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#### **Key indicators**

Single-crystal X-ray study T = 294 K Mean  $\sigma$ (C–C) = 0.009 Å R factor = 0.054 wR factor = 0.095 Data-to-parameter ratio = 7.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 3*a*,20-Bis(dimethylamino)-4,4,14-trimethyl-9,19-cyclo-5*a*-pregnan-16-ol

The title compound,  $C_{28}H_{50}N_2O$ , also known as cyclovirobuxine A, was obtained from *N*-methylation of naturally occurring cyclovirobuxine D. It has a four-ring triterpenoid nucleus in a *trans–cis–trans* configuration. The six-membered rings have chair conformations, while the conformation of the five-membered ring is an envelope. Received 11 September 2006 Accepted 13 September 2006

### Comment

The Buxus plants are a rich source of triterpenoid alkaloids. Previous phytochemical studies on the Buxus species have resulted in the isolation of more than 200 such compounds (Rahman & Choudhary, 1998). In the traditional system of medicine, extracts of genus Buxus have been used for the treatment of various disorders (Cordell, 1981). The title compound, (I), also known as cyclovirobuxine A, possessing potential applications in the treatment of cardiovascular and cerebrovascular diseases (Chiu & Nie, 1992), was isolated and its structure elucidated for the first time by Khuong-Huu-Laine *et al.* (1966). We have semi-synthesized the title compound, (I), from naturally occurring cyclovirobuxine D.



In the molecule of (I) (Fig. 1), the bond lengths and angles are within normal ranges (Allen *et al.*, 1987). The structure of (I) contains the fused four-ring triterpenoid system: A, B, C and D. Similar to cyclovirobuxine D (Choudhary *et al.*, 2003), the triterpenoid nucleus has a *trans–cis–trans* configuration for ring junctions A/B, B/C and C/D.

Rings *A*, *B* and *C* are not planar, having total puckering amplitudes,  $Q_{\rm T}$ , of 0.554 (3), 0.510 (4) and 0.625 (4) Å, respectively, and chair conformations [ $\varphi = -95.77 (10)^{\circ}$ ,  $\theta =$ 3.17 (5)°;  $\varphi = 85.75 (7)^{\circ}$ ,  $\theta = 42.72 (6)^{\circ}$ ; and  $\varphi = -89.92 (5)^{\circ}$ ,  $\theta =$ 110.47 (5)°; Cremer & Pople, 1975]. The conformation of ring *D* is an envelope, with atom C11 at the flap position, 0.701 (5) Å from the mean plane through the other four atoms.

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Figure 1

The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

The dimethylamino substituents are attached to rings A and D; the torsion angles N1-C1-C17-C16  $[-172.4 (5)^{\circ}]$  and C19-N1-C1-C17 [92.9 (6)°] are indicative of a (+)-synclinal conformation. The sums of the bond angles around N1 (335.8°) and N2 (335.4°) indicate  $sp^3$  character of the two N atoms.

## **Experimental**

Wood extract containing cyclovirobuxine D, (1) (65.9%), cyclobuxine D (24.3%) and other Buxus alkaloids (9.84%) was purchased from Jiangsu, China. The crude alkaloid was then purified according to the literature method (Choudhary *et al.*, 2003) followed by recrystallization from acetone. The melting point of purified compound (1) is 493–494 K (acetone, decomp.), similar to that (494-497 K) reported in the literature (Brown & Kupchan, 1964). To a high-pressure autoclave (1000 ml) containing Pd/C (1 g, 10%) and C<sub>2</sub>H<sub>5</sub>OH (300 ml) were added aqueous formaldehyde (15 ml, 37%) and purified cyclovirbuxine D (10 g). The *N*-methylation reaction was performed by stirring with H<sub>2</sub> at 2 atmospheres pressure at 363 K. After 6 h, the mixture was filtered, the solvent was evaporated and the residue was recrystallized from acetone. The melting point is 514–515 K (decomp.), similar to that reported in the literature (Khuong-HuuLaine *et al.*, 1966)

Crystal data

$C_{28}H_{50}N_2O$	V = 635.7 (6) Å <sup>3</sup>
$M_r = 430.70$	Z = 1
Triclinic, P1	$D_x = 1.125 \text{ Mg m}^{-3}$
a = 6.245 (3) Å	Mo $K\alpha$ radiation
b = 7.106 (4)  Å	$\mu = 0.07 \text{ mm}^{-1}$
c = 15.858 (8) Å	T = 294 (2) K
$\alpha = 77.993 \ (9)^{\circ}$	Plate, colorless
$\beta = 87.575 \ (10)^{\circ}$	$0.45 \times 0.38 \times 0.10 \text{ mm}$
$\gamma = 67.574 \ (8)^{\circ}$	

#### Data collection

Refinement

Bruker SMART CCD area-detector	3380 measured reflections
diffractometer	2224 independent reflections
$\omega$ and $\omega$ scans	1002 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan	$R_{\rm int} = 0.042$
(SABABS; Sheldrick, 1996)	$\theta_{\rm max} = 25.0^{\circ}$
$T_{\min} = 0.971, \ T_{\max} = 0.993$	

Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.054$	H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0225P)^2]$
$wR(F^2) = 0.095$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
2224 reflections	$\Delta \rho_{\rm max} = 0.16 \text{ e } \text{\AA}^{-3}$
280 parameters	$\Delta \rho_{\rm min} = -0.16 \text{ e } \text{\AA}^{-3}$

H atoms were positioned geometrically, with O-H = 0.82 Å, and C-H = 0.98, 0.97 and 0.96 Å for methine, methylene and methyl H, respectively, and constrained to ride on their parent atoms, with  $U_{iso}(H) = xU_{eq}(C,O)$ , where x = 1.2 for methine and methylene H, and x = 1.5 for all other H atoms. In the absence of significant anomalous scattering effects, Friedel pairs were merged; the absolute configuration is assigned arbitrarily.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

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