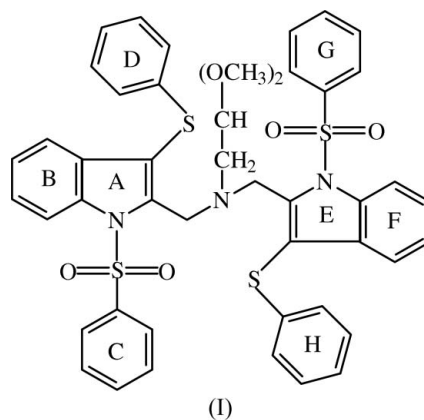


{*N,N*-Bis[3-(phenylsulfanyl)-1-(phenylsulfonyl)-indol-2-ylmethyl]amino}acetaldehyde dimethyl acetal**M. Yogavel,^a D. Velmurugan,^{a*}
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Key indicatorsSingle-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C})$ = 0.007 Å
Disorder in main residue
R factor = 0.074
wR factor = 0.207
Data-to-parameter ratio = 19.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.The structure of the title compound, $\text{C}_{46}\text{H}_{41}\text{N}_3\text{O}_6\text{S}_4$, is stabilized by $\text{C}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{S}$ and $\text{C}-\text{H}\cdots\text{N}$ interactions, and the packing of the molecules is stabilized by several $\text{C}-\text{H}\cdots\pi$ interactions, and face-to-face and edge-to-face $\pi-\pi$ interactions.**Comment**Indole is present in a large number of natural products, many of which have a wide range of biological activities, such as antibacterial (Okabe & Adachi, 1998), antitumour (Schollmeyer *et al.*, 1995), antidepressant (Grinev *et al.*, 1984), antimicrobial (El-Sayed *et al.*, 1986; Gadaginamath & Patil, 1999) and anti-inflammatory (Rodriguez *et al.*, 1985). Indoles have been proved to display high aldose reductase inhibitory activity (Rajeswaran *et al.*, 1999). The interaction of phenylsulfonyl indole with the calf-thymus DNA has also been studied by spectroscopic methods (Sivaraman *et al.*, 1996). The structure determination of the title compound, (I), was undertaken as part of our studies on indole derivatives.

The dihedral angle between the indole groups (*AB* and *EF*) and the phenyl rings (*C*, *D*, *G* and *H*) are $AB/C = 75.0 (1)^\circ$, $EF/G = 84.0 (1)^\circ$, $AB/D = 84.9 (1)^\circ$ and $EF/H = 66.6 (1)^\circ$. The dihedral angle between rings *C/D* and *G/H* are $69.5 (1)$ and $77.8 (1)^\circ$, respectively. Atom N25 deviates by $0.217 (3)$ Å from the plane through atoms C26, C29 and S3, and the sum of the angles around N25 is 353.8° . This slight pyramidalization behaviour is also observed in related indole derivatives (Sankaranarayan *et al.*, 2000, 2001; Sankaranarayan, Yogavel, Velmurugan, Sekar, Babu *et al.*, 2003; Sankaranarayan, Yogavel, Velmurugan, Sekar, Srinivasan *et al.*, 2003; Sethu-Sankar *et al.*, 2002; Ganesh *et al.*, 2003; Krishna *et al.*, 2003).

The molecular structure is stabilized by $\text{C}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{S}$ and $\text{C}-\text{H}\cdots\text{N}$ interactions. The phenyl rings *C* of the

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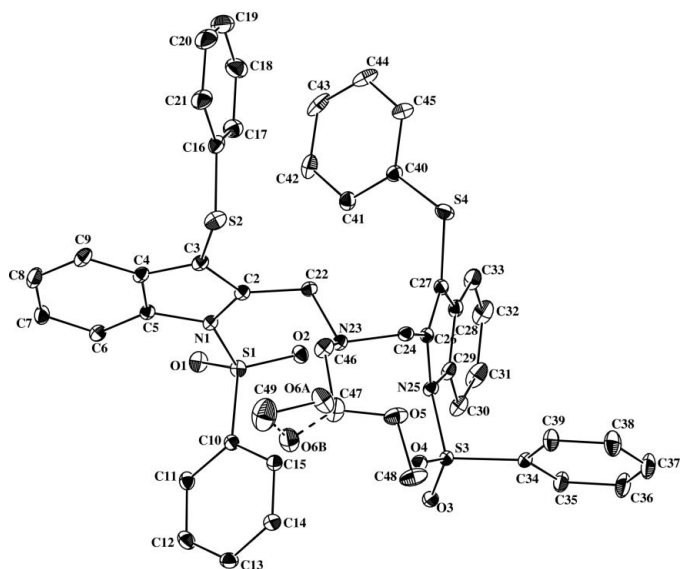


Figure 1
The molecular structure of the title compound, showing 35% probability displacement ellipsoids. H atoms have been omitted. Dashed lines indicate the minor component of the disordered O atom.

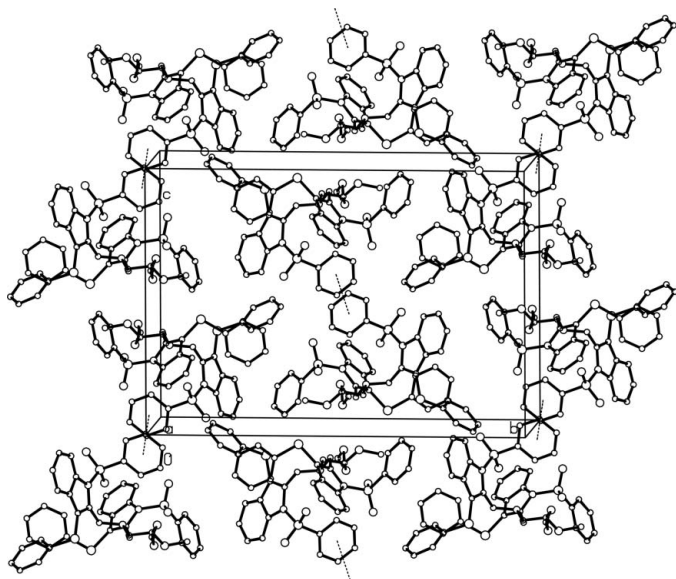


Figure 2
The crystal structure of (I), viewed down the *a* axis. The π - π interactions are indicated as dashed lines. H atoms have been omitted.

molecules at (x, y, z) and $(-x, -y, -z)$ are stacked in a face-to-face manner, with their centroids separated by 3.772 (2) Å. Several intermolecular edge-to-face interactions and several C—H $\cdots\pi$ interactions are also observed (Table 2). Thus, the crystal packing is stabilized by C—H $\cdots\pi$, face-to-face π - π and edge-to-face π - π interactions.

Experimental

To a solution of aminoacetaldehyde dimethyl acetal (4.4 ml, 40 mmol) in chloroform (40 ml) was added a solution of 1-phenylsulfonyl-2-bromomethyl-3-phenylthioindole (10 mmol) in chloroform (15 ml) and the mixture was stirred at room temperature for 4 h.

Work-up of the reaction and crystallization of the residue from methanol afforded the title compound (yield 4.1 g, 85%) as the sole product.

Crystal data

$C_{46}H_{41}N_3O_6S_4$
 $M_r = 857.03$
 Monoclinic, $P2_1/n$
 $a = 10.7146$ (1) Å
 $b = 23.8939$ (1) Å
 $c = 16.8782$ (3) Å
 $\beta = 91.846$ (1)°
 $V = 4318.81$ (9) Å³
 $Z = 4$

$D_x = 1.318$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 8042 reflections
 $\theta = 1.5$ – 28.5°
 $\mu = 0.27$ mm⁻¹
 $T = 293$ (2) K
 Block, colourless
 $0.40 \times 0.36 \times 0.28$ mm

Data collection

Siemens SMART CCD area-detector diffractometer
 ω scans
 Absorption correction: none
 29148 measured reflections
 10689 independent reflections

5223 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.066$
 $\theta_{max} = 28.5^\circ$
 $h = -14 \rightarrow 12$
 $k = -31 \rightarrow 30$
 $l = -15 \rightarrow 22$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.074$
 $wR(F^2) = 0.207$
 $S = 1.01$
 10689 reflections
 543 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0814P)^2 + 1.9503P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.56$ e Å⁻³
 $\Delta\rho_{min} = -0.45$ e Å⁻³

Table 1

Selected interatomic distances (Å).

S1—O2	1.406 (3)	S3—O4	1.426 (3)
S1—O1	1.427 (3)	S3—O3	1.432 (3)
N1—C5	1.415 (4)	N25—C26	1.425 (4)
N1—C2	1.421 (4)	N25—C29	1.435 (4)

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C6—H6 \cdots O1	0.93	2.37	2.923 (5)	118
C11—H11 \cdots O1	0.93	2.57	2.924 (5)	103
C15—H15 \cdots N23	0.93	2.48	3.334 (4)	153
C22—H22A \cdots O2	0.97	2.36	2.913 (4)	115
C22—H22B \cdots S2	0.97	2.85	3.311 (4)	110
C24—H24A \cdots O4	0.97	2.33	2.887 (4)	116
C24—H24A \cdots O5	0.97	2.39	3.090 (5)	129
C24—H24B \cdots S4	0.97	2.84	3.287 (4)	109
C30—H30 \cdots O3	0.93	2.33	2.920 (5)	121
C35—H35 \cdots O3	0.93	2.55	2.904 (5)	103
C32—H32 \cdots CgA ⁱ	0.93	2.95	3.728 (5)	143
C32—H32 \cdots CgB ⁱ	0.93	3.09	3.752 (5)	129
C31—H31 \cdots CgC ⁱ	0.93	2.99	3.680 (6)	132
C7—H7 \cdots CgD ⁱⁱ	0.93	3.09	3.661 (5)	122
C9—H9 \cdots CgH ⁱⁱⁱ	0.93	3.02	3.682 (5)	130

Symmetry codes: (i) $x + 1, y, z$; (ii) $x - \frac{3}{2}, -y - \frac{1}{2}, z - \frac{3}{2}$; (iii) $x - 1, y, z$. CgA, CgB, CgC, CgD and CgH are the centroids of the rings A, B, C, D and H, respectively.

All the H atoms were positioned geometrically and allowed to ride on their parent atoms [C—H = 0.93–0.98 Å, and $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(C_{methyl})$]. Atom O6 was found to be disordered and the site occupancy factors of the major and minor components were refined.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINTE* (Siemens, 1996); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ZORTEP* (Zsolnai, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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