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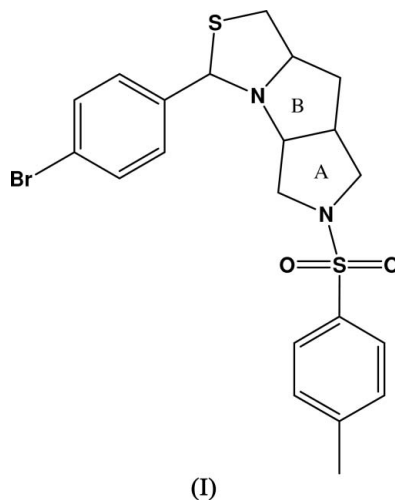
## Key indicators

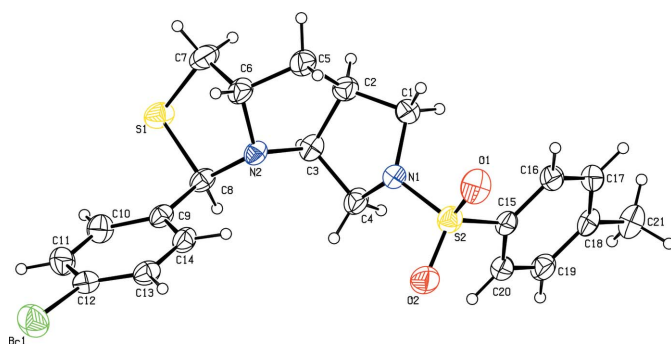
Single-crystal X-ray study  
*T* = 293 K  
Mean  $\sigma(\text{C}-\text{C})$  = 0.004 Å  
*R* factor = 0.043  
*wR* factor = 0.126  
Data-to-parameter ratio = 19.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.3-(4-Bromophenyl)-6-(*p*-tosyl)perhydrothiazolidino[3,4-*a*]pyrrolo[4,5-*c*]pyrroleIn the title compound, C<sub>21</sub>H<sub>23</sub>BrN<sub>2</sub>O<sub>2</sub>S<sub>2</sub>, the two pyrrolidine rings adopt envelope conformations and the thiazolidine ring adopts a twisted conformation with a pseudo-twofold axis passing through the S atom and the C–N ring-fusion bond. The crystal packing is stabilized by a C–H··· $\pi$  interaction between inversion-related molecules.

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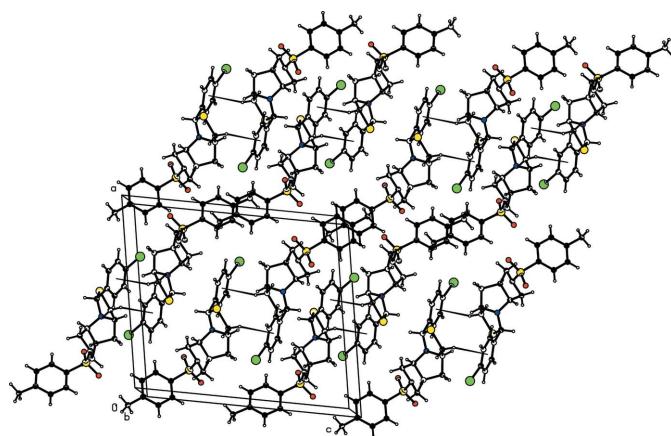
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## Comment

Pyrrole occurs widely in nature and is a structural part of porphyrin heme, chlorophyll and vitamin B12. It has been shown that *N*-substituted pyrrole derivatives inhibit human immunodeficiency virus type-I (HIV-I) (Jiang *et al.*, 2004). Thiazolidine and its derivatives have diverse biological importance, such as local anesthetic, antiseizure, anti-tubercular, antibacterial, anti-amebic, antidiabetic, anti-inflammatory (Chaurasia, 1971; Zimenkovsky *et al.*, 1999), anti-oxidant (Rachinskii, 1964) and flavor-enhancing properties, and the presence of N–C–S in the ring enhances the antifungal activity (Eswaramoorthy *et al.*, 1991).In the title compound, (I), the S1–C8 bond length is long compared with S1–C7, while the N2–C8 bond is short compared with N2–C6 and N2–C3 (Table 1). These long and short bonds may be due to the bromophenyl substitution at C8. The sums of the bond angles around N1 (343.4°) and N2 (333.3°) indicate *sp*<sup>3</sup>-hybridization. Atom Br1 deviates by 0.185 (1) Å from the mean plane of the benzene ring C9–C14. The two fused pyrrolidine rings (N1/C1–C4, *A*, and C2/C3/N2/C5/C6, *B*) adopt envelope conformations; atom N1 in ring *A* lies 0.609 (2) Å above the C1–C4 mean plane and atom C6 in ring *B* lies 0.562 (3) Å below the N2/C3/C2/C5 mean plane.



**Figure 1**  
The molecular structure of (I), showing 30% probability displacement ellipsoids.



**Figure 2**  
The molecular packing of (I), viewed down the *b* axis, showing C—H... $\pi$  interactions as dashed lines.

The thiazolidine ring adopts a twisted conformation with a pseudo-twofold axis passing through atom S1 and the C6—N2 bond. The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) are  $q_2 = 0.409$  (2) Å,  $\varphi = -179.5$  (4)° and  $\Delta_s(\text{N}_1) = 0.003$  (1) for ring A,  $q_2 = 0.366$  (3) Å,  $\varphi = -75$  (1)° and  $\Delta_s(\text{C}_6) = 0.016$  (3) for ring B, and  $q_2 = 0.368$  (3) Å,  $\varphi = 19.9$  (4)° and  $\Delta_s(\text{S}_1) = 0.006$  (1) for the thiazolidine ring.

In the crystal packing, the molecules are linked into centrosymmetric dimers by a pair of C—H... $\pi$  interactions (Fig. 2 and Table 2). Atom C7 at (*x*, *y*, *z*) acts as a donor, through atom H7B, to the benzene ring C9—C14 at (1 - *x*, 2 - *y*, 1 - *z*) with the H7B...Cg1 distance of 2.73 Å (Cg1 is the centroid of the C9—C14 ring).

## Experimental

A solution of *N*-allyl-*N*-(2-oxo-ethyl)-4-methylbenzene-sulfonamide (1 mmol) and 2-(*p*-bromophenyl)thiazolidine-4-carboxylic acid (1.2 mmol) in toluene (20 ml) was refluxed under Dean–Stark conditions for 2 h. After completion of the reaction the mixture was chromatographed over silica gel using a hexane and ethyl acetate (9:1) mixture to yield the title compound. Single crystals were grown by slow evaporation of an ethyl acetate solution.

## Crystal data

$\text{C}_{21}\text{H}_{23}\text{BrN}_2\text{O}_2\text{S}_2$   
 $M_r = 479.44$   
Monoclinic,  $P2_1/c$   
 $a = 15.3215$  (8) Å  
 $b = 8.0676$  (4) Å  
 $c = 17.1663$  (9) Å  
 $\beta = 100.119$  (1)°  
 $V = 2088.88$  (19) Å<sup>3</sup>

$Z = 4$   
 $D_x = 1.525$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
 $\mu = 2.19$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
Block, colorless  
0.23 × 0.22 × 0.20 mm

## Data collection

Bruker SMART CCD area-detector diffractometer  
 $\omega$  scans  
Absorption correction: none  
23302 measured reflections

4943 independent reflections  
3520 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.030$   
 $\theta_{\text{max}} = 28.0^\circ$

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.043$   
 $wR(F^2) = 0.126$   
 $S = 0.96$   
4943 reflections  
254 parameters  
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0658P)^2 + 1.1265P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.89$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.53$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

C1—N1	1.480 (3)	C8—S1	1.869 (2)
C3—N2	1.469 (3)	C12—Br1	1.893 (3)
C4—N1	1.473 (3)	C15—S2	1.763 (3)
C6—N2	1.463 (3)	N1—S2	1.627 (2)
C7—S1	1.795 (3)	O1—S2	1.423 (2)
C8—N2	1.441 (3)	O2—S2	1.428 (2)
C4—N1—C1	105.6 (2)	O1—S2—O2	120.3 (1)
C4—N1—S2	118.9 (2)	O1—S2—N1	106.7 (1)
C1—N1—S2	118.9 (2)	O2—S2—N1	106.6 (1)
C8—N2—C6	109.9 (2)	O1—S2—C15	108.0 (1)
C8—N2—C3	115.8 (2)	O2—S2—C15	108.4 (1)
C6—N2—C3	107.6 (2)	N1—S2—C15	106.0 (1)
C7—S1—C8	92.6 (1)		

**Table 2**

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C7—H7B...Cg1 <sup>i</sup>	0.97	2.73	3.632 (4)	156

Symmetry code: (i)  $-x + 1, -y + 2, -z + 1$ .

All H atoms were positioned geometrically and allowed to ride on their parent C atoms, with C—H distances in the range 0.93–0.98 Å and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

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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