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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.005 \text{ Å}$ R factor = 0.061 wR factor = 0.170 Data-to-parameter ratio = 16.2

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

4'-(4-Chlorophenyl)-1'-methyl-4",5",6",7"tetrahydro-1*H*-indole-3-spiro-2'-pyrrolidine-3'-spiro-2"-(thiazolo[3,2-*a*]pyrimidine)-2(3*H*),-3"(2"*H*)-dione

The title compound, $C_{23}H_{21}ClN_4O_2S$, was synthesized by the intermolecular [3 + 2]-cycloaddition of azomethine ylide, derived from isatin and sarcosine by a decarboxylative route, and 2-(4-chlorobenzylidene)-6,7-dihydro-5*H*-thiazolo[3,2-*a*]-pyrimidin-3-one. In the molecule, the two spiro junctions link a planar 2-oxindole ring, a pyrrolidine ring in an envelope conformation and a 6,7-dihydro-5*H*-thiazolo[3,2-*a*]-pyrimidin-3-one ring. Two molecules are connected into a dimer by two N-H···N hydrogen bonds

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Comment

Spiro-compounds represent an important class of naturally occurring substances, which in many cases exhibit important biological properties (Kobayashi *et al.*, 1991; James *et al.*, 1991). 1,3-Dipolar cycloaddition reactions are widely used for the construction of spiro-compounds (Caramella & Grunanger, 1984). In this paper, the structure of the title compound, (I), is reported.



The compound was synthesized by the intermolecular [3 + 2]-cycloaddition of an azomethine ylide, derived from isatin and sarcosine by a decarboxylative route, and 2-(4-chlorobenzylidene)-6,7-dihydro-5*H*-thiazolo[3,2-*a*]pyrimidin-3-one. The molecular structure of (I) is shown in Fig. 1.

There are two spiro junctions in the molecule, which consists of a planar 2-oxindole ring, a pyrrolidine ring in an envelope conformation and a thiazolo[3,2-*a*]pyrimidine ring. Two molecules are connected into a centrosymmetric dimer by N3-H···N2ⁱ hydrogen bonds [symmetry code: (i) 2 - x, y, $\frac{1}{2} + z$], with an N···N distance of 2.855 (2) Å and an N-H···N angle of 166.9 (3)° (Fig. 2). The structure of 4'-(4-methoxyphenyl)-1'-methyl-4",5",6",7"-tetrahydro-1*H*-indole-3-spiro-2'-pyrrolidine-3'-spiro-2"-(thiazolo[3,2-*a*]pyrimidine)-2(3*H*),-3"(2"*H*)-dione has been reported previously (Li *et al.*, 2003).

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Figure 1

The molecular structure of (I), drawn with 30% probability displacement ellipsoids.



Figure 2

The crystal packing of (I), viewed along the b axis. Intermolecular N- $H \cdot \cdot \cdot N$ hydrogen bonds are shown as dashed lines.

Experimental

A mixture of 2-(4-chlorobenzylidene)-6,7-dihydro-5H-thiazolo[3,2a]pyrimidin-3-one (1 mmol), isatin (1 mmol) and sarcosine (1 mmol) was refluxed in methanol (60 ml) until the disappearance of the starting material, as evidenced by thin-layer chromatography. When the reaction was complete, the solvent was removed in vacuo and the residue was separated by column chromatography (silica gel, petroleum ether/ethyl acetate = 5:1), giving the title compound, (I). IR (KBr): 3351.4 (-NH), 1724.4, 1689.7 (C=O) cm⁻¹; ¹H NMR (δ , p.p.m.): 1.18 (m, 1H, -CH₂), 1.66 (m, 1H, -CH₂), 2.27 (s, 3H, N- CH_3), 3.35 (*m*, 2H, CH_2), 3.38 (*m*, 2H, CH_2), 3.69 (*m*, 1H, $-CH_2$), 4.09 (m, 1H, -CH₂), 4.59 (m, 1H, -CH), 6.82-7.53 (m, 8H, ArH), 8.50 (bs, 1H, -NH); 20 mg of (I) was dissolved in 15 ml dioxane and the solution was kept at room temperature for 15 d; natural evaporation gave colorless single crystals of (I) suitable for X-ray analysis.

> $D_r = 1.386 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation

reflections $\theta = 3.0-26.2^{\circ}$

 $\mu = 0.30 \text{ mm}^{-1}$

T = 293 (2) K

 $R_{\rm int} = 0.054$ $\theta_{\rm max} = 26.4^\circ$ $h = -27 \rightarrow 27$ $k = -15 \rightarrow 16$ $l = -18 \rightarrow 18$

Block, colorless $0.40 \times 0.22 \times 0.20 \text{ mm}$

Cell parameters from 1009

4454 independent reflections

2952 reflections with $I > 2\sigma(I)$

Crystal data

C23H21CIN4O2S $M_r = 452.95$ Monoclinic, C2/c a = 22.308 (8) Å b = 13.224(5) Å c = 15.054 (6) Å $\beta = 102.250 \ (8)^{\circ}$ $V = 4340 (3) \text{ Å}^3$ Z = 8

Data collection

Bruker SMART CCD area-detector
diffractometer
φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 1997)
$T_{\min} = 0.769, \ T_{\max} = 0.942$
17147 measured reflections

Refinement

27

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.061$	$w = 1/[\sigma^2(F_o^2) + (0.084P)^2]$
$WR(F^{-}) = 0.170$	where $P = (F_o^- + 2F_c^-)/3$
S = 1.04	$(\Delta/\sigma)_{\text{max}} < 0.001$
275 parameters	$\Delta \rho_{\rm max} = 0.57 \text{ e A}^{-3}$ $\Delta \rho_{\rm min} = -0.57 \text{ e Å}^{-3}$

H atoms were positioned geometrically, with C-H = 0.93-0.98 Å, and refined with a riding model, with $U_{iso}(H) = 1.2U_{eq}(carrier)$.

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

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