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# 2,3-Bis[(4-methylphenyl)sulfanyl]-4H-1-benzothiopyran-4-one 1,1-dioxide forms a framework built from $\mathbf{C}-\mathbf{H} \cdots \mathrm{O}, \mathrm{C}-\mathbf{H} \cdots \pi$ (arene) and $\pi-\pi$ interactions 

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Molecules of the title comound, $\mathrm{C}_{23} \mathrm{H}_{18} \mathrm{O}_{3} \mathrm{~S}_{3}$, are linked into sheets by a combination of one $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond and two $\mathrm{C}-\mathrm{H} \cdots \pi$ (arene) hydrogen bonds, and these sheets are weakly linked into a three-dimensional structure by means of a single aromatic $\pi-\pi$ stacking interaction.

## Comment

1,4-Naphthoquinone derivatives have been extensively studied as dyes and pigments (Tilak, 1971). Although 4H-1-benzothiopyran-4-one 1,1-dioxide derivatives have not been so well studied, it is known that their absorption spectra exhibit hypsochromic effects compared with the 1,4naphthoquinone analogues (Christie et al., 1980; Watanabe et al., 1989, 1990). The crystal structures of two halogenated derivatives have been investigated as part of a study of nonlinear optical materials (Watanabe et al., 1989, 1992). Accordingly, we have now investigated the title nonhalogenated analogue, (I) (Fig. 1), whose structure exhibits a rich variety of weak direction-specific intermolecular interactions.

The $\mathrm{S}-\mathrm{C}$ distances in (I) (Table 1) show an unusual pattern. For the sulfone atom $S$, formally $S^{V I}$, the $S-C$ distance involving aryl C atoms is less than those involving alkene C atoms, whereas for the sulfide atoms S 2 and S 3 , formally $\mathrm{S}^{-\mathrm{II}}$, the bonds to alkene C atoms are the shorter. Within the heterocyclic ring, there is clear bond fixation with a short C8-C9 bond, while the $\pi$-bonding in each of the arene rings is fully delocalized.

The $\mathrm{O}-\mathrm{S}-\mathrm{O}$ bond angle in the sulfone fragment of (I) is significantly larger than the tetrahedral angle, while the opposed $\mathrm{C}-\mathrm{S}-\mathrm{C}$ angle at S 1 is correspondingly much smaller
than the tetrahedral value. The bond angles at S2 and S3 are typical of those in organosulfides.

(I)

For the heterocyclic ring of (I), the ring-puckering parameters (Cremer \& Pople, 1975) for the atom sequence S1/C1/ C6-C9 are $\theta=85.5(5)^{\circ}$ and $\varphi=18.6(5)^{\circ}$, indicative of a conformation best described as twist-boat (Evans \& Boeyens, 1989). The values of the torsion angles $\mathrm{C} 9-\mathrm{C} 8-\mathrm{S} 2-\mathrm{C} 21$ and $\mathrm{C} 8-\mathrm{C} 9-\mathrm{S} 3-\mathrm{C} 31$ are indicative of entirely different orientations for the two pendant groups (see also Fig. 1).

The molecules of (I) are linked into a three-dimensional framework by a variety of weak but direction-specific intermolecular forces, and the formation of this framework is readily analysed by means of the substructure approach (Gregson et al., 2000). A single $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond (Table 2) links the molecules into chains. These chains are linked into sheets by the combined action of two $\mathrm{C}-\mathrm{H} \cdots \pi$ (arene) hydrogen bonds and the sheets are linked together by a single aromatic $\pi-\pi$ stacking interaction.

Atom C32 in the molecule at $(x, y, z)$ acts as hydrogen-bond donor to sulfone atom O 11 in the molecule at $\left(x, \frac{1}{2}-y, z-\frac{1}{2}\right)$, so producing a zigzag $C(7)$ chain (Bernstein et al., 1995)


Figure 1
The molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $30 \%$ probability level.
running parallel to the [001] direction and generated by the $c$ glide plane at $y=\frac{1}{4}$ (Fig. 2). The formation of this chain is modestly reinforced by a short dipolar contact between the negatively charged sulfone atom O11 in the molecule at $(x, y$, $z$ ) and the positively charged carbonyl atom C 7 in the molecule at $\left(x, \frac{1}{2}-y, \frac{1}{2}+z\right)$. The $\mathrm{O} 11 \cdots \mathrm{C}^{\mathrm{iv}}$ distance is 2.938 (3) $\AA$ and the $\mathrm{S} 1-\mathrm{O} 11 \cdots \mathrm{C}^{\mathrm{iv}}$ angle is $138.6(2)^{\circ}$ [symmetry code: (iv) $\left.x, \frac{1}{2}-y, \frac{1}{2}+z\right]$.

Atoms C5 and C35 in the molecule at $(x, y, z)$ act as hydrogen-bond donors to, respectively, the C31-C36 ring in the molecule at $(x-1, y, z)$ and the C21-C26 ring in the molecule at $(1+x, y, z)$. In this manner, a chain of rings is generated by translation along the [100] direction (Fig. 3). The combination of the [100] and [001] chains produces an (010)


Figure 2
Part of the crystal structure of (I), showing the formation of a $C(7)$ chain along [001]. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (\#) are at the symmetry positions $\left(x, \frac{1}{2}-y, z-\frac{1}{2}\right)$ and $(x, y, z-1)$, respectively.


Figure 3
Part of the crystal structure of (I), showing the formation of a chain of rings along [100]. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (\#) are at the symmetry positions $(1+x, y, z)$ and $(x-1, y, z)$, respectively.


Figure 4
A stereoview of part of the crystal structure of (I), showing the formation of an (010) sheet by the combination of the [100] and [001] chains. For the sake of clarity, H atoms not involved in the motifs shown have been omitted.


Figure 5
Part of the crystal structure of (I), showing the $\pi-\pi$ stacking interaction which links adjacent (010) sheets. For the sake of clarity, H atoms have been omitted. Atoms marked with an asterisk (*) are at the symmetry position $(-x,-y, 2-z)$.
sheet lying in the domain $-0.06<y<0.56$ and generated by the $c$-glide plane at $y=\frac{1}{4}$ (Fig. 4). A similar sheet, generated by the $c$-glide plane at $y=\frac{3}{4}$ and related to the first by inversion, lies in the domain $0.44<y<1.06$.

Adjacent (010) sheets are weakly linked by a $\pi-\pi$ stacking interaction. The C21-C26 rings in the molecules at $(x, y, z)$ and $(-x,-y, 2-z)$ lie in (010) sheets generated by glide planes at $y=\frac{1}{4}$ and $-\frac{1}{4}$, respectively. These two rings are parallel, with an interplanar spacing of 3.548 (2) $\AA$. The ring-centroid separation is $3.876(2) \AA$, corresponding to a centroid offset of 1.561 (2) $\AA$ (Fig. 5). Propagation by the space group of this interaction is sufficient to link all of the (010) sheets into a single framework structure.

## Experimental

A sample of compound (I) was prepared following the procedure of Christie et al. (1980). The crystals used in the present structure determination were grown by slow evaporation of a solution in ethanol [m.p. 432-434 K; literature m.p. 433-433.5 K (Christie et al., 1980)].

## Crystal data

$\mathrm{C}_{23} \mathrm{H}_{18} \mathrm{O}_{3} \mathrm{~S}_{3}$
$M_{r}=438.58$
Monoclinic, $P 2_{\mathrm{a}_{1}} / c$
$a=9.6349$ (3) $\AA$
$b=22.5791$ (7) $\AA$
$c=9.1895$ (4) A
$\beta=91.253(2)^{\circ}$
$V=1998.67(12) \AA^{3}$
$Z=4$

$$
D_{x}=1.458 \mathrm{Mg} \mathrm{~m}^{-3}
$$

Mo $K \alpha$ radiation
Cell parameters from 4564 reflections
$\theta=3.2-27.5^{\circ}$
$\mu=0.39 \mathrm{~mm}^{-1}$
$T=120$ (2) K
Block, orange
$0.30 \times 0.20 \times 0.10 \mathrm{~mm}$

## Data collection

Nonius KappaCCD area-detector diffractometer
$\varphi$ scans, and $\omega$ scans with $\kappa$ offsets
Absorption correction: multi-scan
(SORTAV; Blessing, 1995, 1997)
$T_{\text {min }}=0.891, T_{\text {max }}=0.962$
8989 measured reflections

4564 independent reflections
3002 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.051$
$\theta_{\text {max }}=27.5^{\circ}$
$h=-12 \rightarrow 12$
$k=-27 \rightarrow 29$
$l=-11 \rightarrow 11$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.044$
$w R\left(F^{2}\right)=0.107$
$S=1.00$
4564 reflections
264 parameters

H-atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0507 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }=0.001$
$\Delta \rho_{\text {max }}=0.32 \mathrm{e}^{\circ} \AA^{-3}$
$\Delta \rho_{\min }=-0.52 \mathrm{e}^{-3}$

Table 1
Selected geometric parameters ( $\left(\AA^{\circ}{ }^{\circ}\right)$.

| S1-C1 | $1.753(2)$ | S1-O11 | $1.4363(17)$ |
| :--- | ---: | :--- | :---: |
| S1-C9 | $1.770(2)$ | S1-O12 | $1.4311(16)$ |
| C8-S2 | $1.756(2)$ | C6-C7 | $1.488(3)$ |
| S2-C21 | $1.778(2)$ | C7-C8 | $1.513(3)$ |
| C9-S3 | $1.757(2)$ | C8-C9 | $1.349(3)$ |
| S3-C31 | $1.784(2)$ | C7-O71 | $1.213(3)$ |
|  |  |  |  |
| O11-S1-O12 | $117.38(10)$ | C8-S2-C21 | $101.52(10)$ |
| C1-S1-C9 | $104.88(10)$ | C9-S3-C31 | $103.53(11)$ |
|  |  |  |  |
| S1-C1-C6-C7 | $8.8(3)$ | O71-C7-C8-C9 | $159.0(2)$ |
| C1-C6-C7-C8 | $13.2(3)$ | O71-C7-C6-C1 | $-167.9(2)$ |
| C6-C7-C8-C9 | $-22.1(3)$ | C9-C8-S2-C21 | $133.62(19)$ |
| C7-C8-C9-S1 | $7.4(3)$ | C8-S2-C21-C22 | $-49.7(2)$ |
| C8-C9-S1-C1 | $11.8(2)$ | C8-C9-S3-C31 | $49.6(2)$ |
| C9-S1-C1-C6 | $-19.7(2)$ | C9-S3-C31-C32 | $35.3(2)$ |
|  |  |  |  |

Table 2
Hydrogen-bonding geometry ( $\AA{ }^{\circ},{ }^{\circ}$ ).
$C g 1$ is the centroid of the $\mathrm{C} 31-\mathrm{C} 36$ ring and $C g 2$ is the centroid of the $\mathrm{C} 21-$ C26 ring.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| C32-H32 $\cdots$ O11 $1^{\mathrm{i}}$ | 0.95 | 2.32 | $3.197(3)$ | 154 |
| C5-H5 Cg1 | $1^{\text {(2) }}$ | 0.95 | 2.55 | $3.308(2)$ |
| C35-H35 $\cdots$ Cg2 $^{\text {iii }}$ | 0.95 | 2.73 | $3.658(2)$ | 137 |

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

The space group $P 2_{1} / c$ was uniquely assigned from the systematic absences. All H atoms were located from difference maps and then treated as riding atoms, with $\mathrm{C}-\mathrm{H}$ distances of 0.95 (aromatic) or $0.98 \AA$ (methyl), and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ for aromatic H and $1.5 U_{\text {eq }}(\mathrm{C})$ for methyl H atoms.

Data collection: KappaCCD Server Software (Nonius, 1997); cell refinement: DENZO-SMN (Otwinowski \& Minor, 1997); data reduction: $D E N Z O-S M N$; program(s) used to solve structure: OSCAIL (McArdle, 2003) and SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: OSCAIL and SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and PRPKAPPA (Ferguson, 1999).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1715). Services for accessing these data are described at the back of the journal.

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