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Key indicators

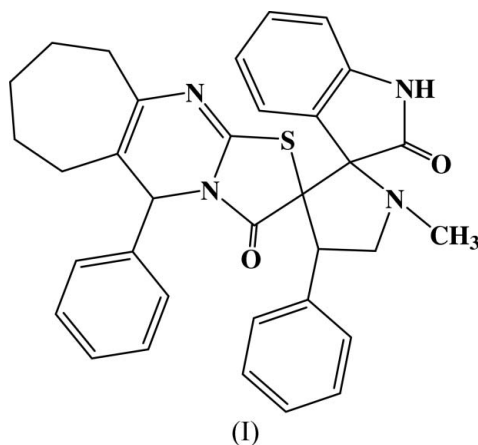
Single-crystal X-ray study
T = 293 K
Mean $\sigma(C-C)$ = 0.002 Å
R factor = 0.042
wR factor = 0.123
Data-to-parameter ratio = 17.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1'-Methyl-4',5-diphenyl-5,6,7,8,9,10-hexa-
hydro-1,3-cycloheptapyrimidino[2,3-*b*]-
thiazole-2-spiro-3'-pyrrolidine-2'-spiرو-
3''-1*H*-indole-2'',3(2*H*,3''*H*)-dioneThe pyrrolidine ring of the title compound, C₃₄H₃₂N₄O₂S,
adopts an envelope conformation. The packing is stabilized by
intermolecular C—H···O and N—H···N interactions.

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Comment

Pyrrolidine derivatives are widely studied for their pharmaco-
logical properties. The pyrrolidine motif occurs in many
families of biologically important compounds. Pyrrolidine
derivatives possess anti-influenza virus (Galeazzi *et al.*, 1999)
and anti-convulsant activities (Obniska *et al.*, 2002). They also
show inhibitory activity towards post-proline cleaving
enzymes and show strong anti-amnesic activities (Saito *et al.*,
1991). Oxindole derivatives help to treat and prevent diabetic
complications arising from elevated levels of sorbitol and act
as aldose reductase inhibitor (Rajeswaran *et al.*, 1999). Thia-
zole derivatives possess anti-inflammatory properties (Köysal
et al., 2004) and thiazole naphthyridine derivatives exhibit
good antibacterial activity (Kondo *et al.*, 1990). A series of
thiazolo[3,4-*a*]benzimidazole derivatives have been evaluated
in vitro as antitumor agents against 60 human tumor cell-lines
(Chimirri *et al.*, 1994).The two spiro junctions in the title molecule, (I), are formed
by a pyrrolidine ring, an oxindole ring and a hexahydro-1,3-
cycloheptapyrimidino[2,3-*b*]thiazol-3-one ring. The sum of the
bond angles at atom N1 of the pyrrolidine ring (337°) indicates
*sp*³ hybridization. The bond lengths and angles of the pyr-
rolidine ring are somewhat distorted, which may be due to the
spiro fusion. Selected geometric parameters are given in
Table 1. The pyrrolidine ring adopts an envelope conforma-
tion, with atom N1 deviating by −0.587 (1) Å from the plane
composed of atoms C1, C2, C3 and C4.Intermolecular C—H···O and N—H···N interactions
stabilize the crystal packing. Atom C1 acts as a donor to O2 at

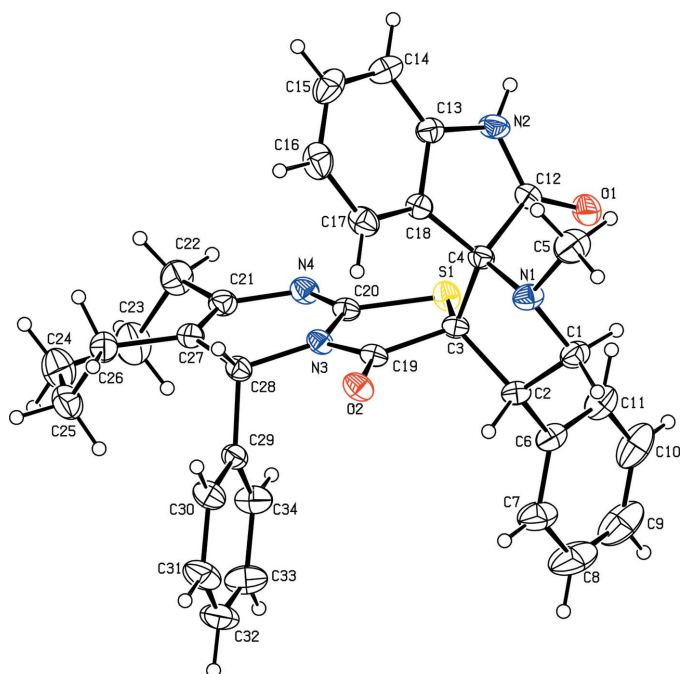


Figure 1
The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

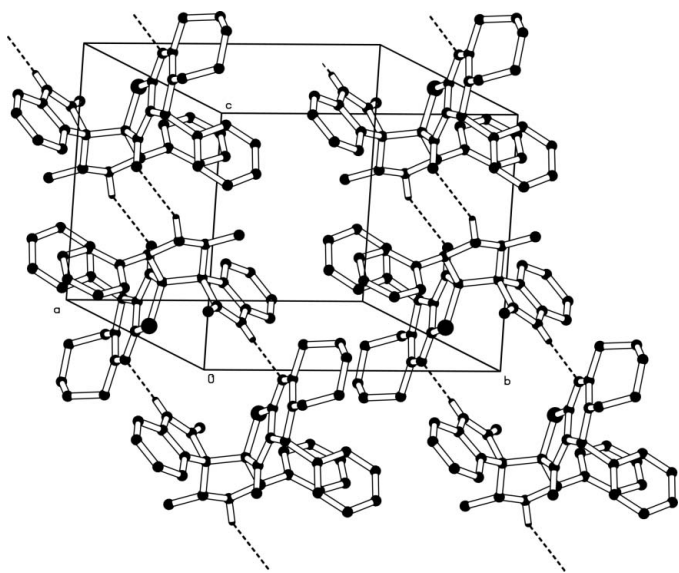


Figure 2
The molecular packing of (I). For the sake of clarity, H atoms not involved in the hydrogen bonds (dashed lines) have been omitted.

$(1-x, 2-y, 1-z)$, generating a centrosymmetric $R_2^2(12)$ ring centred at $(\frac{1}{2}, 1, \frac{1}{2})$ and amine atom N2 acts as a donor to N4 at $(-x, 2-y, -z)$, generating a centrosymmetric $R_2^2(16)$ ring centred at $(0, 1, 0)$. The propagation of these two hydrogen bonds generates a chain running along $[001]$.

Experimental

A mixture of 2-(phenylmethylene)-5-phenyl-5,6,7,8,9,10-hexahydro-1,3-cycloheptapyrimidino[2,3-*b*]thiazol-3-one (1 mmol), isatin (1 mmol) and sarcosine (1 mmol) was refluxed in 20 ml of methanol-

dioxane solvent (1:1) for 5 h. After completion of the reaction, as evidenced by thin-layer chromatography, the residue was chromatographed with a hexane-ethyl acetate mixture (4:1) to get obtain the product. Diffraction quality crystals were obtained from a methanol solution.

Crystal data

$C_{34}H_{32}N_4O_2S$
 $M_r = 560.70$
Triclinic, $P\bar{1}$
 $a = 10.483$ (1) Å
 $b = 12.133$ (1) Å
 $c = 12.870$ (1) Å
 $\alpha = 96.24$ (1)°
 $\beta = 108.27$ (1)°
 $\gamma = 107.14$ (1)°
 $V = 1448.5$ (3) Å³

$Z = 2$
 $D_x = 1.286$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 5673 reflections
 $\theta = 1.7$ – 25.0 °
 $\mu = 0.15$ mm⁻¹
 $T = 293$ (2) K
Block, colourless
 $0.24 \times 0.21 \times 0.20$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω scans
Absorption correction: none
16755 measured reflections
6629 independent reflections

5676 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.018$
 $\theta_{max} = 28.0$ °
 $h = -13 \rightarrow 13$
 $k = -16 \rightarrow 15$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.123$
 $S = 1.02$
6629 reflections
370 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0697P)^2 + 0.4648P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.33$ e Å⁻³
 $\Delta\rho_{min} = -0.16$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

S1–C20	1.743 (1)	N3–C20	1.377 (2)
S1–C3	1.831 (1)	N3–C28	1.477 (2)
N1–C5	1.455 (2)	N4–C20	1.277 (2)
N1–C1	1.455 (2)	N4–C21	1.429 (2)
N1–C4	1.460 (2)	C2–C3	1.582 (2)
N2–C12	1.351 (2)	C3–C4	1.581 (2)
N2–C13	1.401 (2)	C4–C12	1.570 (2)
N3–C19	1.373 (2)		
C20–S1–C3	93.4 (1)	C19–N3–C28	121.1 (1)
C1–N1–C4	107.2 (1)	C20–N3–C28	121.6 (1)
C12–N2–C13	112.2 (1)	C20–N4–C21	116.5 (1)
C19–N3–C20	117.1 (1)		

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2–H2 ⁱ ⋯N4 ⁱ	0.86	2.11	2.964 (2)	175
C1–H1A ⁱ ⋯O2 ⁱⁱ	0.97	2.55	3.415 (2)	148

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

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with $N-H = 0.86$ Å and $C-H$ distances in the range 0.93 – 0.97 Å and with $U_{iso}(H) = 1.2 U_{eq}(C,N)$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINTE* (Bruker, 2001); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine

structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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