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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.046 wR factor = 0.130 Data-to-parameter ratio = 19.6

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(2*R*,3*aR*,4*S*,6*aR*,7*aR*)-*cis*-4-Benzyl-2-(4-bromophenyl)-5-(4-methylphenylsulfonyl)perhydrothiazolidino[3,4-*a*]pyrrolo[4,5-c]pyrrole

In the title compound, $C_{28}H_{29}BrN_2O_2S_2$, the thiazolidine and the two pyrrolidine rings adopt envelope conformations. The crystal packing is stabilized by weak $C-H\cdots\pi$ interactions.

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Comment

Pyrrolidine derivatives are extensively studied for their important medicinal properties. Pyrrolidine occurs widely in nature and is a structural component of porphyrin heme, chlorophyll and vitamin B12. Pyrrolidine compounds have antifungal and antimicrobial activities (Amal Raj *et al.*, 2003). Thiazolidine derivatives possess antidiabetic and adipogenic properties (Norisada *et al.*, 2004).



In the title compound, (I), the S2–C6 bond length is long compared with S2–C5, while the N2–C6 bond is short compared with N2–C4 and N2–C7 (Table 1). These long and short bonds may be a result of the bromophenyl substituent at C6. The sums of the bond angles around N1 (353.8°) and N2 (334.9°) indicate sp^2 - and sp^3 -hybridization, respectively. The C23–C28 phenyl ring makes dihedral angles of 21.1 (2) and 79.6 (2)°, respectively, with the C9–C14 and C15–C20 benzene rings. Atom Br1 deviates by 0.094 (1) Å from the mean plane of the benzene ring C9–C14.

The two fused pyrrolidine rings (N1/C1/C2/C7/C8, A and C2-C4/N2/C7, B) adopt envelope conformations, with flap atom C2 deviating by 0.506 (2) and 0.552 (3) Å, respectively, from the N1/C1/C7/C8 and C3/C4/N2/C7 planes. The thiazolidine ring also adopts an envelope conformation, with atom S2 deviating by 0.888 (1) Å from the other atoms in the ring, whereas in a similar structure, 3-(4-bromophenyl)-6-(*p*-tos-yl)perhydrothiazolidino[3,4-*a*]pyrrolo[4,5-*c*]pyrrole (Kavitha *et al.*, 2006), the thiazolidine ring adopts a twist conformation.



Figure 1

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



Figure 2

The molecular packing of (I), viewed approximately down the *a* axis. For clarity, H atoms not involved in the $C-H\cdots\pi$ interactions (dashed lines) have been omitted.

The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) are $q_2 = 0.326$ (3) Å, $\varphi = 257.4$ (4)° and $\Delta_s(C_2) = 3.4$ (2)° for ring A, $q_2 = 0.358$ (3) Å, $\varphi = 105.9$ (4)° and $\Delta_s(C_2) = 1.5$ (3)° for ring B, and $q_2 = 0.522$ (2) Å, $\varphi = 357.6$ (3)° and $\Delta_s(S_2) =$ $3.8(2)^{\circ}$ for the thiazolidine ring.

The crystal packing is stabilized by van der Waals and weak $C-H\cdots\pi$ intermolecular interactions. Atom C27 acts as a donor to the C9–C14 benzene ring at $(x, \frac{1}{2} - y, -\frac{1}{2} + z)$

(centroid Cg), with $H \cdots Cg$ and $C \cdots Cg$ distances of 2.85 and 3.716 (4) Å and a C-H···Cg angle of 156° (Fig. 2).

Experimental

of (S)-2-(N-allyl-N-tosylamino)-3-phenylpropanal А mixture (1.0 mmol) and 2-(p-bromophenyl)thiazolidine-4-carboxylic acid (1.5 mmol) in toluene (30 ml) was refluxed under Dean-Stark conditions till the completion of the reaction. The reaction mixture was then concentrated in vacuo and extracted with dichloromethane $(2 \times 20 \text{ ml})$ and water $(2 \times 20 \text{ ml})$. The organic layer was washed with brine $(2 \times 20 \text{ ml})$, dried with anhydrous sodium sulfate and concentrated in vacuo. The residue was then subjected to column chromatography (silica gel, 100-200 mesh), eluting with a hexaneethyl acetate (8:2) mixture, to give the title compound. The title compound was crystallized from a hexane-ethyl acetate (4:1) solution by slow evaporation.

Crystal data

$C_{28}H_{29}BrN_2O_2S_2$	V = 2632.1 (2) Å ³
$M_r = 569.56$	Z = 4
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 14.7207 (8) Å	$\mu = 1.75 \text{ mm}^{-1}$
b = 9.1692 (5) Å	T = 293 (2) K
c = 20.6819 (11) Å	$0.25 \times 0.24 \times 0.22$ mm
$\beta = 109.461 \ (1)^{\circ}$	

Data collection

Bruker SMART APEX CCD area-	6207 independent reflections
detector diffractometer	4428 reflections with $I > 2\sigma(I)$
Absorption correction: none 29550 measured reflections	$R_{\rm int} = 0.033$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$	317 parameters
$wR(F^2) = 0.130$	H-atom parameters constrained
S = 0.97	$\Delta \rho_{\rm max} = 0.80 \ {\rm e} \ {\rm \AA}^{-3}$
6207 reflections	$\Delta \rho_{\rm min} = -0.53 \text{ e} \text{ \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

.47 (2)
78 (3)
4.8 (2)
0.7(2)
9.4 (2)
;

H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H = 0.93-0.98 Å and $U_{iso}(H)$ = $1.5U_{eq}$ (methyl C) or $1.2U_{eq}$ (C).

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; 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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