

A 1:1 molecular complex of bis(4-aminophenyl) disulfide and 4-aminothiophenol

Venu R. Vangala,^a Gautam R. Desiraju,^{a*} Ram K. R. Jetti,^b Dieter Bläser^b and Roland Boese^b

^aSchool of Chemistry, University of Hyderabad, Hyderabad 500 046, India, and

^bInstitut für Anorganische Chemie der Universität-GH Essen, Universitätsstraße 3-5, D-45117 Essen, Germany

Correspondence e-mail: desiraju@uohyd.ernet.in

Received 18 June 2002

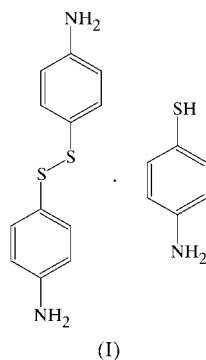
Accepted 18 July 2002

Online 30 September 2002

The centrosymmetric crystal structure of the title complex, $C_{12}H_{12}N_2S_2 \cdot C_6H_7NS$, is built up of dimers of the constituent molecules and stabilized by a herring-bone geometry between the phenyl rings. The structure reveals an $N-H \cdots N-H \cdots N-H \cdots S$ co-operative hydrogen-bonded chain, and $C-H \cdots S$ and $N-H \cdots \pi$ hydrogen bonds. The $S-H$ group forms an uncommon $S-H \cdots \pi$ interaction.

Comment

Crystals of the title complex, (I), were obtained during an investigation of 4-aminothiophenol. Aerial oxidation of this compound leads to the disulfide, which cocrystallizes with the original thiophenol to give the present 1:1 molecular complex.



Complex (I) illustrates a situation wherein a crystal structure is stabilized by the interplay of strong and also weak intermolecular interactions. The molecular geometry and numbering of (I) are given in Fig. 1. The $S-S$ distance of 2.0645 (7) Å is normal and is comparable with the length of 2.074 (1) Å reported for bis(4-aminophenyl) disulfide (Vittal & Anjali, 1998). The $C-S-S-C$ torsion angle in (I) is 94.8 (1)° [80.2 (1)° in bis(4-aminophenyl) disulfide]. Fig. 2 shows a packing diagram of (I) viewed down the [010] axis.

The geometry of the intermolecular interactions is given in Table 1.

The structure of (I) is stabilized by a finite co-operative chain of $N3-H3A \cdots N2-H2B \cdots N1-H1A \cdots S2$ hydrogen bonds in the ac plane. The weak $C2-H2 \cdots S2$ interaction is antico-operative with respect to the above set of interactions. Other antico-operative interactions, namely $N3-H3B \cdots \pi$ and $C3-H3 \cdots \pi$, are also seen. A weak $C15-H15 \cdots \pi$ interaction exists between screw-related molecules.

Although the $S-H$ group is considered to be one of the classical hydrogen-bonding groups, it does not form $S-H \cdots X$ interactions very often (Desiraju & Steiner, 1999; Allen *et al.*,

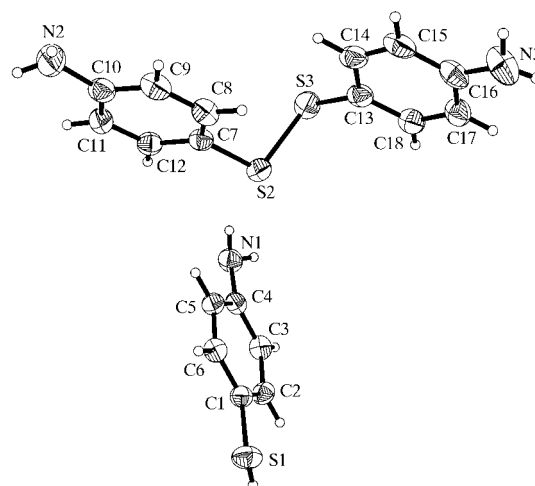


Figure 1

A view of the molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

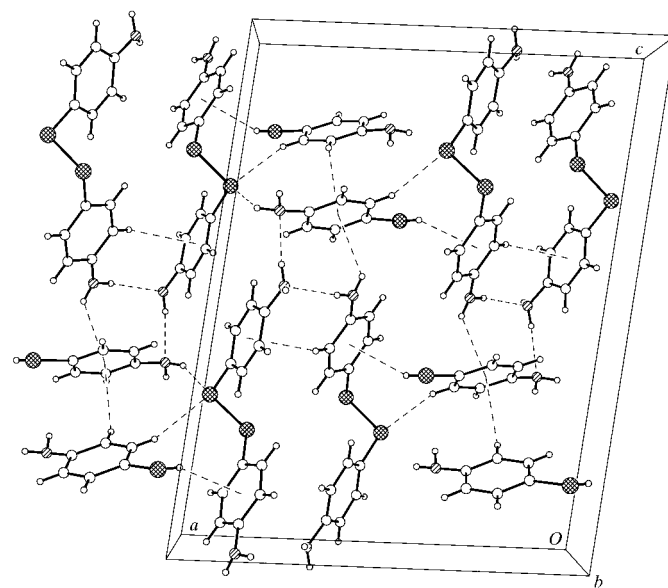


Figure 2

The packing of (I), viewed down [010]. N and S atoms are shaded differently for clarity, and hydrogen bonds are shown as dotted lines. Note the $N-H \cdots N-H \cdots N-H \cdots S$ chain and the $S-H \cdots \pi$ interactions.

1997; Steiner, 2000). In (I), there is a well characterized S—H···π interaction, which is uncommon. In the Cambridge Structural Database (CSD, Version 5.23; Allen & Kennard, 1993), there are only eight crystal structures [refcodes HIPMUO (Zhu-Ohlbach *et al.*, 1998), SIZBAE (Bernardinelli *et al.*, 1991), TASPOS (Slusarchyk *et al.*, 1995), TAXMUA (Nishio *et al.*, 1996), VOPBEH (Garbarczyk & Krolikowska, 1992), YOMWAY01 (Gibbs *et al.*, 1995), and YULZOU and YULZUA (Sellmann *et al.*, 1995)], containing 12 thiol groups, wherein such S—H···π interactions are seen. Not many crystal structures of aminothiols have been reported to date. The present structure is in contrast with that of 6-amino-pyridine-1-thiol (Sabino *et al.*, 2002), where N—H···N hydrogen bonds are preferentially formed.

Experimental

Aerial oxidation of 4-aminothiophenol leads to the disulfide, which cocrystallizes with the original thiophenol, leading to yellow crystals of the 1:1 molecular complex, (I).

Crystal data

C₁₂H₁₂N₂S₂·C₆H₇NS
M_r = 373.54
 Monoclinic, *P*₂₁/*n*
a = 15.210 (3) Å
b = 6.0099 (11) Å
c = 19.683 (4) Å
 β = 96.814 (3)°
V = 1786.6 (6) Å³
Z = 4

D_x = 1.389 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 5202 reflections
 θ = 2.7–28.3°
 μ = 0.42 mm⁻¹
T = 203 (2) K
 Rod, yellow
 0.28 × 0.08 × 0.05 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω and φ scans
 Absorption correction: empirical (*SADABS*; Blessing, 1995)
T_{min} = 0.89, *T_{max}* = 0.99
 8776 measured reflections

4410 independent reflections
 3497 reflections with *I* > 2σ(*I*)
R_{int} = 0.081
 θ_{max} = 28.3°
h = -20 → 20
k = 0 → 8
l = -26 → 26

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.050
wR (*F*²) = 0.128
S = 1.04
 4410 reflections
 217 parameters
 H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.070*P*)² + 0.063*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} = 0.001
 Δρ_{max} = 0.79 e Å⁻³
 Δρ_{min} = -0.25 e Å⁻³

H atoms on C atoms were refined using a riding model starting from idealized geometries, with *U*_{iso}(H) = 1.5*U*_{eq}(C). The positions of H atoms on N and S atoms were taken from a difference Fourier map and these H atoms were also refined as riding, with *U*_{iso}(H) = 1.2*U*_{eq}(N) or 1.5*U*_{eq}(S). We note that the geometry around the N1-amino group is pyramidal and this is in accord with previous observations in crystal structures of aromatic amines where the N atom is a hydrogen-bond acceptor (Allen *et al.*, 1997). The maximum/minimum residual electron densities are located close to the S atoms, obviously also influencing the position of the attached H1 atom.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve

Table 1

Hydrogen-bonding geometry (Å, °).

Cg1, Cg2 and Cg3 are the centroids of rings C–C6, C7–C12 and C13–C18, respectively.

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
N2—H2B···N1 ⁱ	0.87	2.34	3.199 (2)	168
N1—H1A···S2	0.88	2.99	3.7452 (16)	144
N3—H3A···N2 ⁱⁱ	0.91	2.52	3.410 (3)	168
C2—H2···S2 ⁱⁱⁱ	0.94	2.84	3.7387 (18)	162
S1—H1···Cg3 ⁱⁱⁱ	0.90	2.89	3.7310 (12)	155
N3—H3B···Cg1 ^{iv}	0.86	2.65	3.312 (2)	135
C3—H3···Cg1 ⁱⁱⁱ	0.94	2.88	3.544 (2)	129
C15—H15···Cg2 ⁱⁱ	0.94	2.83	3.614 (2)	142

Symmetry codes: (i) 1 - *x*, 1 - *y*, -*z*; (ii) $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$; (iii) $\frac{1}{2} - x, y - \frac{1}{2}, \frac{1}{2} - z$; (iv) $x - \frac{1}{2}, \frac{1}{2} - y, z - \frac{1}{2}$.

structure: *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

VRV thanks the CSIR for fellowship support. RKRJ and RB are grateful to the DFG for financial assistance. GRD is a recipient of the Alexander von Humboldt Research Award.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1383). Services for accessing these data are described at the back of the journal.

References

- Allen, F. H., Bird, C. M., Rowland, R. S. & Raithby, P. R. (1997). *Acta Cryst.* **B53**, 696–701.
- Allen, F. H., Hoy, V. J., Howard, J. A. K., Thalladi, C. R., Desiraju, G. R., Wilson, C. C. & McIntyre, G. J. (1997). *J. Am. Chem. Soc.* **119**, 3477–3480.
- Allen, F. H. & Kennard, O. (1993). *Chem. Des. Autom. News*, **8**, 1, 31–37.
- Bernardinelli, G., Geoffroy, M. & Franz, R. (1991). *Z. Kristallogr.* **195**, 147–149.
- Blessing, R. H. (1995). *Acta Cryst.* **A51**, 33–38.
- Bruker (1997). *SMART*. Version 5.054. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2000). *SAINT*. Version 6.01. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2001). *SHELXTL*. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
- Desiraju, G. R. & Steiner, T. (1999). In *The Weak Hydrogen Bond in Structural Chemistry and Biology*. Oxford University Press.
- Garbarczyk, J. & Krolikowska, M. (1992). *Z. Kristallogr.* **198**, 322–324.
- Gibbs, C. G., Sujeeth, P. K., Rogers, J. S., Stanley, G. G., Krawiec, M., Watson, W. H. & Gutsche, C. D. (1995). *J. Org. Chem.* **60**, 8394–8402.
- Nishio, T., Mori, Y.-I., Iida, I. & Hosomi, A. (1996). *J. Chem. Soc. Perkin Trans. 1*, pp. 921–926.
- Sabino, J. R., da Silva, C. H. T. P. & Yonashiro, M. (2002). *Acta Cryst.* **C58**, o78–o79.
- Sellmann, D., Ruf, R., Knoch, F. & Moll, M. (1995). *Z. Naturforsch. Teil B*, **50**, 791–801.
- Slusarchyk, W. A., Robl, J. A., Taunk, P. C., Asaad, M. M., Bird, J. E., DiMarco, J. & Pan, Y. (1995). *Bioorg. Med. Chem. Lett.* **5**, 753–758.
- Steiner, T. (2000). *Acta Cryst.* **C56**, 876–877.
- Vittal, J. J. & Anjali, K. S. (1998). *Cryst. Eng.* **1**, 147–152.
- Zhu-Ohlbach, Q., Gleiter, R., Rominger, F., Schmidt, H.-L. & Reda, T. (1998). *Eur. J. Org. Chem.* pp. 2409–2416.