

Bis(*N,N*-dimethylthiocarbamoylthio)-
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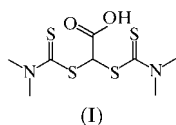
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Bis(*N,N*-dimethylthiocarbamoylthio)acetic acid, [(CH₃)₂N-
C(=S)S]₂CHC(=O)OH or C₈H₁₄N₂O₂S₄, exists as a centro-
symmetric hydrogen-bonded dimer [O···O 2.661 (3) Å].

Comment

The thiocarbamoylthioacetic acids, R₂NC(S)SCH₂C(O)OH, are plant growth-regulating compounds (Fawcett *et al.*, 1956; van der Kerk *et al.*, 1995) that are readily synthesized by displacing the chloride of chloroacetic acid by the dithiocarbamate nucleophile (Nachmias, 1952). Their acid dissociation constants (Janssen, 1963) have been explained in terms of extensive mesomeric electron displacements of the thio-carbonyl group; the R₂NC(S)S portion also exerts an appreciable electron-withdrawing effect (Janssen, 1962). The analogous [R₂NC(S)S]₂CHC(O)OH acid is known only for the R = CH₃ homolog as other homologs cannot be synthesized by the reaction of the dithiocarbamate ion with dichloroacetic acid (Pluijgers, 1959). The title compound, (I), crystallizes as a centrosymmetric dimer that is held together by a short hydrogen bond [O···O 2.664 (3) Å]. Bond dimensions in the acid are similar to those found in the dicyclohexylammonium (Ng, 1996) and tetramethylammonium (Ng, 1997) salts. The tetramethylammonium salt crystallizes with a molecule of the acid, and the carboxylate and acid moieties are linked across a center of inversion by the acid H atom [O···O 2.486 (3) Å].



An examination of the bond dimensions found in [(CH₃)₂NC(S)S]₂CHC(O)OH acid does not suggest any particularly abnormal values. The bond dimensions that are calculated by the GAUSSIAN suite (Frisch *et al.*, 1995) compare well with those found experimentally. Similar calculations were also performed on [(C₂H₅)₂N-C(S)S]₂CHC(O)OH to ascertain why this and other homologs could not be isolated. The calculations, which used the same model chemistry (HF/3-21G), gave insignificant differences, other than for somewhat longer carbon–sulfur double-bonds, *i.e.* this acid should be thermodynamically stable (Table 2).

Experimental

The title acid was synthesized from dimethylamine, carbon disulfide and dichloroacetic acid (Nachmias, 1952; Pluijgers & van der Kerk, 1961), and crystals were obtained upon recrystallization of the compound from ethanol.

Crystal data

C₈H₁₄N₂O₂S₄
M_r = 298.45
Triclinic, P $\bar{1}$
a = 7.3275 (1) Å
b = 7.6321 (2) Å
c = 12.6379 (4) Å
α = 95.809 (2)°
β = 96.991 (1)°
γ = 92.244 (2)°
V = 696.96 (3) Å³

Z = 2
D_x = 1.422 Mg m⁻³
Mo Kα radiation
Cell parameters from 3176
reflections
θ = 2.80–28.28°
μ = 0.669 mm⁻¹
T = 293 (2) K
Block, faint tan
0.38 × 0.30 × 0.16 mm

Data collection

Siemens CCD area-detector
diffractometer
ω scans
Absorption correction: empirical
(SADABS; Sheldrick, 1996)
T_{min} = 0.785, T_{max} = 0.900
5032 measured reflections

3349 independent reflections
2292 reflections with I > 2σ(I)
R_{int} = 0.037
θ_{max} = 28.28°
h = -9 → 9
k = -10 → 10
l = -16 → 9

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.053
wR(F²) = 0.139
S = 0.946
3349 reflections
151 parameters
H-atom parameters constrained

w = 1/[σ²(F_o²) + (0.0705P)²]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} = 0.001
Δρ_{max} = 0.38 e Å⁻³
Δρ_{min} = -0.39 e Å⁻³
Extinction correction: SHELXL97
Extinction coefficient: 0.030 (5)

Table 1

Hydrogen-bonding geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···O2 ⁱ	0.82	1.84	2.661 (3)	178

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

Table 2

Bond distances (Å) and angles (°) derived from *ab initio* (HF/3-21G) calculations.

	(I)	HF/3-21G
S1—C3	1.790 (3)	1.856
S1—C2	1.812 (3)	1.873
S2—C3	1.655 (3)	1.713
S3—C6	1.783 (3)	1.841
S3—C2	1.799 (3)	1.856
S4—C6	1.661 (3)	1.720
O1—C1	1.313 (3)	1.312
O2—C1	1.214 (3)	1.214
N1—C3	1.332 (3)	1.332
N1—C5	1.463 (4)	1.461
N1—C4	1.467 (4)	1.469
N2—C6	1.326 (3)	1.327
N2—C8	1.462 (4)	1.462
N2—C7	1.465 (3)	1.462
C1—C2	1.531 (4)	1.529
C3—S1—C2	102.2 (1)	103.1
C6—S3—C2	102.4 (1)	101.5
C3—N1—C5	123.8 (2)	123.0
C3—N1—C4	119.7 (2)	118.8
C5—N1—C4	116.5 (2)	118.2
C6—N2—C8	123.4 (2)	122.4
C6—N2—C7	120.7 (2)	119.1
C8—N2—C7	115.8 (2)	118.5
O2—C1—O1	124.9 (3)	123.8
O2—C1—C2	123.0 (2)	126.8
O1—C1—C2	112.0 (2)	109.3
C1—C2—S3	113.9 (2)	113.0
C1—C2—S1	105.0 (2)	106.7
S3—C2—S1	106.4 (1)	107.1
N1—C3—S2	124.9 (2)	124.9
N1—C3—S1	111.4 (2)	112.5
S2—C3—S1	123.7 (2)	122.5
N2—C6—S4	125.6 (2)	124.8
N2—C6—S3	113.0 (2)	114.0
S4—C6—S3	121.4 (1)	121.2

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); software used to prepare material for publication: *SHELXL97*.

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