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Key indicators

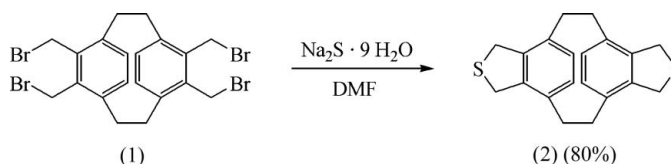
Single-crystal X-ray study
 $T = 143\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.044
 wR factor = 0.107
Data-to-parameter ratio = 13.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

[2.2]Dihydroisobenzothiophene

In the unit cell of the title compound, 1¹,1³,4¹,4³-tetrahydro-1,4(4,7)-di(benzo[*c*]thiophena)cyclohexaphane, $\text{C}_{20}\text{H}_{20}\text{S}_2$, there are four independent but closely similar molecules, each with crystallographic inversion symmetry. The packing is interpreted in terms of layers formed *via* C—H... π interactions.

Comment

2,5-Dihydroisobenzothiophene and its derivatives are useful substrates for the generation of *o*-xylylene intermediates, which can subsequently be trapped by various double- and triple-bond dienophiles to yield six-membered ring systems (Gajewski, 2004). In order to produce such intermediates in the cyclophane series, we required the title [2.2]cyclophane, (2), as a precursor. This is easily obtained from the tetrabromide, (1) (El-Tamany & Hopf, 1980; see *Experimental* section). Here, we report the structure determination of (2) by single-crystal X-ray crystallography.



Compound (2) crystallizes with four independent molecules in the unit cell, each with inversion symmetry (Fig. 1). Molecule 1 is centred at $(\frac{1}{2}, 0, 0)$, molecule 2 (primed atom names) at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$, molecule 3 (doubly primed atom names) at $(0, \frac{1}{2}, 0)$,

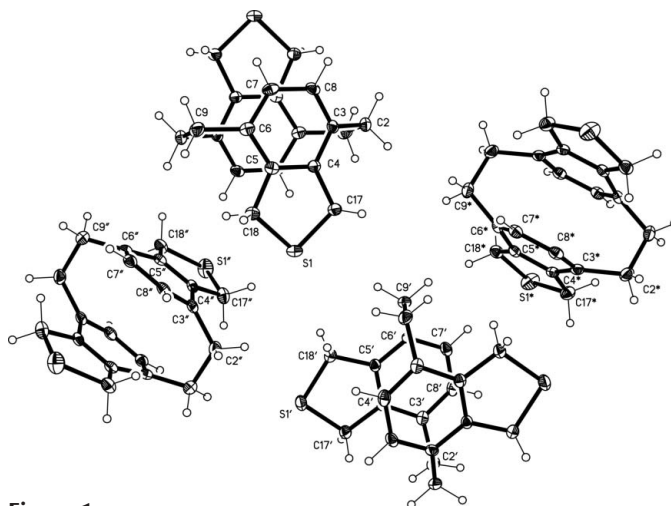


Figure 1

The four independent molecules of the title compound in the crystal structure. Ellipsoids are drawn at the 30% probability level. Only the asymmetric unit is numbered.

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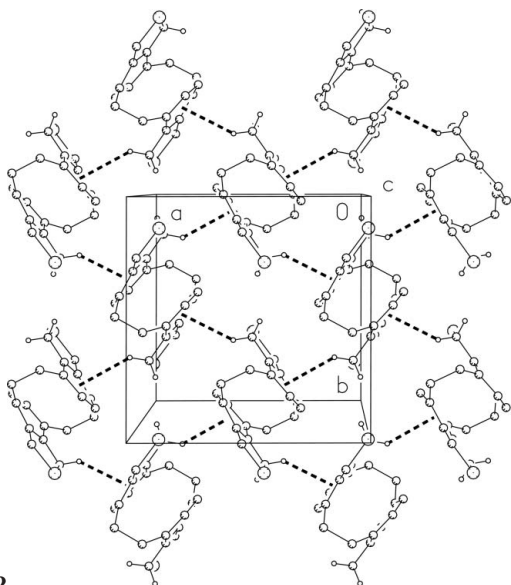


Figure 2

A packing diagram of the title compound, showing a view of one layer at $z \approx 0$. C—H... π contacts (see text) are indicated by dashed lines; H atoms not involved in these interactions have been omitted.

and molecule 4 (starred atom names) at $(1, 0, \frac{1}{2})$. IUPAC-consistent numbering has been chosen as far as possible consistent with the crystallographic symmetry.

All four molecules are closely similar. Bond lengths and angles, *e.g.* those at the S atoms (Table 1), may be considered normal. The six-membered rings show the flattened boat conformation typical of [2.2]cyclophanes, with the bridgehead atoms lying 0.15–0.16 Å out of the plane of the other four atoms. The C—C bonds in the bridges are lengthened (Table 1), also typical of cyclophanes. The thiophenyl rings are planar (r.m.s. deviations < 0.01 Å) and subtend angles of 0–2° with the above planes. Full numerical details are given in the deposited CIF.

The molecular packing involves layers parallel to (001). Molecules 1 and 3 form a layer at $z \approx 0$ (Fig. 2), and molecules 2 and 4 a layer at $z \approx \frac{1}{2}$ (Fig. 3). Both within and between the layers there are a number of C—H...S contacts (Table 2), but these are long (> 2.9 Å) and have no clear upper length limit. For this reason, we prefer to analyse the packing in terms of C—H... π interactions, as we have done previously for other cyclophane systems (*e.g.* El Shaieb *et al.*, 2003). There are four such interactions, listed in Table 2 for normalized C—H bond lengths of 1.08 Å (Steiner, 1998) and for centroids Cg1–Cg4 calculated from the four non-bridgehead atoms C4, C5, C7 and C8 of the six-membered rings. However, we note that all four contacts display either narrow angles or long distances.

Experimental

A solution of the tetrabromide, (1), (1.74 g, 3.0 mmol) in dimethylformamide (DMF; 100 ml) was added to a well stirred suspension of sodium sulfide enneahydrate (1.44 g, 6.0 mmol) in DMF (350 ml) at room temperature over 8 h. After the addition was complete, the mixture was stirred for another 12 h. Extraction with dichloromethane, drying of the organic phase over sodium sulfate and solvent

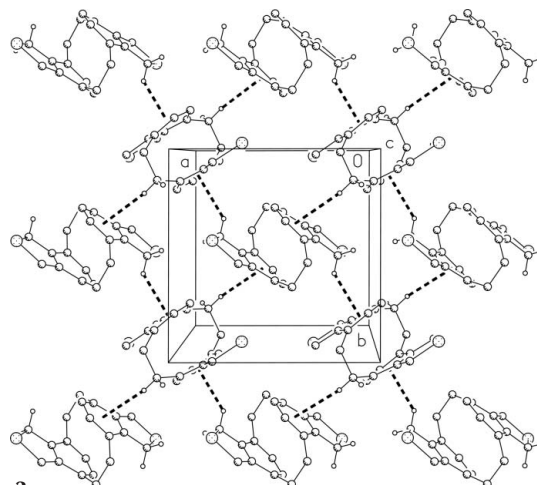


Figure 3

A packing diagram of the title compound, showing a view of one layer at $z \approx \frac{1}{2}$. C—H... π contacts (see text) are indicated by dashed lines; H atoms not involved in these interactions have been omitted.

removal provided (2) (0.7 g, 80%) after silica-gel chromatography. The structure of (2) is also supported by all other spectroscopic and analytical data (Kannan & Hopf, unpublished results).

Crystal data

$C_{20}H_{20}S_2$	$Z = 4$
$M_r = 324.48$	$D_x = 1.379 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 10.088 (2) \text{ \AA}$	Cell parameters from 52 reflections
$b = 10.182 (2) \text{ \AA}$	$\theta = 10\text{--}11.5^\circ$
$c = 15.247 (2) \text{ \AA}$	$\mu = 0.33 \text{ mm}^{-1}$
$\alpha = 86.53 (2)^\circ$	$T = 143 (2) \text{ K}$
$\beta = 88.44 (2)^\circ$	Prism, colourless
$\gamma = 89.81 (2)^\circ$	$0.6 \times 0.35 \times 0.3 \text{ mm}$
$V = 1562.7 (5) \text{ \AA}^3$	

Data collection

Stoe STADI-4 diffractometer	$\theta_{\text{max}} = 25.0^\circ$
θ/ω scans	$h = -12 \rightarrow 6$
Absorption correction: none	$k = -12 \rightarrow 12$
9570 measured reflections	$l = -18 \rightarrow 18$
5496 independent reflections	3 standard reflections
4469 reflections with $I > 2\sigma(I)$	frequency: 60 min
$R_{\text{int}} = 0.026$	intensity decay: 2%

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0354P)^2 + 1.5793P]$
$R[F^2 > 2\sigma(F^2)] = 0.044$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.107$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.01$	$\Delta\rho_{\text{max}} = 0.54 \text{ e \AA}^{-3}$
5496 reflections	$\Delta\rho_{\text{min}} = -0.38 \text{ e \AA}^{-3}$
397 parameters	
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

S1—C18	1.814 (3)	S1''—C18''	1.807 (3)
S1—C17	1.824 (3)	S1''—C17''	1.826 (3)
C2—C9 ⁱ	1.589 (4)	C2''—C9'' ⁱⁱⁱ	1.590 (4)
S1'—C17'	1.813 (2)	S1*—C18*	1.811 (3)
S1'—C18'	1.819 (3)	S1*—C17*	1.818 (3)
C2'—C9'' ⁱⁱ	1.585 (4)	C2*—C9* ^{iv}	1.589 (4)
C18—S1—C17	95.62 (11)	C18''—S1''—C17''	95.74 (12)
C17'—S1'—C18'	95.44 (11)	C18*—S1*—C17*	95.87 (12)

Symmetry codes: (i) $-x + 1, -y, -z$; (ii) $-x + 1, -y + 1, -z + 1$; (iii) $-x, -y + 1, -z$; (iv) $-x + 2, -y, -z + 1$.

Table 2
Hydrogen-bond geometry (Å, °).

<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>
C2''—H2''2...S1'	0.99	2.95	3.723 (3)	136
C9''—H9''2...S1'''	0.99	3.02	3.732 (3)	130
C9*—H9*1...S1''v	0.99	2.96	3.937 (3)	168
C7—H7...S1*vi	0.95	2.94	3.854 (3)	162
C9'—H9'1...S1	0.99	3.07	3.907 (3)	143
C8''—H8''...S1''ii	0.95	3.08	3.896 (3)	145
C8*—H8*...S1''v	0.95	3.10	3.879 (3)	140
C17—H17B...S1*viiii	0.99	3.07	3.974 (3)	153
C18''—H18F...Cg1 ^{ix}	1.08	2.74	3.63	140
C18—H18A...Cg3	1.08	2.78	3.66	111
C17'—H17C...Cg4 ⁱⁱ	1.08	2.61	3.45	135
C2*—H2*2...Cg2 ^v	1.08	2.98	4.04	166

Symmetry codes: (ii) $-x + 1, -y + 1, -z + 1$; (iii) $-x, -y + 1, -z$; (v) $x + 1, y, z$; (vi) $x, y, z - 1$; (vii) $-x + 1, -y + 1, -z$; (viii) $-x + 1, -y, -z + 1$; (ix) $x - 1, y, z$.

H atoms were included in calculated positions and refined using a riding model, with fixed C—H bond lengths of 0.95 Å (sp^2 CH) or 0.99 Å (CH₂). $U_{iso}(H)$ values were fixed at 1.2 times U_{eq} of the parent atom.

Data collection: *DIF4* (Stoe & Cie, 1992); cell refinement: *DIF4*; data reduction: *REDU4* (Stoe & Cie, 1992); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

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