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## Key indicators

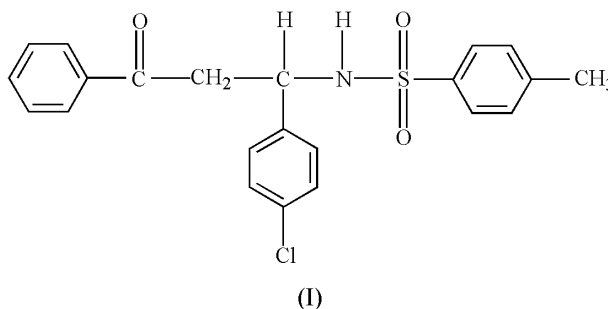
Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å  
 $R$  factor = 0.049  
 $wR$  factor = 0.148  
Data-to-parameter ratio = 16.5For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.3-(4-Chlorophenyl)-1-phenyl-3-(*p*-tolylsulfonyl-  
amino)propan-1-one

The title compound,  $\text{C}_{22}\text{H}_{20}\text{ClNO}_3\text{S}$ , was synthesized by the Reformatsky reaction of *N*-[(4-chlorophenyl)methylene]-4-methylbenzenesulfonamide and 2-bromoacetophenone. The S atom in the sulfonyl group has a distorted tetrahedral geometry. Two intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds are formed between the amine and sulfonyl groups of two molecules related by a center of symmetry.

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## Comment

The addition of nucleophilic reagents to unsaturated carbon–nitrogen bonds is a common practice in the synthesis of amine derivatives (Kobayashi & Ishitani, 1999). In our laboratory, we have recently investigated a new Reformatsky addition reaction to *N*-sulfonylimines with zinc enolates of ketones to give the corresponding  $\beta$ -sulfonaminoketones. A new compound, namely the title compound, (I), has been synthesized by the reaction of *N*-[(4-chlorophenyl)methylene]-4-methylbenzenesulfonamide (Shim & Yamamoto, 2000) with 2-bromoacetophenone in the presence of active zinc powder. An X-ray crystal structure determination of (I) was carried out in order to elucidate the structure, and the results are presented here.



The molecular structure of (I) is illustrated in Fig. 1. The S atom has a distorted tetrahedral geometry, with the  $\text{O}3-\text{S}1-\text{O}2$  [ $119.2(1)^\circ$ ] and  $\text{O}2-\text{S}1-\text{N}1$  [ $105.0(1)^\circ$ ] angles deviating significantly from the regular tetrahedral value. In this structure, the ring attached to the sulfonyl group and the 4-chlorophenyl ring make dihedral angles of  $94.5(3)^\circ$  and  $103.2(3)^\circ$ , respectively, with the carbonylphenyl ring. Two molecules related by a center of symmetry are linked by two intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds (Table 2), resulting in the formation of a dimer.

## Experimental

To a solution of *N*-[(4-chlorophenyl)methylene]-4-methylbenzenesulfonamide (1.0 mmol) in dichloromethane (5.0 ml) was added

2-bromoacetophenone (1.5 mmol). Zinc powder (3.0 mmol) and a trace amount of iodine were added to the mixture. After the reaction mixture was refluxed with stirring for 11 h and quenched with a saturated aqueous solution of  $\text{NH}_4\text{Cl}$  (5.0 ml) and 25% ammonia (5.0 ml), the mixture was extracted with dichloromethane. The extract was washed with water and dried over magnesium sulfate. After the solvent was evaporated, a white powder was obtained (yield 86%) by flash chromatography (ethyl acetate–trichloromethane). Slow evaporation of an ethyl acetate–petroleum ether mixture afforded the title compound as a crystalline solid (m.p. 401–403 K). IR (KBr,  $\text{cm}^{-1}$ ):  $\nu$  3261, 1690, 1318, 1154;  $^1\text{H}$  NMR ( $\text{CDCl}_3$ , p.p.m.):  $\delta$  7.81–7.07 (*m*, 13H), 5.73 (*br*, 1H), 4.81 (*m*, 1H), 3.52 (*dd*, 1H), 3.40 (*dd*, 1H), 2.36 (*s*, 3H). Analysis calculated for  $\text{C}_{22}\text{H}_{20}\text{ClNO}_3\text{S}$ : C 63.78, H 4.83, N 3.38%; found: C 63.84, H 4.87, N 3.35%.

#### Crystal data

$\text{C}_{22}\text{H}_{20}\text{ClNO}_3\text{S}$   
 $M_r = 413.91$   
 Monoclinic,  $P2_1/c$   
 $a = 15.975$  (5) Å  
 $b = 9.654$  (3) Å  
 $c = 14.566$  (4) Å  
 $\beta = 115.752$  (4) $^\circ$   
 $V = 2023.3$  (11) Å $^3$   
 $Z = 4$

$D_x = 1.359$  Mg m $^{-3}$   
 Mo  $K\alpha$  radiation  
 Cell parameters from 923 reflections  
 $\theta = 3.1$ – $25.0$  $^\circ$   
 $\mu = 0.32$  mm $^{-1}$   
 $T = 293$  (2) K  
 Prism, colorless  
 $0.40 \times 0.34 \times 0.18$  mm

#### Data collection

Bruker SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.882$ ,  $T_{\max} = 0.945$   
 11 492 measured reflections

4185 independent reflections  
 2521 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.042$   
 $\theta_{\max} = 26.5$  $^\circ$   
 $h = -17 \rightarrow 20$   
 $k = -12 \rightarrow 6$   
 $l = -18 \rightarrow 17$

#### Refinement

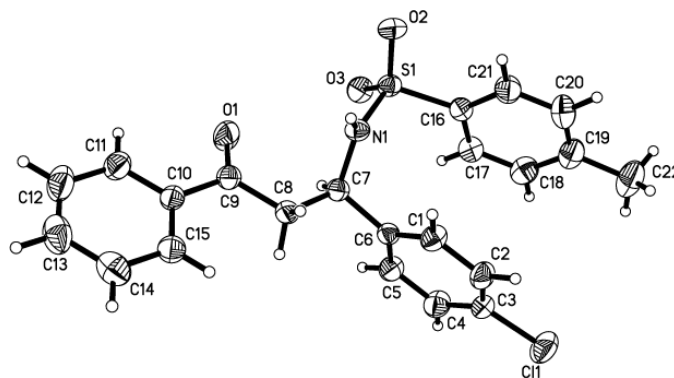
Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.148$   
 $S = 1.06$   
 4185 reflections  
 253 parameters

H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.077P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.29$  e Å $^{-3}$   
 $\Delta\rho_{\min} = -0.47$  e Å $^{-3}$

**Table 1**

Selected geometric parameters (Å,  $^\circ$ ).

S1–O3	1.426 (2)	S1–C16	1.758 (3)
S1–O2	1.440 (2)	O1–C9	1.212 (3)
S1–N1	1.608 (2)	N1–C7	1.475 (3)
O3–S1–O2	119.2 (1)	O2–S1–C16	107.3 (1)
O3–S1–N1	107.9 (1)	N1–S1–C16	108.6 (1)
O2–S1–N1	105.0 (1)	C7–N1–S1	123.0 (2)
O3–S1–C16	108.5 (1)	S1–N1–C7–C6	78.1 (2)
O3–S1–N1–C7	42.4 (2)	N1–C7–C8–C9	67.8 (3)
O2–S1–N1–C7	170.5 (2)	C7–C8–C9–O1	−0.3 (4)
C16–S1–N1–C7	−74.9 (2)	C7–C8–C9–C10	178.4 (2)
S1–N1–C7–C8	−157.12 (17)		



**Figure 1**

View of the molecule of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 35% probability level.

**Table 2**

Hydrogen-bonding geometry (Å,  $^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O2}^i$	0.86	2.18	2.951 (3)	149

Symmetry code: (i)  $-x, 1-y, 2-z$ .

All H atoms were positioned geometrically and refined as riding ( $\text{N}-\text{H} = 0.86$  Å and  $\text{C}-\text{H} = 0.93$ – $0.98$  Å). For NH, CH and  $\text{CH}_2$  groups,  $U_{\text{iso}}(\text{H})$  values were set at  $1.2U_{\text{eq}}$  (carrier atom), and for the methyl groups they were set at  $1.5U_{\text{eq}}$  (carrier atom).

Data collection: SMART (Bruker, 1997); cell refinement: SMART; data reduction: SAINT (Bruker, 1997); 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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