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6a,7-Dihydro-2,6-dimethyl-7-phenyl-6*H*chromeno[4,3-*d*]-1,2,4-triazolo[1,5-*a*]pyrimidin-6a-ol†

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Abstract

In the title compound, $C_{20}H_{18}N_4O_2$, the dihydropyrimidine and dihydropyran rings adopt a distorted sofa conformation. The phenyl substituent at C7 and the hydroxy group at C6a adopt axial positions, whereas the methyl group at C6 is equatorial.

Comment

Derivatives of 1,2,4-triazolo[1,5-a]pyrimidine possess high biological activity, for example, as cardiovascular agents (Tsuda *et al.*, 1986). The chemical and physiological characteristics of these compounds are directly related to their conformations. However, relevant data on

© 1997 International Union of Crystallography Printed in Great Britain – all rights reserved the structures of dihydro-1,2,4-triazolo[1,5-a]pyrimidines condensed with pyran rings, (I), or other heterocycles have not previously been available.



Torsion angles N7a—C10a—N11—C11a and C1— C1a—C11a—N11 are 15.4 (4) and 9.4 (4)°, respectively. This indicates some loss of planarity of the conjugated system of the molecule, although the C1a—C11a, N11—C11a and C10a—N11 bond lengths [1.453 (4), 1.294 (3) and 1.381 (3) Å, respectively] are evidence of the conjugation between the benzene ring, azomethine group and 1,2,4-triazole ring (Fig. 1).



Fig. 1. A view of (I) with displacement ellipsoids drawn at the 50% probability level. The H atoms are omitted for clarity.

The conformations of both the pyrimidine and pyran rings can be described as distorted sofas. The puckering coordinates (Zefirov, Palyulin & Dashevskaya, 1990) are S = 0.63, $\theta = 50.9$ and $\varphi = 18.3$ for the pyrimidine, and S = 0.72, $\theta = 44.7$ and $\varphi = 10.2$ for the pyran moiety, where S is the puckering degree, and θ and φ are polar angles which describe the ring conformation type (values of these angles for ideal conformations are: sofa $\theta = 45$, $\varphi = 0$; half chair $\theta = 45$, $\varphi = 30$). Comparison of the title compound with *cis*-3-bromo-2,3-dihydro-2-phenyl-4H-1-benzopyran-4-one and (2R,3R)-3,5,7-trihydroxy-2-(2R*,3R*)-3-(4-hydroxy-3-methoxyphenyl)-2-hydroxymethyl-(1,4-benzodioxan-6-yl)-4-chromanone

[†] Alternative name: 6a,7-dihydro-2,6-dimethyl-7-phenyl-6*H*-[1,2,4]-triazolo[2',3':1,2]pyrimidino[5,4-*c*]chromen-6a-ol.

shows that the dihydropyranone ring adopts a distorted half-chair conformation and annelation of the ring concerned leads to the change in its conformation (Cantrell & Hockstein, 1982; Lotter & Wagner, 1983). An alkyl substituent in the second position has an equatorial orientation (Geddes, Sheldrick & Akrigg, 1980; Chambers & Marfat, 1994). Comparison of (I) with 5,7-diphenyl-6-oxy-6,7-dihydropyrazolo[1,5-*a*]pyrimidine, the puckering coordinates for which are S = 0.61, $\theta = 46.2$ and $\varphi = 28.9$, indicates that annelation of the dihydropyrimidine ring results in a decrease in its twisting (Desenko *et al.*, 1993).

The crystal is composed of a racemic mixture of the (6R, 6aS, 7R) and (6S, 6aR, 7S) isomers. The phenyl and hydroxyl substituents on C7 and C6a are both in axial positions [the C10a—N7a—C7—C15 and O12—C6a—C11a—C1a torsion angles are 95.7 (3) and -104.2 (3)°, respectively]. The methyl group is equatorial; the C4a—O5—C6—C14 and C11a—C6a—C6—C14 torsion angles are -178.7 (2) and -168.1 (2)°, respectively.

In the crystal, the molecules of (I) form chains due to intermolecular hydrogen bonds O12—H12O···N10ⁱ, characterized by H···N 1.92 (4) Å and O—H···N 174 (4)° [symmetry code: (i) $\frac{1}{2} - x$, $\frac{1}{2} + y$, $\frac{3}{2} - z$].

Experimental

Compound (I) was obtained from 2,9-dimethyl-3-phenyl-3,7-dihydro-2*H*-1,2,4-triazolo[1',5'-a']pyrimido[4,5-d]benzo-[b]pyran by dissolving the latter in DMSO and allowing slow aerial oxidation to occur over a period of two weeks. Spectral characteristics of (I) are as follows: UV spectrum (isopropanole, Specord M40 spectrometer): Λ max (ε /1000): 313 (20.7), 376 (20.8); IR spectrum (KBr tablets, Specord 75 IR spectrometer): ν (C=N) 1618 cm⁻¹; PMR spectrum (DMSO-d₆, Varian VXR-300 spectrometer): δ (p.p.m.): 6.8– 7.9 (8H, m), 7.94 (1H, s, 6-H), 7.18 (1H, s, OH), 5.59 (1H, s, 3-H), 3.68 (1H, q, J = 6.4 Hz, 2-H), 2.32 (3H, s, 13-CH₃), 1.41 (3H, d, J = 6.4 Hz, 14-CH₃). Single crystals were grown by diffusion of benzene into a solution of (I) in DMSO.

Crystal data

Siemens P3/PC diffractom-

eter

$C_{20}H_{18}N_4O_2$	Mo $K\alpha$ radiation
$M_r = 346.38$	$\lambda = 0.71073 \text{ Å}$
Monoclinic	Cell parameters from 24
$P2_{1}/n$	reflections
a = 10.770(2) Å	$\theta = 11 - 12^{\circ}$
b = 10.045(2) Å	$\mu = 0.089 \text{ mm}^{-1}$
c = 16.499 (4) Å	T = 293 (2) K
$\beta = 105.07(2)^{\circ}$	Plate
V = 1723.6 (6) Å ³	$0.40 \times 0.20 \times 0.10 \text{ mm}$
Z = 4	Yellow
$D_x = 1.335 \text{ Mg m}^{-3}$	
D_m not measured	
Data collection	

 $R_{\rm int} = 0.023$

 $\theta_{\rm max} = 25.04^{\circ}$

θ –2 θ scans	$h = 0 \rightarrow 12$
Absorption correction: none	$k = -11 \rightarrow 0$
2618 measured reflections	$l = -19 \rightarrow 18$
2467 independent reflections	2 standard refle
1685 reflections with	every 98 refle
$I > 2\sigma(I)$	intensity deca

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.201$ S = 1.1522365 reflections 241 parameters H atoms riding $w = 1/[\sigma^2(F_o^2) + (0.0575P)^2 + 0.7176P]$ where $P = (F_o^2 + 2F_c^2)/3$ $k = -11 \rightarrow 0$ $l = -19 \rightarrow 18$ 2 standard reflections every 98 reflections intensity decay: 5%

 $(\Delta/\sigma)_{max} = 0.037$ $\Delta\rho_{max} = 0.17 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{min} = -0.19 \text{ e } \text{\AA}^{-3}$ Extinction correction: none Scattering factors from International Tables for Crystallography (Vol. C)

Refinement on F^2 was carried out on all reflections except for 66 with very negative F^2 or having $|(F_o - F_c)| > 4\sigma(F_o)$ and therefore flagged by the user for potential systematic errors. Standard uncertainties on C—C bond distances are 0.004 Å.

Data collection: P3/PC Diffractometer Program (Siemens, 1989). Cell refinement: P3/PC Diffractometer Program. Data reduction: SHELXTL (Sheldrick, 1994) . Program(s) used to solve structure: SHELXTL. Program(s) used to refine structure: SHELXTL. Molecular graphics: SHELXTL. Software used to prepare material for publication: SHELXTL.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BM1134). Services for accessing these data are described at the back of the journal.

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