

2-(2,5-Dimethylbenzyl)-3-phenylsulfanyl-1-phenylsulfonyl-1H-indole

G. Usha,^a S. Selvanayagam,^a
D. Velmurugan,^{a*}
K. Ravikumar,^b N. Sureshbabu^c
and P. C. Srinivasan^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

Correspondence e-mail: d_velu@yahoo.com

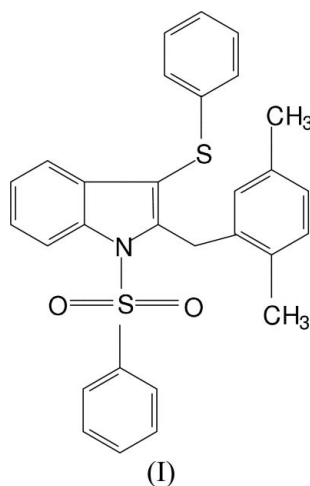
Key indicators

Single-crystal X-ray study
 $T = 273$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.049
 wR factor = 0.143
Data-to-parameter ratio = 16.9For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

In the title compound, $\text{C}_{29}\text{H}_{25}\text{NO}_2\text{S}_2$, the benzene rings of the phenylsulfanyl and the dimethylbenzyl substituents are almost perpendicular to the indole unit, whereas the dihedral angle between the phenyl rings of the phenylsulfanyl and phenylsulfonyl groups is $71.2(1)^\circ$. The molecules in the crystal structure are held together by van der Waals, $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{S}$ interactions.

Comment

Indoles and their derivatives have been of interest for many years, since a large number of natural products contain indole systems, and they are found in a number of pharmaceutical products, fragrances and dyes (Padwa *et al.*, 1999). Spiroindole derivatives exhibit antibacterial and antifungal properties (Sehgal *et al.*, 1994). The sulfonamide-containing drugs inhibit the growth of bacterial organisms and are also used for treating urinary and gastrointestinal infections. The wide range of biological activities of indole and its derivatives prompted us to undertake the crystal structure analysis of the title compound, (I).



The $\text{S}-\text{O}$, $\text{S}-\text{C}$ and $\text{S}-\text{N}$ bond distances are in good agreement with the related reported values of 1.435 (5), 1.767 (7) and 1.685 (5) Å, respectively (Govindasamy *et al.*, 1998). The electron-withdrawing character of the phenylsulfonyl group affects the $\text{C}-\text{N}$ distances in the indole ring system [$\text{C}5-\text{N}1 = 1.417(2)$ Å and $\text{C}2-\text{N}1 = 1.416(2)$ Å] and this is observed in similar reported structures (Rodriguez *et al.*, 1995; Govindasamy *et al.*, 1999). The sum of the angles around atom N1 (358.4°) indicates sp^2 hybridization. In the dimethylphenyl ring, the endocyclic angles at C19 and C22 are $117.4(2)$ and $117.6(2)^\circ$, respectively, and this decrease can be

Received 3 June 2005
Accepted 14 June 2005
Online 24 June 2005

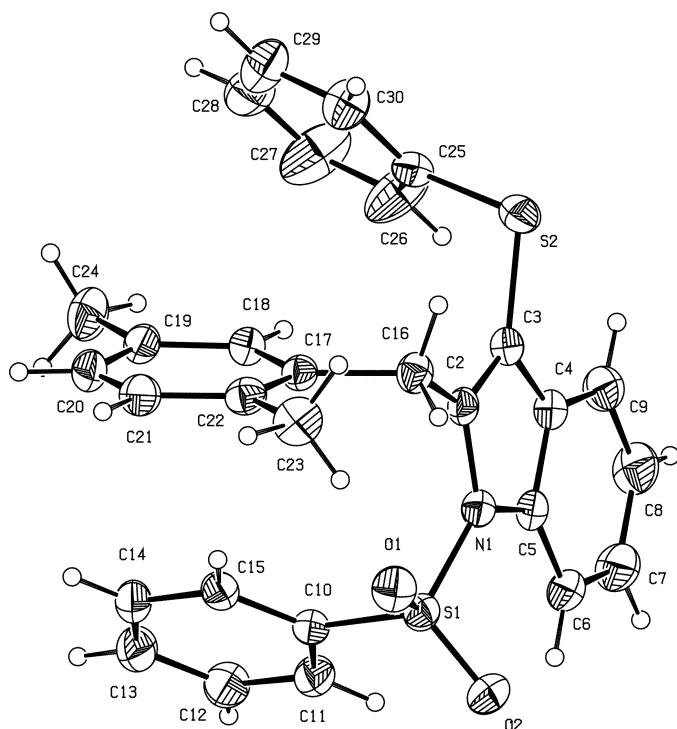


Figure 1
The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

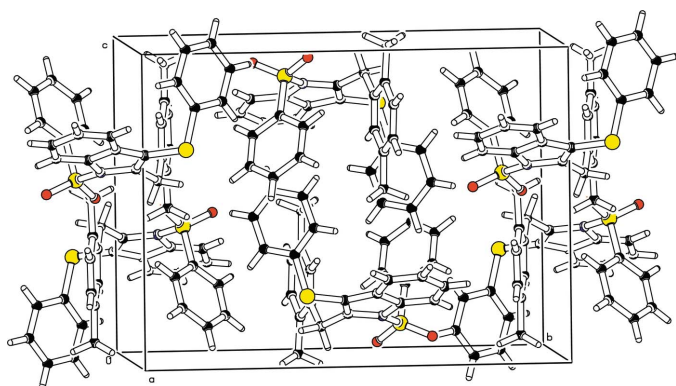


Figure 2
The molecular packing of (I), viewed approximately down the *c* axis.

attributed to the steric hindrance caused by the methyl groups. The bond angle C2–C16–C17 [115.5 (2)°] is widened.

The dihedral angles between the indole ring system and the mean planes of the phenylsulfonyl, dimethylphenyl and the phenylsulfanyl rings are 85.1 (1), 82.6 (1) and 71.2 (1)°, respectively, and show that the substituent rings are almost perpendicular to the indole system.

The packing of the molecules in the unit cell is governed by van der Waals forces and the crystal structure is stabilized by C–H···O and C–H···S interactions.

Experimental

The title compound was prepared by the reaction of 2-hydroxy-methyl-3-phenylsulfanyl-1-phenylsulfonyl-1*H*-indole with *p*-xylene in

the presence of a catalytic amount of boron trifluoride etherate in boiling chloroform, following a published procedure (Rajeswaran & Srinivasan, 1992). The crude product was purified by silica-gel column chromatography, eluting with hexane–ethyl acetate (9:1). Diffraction quality crystals were obtained from a hexane/ethyl acetate (1:1) solution.

Crystal data

C₂₉H₂₅NO₂S₂
M_r = 483.62
 Monoclinic, *P*2₁/*n*
a = 10.8372 (6) Å
b = 17.2384 (10) Å
c = 12.7220 (7) Å
 β = 90.269 (1)°
V = 2376.6 (2) Å³
Z = 4

D_x = 1.352 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 5656 reflections
 θ = 2.2–27.5°
 μ = 0.25 mm⁻¹
T = 273 (2) K
 Block, colourless
 0.23 × 0.22 × 0.20 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω scans
 Absorption correction: none
 14279 measured reflections
 5221 independent reflections

4227 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.020
 θ_{\max} = 28.0°
h = -14 → 14
k = -22 → 21
l = -14 → 16

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.049
wR(*F*²) = 0.143
S = 1.01
 5221 reflections
 309 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0882P)^2 + 0.5156P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.50 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

S1–O1	1.414 (2)	N1–C5	1.417 (2)
S1–O2	1.422 (2)	N1–C2	1.416 (2)
S1–N1	1.682 (2)	C2–C16	1.496 (3)
S1–C10	1.760 (2)	C16–C17	1.522 (2)
S2–C3	1.759 (2)	C19–C24	1.499 (3)
S2–C25	1.780 (2)	C22–C23	1.501 (3)
O1–S1–O2	120.6 (1)	C3–S2–C25	103.0 (1)
O1–S1–N1	106.7 (1)	C5–N1–C2	108.5 (2)
O2–S1–N1	105.5 (1)	C5–N1–S1	122.1 (1)
O1–S1–C10	109.2 (1)	C2–N1–S1	127.8 (1)
O2–S1–C10	108.7 (1)	C2–C16–C17	115.5 (2)
N1–S1–C10	105.0 (1)		
O2–S1–C10–C15	149.7 (2)		

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>
C6–H6···O2	0.93	2.31	2.900 (3)	121
C15–H15···O1	0.93	2.55	2.914 (3)	104
C16–H16 <i>A</i> ···S2	0.97	2.83	3.294 (2)	110
C16–H16 <i>B</i> ···O1	0.97	2.38	2.881 (3)	112

The H atoms were positioned geometrically and were treated as riding on their parent C atoms, with aromatic C–H = 0.93 Å, methyl

C–H = 0.96 Å and methylene C–H = 0.97 Å, and with N–H = 0.86 Å, and with $U_{\text{iso}} = 1.5U_{\text{eq}}(\text{C})$ for methyl H and $1.2U_{\text{eq}}(\text{N,C})$ for the remaining H atoms.

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: *ORTEP3* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

GU thanks the University Grants Commission (UGC) for the award of the Faculty Improvement Programme (FIP). SS thanks the Council of Scientific and Industrial Research (CSIR) for providing a Senior Research Fellowship. DV acknowledges the UGC and the Department of Bio-Technology (DBT) for providing computing facilities under Major Research Projects and also acknowledges financial support to

the Department under UGC–SAP and DST–FIST programmes.

References

- Bruker (2001). *SAINTE* (Version 6.28a) and *SMART* (Version 5.625). Bruker AXS Inc., Madison, Wisconsin, USA.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Govindasamy, L., Velmurugan, D., Ravikumar, K. & Mohanakrishnan, A. K. (1998). *Acta Cryst.* **C54**, 635–637.
- Govindasamy, L., Velmurugan, D., Shanmuga Sundara Raj, S. & Fun, H. K. (1999). *Acta Cryst.* **C55**, 1315–1317.
- Nardelli, M. (1995). *J. Appl. Cryst.* **28**, 659.
- Padwa, A., Brodney, M. A., Liu, B., Stake, K. & Wu, T. (1999). *J. Org. Chem.* **64**, 3595–3607.
- Rajeswaran, W. G. & Srinivasan, P. C. (1992). *Ind. J. Heterocyclic Chem.* **2**, 89–90.
- Rodriguez, J. G., del Valle, C., Calderon, C. E. & Ripoll, M. M. (1995). *J. Chem. Crystallogr.* **25**, 249–257.
- Sehgal, V., Singh, P., Dandia, A. & Bohra, R. (1994). *Acta Cryst.* **C50**, 1156–1159.
- Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
- Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.