

3-(2,4-Dichlorophenyl)-*N,N*-diethyl-3-(4-methylphenylsulfonylamino)propanamide

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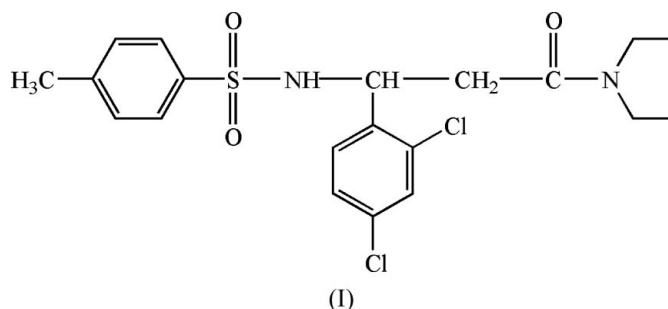
Key indicators

Single-crystal X-ray study
 $T = 294$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.053
 wR factor = 0.132
Data-to-parameter ratio = 14.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The S atom of the sulfonyl group in the title compound, $\text{C}_{20}\text{H}_{24}\text{Cl}_2\text{N}_2\text{O}_3\text{S}$, has a distorted tetrahedral geometry. Two intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds between the amino groups and the carbonyl groups link the molecules into a centrosymmetric dimer.

Comment

The addition of a nucleophile to unsaturated carbon–nitrogen bonds is a convenient method of synthesizing amine derivatives (Kobayashi & Ishitani, 1999). In our laboratory, we have recently investigated a new metal-mediated reaction for the preparation of β -sulfonylaminoamides. The title compound, (I), has been synthesized by the reaction of *N*-(2,4-dichlorobenzylidene)-4-methylbenzenesulfonamide and 2-bromo-*N,N*-diethylacetamide in the presence of zinc powder.



The molecular structure of (I) and the atom-numbering scheme are shown in Fig. 1. In this structure, the benzene ring attached to the sulfonyl group makes a dihedral angle of $11.7(3)^\circ$ with the dichlorophenyl ring. The S atom has a distorted tetrahedral geometry, with the $\text{O}2-\text{S}1-\text{O}1$ and $\text{O}2-\text{S}1-\text{N}1$ angles (Table 1) deviating most from the regular tetrahedral value. Intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds are formed between the amino groups and the carbonyl groups of two molecules, forming a centrosymmetric dimer.

Experimental

The title compound was synthesized by adding 2-bromo-*N,N*-diethylacetamide (3.0 mmol) to a solution of *N*-(2,4-dichlorobenzylidene)-4-methylbenzenesulfonamide (1.0 mmol) in dichloromethane (5.0 ml). Zinc powder (6.0 mmol) and a trace amount of iodine were added to the mixture. After the mixture had been refluxed for 6 h, the reaction was quenched with a saturated solution of ammonium chloride (8.0 ml). The mixture was extracted with dichloromethane, evaporated and separated by flash chromatography (using ethyl acetate–petroleum ether). A white powder (yield 64%) was obtained. Slow evaporation of a dichloromethane–petroleum ether (1:7) solution afforded the title compound as single crystals.

Crystal data

C₂₀H₂₄Cl₂N₂O₃S
M_r = 443.38
 Triclinic, *P* $\bar{1}$
a = 7.895 (3) Å
b = 10.704 (4) Å
c = 14.046 (4) Å
 α = 105.881 (5)°
 β = 100.025 (4)°
 γ = 99.190 (4)°

V = 1097.0 (7) Å³
Z = 2
D_x = 1.342 Mg m⁻³
 Mo *K*α radiation
 μ = 0.41 mm⁻¹
T = 294 (2) K
 Block, colourless
 0.36 × 0.22 × 0.14 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.896, *T_{max}* = 0.944

5569 measured reflections
 3833 independent reflections
 3131 reflections with *I* > 2σ(*I*)
R_{int} = 0.016
 θ_{max} = 25.0°

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.053
wR(*F*²) = 0.132
S = 1.06
 3833 reflections
 258 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0549P)^2 + 0.8034P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.60 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{min} = -0.71 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

S1—O2	1.426 (2)	S1—N1	1.606 (2)
S1—O1	1.427 (2)	S1—C8	1.766 (3)
O2—S1—O1	120.48 (15)	O2—S1—C8	107.45 (13)
O2—S1—N1	105.55 (14)	O1—S1—C8	107.80 (14)
O1—S1—N1	107.42 (13)	N1—S1—C8	107.55 (12)

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>
N1—H1...O3 ⁱ	0.79 (3)	2.04 (3)	2.806 (3)	166 (3)

Symmetry code: (i) $-x + 1, -y + 1, -z + 1$.

All carbon-bound H atoms were positioned geometrically and refined as riding (C—H = 0.93–0.98 Å), with *U_{iso}*(H) set at 1.2*U_{eq}*(C) or 1.5*U_{eq}*(methyl C). The coordinates of atom H1 were refined. Methyl groups C14 and C18 were allowed to rotate but not to tip.

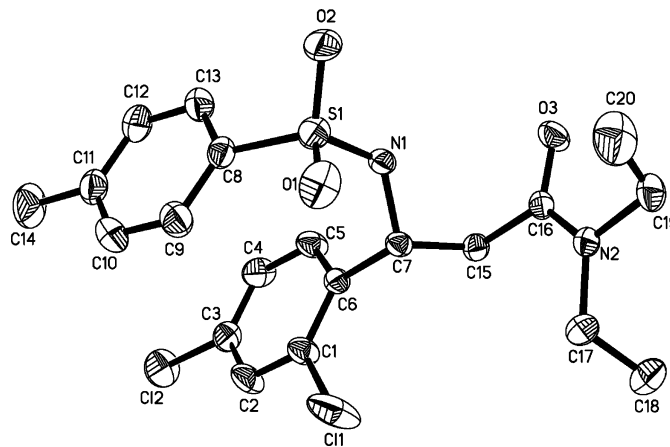


Figure 1

View of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms have been omitted for clarity.

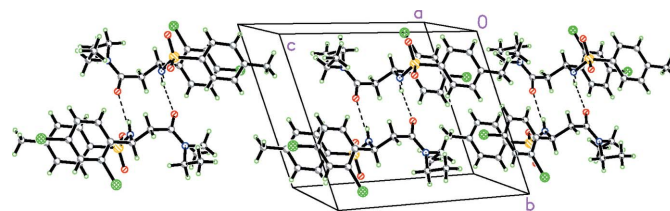


Figure 2

View of the molecular structure of (I). Dashed lines indicate N—H...O interactions.

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*.

References

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