organic papers

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Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.004 Å R factor = 0.045 wR factor = 0.107 Data-to-parameter ratio = 16.3

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

[4-(2,3-Dichlorophenyl)piperazin-1-yl][2-(2-methoxyphenylsulfanyl)phenyl]methanone

In the title compound, $C_{24}H_{22}Cl_2N_2O_2S$, synthesized from 2-(2-methoxyphenylsulfanyl)benzoyl chloride and 1-(2,3dichlorophenyl)piperazine, the piperazine ring adopts a normal chair conformation. Received 3 January 2006 Accepted 8 February 2006

Comment

Substituted diphenyl sulfide derivatives which display high *in vitro* and *in vivo* affinities for serotonin transporter (SERT), high selectivity for the dopamine transporter (DAT) and partial selectivity for norepinephrine transporter sites (NET) have been described as potent and selective SERT ligands (Mehta & Brieaddy, 1997; Wilson & Houle, 1999; Younes *et al.*, 2000).



The structure of (I) is illustrated in Fig. 1. The piperazine ring has a normal chair conformation. The dihedral angle relating the two rings bonded to the S atom is $102.28 (12)^{\circ}$. The dichlorophenyl ring is also twisted with respect to the piperazine ring [C19-N2-C17-C18 = 164.8 (2)°].



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The molecular structure of (I), drawn with 30% probability ellipsoids.

Experimental

2-(2-Methoxyphenylsulfanyl)benzoyl chloride (8 mmol), triethylamine (20 mmol) and 1-(2,3-dichlorophenyl)piperazine (8 mmol) in CHCl₃ (60 ml) were stirred at room temperature for 3 h. The mixture was then washed with 2 M sodium hydroxide. The organic layer was dried and evaporated in vacuo to dryness, giving a yellow oil, which solidified at room temperature. After filtration through active charcoal and recrystallization from 60% aqueous ethanol (70 ml), the title compound was obtained as white crystals. Crystals suitable for X-ray analysis were grown by slow evaporation of an absolute methanol solution at room temperature over a period of 15 d.

Crystal data

CarHanClaNaOaS	$D = 1.412 \text{ Mg m}^{-3}$
$M_r = 473.40$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2087
a = 17.034 (3) Å	reflections
b = 7.6898 (14) Å	$\theta = 2.4 - 21.9^{\circ}$
c = 17.229 (3) Å	$\mu = 0.41 \text{ mm}^{-1}$
$\beta = 99.344 \ (3)^{\circ}$	T = 294 (2) K
V = 2226.9 (7) Å ³	Block, colourless
Z = 4	$0.24 \times 0.22 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer φ and φ scans Absorption correction: multi-scan (SADABS; Bruker, 1997) $T_{\min} = 0.900, T_{\max} = 0.921$ 12169 measured reflections

4587 independent reflections 2529 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.054$ $\theta_{\rm max} = 26.5^{\circ}$ $h = -21 \rightarrow 19$ $k = -7 \rightarrow 9$

 $l = -21 \rightarrow 18$

Refinement Refinement on F^2

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.041P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	+ 0.1467P]
$wR(F^2) = 0.107$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.99	$(\Delta/\sigma)_{\rm max} = 0.003$
4587 reflections	$\Delta \rho_{\rm max} = 0.22 \ {\rm e} \ {\rm \AA}^{-3}$
282 parameters	$\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

All H atoms were positioned geometrically and refined as riding, with C-H = 0.93–0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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