

**5 α ,8 α -Epidioxyergosta-6,22-dien-3 β -ol
(ergosterol peroxide) methanol solvate**

Jian-Feng Wang,^{a*} Yao-Jian Huang,^b Mei-Juan Fang,^a Wen-Ling Xie,^b Wen-Jin Su^b and Yu-Fen Zhao^a

^aThe Key Laboratory for Chemical Biology of Fujian Province, Xiamen University, Xiamen 361005, China, and ^bDepartment of Biology, Xiamen University, Xiamen 361005, China

Correspondence e-mail:
jfwang@yanan.xmu.edu.cn

Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.009\text{ \AA}$
 R factor = 0.068
 wR factor = 0.170
Data-to-parameter ratio = 8.6

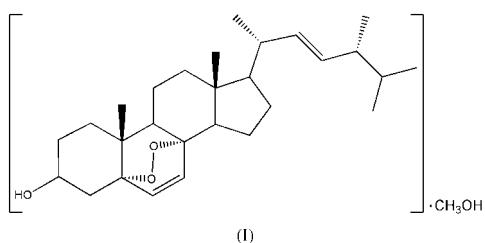
For details of how these key indicators were automatically derived from the article, see
<http://journals.iucr.org/e>.

The title compound was isolated from *Tubercularia sp.*, an endophytic fungus of *Taxus mairei*, and crystallizes as a methanol solvate, $\text{C}_{28}\text{H}_{44}\text{O}_3\cdot\text{CH}_3\text{OH}$. The crystal structure shows that the OH group complexes with the methanol solvent molecule via intermolecular hydrogen bonds, and that the peroxy unit has an O—O bond length of 1.482 (5) Å and a C—O—O—C torsion angle of $-6.0(6)^\circ$.

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Comment

5 α ,8 α -Epidioxyergosta-6,22-dien-3 β -ol (ergosterol peroxide), was isolated from *Tubercularia sp.*, an endophytic fungus with antitumor activity (Wang *et al.*, 2000). It is found that ergosterol peroxide has various important biological activities, including immunosuppressive (Fujimoto *et al.*, 1994), anti-inflammatory (Yasukawa *et al.*, 1996), antiviral (Lindequist *et al.*, 1989; Nakanishi *et al.*, 1998), antiplasmodial (Kuria *et al.*, 2002) and antitumor (Bok *et al.*, 1999; Nam *et al.*, 2001) activities.



The title compound crystallizes as a methanol solvate, (I) (Fig. 1), and the OH group complexes with the methanol solvent molecule via intermolecular hydrogen bonds, with an $\text{O}1\text{S}\cdots\text{O}1(-x + 1, y - \frac{1}{2}, -z + 2)$ distance of 2.666 (9) Å and an $\text{O}1\cdots\text{O}1\text{S}(x, y + 1, z)$ distance of 2.695 (12) Å. The structure of (I) is similar to that of 5 α ,8 α -epidioxyergosta-6,22-

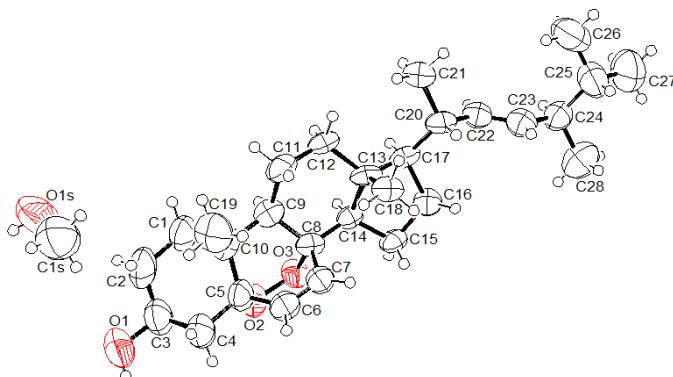


Figure 1
ORTEP-3 (Farrugia, 1997) plot of (I) at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.

dien-3 β -acetate, (II) (Kutshabsky *et al.*, 1990). The two structures differ in that (I) contains a hydroxy group instead of an acetate group. The peroxy distance [O—O = 1.482 (5) Å] in (I) is close to that in (II) [O—O = 1.48 (1) Å]. The torsion angle [C—O—O—C = -6.0 (6) $^\circ$] of the peroxy unit in (I) is very similar to that in (II) [C—O—O—C = -5.6 (7) $^\circ$].

Experimental

The title compound was isolated from an endophytic fungus, *Tuberularia sp.*, which was found in the inner bark of *Taxus mairei* of Fujian Province, China. Crystals were grown from methanol. The molecular formula of the title compound was deduced from the high resolution EI-MS spectrum as C₂₈H₄₄O₃, showing an accurate mass of *m/z* 428.3304 (M^+ , Δ = 1.4 mmu).

Crystal data

C ₂₈ H ₄₄ O ₃ ·CH ₄ O	$D_x = 1.081 \text{ Mg m}^{-3}$
$M_r = 460.67$	Mo K α radiation
Monoclinic, $P2_1$	Cell parameters from 684
$a = 10.022 (5) \text{ \AA}$	reflections
$b = 7.373 (5) \text{ \AA}$	$\theta = 2.9\text{--}24.4^\circ$
$c = 19.156 (5) \text{ \AA}$	$\mu = 0.07 \text{ mm}^{-1}$
$\beta = 91.556 (5)^\circ$	$T = 298 (2) \text{ K}$
$V = 1415.0 (12) \text{ \AA}^3$	Needle, colorless
$Z = 2$	$0.43 \times 0.18 \times 0.11 \text{ mm}$

Data collection

Bruker SMART APEX area-detector diffractometer	2692 independent reflections
φ and ω scans	1468 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2001)	$R_{\text{int}} = 0.079$
$T_{\min} = 0.909$, $T_{\max} = 0.992$	$\theta_{\max} = 25.0^\circ$
7222 measured reflections	$h = -11 \rightarrow 10$
	$k = -8 \rightarrow 8$
	$l = -14 \rightarrow 22$

Refinement

Refinement on F^2	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.069$	$w = 1/[\sigma^2(F_o^2) + (0.0389P)^2]$
$wR(F^2) = 0.170$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.09$	$(\Delta/\sigma)_{\max} = 0.013$
2692 reflections	$\Delta\rho_{\max} = 0.13 \text{ e \AA}^{-3}$
312 parameters	$\Delta\rho_{\min} = -0.12 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bonding geometry (Å, $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1S—H1S \cdots O1 ⁱ	0.82	1.89	2.666 (9)	158
O1—H1 \cdots O1S ⁱⁱ	0.79 (8)	1.94 (8)	2.695 (12)	160 (9)

Symmetry codes: (i) $1 - x, y - \frac{1}{2}, 2 - z$; (ii) $x, 1 + y, z$.

The H atom on atom O1 was located in a difference Fourier synthesis and refined freely. All other H atoms were positioned geometrically (C—H = 0.93, 0.96, 0.97 and 0.98 Å, and O—H = 0.82 Å) and were included in the refinement in the riding-model approximation. The displacement parameters of H atoms were set to 1.2U_{eq} of their parent atoms. The absolute configuration was taken as that of 5 α ,8 α -epidioxyergosta-6,22-dien-3 β -acetate (Kutshabsky *et al.*, 1990). In the absence of significant anomalous scattering effects, Friedel pairs were merged.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SIR97 (Altomare *et al.*, 1999); program(s) used to refine

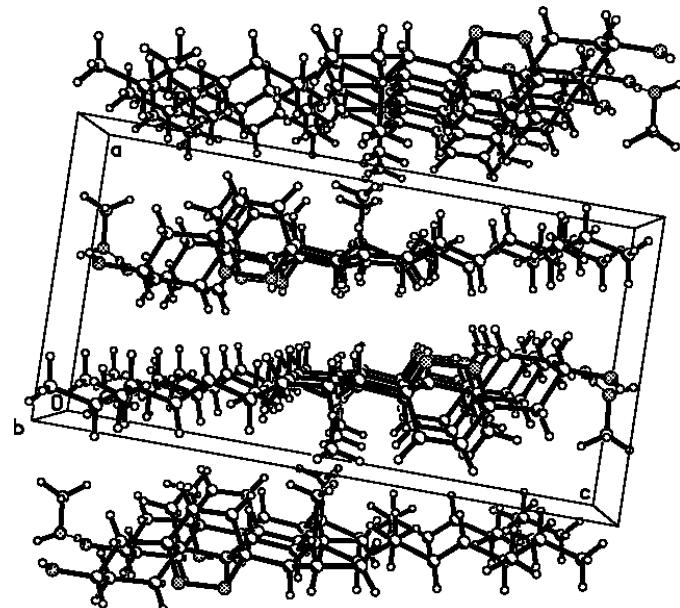


Figure 2
Packing diagram (ORTEPII; Johnson, 1976) of (I).

structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997) and ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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