydrogen, H2A, were not determined. The final refinement converged (max. $\Delta/\sigma = 0.02$) to values of the standard crystallographic agreement factors of $R = 0.058$, $wR = 0.065$ and $S = 1.179$ for 2113 reflections with $I \geq 2\sigma(I)$ and 569 parameters. The function minimized was $\sum w(|F_o| - |F_c|)^2$. Weights were assigned to the data as $w = 4F_o^2/\sigma^2(I)$ where $\sigma(I) = [\sigma(I)^2 + (0.07F_o)^2]^{1/2}$. An extinction coefficient of the form proposed by Zachariasen (1963) was applied and refined: $g = 1.80(1) \times 10^{-7}$. OW1 and OW2 were located on special crystallographic positions with occupancies of 0.5. Neutral atom scattering factors were taken from *International Tables for X-ray Crystallography* (1974, Vol. IV) except for H atoms (Stewart, Davidson & Simpson, 1965). The effects of anomalous dispersion for non-H were included. A final difference Fourier map showed max. excursions of $(\Delta\rho)_{\text{max}} = 0.253$ to $(\Delta\rho)_{\text{min}} = -0.238$ e Å⁻³. The atomic coordinates and equivalent isotropic thermal parameters are collected in Table 1.* All programs

* Lists of anisotropic displacement parameters, selected torsion angles, least-squares planes, H-atom positions and structure factors have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 54288 (22 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

Table 1. Positional and equivalent isotropic thermal parameters with e.s.d.'s

$$B_{eq} = (8\pi^2/3)\sum_i \sum_j U_{ij} a_i^* a_j^* a_i \cdot a_j$$

	x	y	z	B (Å ²)
O1A	0.2695 (2)	0.8396 (9)	-0.1610 (2)	3.2 (1)
O3''A	0.1033 (2)	0.6017 (9)	-0.0495 (2)	3.7 (1)
O4A	0.4627 (2)	0.837 (1)	-0.1602 (2)	6.2 (2)
O4'A	0.0913 (3)	0.878 (1)	-0.3664 (2)	7.3 (2)
O4''A	0.0129 (2)	0.832 (1)	-0.1005 (2)	3.5 (1)
O5A	0.4830 (2)	0.803 (1)	-0.0678 (2)	6.8 (2)
O7A	0.2994 (2)	0.796 (1)	-0.0113 (2)	4.1 (2)
O7''A	0.1812 (2)	0.9665 (8)	-0.0952 (2)	2.9 (1)
C1'A	0.2407 (4)	0.864 (1)	-0.2467 (3)	3.0 (2)
C1''A	0.2111 (3)	0.801 (1)	-0.0750 (3)	2.6 (2)
C2A	0.2930 (3)	0.857 (1)	-0.2037 (3)	2.7 (2)
C2'A	0.1769 (4)	0.892 (1)	-0.2405 (5)	3.9 (2)
C2''A	0.1774 (3)	0.642 (1)	-0.1039 (3)	2.8 (2)
C3A	0.3560 (4)	0.864 (2)	-0.2043 (3)	4.3 (3)
C3'A	0.1281 (4)	0.900 (2)	-0.2812 (3)	4.5 (3)
C3''A	0.1061 (3)	0.644 (1)	-0.0995 (3)	2.3 (2)
C4A	0.4021 (4)	0.843 (2)	-0.1600 (3)	4.7 (3)
C4'A	0.1412 (4)	0.872 (2)	-0.3270 (3)	4.7 (3)
C4''A	0.0759 (3)	0.827 (1)	-0.1143 (3)	2.6 (2)
C5A	0.4184 (4)	0.808 (2)	-0.0692 (3)	4.5 (2)
C5'A	0.2044 (4)	0.844 (2)	-0.3345 (3)	4.2 (2)
C5''A	0.1147 (4)	0.980 (1)	-0.0869 (3)	3.5 (2)
C6A	0.3934 (4)	0.795 (2)	-0.0261 (3)	4.3 (2)
C6'A	0.2533 (4)	0.839 (2)	-0.2936 (3)	4.1 (2)
C6''A	0.0900 (4)	1.164 (1)	-0.1063 (4)	4.8 (3)
C7A	0.3266 (3)	0.802 (1)	-0.0303 (3)	3.2 (2)
C8A	0.2834 (3)	0.812 (1)	-0.0750 (3)	3.1 (2)
C9A	0.3103 (3)	0.829 (1)	-0.1170 (3)	2.8 (2)
C10A	0.3778 (3)	0.827 (2)	-0.1143 (3)	3.7 (2)
C11A	0.3420 (4)	0.787 (2)	0.0585 (3)	6.0 (3)
O1B	0.2703 (2)	0.2042 (9)	0.2072 (2)	3.0 (1)
O3''B	0.1945 (2)	0.4285 (8)	0.0224 (2)	2.9 (1)
O4B	0.4239 (2)	0.194 (1)	0.3198 (2)	4.5 (2)
O4'B	-0.0039 (2)	0.213 (1)	0.2567 (2)	6.5 (2)
O4''B	0.0892 (2)	0.2142 (9)	0.0248 (2)	3.4 (1)
O5B	0.4994 (2)	0.208 (1)	0.2543 (2)	5.5 (2)
O7B	0.3975 (2)	0.256 (1)	0.0858 (2)	5.1 (2)
O7''B	0.2382 (2)	0.0709 (8)	0.1076 (2)	2.6 (1)
C1'B	0.1920 (3)	0.202 (1)	0.2561 (3)	2.9 (2)
C1''B	0.2747 (3)	0.231 (1)	0.1051 (2)	2.5 (2)
C2B	0.2610 (4)	0.202 (1)	0.2542 (3)	3.0 (2)
C2'B	0.1474 (4)	0.159 (1)	0.2144 (3)	3.2 (2)
C2''B	0.2319 (4)	0.399 (1)	0.1087 (3)	3.1 (2)
C3B	0.3116 (4)	0.198 (1)	0.2925 (3)	3.3 (2)
C3'B	0.0819 (4)	0.162 (1)	0.2148 (3)	3.9 (2)
C3''B	0.1719 (4)	0.393 (1)	0.0679 (3)	2.8 (2)
C4B	0.3772 (4)	0.198 (1)	0.2836 (3)	3.7 (2)
C4'B	0.0609 (4)	0.215 (2)	0.2583 (3)	4.2 (2)
C4''B	0.1392 (3)	0.210 (1)	0.0661 (3)	2.7 (2)
C5B	0.4457 (3)	0.216 (2)	0.2196 (3)	3.6 (2)
C5'B	0.1050 (4)	0.254 (2)	0.3006 (3)	4.5 (3)
C5''B	0.1864 (3)	0.056 (1)	0.0657 (3)	2.3 (2)
C6B	0.4516 (4)	0.231 (2)	0.1704 (3)	4.5 (2)
C6'B	0.1703 (4)	0.247 (1)	0.2990 (3)	3.3 (2)
C6''B	0.1569 (4)	-0.127 (1)	0.0696 (3)	3.2 (2)
C7B	0.3955 (3)	0.241 (1)	0.1342 (3)	3.2 (2)
C8B	0.3340 (3)	0.224 (1)	0.1462 (3)	2.8 (2)
C9B	0.3305 (3)	0.209 (1)	0.1952 (3)	2.7 (2)
C10B	0.3852 (3)	0.210 (1)	0.2330 (3)	2.8 (2)
C11B	0.4590 (4)	0.282 (2)	0.0709 (3)	7.8 (4)
OW1	0.000	0.500	0.000	3.8 (2)
OW2	0.500	0.436 (1)	0.500	3.6 (2)
OW3	0.2616 (3)	1.252 (1)	-0.0418 (2)	5.9 (2)
OW4	0.3835 (3)	1.302 (2)	-0.0466 (2)	12.5 (4)

The flavone distances and angles are comparable to those in 4'-bromo-5-hydroxyflavone or 4'-bromo-3-hydroxyflavone (Hayashi, Kawai, Ohno, Iitaka & Akimoto, 1974), hymenoxin (Watson, Kashyap, Gao & Mabry, 1991), glabratephrin (Vleggaar, Kruger, Smalberger & van den Berg, 1978), 3-chloroflavone (Tomlin & Cantrell, 1990), 3-hydroxyflavone (Etter, Urbanczyk-Lipkowska, Baer & Barbara, 1986) or 5-hydroxyflavone (Shoja, 1990), 5-hydroxy-7-methoxyflavone (Shoja, 1989), 5,4'-dihydroxy-3,6,7,8-tetramethoxyflavone (Vijayalakshmi, Rajan, Srinivasan & Ramachandran Nair, 1986), 3',5,5',6'-tetramethoxyflavone (Ting, Watson & Dominguez, 1972), 4',5,7-trihydroxyisoflavone (Breton, Precigoux, Courseille & Hospital, 1975), 5,7,4'-trimethoxyflavanone (Mariezcurrena, 1978), and 5-hydroxy-6-bromo-2'',3'',4',4'',6'',7-hexaacetyl-vitexin or 5-hydroxy-3',6-dibromo-2'',3'',4',4'',6'',7-hexaacetyl-vitexin (Jurnak & Templeton, 1975).

The benzopyran is virtually planar; the dihedral angle between planar ring halves is 1 (2)° for molecule *A* and 3.1 (6)° for molecule *B*. The phenolic substituent is planar, but is not coplanar to the benzopyran in either molecule as evidenced by the C3—C2—C1'—C6' torsion angles of 14 (2) and 18 (2)° for molecules *A* and *B*, respectively. The twisting of the phenolic ring is a result of steric and electronic effects. Theoretical calculations (Glusker & Rossi, 1986; Rossi, Cantrell, Farber, Dyott, Carrell & Glusker, 1980) suggest that for flavones with an H atom at position 3 a torsion angle of 22.8° results in a minimum-energy conformation.

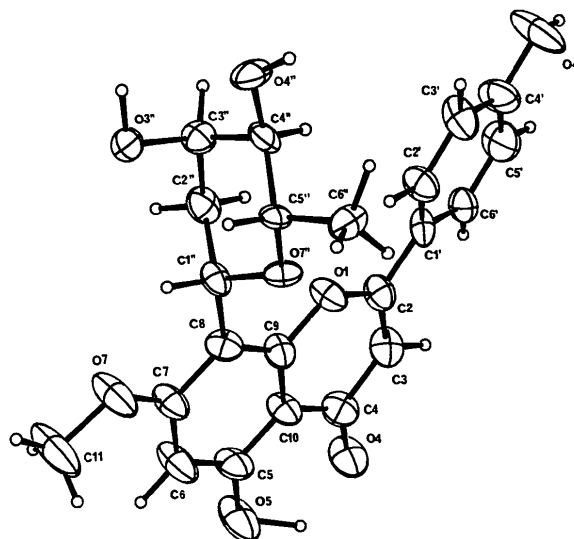


Fig. 1. Molecular structure of aciculatin (molecule *B*) showing labelling scheme. Non-H atoms are drawn with ellipsoids of 50% probability, H atoms are represented as spheres of arbitrary size.

used were from the locally modified Enraf–Nonius (1979) *Structure Determination Package*.

Discussion. Aciculatin crystallizes as a sesquihydrate with two independent molecules per asymmetric unit. The molecular structure of molecule *B* is presented in Fig. 1. Selected bond distances and bond angles are listed in Table 2. Intermolecular hydrogen bonding interactions are collected in Table 3.

Table 2. Selected bond distances (Å) and angles (°) with e.s.d.'s

O1A	C2A	1.361 (9)
O1A	C9A	1.353 (9)
O3''A	C3''A	1.424 (9)
O4A	C4A	1.28 (1)
O4'A	C4'A	1.37 (1)
O4''A	C4''A	1.443 (9)
O5A	C5A	1.35 (1)
O7A	C7A	1.370 (9)
O7A	C11A	1.441 (9)
O7''A	C1''A	1.44 (1)
O7''A	C5''A	1.46 (1)
C1'A	C2A	1.46 (1)
C1'A	C2'A	1.40 (1)
C1'A	C6'A	1.38 (1)
C1''A	C2''A	1.52 (1)
C1''A	C8A	1.52 (1)
C2A	C3A	1.33 (1)
C2'A	C3'A	1.38 (1)
C2''A	C3''A	1.53 (1)
C3A	C4A	1.42 (1)
C3'A	C4'A	1.35 (1)
C3''A	C4''A	1.51 (1)
C4A	C10A	1.44 (1)
C4'A	C5'A	1.40 (1)
C4''A	C5''A	1.51 (1)
C5A	C6A	1.38 (1)
C5A	C10A	1.38 (1)
C5'A	C6'A	1.38 (1)
C5''A	C6''A	1.51 (1)
C6A	C7A	1.39 (1)
C7A	C8A	1.40 (1)
C8A	C9A	1.38 (1)
C9A	C10A	1.41 (1)

C2A	O1A	C9A	120.5 (6)
C7A	O7A	C11A	118.0 (6)
C1''A	O7''A	C5''A	111.2 (7)
C2A	C1'A	C2'A	120.4 (8)
C2A	C1'A	C6'A	120.6 (8)
C2'A	C1'A	C6'A	119.0 (8)
O7''A	C1''A	C2''A	108.7 (6)
O7''A	C1''A	C8A	108.9 (8)
C2''A	C1''A	C8A	114.3 (8)
O1A	C2A	C1'A	111.3 (7)
O1A	C2A	C3A	122.0 (8)
C1'A	C2A	C3A	126.6 (8)
C1'A	C2'A	C3'A	119.9 (9)
C1''A	C2''A	C3''A	109.0 (7)
C2A	C3A	C4A	121.0 (8)
C2'A	C3'A	C4'A	120.3 (9)
O3''A	C3''A	C2''A	106.7 (7)
O3''A	C3''A	C4''A	111.3 (7)
C2''A	C3''A	C4''A	110.9 (7)
O4A	C4A	C3A	121.9 (8)
O4'A	C4'A	C10A	120.7 (8)
C3A	C4A	C10A	117.4 (8)
O4'A	C4'A	C3'A	118.6 (9)
O4''A	C4''A	C5''A	120.2 (9)
C3'A	C4'A	C5'A	121.3 (9)
O4''B	C4''B	C5''B	114.2 (8)
C3''B	C4''B	C5''B	112.2 (6)
O5B	C5B	C6B	119.1 (7)
O5B	C5B	C10B	120.3 (8)
C6B	C5B	C10B	120.6 (8)
C4''B	C5''B	C6''B	119.0 (9)
O7''B	C5''B	C4''B	109.8 (7)
O7''B	C5''B	C6''B	106.1 (7)
O4'A	C4'A	C3'A	108.3 (8)
O4''A	C4''A	C5''A	106.8 (7)
C3''A	C4''A	C5''A	111.6 (7)
O5A	C5A	C6A	120.6 (8)
O5A	C5A	C10A	118.9 (8)
C6A	C5A	C10A	120.6 (7)
C4'A	C5'A	C6'A	118.2 (9)
O7''A	C5''A	C4''A	108.6 (7)
O7''A	C5''A	C6''A	106.3 (8)
C4''A	C5''A	C6''A	111.9 (8)
C5A	C6A	C7A	117.5 (8)
C1'A	C6'A	C5'A	121.3 (8)
O7A	C7A	C6A	119.9 (7)
O7A	C7A	C8A	115.7 (7)
C6A	C7A	C8A	124.3 (8)
C1''A	C8A	C7A	119.3 (8)

O1B	C2B	1.34 (1)
O1B	C9B	1.367 (9)
O3''B	C3''B	1.44 (1)
O4B	C4B	1.267 (9)
O4'B	C4'B	1.356 (9)
O4''B	C4''B	1.404 (9)
O5B	C5B	1.343 (9)
O7B	C7B	1.346 (9)
O7B	C11B	1.44 (1)
O7''B	C1''B	1.42 (1)
O7''B	C5''B	1.442 (9)
C1'B	C2B	1.46 (1)
C1'B	C2'B	1.38 (1)
C1'B	C6'B	1.38 (1)
C1''B	C2''B	1.54 (1)
C1''B	C8B	1.53 (1)
C2B	C3B	1.36 (1)
C2'B	C3'B	1.38 (1)
C2''B	C3''B	1.53 (1)
C3B	C4B	1.45 (1)
C3'B	C4'B	1.40 (1)
C3''B	C4''B	1.51 (1)
C4B	C10B	1.43 (1)
C4'B	C5'B	1.38 (1)
C4''B	C5''B	1.51 (1)
C5B	C6B	1.38 (1)
C5B	C10B	1.39 (1)
C5'B	C6'B	1.38 (1)
C5''B	C6''B	1.49 (1)
C6B	C7B	1.40 (1)
C7B	C8B	1.40 (1)
C8B	C9B	1.37 (1)
C9B	C10B	1.40 (1)

C1''A	C8A	C9A	124.5 (7)
C7A	C8A	C9A	116.2 (7)
O1A	C9A	C8A	117.5 (7)
O1A	C9A	C10A	121.3 (7)
C8A	C9A	C10A	121.0 (7)
C4A	C10A	C5A	122.2 (8)
O7''A	C1''A	C9A	117.6 (8)
C5A	C10A	C9A	120.2 (7)
C2B	O1B	C9B	122.3 (6)
C4''B	C5B	C6''B	113.2 (7)
C5B	C6B	C7B	119.1 (8)
C1'B	C6'B	C5'B	121.3 (9)
O7B	C7B	C6B	122.4 (7)
O7B	C7B	C8B	115.8 (7)
C6B	C7B	C8B	121.6 (8)
C1''B	C8B	C7B	119.4 (7)
C1''B	C8B	C9B	123.5 (7)
C7B	O7B	C11B	118.6 (6)
C1''B	O7''B	C5''B	111.3 (6)
C2B	C1'B	C2'B	120.1 (8)
C2B	C1'B	C6''B	120.9 (8)
C2'B	C1'B	C6'B	119.1 (8)
O7''B	C1''B	C2''B	109.7 (6)
O7''B	C1''B	C8B	108.2 (8)
C2''B	C1''B	C8B	112.9 (8)
O1B	C2B	C1'B	110.6 (7)
O1B	C2B	C3B	121.3 (8)
C1'B	C2B	C3B	128.1 (8)
C1'B	C2'B	C3'B	121.3 (8)
C1''B	C2''B	C3''B	110.2 (8)
C2B	C3B	C4B	120.6 (8)
C2'B	C3'B	C4'B	118.6 (8)
O3''B	C3''B	C2''B	106.0 (7)
O3''B	C3''B	C4''B	110.6 (7)
C2''B	C3''B	C4''B	110.9 (8)
O4B	C4B	C3B	119.7 (8)
O4B	C4B	C10B	123.7 (8)
C3B	C4B	C10B	116.5 (7)
O4'B	C4'B	C3'B	116.0 (9)
O4''B	C4''B	C5''B	123.2 (9)
C3'B	C4'B	C5'B	120.6 (8)
O4''B	C4''B	C3''B	106.0 (7)
C7B	C8B	C9B	117.1 (7)
O1B	C9B	C8B	117.2 (7)
O1B	C9B	C10B	119.5 (7)
C8B	C9B	C10B	123.2 (8)
C4B	C10B	C5B	122.2 (7)
C4B	C10B	C9B	119.7 (7)
C5B	C10B	C9B	118.1 (8)

Table 3. Intermolecular hydrogen-bonding interactions

D—H...A	D...A (Å)	D—H...A (°)	Symmetry operation*
O4''B—H6B...O4A	2.67 (1)	179	(iii) 00 - 1
O4''A—H15A...O4B	2.81 (1)	150	(iii) - 10 - 1
O4''B—H13B...O3''A	2.93 (1)	176	(i) 000
O4''B—H13B...O3''A	2.928 (6)	145	(iii) - 10 - 1
O3''B—H13B...O3''A	2.80 (1)	150	(i) 000
O4''B—H15B...O4''B	2.77 (1)	177	(i) 0 - 10
O4''A—H6A...O4''A	2.60 (1)	163	(iv) 0 1 - 1
O4''B—H15B...O4''B	2.62 (1)	165	(i) 000
O4''B—H15B...O4''B	2.78 (1)	174	(iii) - 1 - 1 - 1
O4''B—H15B...O4''B	2.82 (1)	140	(i) 000
O3''A—H13A...O4''B	2.863 (6)	143	(i) 000

* Translations are along x , y and z , respectively, with the symmetry operators defined as: (i) x, y, z ; (ii) $-x, y, -z$; (iii) $\frac{1}{2} + x, \frac{1}{2} + y, \frac{1}{2} + z$; (iv) $\frac{1}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$.

Intramolecular hydrogen bonding occurs between the hydroxyl and ketonic O atoms of the flavone ring system. In molecule *B*, the hydroxyl H atom, H2B, located directly from the difference Fourier map, shows hydrogen bonding to the ketonic O atom, O4B (O5B—H2B = 1.21, O5B...O4B = 2.61 (1), H2B...O4B = 1.76 Å and O5B—H2B...O4B = 122°). The related H atom was not located for molecule *A*; however, its presence is inferred from the distance of O5A...O4A = 2.514 (9) Å and the absence of other short contacts to O5A. The intramolecular hydrogen bonding of aciculatin is similar to that exhibited in the related hydroxyflavones, hymenoxin (Watson, Kashyap, Gao & Mabry, 1991), 3-hydroxyflavone (Etter, Urbanczyk-Lipkowska, Baer & Barbara, 1986) or 5-hydroxyflavone (Shoja, 1990), 5-hydroxy-7-methoxyflavone (Shoja, 1989), 5,4'-dihydroxy-3,6,7,8-tetramethoxyflavone (Vijayalakshmi, Rajan, Srinivasan & Ramachandran Nair, 1986) and 4',5,7-trihydroxyisoflavone (Breton, Precigoux, Courseille & Hospital, 1975).

The 2,6-dideoxy- β -ribo-hexopyranosyl ring is linked to C8 of the flavone through a β -C-glycosidic bond. The hexopyranosyl ring adopts a chair conformation with C1'' and C4'' located on opposite sides of the plane generated by C2''—C3''—C5''—O7''. This is similar to both pyran rings in isokidamycin bis(*m*-bromobenzoate) (Furukawa & Iitaka, 1974, 1980) or one of the pyran rings (ring *E*) in triacetyl-methoxykidamycin bis(trimethylammonium) iodide (Furukawa, Itai & Iitaka, 1973; Furukawa & Iitaka, 1974) and the β -D-glucosyl ring in 5-hydroxy-3',6'-dibromo-2'',3'',4'',4'',6'',7'-hexaacetylvitexin (Jur-nak & Templeton, 1975). The 4''-OH and 5''-CH₃ substituents adopt equatorial positions while the 3''-OH substituent adopts an axial position. As with the vitexin derivative, the plane of the hexopyranosyl ring is essentially perpendicular to the benzopyran ring [molecule *A* 91.4 (3), molecule *B* 88.9 (3)°] in order to minimize steric interactions.

The two independent molecules of aciculatin also differ in the magnitude of the torsion angles associated with the hydroxyl groups on the phenolic and

hexopyranosyl rings. The 4'-OH [C5'A—C4'A—O4'A—H6A = -8 (1) and C5'B—C4'B—O4'B—H6B = 30 (2)°] and 3''-OH (C4''A—C3''A—O3''A—H13A = 93.1 (8) and C4''B—C3''B—O3''B—H13B = 74.9 (7)°] torsion angles are moderately different. The largest difference in torsion angles occurs with the 4''-hydroxyls [C5''A—C4''A—O4''A—H15A = 128.4 (7) and C5''B—C4''B—O4''B—H15B = 64.6 (8)°].

Intermolecular hydrogen bonding (Table 3) occurs between O3''A and O3''B [O3''B...O3''A = 2.80 (1) Å and 150°] of the two independent molecules. In addition, hydrogen bonding occurs between the ketonic O atom O4A and the phenolic O atom O4'B [O4'B...O4A = 2.67 (1) Å and 179°] and between the ketonic O atom O4B and the hydroxyl O atom O4''A [O4''A...O4B = 2.81 (1) Å and 150°]. There is also a network of hydrogen-bonding interactions between the aciculatin molecules and the water molecules which stabilizes the crystal lattice.

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Structure of Cortivazol, 11 β ,17 α ,21-Trihydroxy-6,16 α -dimethyl-2'-phenyl-2'H-pregna-2,4,6-trieno[3,2-c]pyrazol-20-one 21-Acetate

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Abstract. C₃₂H₃₈N₂O₅, $M_r = 530.7$, monoclinic, C_2 , $a = 30.625$ (5), $b = 6.229$ (2), $c = 15.289$ (2) Å, $\beta = 93.86$ (2)°, $V = 2909.8$ Å³, $Z = 4$, $D_x = 1.211$ g cm⁻³, $\lambda(\text{Cu } K\alpha) = 1.5418$ Å, $\mu = 6.2$ cm⁻¹, $F(000) = 1136$, $T = 292$ K, final $R = 0.047$ for 2415 reflections with $I > 2.5\sigma(I)$. All bond lengths and angles are within normal limits. Ring *A* with two double bonds is not

planar, but is in the 1 α ,10 β half-chair conformation. Ring *B* is in a 9 α ,10 β half-chair conformation distorted towards a 9 α sofa. Ring *C* is in the expected chair conformation, whereas ring *D* is in the 13 β envelope conformation. The C20, C26, C27 and N29 substituents are equatorial, O11, C18 and C19 are β axial, and O17 and C25 are α axial. Rings *C* and *D*