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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.057 wR factor = 0.150 Data-to-parameter ratio = 18.7

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

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# 6-Ethyl-1-(4-fluorophenyl)-5-(phenylsulfonyl)perhydro-cis-pyrrolo[3,4-b]pyrrole

In the title compound,  $C_{20}H_{23}FN_2O_2S$ , one of the pyrrolidine rings adopts an envelope conformation, while the other is in a twist conformation. The molecules are linked into C(6) chains by  $C-H\cdots O$  hydrogen bonds.

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## Comment

The pyrrolidine ring occurs in many families of biologically important compounds. Many pyrrolidine derivatives with varying properties have been synthesized as a result of the ease of substitution and modifications at several positions (Baldwin *et al.*, 1994). Pyrrolidine derivatives have been found to exhibit antimicrobial, antifungal (Amal Raj *et al.*, 2003), anti-inflammatory (Fernandes *et al.*, 2004) and antiviral (Borthwick *et al.*, 2003) activities. We report here the crystal structure of the title compound, (I).



The bond lengths in (I) (Fig. 1) show normal values (Allen *et al.*, 1987). The sums of the bond angles (Table 1) around atoms N1 (351.8°) and N2 (357.8°) indicate that they are  $sp^2$ -hybridized. Atom F1 deviates from the plane of the C15–C20 benzene ring by 0.026 (3) Å.

The N1/C7–C10 pyrrolidine ring adopts an envelope conformation, with atom C9 deviating by 0.452 (3) Å from the mean plane of other ring atoms. The other pyrrolidine ring (C8/C9/N2/C14/C13) adopts a twist conformation with a pseudo-twofold axis passing through atom N2 and the C8–C13 bond. The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) are  $q_2 = 0.284$  (3) Å and  $\varphi = 295.5$  (6)° and  $\Delta_s$ (C9) = 5.3 (3)° for the N1/C7–C10 pyrrolidine ring, and  $q_2 = 0.325$  (3) Å,  $\varphi = 91.1$  (5)° and  $\Delta_2$ (N2) = 1.3 (3)° for the other pyrrolidine ring. The dihedral angle between the C1–C6 and C15–C20 rings is 11.0 (2)°.



#### 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 hydrogen bonds (dashed lines) have been omitted.

The C3-H3···O2(2 - x,  $-\frac{1}{2} + y, \frac{1}{2} - z$ ) hydrogen bond (Table 2) links the molecules into a C(6) chain running along the *b* axis.

## **Experimental**

A solution of 2-(N-allyl-N-benzenesulfonylamino)butanal (1 mmol) and p-fluorophenylglycine (1.2 mmol) in dry toluene (20 ml) was refluxed until completion of the reaction as evidenced by thin-layer

chromatography. The solvent was evaporated under vacuum and the residue was column chromatographed (silica gel, 100–200 mesh) using a hexane–ethyl acetate (9:1) mixture, yielding the title compound which was recrystallized by slow evaporation of an ethyl acetate solution.

V = 1869.47 (17) Å<sup>3</sup>

 $0.23 \times 0.22 \times 0.21$  mm

4395 independent reflections 3627 reflections with  $I > 2\sigma(I)$ 

Absolute structure: Flack (1983);

Mo  $K\alpha$  radiation  $\mu = 0.20 \text{ mm}^{-1}$ 

T = 293 (2) K

 $R_{\rm int} = 0.024$ 

 $\Delta \rho_{\text{max}} = 0.38 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\text{min}} = -0.22 \text{ e } \text{\AA}^{-3}$ 

1847 Friedel pairs

Flack parameter: 0.41 (9)

Z = 4

Crystal data

 $C_{20}H_{23}FN_2O_2S$   $M_r = 374.46$ Orthorhombic,  $P2_12_12_1$  a = 7.8671 (4) Å b = 13.6413 (7) Å c = 17.4200 (9) Å

## Data collection

Bruker SMART area-detector diffractometer Absorption correction: none 16247 measured reflections

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.057$   $wR(F^2) = 0.150$  S = 1.05 4395 reflections 235 parametersH-atom parameters constrained

### Table 1

Selected geometric parameters (Å, °).

C1-S1	1.770 (3)	C15-N2	1.379 (4)
C7-N1	1.481 (3)	C18-F1	1.383 (4)
C9-N2	1.450 (4)	N1-S1	1.617 (2)
C10-N1	1.481 (3)	O1-S1	1.434 (2)
C14-N2	1.469 (3)	O2-S1	1.422 (2)
C7-N1-C10	110.9 (2)	O2-S1-O1	120.2 (2)
C7-N1-S1	117.8 (2)	O2-S1-N1	106.9 (1)
C10-N1-S1	123.1 (2)	O1-S1-N1	106.4 (1)
C15-N2-C9	123.2 (2)	O2-S1-C1	107.4 (1)
C15-N2-C14	122.1 (2)	O1-S1-C1	106.6 (1)
C9-N2-C14	112.5 (2)	N1-S1-C1	108.9 (1)

Table 2Hydrogen-bond geometry (Å,  $^{\circ}$ ).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$		
$C3-H3\cdots O2^i$	0.93	2.59	3.504 (4)	168		
Symmetry code: (i) $-x + 2$ , $y - \frac{1}{2}$ , $-z + \frac{1}{2}$ .						

H atoms were positioned geometrically and allowed to ride on their parent atoms, with C–H = 0.93–0.98 Å and  $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm methyl}\ {\rm C})$  or  $1.2U_{\rm eq}({\rm C})$ . The value of the Flack parameter indicates inversion twinning.

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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