organic compounds

Bis(N-methyl-N-phenylcarbamoyl)-disulfane

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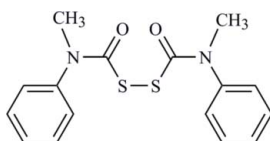Correspondence e-mail: barany@umn.edu

The title compound, $C_{16}H_{16}N_2O_2S_2$, has been synthesized by several different high-yield routes, and has been encountered as a co-product in a number of reaction pathways, ever since it became of interest to our research program over 30 years ago. We now confirm the proposed molecular structure in which the molecule exhibits a twofold axis of symmetry through the mid-point of the S—S bond and the two planes defined by the (carbamoyl)sulfenyl moieties are essentially perpendicular to each other [dihedral angle = 81.55 (14)°].

Related literature

For the preparation of the title compound, and of very closely related chemical structures, see: Kobayashi *et al.* (1973); Barany *et al.* (1983); Schroll & Barany (1986); Schrader *et al.* (2011). For related structures, see: CSD refcodes BOWGAV (Bereman *et al.*, 1983), DBZOSS01&03 (Rout *et al.*, 1983; Paul & Srikrishnan, 2004), METHUS03 (Wang & Liao, 1989), NELTUT (Fun *et al.*, 2001), JAXPOO (Raya *et al.*, 2005), UDALER (Li *et al.*, 2006) and EMASIV (Singh *et al.*, 2011). For the theoretical optimum torsion angle about the disulfane, see: Pauling (1949); Torrico-Vallejos *et al.* (2010) and references cited therein.

Experimental

Crystal data

$C_{16}H_{16}N_2O_2S_2$	$V = 1654.9$ (5) Å^3
$M_r = 332.43$	$Z = 4$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 15.286$ (3) Å	$\mu = 0.33$ mm^{-1}
$b = 9.7849$ (18) Å	$T = 296$ K
$c = 11.597$ (2) Å	$0.40 \times 0.16 \times 0.13$ mm
$\beta = 107.433$ (3)°	

Data collection

Bruker SMART CCD diffractometer	5726 measured reflections
Absorption correction: multi-scan (*SADABS*; Bruker, 2010)	1468 independent reflections
$T_{min} = 0.880$, $T_{max} = 0.958$	1140 reflections with $I > 2\sigma(I)$
	$R_{int} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$	101 parameters
$wR(F^2) = 0.110$	H-atom parameters constrained
$S = 1.05$	$\Delta\rho_{max} = 0.21$ e Å^{-3}
1468 reflections	$\Delta\rho_{min} = -0.18$ e Å^{-3}

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: QM2060).

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supporting information

Bis(*N*-methyl-*N*-phenylcarbamoyl)disulfane

Alayne L. Schroll, Maren Pink and George Barany

S1. Comment

Bis(*N*-methyl-*N*-phenylcarbamoyl)disulfane ($C_{16}H_{16}N_2O_2S_2$) was first reported by Kobayashi *et al.* (1973). The compound became of interest to our research program over thirty years ago (Barany *et al.*, 1983; Schroll and Barany, 1986) and has been synthesized by several different high-yield routes, as well as encountered as a co-product in a number of reaction pathways (Barany *et al.*, 1983; Schroll and Barany, 1986; Schrader *et al.*, 2011). We now confirm the molecular structure of the title compound by single-crystal X-ray analysis. The disulfane reported herein is the flagship of the homologous series of bis(*N*-methyl-*N*-phenylcarbamoyl)polysulfanes, $C_{16}H_{16}N_2O_2S_n$, which have been prepared and structurally characterized for *n* = 1–6.

The title compound exhibits a twofold axis of symmetry through the center of the S–S bond, and all bond distances and angles are within expected ranges. The N–C bond distance is 1.35 Å, consistent with ~60% double bond character, with the consequence that the (carbamoyl)sulfenyl atoms (S1,C1,O1,N1,C2,C3) are in a plane. The aromatic ring is nearly perpendicular to the (carbamoyl)sulfenyl plane, with a torsion angle of 92.5° (C2–N1–C3–C4). The S–S bond length of 2.03 Å is slightly shorter than the 2.07 Å reported for the S–S bond length in elemental sulfur (S_8), suggesting that some partial double bond character extends through the S–S bond due to its adjacency to carbonyl groups on both sides. Several other reference compounds also have an S–S bond length of 2.01–2.03 Å (Bereman *et al.*, 1983; Rout *et al.*, 1983; Paul and Srikrishnan, 2004; Fun *et al.*, 2001; Raya *et al.*, 2005; Li *et al.*, 2006; Singh *et al.*, 2011). The most noteworthy feature of the title compound is the torsion angle about the disulfane, which is 81.6° and as such is somewhat smaller than the theoretical optimum of 90.0° (Pauling, 1949; Torrico-Vallejos *et al.*, 2010) that has been explained as allowing for minimal mutual repulsion of pπ orbital electron lone pairs in sulfur. A comparable deviation from theory was reported for dibenzoyl disulfide (Rout *et al.*, 1983; Paul & Srikrishnan, 2004), where the torsion angle is 80.8°. Bis(*N*-methyl-*N*-phenylthiocarbamoyl)disulfane, which only differs from the title compound by two thiocarbonyls in place of two carbonyls, has a torsion angle about the disulfane of 89.8° and shows a conformation that is not completely superimposable on the title compound (Fun *et al.*, 2001).

Note regarding nomenclature: The title compound is named in a manner that is consistent with our prior publications. The closely related $C_{16}H_{16}N_2S_4$ was named bis(*N*-methyl-*N*-phenylthiocarbamoyl) disulfide by Fun *et al.* (2001), but we have chosen the "disulfane" revised name for consistency.

Table 1 Selected geometric parameters (Å, °)

N1–C1 1.345 (3)

N1–C2 1.461 (3)

N1–C3 1.442 (2)

C1–O1 1.209 (2)

C1–S1 1.825 (2)

S1–S1 2.0262 (11)

C2–N1–C3–C4 92.5 (3)

C1–S1–S1–C1 81.55 (14)

Symmetry operator (*a*): -*x* + 1, *y*, -*z* + 1/2

S2. Experimental

The title compound was prepared in high yield from the reaction of *N*-methylaniline with bis(chlorocarbonyl)disulfane, and recrystallized from hot carbon tetrachloride/chloroform (3:2) in 60–85% recovery or from hot acetone in 75% recovery (Barany *et al.*, 1983).

S3. Refinement

H atoms were positioned geometrically and refined using a riding model with C—H = 0.93 Å (aromatic) or 0.96 Å (methyl), and $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic and $1.5U_{eq}(C)$ for methyl H atoms.
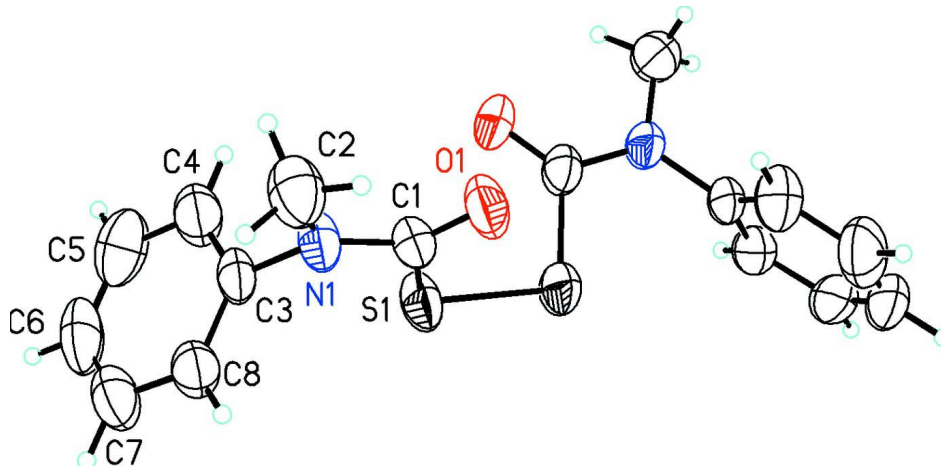
Figure 1

Crystallographic structure of the title compound showing 50% probability displacement ellipsoids and with all non-hydrogen atoms labelled and numbered.

Bis(*N*-methyl-*N*-phenylcarbamoyl)disulfane

Crystal data

$C_{16}H_{16}N_2O_2S_2$	$F(000) = 696$
$M_r = 332.43$	$D_x = 1.334$ Mg m^{-3}
Monoclinic, $C2/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
$a = 15.286$ (3) Å	Cell parameters from 1966 reflections
$b = 9.7849$ (18) Å	$\theta = 2.5–24.4°$
$c = 11.597$ (2) Å	$\mu = 0.33$ mm^{-1}
$\beta = 107.433$ (3)°	$T = 296$ K
$V = 1654.9$ (5) Å^3	Needle, colorless
$Z = 4$	$0.40 \times 0.16 \times 0.13$ mm

Data collection

Bruker SMART CCD diffractometer	5726 measured reflections
Radiation source: sealed tube	1468 independent reflections
Graphite monochromator	1140 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{int} = 0.031$
Absorption correction: multi-scan (*SADABS*; Bruker, 2010)	$\theta_{max} = 25.1°$, $\theta_{min} = 2.5°$
	$h = -18 \rightarrow 17$
	$k = 0 \rightarrow 11$
$T_{min} = 0.880$, $T_{max} = 0.958$	$l = 0 \rightarrow 13$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	
$R[F^2 > 2\sigma(F^2)] = 0.037$	Hydrogen site location: inferred from neighbouring sites
$wR(F^2) = 0.110$	
$S = 1.05$	H-atom parameters constrained
1468 reflections	$w = 1/[\sigma^2(F_o^2) + (0.0553P)^2 + 1.0043P]$
101 parameters	where $P = (F_o^2 + 2F_c^2)/3$
0 restraints	$(\Delta/\sigma)_{max} = 0.001$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{max} = 0.21$ e Å^{-3}
	$\Delta\rho_{min} = -0.18$ e Å^{-3}

Special details

Refinement. Refinement of F^2 against all reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2, conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2. The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on all data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	x	*y*	*z*	U_{iso}*/U_{eq}
S1	0.56058 (4)	0.49722 (6)	0.73116 (5)	0.0567 (2)
O1	0.47025 (9)	0.69997 (17)	0.58995 (15)	0.0626 (5)
N1	0.61329 (11)	0.65722 (19)	0.57979 (17)	0.0514 (5)
C1	0.54116 (13)	0.6361 (2)	0.62131 (19)	0.0481 (5)
C2	0.60771 (17)	0.7649 (3)	0.4905 (2)	0.0697 (7)
H2A	0.5446	0.7887	0.4530	0.105*
H2B	0.6341	0.7328	0.4301	0.105*
H2C	0.6407	0.8438	0.5298	0.105*
C3	0.69962 (13)	0.5872 (2)	0.62742 (19)	0.0466 (5)
C4	0.76230 (16)	0.6332 (3)	0.7319 (2)	0.0681 (7)
H4A	0.7486	0.7071	0.7739	0.082*
C5	0.84668 (17)	0.5672 (4)	0.7739 (3)	0.0852 (9)
H5A	0.8893	0.5962	0.8451	0.102*
C6	0.86714 (18)	0.4597 (3)	0.7107 (3)	0.0806 (9)
H6A	0.9235	0.4159	0.7393	0.097*
C7	0.80540 (18)	0.4168 (3)	0.6063 (3)	0.0716 (7)
H7A	0.8202	0.3449	0.5631	0.086*
C8	0.72070 (16)	0.4796 (2)	0.5640 (2)	0.0562 (6)
H8A	0.6782	0.4493	0.4932	0.067*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0465 (3)	0.0663 (4)	0.0681 (4)	0.0096 (3)	0.0337 (3)	0.0104 (3)
O1	0.0407 (8)	0.0691 (10)	0.0836 (12)	0.0153 (7)	0.0269 (8)	0.0072 (8)
N1	0.0395 (9)	0.0581 (11)	0.0632 (11)	0.0094 (8)	0.0255 (8)	0.0135 (9)
C1	0.0387 (11)	0.0530 (12)	0.0566 (13)	0.0036 (9)	0.0205 (9)	−0.0051 (10)
C2	0.0644 (15)	0.0712 (16)	0.0834 (18)	0.0110 (13)	0.0370 (14)	0.0239 (14)
C3	0.0349 (10)	0.0546 (12)	0.0573 (13)	0.0039 (9)	0.0244 (9)	0.0078 (10)
C4	0.0496 (13)	0.0865 (18)	0.0723 (16)	0.0010 (13)	0.0244 (12)	−0.0099 (14)
C5	0.0468 (14)	0.120 (3)	0.0796 (19)	−0.0082 (16)	0.0043 (13)	0.0141 (19)
C6	0.0438 (14)	0.092 (2)	0.114 (2)	0.0204 (14)	0.0361 (16)	0.0376 (19)
C7	0.0624 (15)	0.0610 (15)	0.105 (2)	0.0186 (13)	0.0467 (16)	0.0171 (15)
C8	0.0508 (13)	0.0567 (14)	0.0675 (14)	0.0032 (10)	0.0274 (11)	0.0033 (11)

Geometric parameters (Å, º)

S1—O1^i	3.0078 (18)	C3—C8	1.377 (3)
S1—C1	1.825 (2)	C4—C5	1.393 (4)
S1—S1^i	2.0262 (11)	C4—H4A	0.9300
O1—C1	1.209 (2)	C5—C6	1.371 (4)
N1—C1	1.345 (3)	C5—H5A	0.9300
N1—C3	1.442 (2)	C6—C7	1.359 (4)
N1—C2	1.461 (3)	C6—H6A	0.9300
C2—H2A	0.9600	C7—C8	1.384 (3)
C2—H2B	0.9600	C7—H7A	0.9300
C2—H2C	0.9600	C8—H8A	0.9300
C3—C4	1.376 (3)		
C1—S1—S1^i	100.51 (7)	C3—C4—C5	118.9 (3)
C1—N1—C3	122.98 (17)	C3—C4—H4A	120.5
C1—N1—C2	118.97 (17)	C5—C4—H4A	120.5
C3—N1—C2	117.79 (17)	C6—C5—C4	120.3 (3)
O1—C1—N1	124.8 (2)	C6—C5—H5A	119.8
O1—C1—S1	122.58 (16)	C4—C5—H5A	119.8
N1—C1—S1	112.64 (14)	C7—C6—C5	120.3 (2)
N1—C2—H2A	109.5	C7—C6—H6A	119.8
N1—C2—H2B	109.5	C5—C6—H6A	119.8
H2A—C2—H2B	109.5	C6—C7—C8	120.3 (3)
N1—C2—H2C	109.5	C6—C7—H7A	119.8
H2A—C2—H2C	109.5	C8—C7—H7A	119.8
H2B—C2—H2C	109.5	C3—C8—C7	119.6 (2)
C4—C3—C8	120.5 (2)	C3—C8—H8A	120.2
C4—C3—N1	119.9 (2)	C7—C8—H8A	120.2
C8—C3—N1	119.5 (2)		
C3—N1—C1—O1	174.2 (2)	C8—C3—C4—C5	−1.3 (4)
C2—N1—C1—O1	0.2 (3)	N1—C3—C4—C5	−177.6 (2)

C3—N1—C1—S1	−6.4 (3)	C3—C4—C5—C6	1.0 (4)
C2—N1—C1—S1	179.66 (17)	C4—C5—C6—C7	0.2 (4)
S1i—S1—C1—O1	−0.1 (2)	C5—C6—C7—C8	−1.2 (4)
S1i—S1—C1—N1	−179.48 (15)	C4—C3—C8—C7	0.4 (3)
C1—N1—C3—C4	−81.5 (3)	N1—C3—C8—C7	176.7 (2)
C2—N1—C3—C4	92.5 (3)	C6—C7—C8—C3	0.9 (4)
C1—N1—C3—C8	102.2 (3)	C1—S1—S1i—C1i	−81.55 (14)
C2—N1—C3—C8	−83.8 (3)		

Symmetry code: (i) −x+1, y, −z+3/2.