organic compounds

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Hydrogen-bonded molecular ladders in 1,4-bis(4-chlorophenylsulfonyl)-2,5dimethylbenzene

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In the title compound, $C_{20}H_{16}Cl_2O_4S_2$, the molecules lie across centres of inversion. A single type of intermolecular C– H···O hydrogen bond, with a C···O distance of 3.254 (3) Å and a C–H···O angle of 132°, links the molecules into ladders whose uprights form C(6) chains and whose rungs enclose centrosymmetric $R_2^2(22)$ rings.

Comment

In molecular solids which contain an excess of hard hydrogenbond acceptors over hard hydrogen-bond donors (Braga *et al.*, 1995), it is to be expected that C—H bonds, particularly those in aromatic systems, will participate as donors in hydrogen bonds of the types C—H···O or C—H···N (Hanton *et al.*, 1992). Aromatic sulfones are particularly effective in C— H···O hydrogen-bond formation between aromatic C—H bonds acting as hydrogen-bond donors and sulfone S==O bonds acting as acceptors (Glidewell *et al.*, 1995; Meehan *et al.*, 1997; Ferguson *et al.*, 1999). In particular, the bis(sulfones) (PhSO₂)₂CH₂ (Glidewell *et al.*, 1995) and (4-MeOC₆H₄-SO₂)₂CH₂ (Meehan *et al.*, 1997) form one-dimensional chains and a three-dimensional framework, respectively, as a result of



such hydrogen-bond formation. As an example of a bis(sulfone) in which a further aromatic nucleus has been incorporated between the two sulfone groups, we now report the structure of the title compound, (I), wherein the molecules are linked into a molecular ladder by $C-H\cdots O$ hydrogen bonds.

Compound (I) crystallizes in the space group $P\overline{1}$ with Z = 1; hence the molecules lie across centres of inversion and, for the sake of convenience, the reference molecule has been placed across the inversion centre at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$. The molecular conformation of (I) (Fig. 1) is dominated by the near coplanarity of the O-S-C-C fragments (Table 1). As commonly observed in diaryl sulfones (Ferguson *et al.*, 1999), there is an electrostatic attraction between the S=O bond, which is polarized in the sense S⁺-O⁻, and an *ortho* C-H bond in the adjacent ring, polarized in the sense C⁻-H⁺. In the most common arrangement, as also found here, each O atom of the sulfone unit interacts with a different ring, so that the local symmetry around the S atom is close to C_2 .

Both of the S1-C7-C9ⁱ and C7ⁱ-C9-C91 bond angles [symmetry code: (i) 1 - x, 1 - y, 1 - z] are significantly greater than 120° (Table 1). This type of angular distortion has been observed in several diaryl sulfones where there is a methyl group adjacent to the SO₂ unit (Jeyakanthan *et al.*, 1998), and it may plausibly be interpreted as arising from repulsive interactions between the methyl group and the adjacent sulfone unit. The O-S-O bond angle is much larger than tetrahedral. All the bond lengths are typical of their types (Allen *et al.*, 1987).

There is a single type of intermolecular hydrogen bond (Table 2) linking the molecules. Atom C2 at (x, y, z), a component of the molecule centred at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$, acts as hydrogen-bond donor to O2 at (x, y - 1, z), which is a component of the molecule centred at $(\frac{1}{2}, -\frac{1}{2}, \frac{1}{2})$; repetition of this hydrogen bond generates a C(6) chain running parallel to [010] and generated by translation (Fig. 2). The symmetryrelated C2 atom in the original $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$ molecule is at (1 - x, 1)-y, 1-z), and this acts as donor to O2 at (1-x, 2-y, 1-z)z), which is a component of the molecule centred at $(\frac{1}{2}, \frac{3}{2}, \frac{1}{2})$; hence, another C(6) chain is generated, running antiparallel to the first. The overall supramolecular structure (Fig. 2) thus takes the form of a molecular ladder, in which the uprights are formed by a pair of antiparallel C(6) chains, while the rungs are provided by the central aryl rings of successive molecules. Between the rungs are $R_2^2(22)$ rings centred at $(\frac{1}{2}, n, \frac{1}{2})$ (n = zero)or integer), but the aromatic and methyl H atoms preclude the presence of any void space. A single ladder runs through each unit cell, and there are no significant interactions between



Figure 1

The molecule of (I) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of an arbitrary radius [symmetry code: (i) 1 - x, 1 - y, 1 - z].

neighbouring ladders; in particular, there are no aromatic $\pi \cdots \pi$ stacking interactions.

It is notable that the intermolecular hydrogen bonding does not involve the central aryl ring. By contrast, in each of (PhSO₂)₂CH₂ (Glidewell et al., 1995) and (4-MeOC₆H₄₋ SO₂)₂CH₂ (Meehan et al., 1997), both of the C-H bonds of the central methylene unit are engaged in hydrogen-bond formation. It is thus of interest to consider some related compounds retrieved from the Cambridge Structural Database (CSD; Allen & Kennard, 1993). In (PhSO₂CH₂)₂ (FAHNOR; Hauback & Mo, 1990), although the molecules could lie across centres of inversion, as in (I), they do not in



Figure 2

Part of the crystal structure of (I) showing the formation of a molecular ladder. For the sake of clarity, the atom labels are applied to the unit at (x, x)y - 1, z) and H atoms not involved in the supramolecular structure have been omitted.

fact do so; the molecules are again linked into molecular ladders, but the C-H···O hydrogen bonds forming the uprights use only H atoms from the central -CH₂-CH₂- unit. The ladders are linked into a two-dimensional array by further C-H...O hydrogen bonds involving the aromatic rings. For neither the Z isomer of PhSO₂CH=CHSO₂Ph (DEBYIS; De Lucchi et al., 1985) nor the centrosymmetric E isomer (DEBYOY; De Lucchi et al., 1985) are there any H-atom coordinates in the CSD; nonetheless, it is clear from the short intermolecular $C \cdots O$ distances that the intermolecular aggregation is dominated by the aromatic C-H bonds rather than those in the alkene fragment. The factors which determine the involvement of aromatic versus other C-H bonds in these systems are as yet not fully clear. However, it may be expected that aliphatic C-H bonds adjacent to SO₂ units are more acidic than is typical for aliphatic C-H bonds, and thus more like aromatic C-H bonds in their hydrogen-bonding behaviour.

Experimental

A sample of 2,5-bis(4-chlorophenylthio)-1,4-dimethylbenzene (Grant et al., 1987) was oxidized using sodium periodate in CH₂Cl₂ to yield (I). Crystals suitable for single-crystal X-ray diffraction were grown from an ethanol solution (m.p. 506–508 K).

$C_{20}H_{16}Cl_2O_4S_2$	Z = 1		
$M_r = 455.35$	$D_{\rm x} = 1.605 {\rm Mg} {\rm m}^{-3}$		
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation		
a = 5.4859 (2) Å	Cell parameters from 2112		
b = 6.9595 (4) Å	reflections		
c = 12.5211 (7) Å	$\theta = 1.65 - 27.45^{\circ}$		
$\alpha = 98.6245 \ (17)^{\circ}$	$\mu = 0.592 \text{ mm}^{-1}$		
$\beta = 93.561 \ (4)^{\circ}$	T = 150 (2) K		
$\gamma = 92.174 \ (3)^{\circ}$	Plate, colourless		
$V = 471.18 (4) \text{ Å}^3$	$0.10 \times 0.10 \times 0.03 \text{ mm}$		

Data collection

Crystal data

Nonius KappaCCD diffractometer 1689 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.039$ φ and ω scans with κ offsets $\theta_{\rm max} = 27.45^{\circ}$ Absorption correction: multi-scan $h = -7 \rightarrow 7$ (SORTAV; Blessing, 1995, 1997) $k = -9 \rightarrow 8$ $T_{\min} = 0.943, T_{\max} = 0.985$ 5965 measured reflections $l = -16 \rightarrow 16$ 2112 independent reflections

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0411P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.039$ + 0.1338P] $wR(F^2) = 0.104$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ S = 1.073 $\Delta \rho_{\rm max} = 0.39 \; {\rm e} \; {\rm \AA}^{-3}$ 2112 reflections $\Delta \rho_{\rm min} = -0.53 \text{ e } \text{\AA}^{-3}$ 128 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

S1-O1	1.4322 (14)	S1-C7	1.782 (2)
S1-O2	1.4416 (14)	Cl1-C1	1.738 (2)
S1-C4	1.772 (2)		
01 51 02	119 (7 (0)		101 76 (19)
01-51-02	118.67 (9)	08-07-09	121.70 (18)
01 - 81 - C4	108.12 (9)	C8_C/_S1	115.96 (15)
O2 - S1 - C4	107.72 (9)	$C9^{1}-C7-S1$	122.24 (14)
O1-S1-C7	110.17 (9)	C8-C9-C7 ⁱ	115.95 (17)
O2-S1-C7	107.60 (9)	C7 ⁱ -C9-C91	124.97 (18)
C4-S1-C7	103.49 (9)	C8-C9-C91	119.06 (18)
O1-S1-C4-C3	-4.3 (2)	O2-S1-C7-C8	17.70 (17)

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

Table 2

Hydrogen-bonding geometry (Å, °).

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C2 - H2 \cdots O2^{ii}$	0.95	2.54	3.254 (3)	132

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

Compound (I) crystallized in the triclinic system; space group $P\overline{1}$ was assumed and confirmed by the analysis. H atoms were treated as riding, with C-H 0.95 (aromatic) and 0.98 Å (methyl). Examination of the structure with PLATON (Spek, 1999) showed that there were no solvent-accessible voids in the crystal lattice.

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Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO* (Otwinowski & Minor, 1997); data reduction: *DENZO*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 1999); software used to prepare material for publication: *SHELXL*97 and *PRPKAPPA* (Ferguson, 1999).

X-ray data were collected at the EPSRC X-ray Crystallographic Service, University of Southampton, using a Nonius KappaCCD diffractometer. The authors thank the staff for all their help and advice.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GG1006). Services for accessing these data are described at the back of the journal.

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