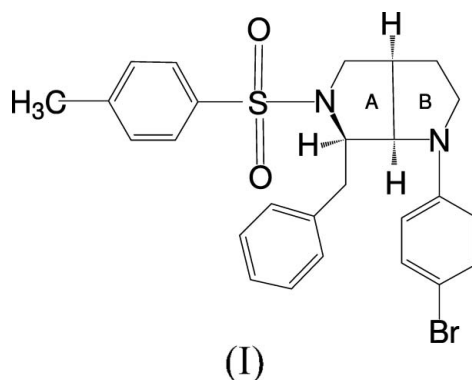


(3*aR*,6*S*,6*aR*)-6-Benzyl-1-(4-bromophenyl)-5-tosyl-*cis*-perhydropyrrolo[3,4-*b*]pyrroleK. N. Vennila,^a D. Gayathri,^a D. Velmurugan,^{a*} K. Ravikumar^b and M. Poornachandran^c^aDepartment of Crystallography and Biophysics, University of Madras, Guindy Campus, Chennai 600 025, India, ^bLaboratory of X-ray Crystallography, Indian Institute of Chemical Technology, Hyderabad 500 007, India, and ^cDepartment of Organic Chemistry, University of Madras, Guindy Campus, Chennai 600 025, India

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Key indicatorsSingle-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.006 \text{ \AA}$
R factor = 0.048
wR factor = 0.153
Data-to-parameter ratio = 19.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.The title compound, C₂₆H₂₇BrN₂O₂S, has two fused pyrrolidine rings, each in a twist conformation.Received 22 March 2007
Accepted 2 April 2007**Comment**In recent years, polyhydroxylated piperidine, pyrrolidine, indolizidine and pyrrolizidine alkaloids have attracted much attention, owing to their ability to act as selective glycosidase inhibitors. Such inhibition may be useful for the treatment of diabetes, cancer, malaria and some viral infections including AIDS (Kilonda *et al.*, 1995). Pyrrolidine compounds are capable of exhibiting antimicrobial and antifungal activity (Amal Raj *et al.*, 2003).The title compound, (I), has two fused pyrrolidine rings, each in a twist conformation. The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) for the pyrrolidine ring A (N1/C1–C4) are $q_2 = 0.286(4) \text{ \AA}$, $\varphi = -85.4(8)^\circ$, $\Delta C_2(\text{N1}) = 3.4(5)$, and for the pyrrolidine ring B (N2/C3/C2/C5/C6) are $q_2 = 0.345(4) \text{ \AA}$, $\varphi = 84.0(6)^\circ$, $\Delta C_2(\text{N2}) = 5.2(6)$. The geometric parameters are in the normal ranges (Allen *et al.*, 1987). The sums of the bond angles around atoms N1 and N2 of the pyrrolidine rings (349.1 and 356.1°, respectively) indicate that N1 and N2 have almost trigonal planar coordination. The dihedral angle between the bromophenyl and phenyl rings is 79.1(1)°.**Experimental**A mixture of (*S*)-2-(*N*-allyl-*N*-tosylamino)-3-phenylpropanal (1 mmol) and *N*-(*p*-bromophenyl)glycine (1 mmol) in 20 ml of toluene was refluxed under Dean–Stark conditions, until the disappearance of the starting materials, as evidenced by thin-layer chromatography. The solvent was then evaporated in vacuum and the residue was column chromatographed using a hexane–ethyl acetate

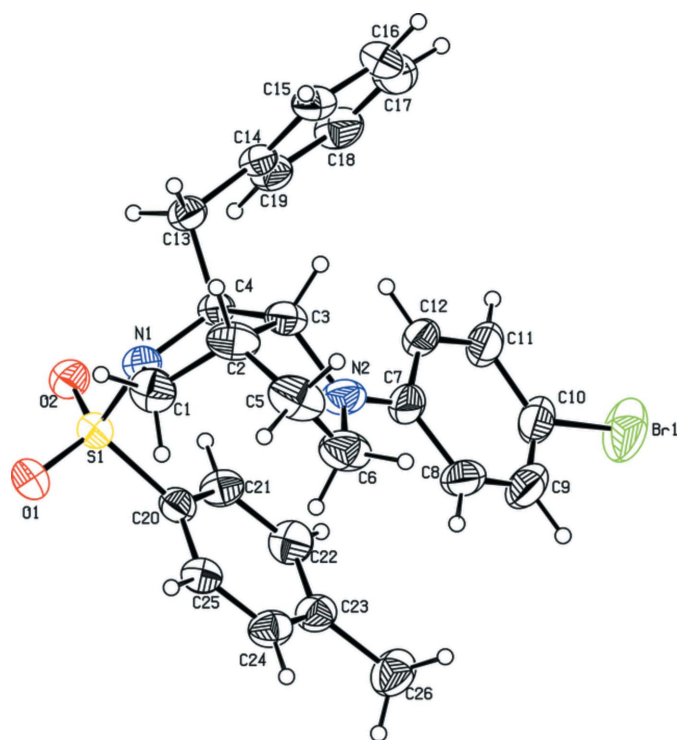


Figure 1
The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

mixture (8:2) to obtain the title compound. The title compound was crystallized from an ethyl acetate solution by slow evaporation.

Crystal data

$C_{26}H_{27}BrN_2O_2S$	$V = 2440.6 (2) \text{ \AA}^3$
$M_r = 511.47$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 20.9376 (11) \text{ \AA}$	$\mu = 1.80 \text{ mm}^{-1}$
$b = 10.5821 (6) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 11.0154 (6) \text{ \AA}$	$0.24 \times 0.23 \times 0.21 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer	5499 independent reflections
Absorption correction: none	3825 reflections with $I > 2\sigma(I)$
20524 measured reflections	$R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$	$\Delta\rho_{\text{max}} = 0.80 \text{ e \AA}^{-3}$
$wR(F^2) = 0.153$	$\Delta\rho_{\text{min}} = -0.67 \text{ e \AA}^{-3}$
$S = 1.07$	Absolute structure: Flack (1983),
5499 reflections	2297 Friedel pairs
289 parameters	Flack parameter: 0.031 (13)
H-atom parameters constrained	

Table 1

Selected bond angles ($^\circ$).

C7–N2–C6	122.0 (4)	C4–N1–C1	111.9 (3)
C7–N2–C3	123.7 (3)	C4–N1–S1	118.7 (2)
C6–N2–C3	111.3 (4)	C1–N1–S1	119.5 (2)

The H atoms were positioned geometrically and were treated as riding on their parent C atoms, with aromatic C–H distances of 0.93 Å, methyl C–H distances of 0.96 Å and methylene C–H distances of 0.97 Å, and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H and $1.2U_{\text{eq}}(\text{C})$ for other H.

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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