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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.027 wR factor = 0.096 Data-to-parameter ratio = 14.5

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

# 2-Oxo-2-phenylethyl morpholine-4-carbodithioate

The title compound,  $C_{13}H_{15}NO_2S_2$ , the morpholine ring adopts the usual chair conformation and the four C atoms deviate only slightly from coplanarity. The crystal structure is stabilized by some weak intramolecular  $C-H\cdots S$  hydrogenbond interactions.

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#### Comment

Morpholine is a versatile chemical which is used as a solvent for resins, dyes and waxes. One of its most important uses is as a chemical intermediate in the preparation of pesticides (Li *et al.*, 1998). Due to their important biological activities, morpholines have attracted a great deal of attention in terms of their synthesis and crystal structures. In a search for new morpholine compounds with higher biological activities, the title compound, (I), was synthesized and we report here its crystal structure.



In the morpholine ring, the average C–N, C–C and C–O bond distances [1.467 (3), 1.498 (3) and 1.418 (3) Å, respectively] are in good agreement with the values in (4-fluoro-benzoyl)(1H-1,2,4-triazol-1-yl)methylmorpholine-4-carbodi-

thioate (Xu *et al.*, 2004), *N*-phenyl-4-morpholinecarbothioamide and *N*-(2-tolyl)-4-morpholinecarbothioamide (Ramnathan *et al.*, 1996). The C1=S1 bond length is 1.653 (3) Å, in good agreement with the mean value of 1.660 Å reported by Allen *et al.* (1987). The morpholine ring adopts the usual chair conformation and its four C atoms deviate only slightly from coplanarity, in agreement with structural data available from the Cambridge Structural Database (Version 5.14; Allen, 2002). Atoms S1, S2, N1, C1, C2, C5 and C6 are coplanar (plane *p*1). The dihedral angle between the benzene ring and *p*1 is 70.4 (6)°. The crystal structure is stabilized by some weak intramolecular C-H···S hydrogen-bond interactions (Table 2).

#### **Experimental**

A mixture of 2-bromo-1-phenylethanone (3.98 g, 0.02 mol), sodium morpholine-4-carbodithioate (3.54 g, 0.02 mol) and acetone (50 ml)

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**Figure 1** View of the title compound, (I), with displacement ellipsoids drawn at the 40% probability level.

was stirred for 1.5 h at about 273 K. The solution was then filtered, concentrated and purified by flash chromatography (silica gel, chloroform/cyclohexane 5:1) to afford the title compound (yield 4.78 g, 85%). Single crystals of the title compound suitable for X-ray measurements were obtained by recrystallization from ethyl acetate at room temperature.

#### Crystal data

$C_{13}H_{15}NO_2S_2$	Mo $K\alpha$ radiation	
$M_r = 281.38$	Cell parameters from 2566	
Orthorhombic, Pbca	reflections	
a = 9.5294 (9)  Å	$\theta = 2.4-20.1^{\circ}$	
b = 8.4805(8)  Å	$\mu = 0.39 \text{ mm}^{-1}$	
c = 33.362 (3) Å	T = 293 (2)  K	
$V = 2696.1 (4) \text{ Å}^3$	Block, colorless	
Z = 8	$0.54 \times 0.34 \times 0.18 \text{ mm}$	
$D_x = 1.386 \text{ Mg m}^{-3}$		

#### Data collection

Bruker SMART 1000 CCD areadetector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\min} = 0.801, T_{\max} = 0.933$ 13423 measured reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.027$   $wR(F^2) = 0.096$  S = 0.872367 reflections 163 parameters 2367 independent reflections 1349 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.053$   $\theta_{max} = 25.0^{\circ}$   $h = -11 \rightarrow 7$   $k = -10 \rightarrow 9$  $l = -38 \rightarrow 39$ 

H-atom parameters constrained $w = 1/[\sigma^2(F^2) + (0.0531P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.16 \text{ e} \text{ Å}^{-3}$
$\Delta \rho_{\rm min} = -0.16 \text{ e} \text{ Å}^{-3}$

## Table 1

Selected geometric parameters (Å, °).

S1-C1	1.653 (3)	O1-C3	1.423 (3)
S2-C1	1.776 (3)	O2-C7	1.208 (3)
S2-C6	1.790 (2)	C2-C3	1.493 (3)
N1-C1	1.339 (3)	C4-C5	1.503 (4)
N1-C2	1.462 (3)	C6-C7	1.513 (3)
N1-C5	1.471 (3)	C7-C8	1.490 (3)
O1-C4	1.413 (3)		
C1-S2-C6	101.24 (12)	N1-C1-S1	124.42 (19)
C2-N1-C5	112.0 (2)	N1-C1-S2	113.49 (18)
C4-O1-C3	109.21 (18)	S1-C1-S2	122.09 (15)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C2-H2B\cdots$ S1	0.97	2.57	3.065 (3)	112
$C5-H5A\cdots S2$	0.97	2.38	2.928 (6)	115
$C6-H6B\cdots S1$	0.97	2.63	3.024 (3)	104

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: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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