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Key indicators

Single-crystal X-ray study $T=293~{\rm K}$ Mean $\sigma({\rm C-C})=0.003~{\rm \AA}$ R factor = 0.035 wR factor = 0.088 Data-to-parameter ratio = 16.6

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

(4-Chlorobenzoyl)methyl morpholine-4-carbodithionate

The title compound, $C_{13}H_{14}CINO_2S_2$, the morpholine crystallizes in space group $P2_1/c$. The morpholine ring adopts the usual chair conformation and the four C atoms deviate only slightly from coplanarity. There are some weak inter- and intramolecular hydrogen-bond interactions in the crystal structure, providing stabilization. Received 18 November 2004 Accepted 9 December 2004 Online 18 December 2004

Comment

Morpholine is a versatile chemical. It is used as a solvent for resins, dyes and waxes. Its alkyl derivatives (e.g. N-methylmorpholine and N-ethylmorpholine) are used as catalysts for the production of polyurethane foams. The most important use is as a chemical intermediate to prepare pharmaceuticals. Drugs containing the morpholine ring have established activities that include the reduction of blood sugar and lipid levels (Yoshioka, 1995), and the amelioration of obesity and insulin resistance (Fisher & Wyvratt, 1990). Owing to their important pharmacological activities, these compounds have received a great deal of attention in respect of their syntheses and in the elucidation of their crystal structures a search of new morpholine compounds with higher pharmacological activities, the title compound, (I), was synthesized.

In the morpholine ring, the average C-N, C-C and C-Obond distances [1.472 (2), 1.506 (3) and 1.421 (2) Å, respectively are also in good agreement with earlier reports (Ramnathan et al., 1996; Yavuz et al., 2004). The C5=S2 distance is 0.02 Å shorter than the mean value of 1.681 Å found in an earlier report (Allen et al., 1987). The C-Cl bond length [1.740 (2) Å] is comparable to the value found by Zucco et al., 1999). The morpholine ring adopts the usual chair conformation and the four C atoms deviate only slightly from coplanarity, in agreement with the structural data available from Version 5.14 of the Cambridge Structural Database (Allen, 2002). Atoms S1, S2, N1, C1, C4, C5 and C6 are coplanar (p1). The dihedral angle between the C8-C13 benzene ring and p1 is 70.4 (6)°. Weak intermolecular and intramolecular hydrogen-bond interactions stabilize the structure (Table 2).

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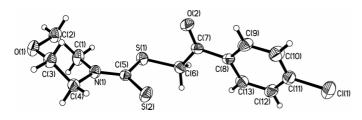


Figure 1
View of (I), with displacement ellipsoids drawn at the 35% probability level

Experimental

A mixture of 2-bromo-1-(4-chlorophenyl)ethanone (4.67 g, 0.02 mol), sodium morpholine-4-carbodithioate (3.54 g, 0.02 mol) and acetone (50 ml) was stirred for 1 h at around 273 K. The solution was then filtered, concentrated and purified by flash chromatography (silica gel, chloroform–cyclohexane 5:1 ν/ν) to afford the title compound (yield 5.05 g, 80%). Single crystals suitable for X-ray measurements was obtained by recrystallization from ethyl acetate at room temperature.

Crystal data

$C_{13}H_{14}CINO_2S_2$	$D_x = 1.469 \text{ Mg m}^{-3}$
$M_r = 315.82$	Mo $K\alpha$ radiation
Monoclinic, P2 ₁ /c	Cell parameters from 967
a = 18.092 (5) Å	reflections
b = 9.471 (2) Å	$\theta = 3.1 - 25.7^{\circ}$
c = 8.464 (2) Å	$\mu = 0.56 \text{ mm}^{-1}$
$\beta = 100.127 (8)^{\circ}$	T = 293 (2) K
$V = 1427.7 (6) \text{ Å}^3$	Block, colorless
Z = 4	$0.48 \times 0.40 \times 0.32 \text{ mm}$

Data collection

 Bruker SMART CCD area-detector diffractometer
 2876

 φ and ω scans
 $R_{\rm int}$

 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $h = T_{\rm min} = 0.761, T_{\rm max} = 0.837$
 $k = T_{\rm min} = 0.761, T_{\rm max} = 0.837$ $k = T_{\rm min} = 0.761, T_{\rm max} = 0.837$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.035$ $wR(F^2) = 0.088$ S = 1.062876 reflections 173 parameters H-atom parameters constrained 2876 independent reflections 2096 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.030$ $\theta_{\rm max} = 26.3^{\circ}$ $h = -22 \rightarrow 20$ $k = -11 \rightarrow 11$ $l = -10 \rightarrow 6$

$$\begin{split} w &= 1/[\sigma^2(F_{\rm o}^2) + (0.0352P)^2 \\ &+ 0.3711P] \\ \text{where } P &= (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} &= 0.002 \\ \Delta\rho_{\rm max} &= 0.21 \text{ e Å}^{-3} \\ \Delta\rho_{\rm min} &= -0.21 \text{ e Å}^{-3} \\ \text{Extinction correction: } SHELXL97 \\ \text{Extinction coefficient: } 0.0260 \ (16) \end{split}$$

Table 1 Selected geometric parameters (Å, °).

S1-C5	1.780 (2)	N1-C4	1.469 (3)
S1-C6	1.787 (2)	N1-C1	1.475 (2)
S2-C5	1.657 (2)	O1-C2	1.416 (3)
Cl1-C11	1.740 (2)	O1-C3	1.426 (2)
N1-C5	1.341 (2)	C3-C4	1.502 (3)
C5-S1-C6	101.47 (10)	C2-O1-C3	109.12 (15)
C4-N1-C1	112.36 (16)	N1-C5-S1	113.54 (14)

Table 2 Hydrogen-bond geometry (Å, °).

$D-H\cdot\cdot\cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$C1-H1B\cdots S1$	0.97	2.39	2.932 (3)	115
$C4-H4A\cdots S2$	0.97	2.57	3.068 (3)	112
$C4-H4B\cdots O2^{i}$	0.97	2.58	3.544 (4)	172
$C6-H6A\cdots S2$	0.97	2.64	3.032 (2)	104

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

All H atoms were placed in calculated positions, with C—H = 0.93 or 0.97 Å, and included in the final cycles of refinement using a riding model, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); 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, 1999); software used to prepare material for publication: *SHELXTL*.

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References

Allen, F. H. (2002). Acta Cryst. B58, 380-388.

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans.* 2, pp. 1–19.

Bruker (1998). SMART. Bruker AXS Inc., Madison, Wisconsin, USA.

Bruker (1999). SAINT and SHELXTL. Bruker AXS Inc., Madison, Wisconsin, USA.

Fisher, M. H. & Wyvratt., M. J. (1990). US Patent Application 3 7292 85.

Rampathan A. Siyakumar K. Sriniyasan N. Janarthanan N. Ramadas

Ramnathan, A., Sivakumar, K., Srinivasan, N., Janarthanan, N., Ramadas, K. & Fun, H.-K. (1996). *Acta Cryst.* C52, 1285–1288.

Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Köysal, Y., Işik, , Septioglu, E. & Çaliş, Ü. (2004). *Acta Cryst.* C**60**, o757–o758. Yoshioka, T. (1995). Japanese Patent 7002824.

Zucco, C., Neves, A., Vencato, I., Szpoganicz, B. & Bertoldi, F. C. (1999). Acta Cryst. C55, 654–656.