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#### **Key indicators**

Single-crystal X-ray study T = 298 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.050 wR factor = 0.147 Data-to-parameter ratio = 17.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 1,10-Bis(phenylsulfanyl)decane

In the title structure,  $C_{22}H_{30}S_2$ ,  $L^{10}$ , the asymmetric unit contains two independent molecules, both of which lie on crystallographic inversion centers, and the aliphatic chains are in the all-trans conformation. The crystal structure is stabilized by a combination of van der Waals forces and weak  $C - H \cdots \pi$ (arene) interactions.

### Comment

The self-assembly of metal-organic coordination polymers has attracted a great deal of attention because of their potential application as functional materials (Yaghi et al., 1995). Bifunctional organic ligands are usually used as spacers for the building of metal-organic framework materials (MOF) (Carlucci et al., 2002). The S atom is a soft base and has a good ability to coordinate to Ag, which is a soft acid. In addition, the dithiolate building block can afford two coordination sites to a metal center and can be used to construct supramolecular architectures (Black et al., 1995; Bu et al., 2002). When the length of the aliphatic segment between the S atoms is increased, the number of possible conformations and the flexibility of the spacer are augmented. As a result, in metalorganic crystal engineering, when flexible ligands are involved in supramolecular networks, the prediction of the topology of the polymer coordination is more difficult, since there are several other factors affecting the formation of the framework (Withersby et al., 1999, 1997; Noro et al., 2002; Blake et al., 1999). In order to study the effect of the length of the building block on the topology of metal-organic polymer coordination, the title compound,  $L^{10}$ , (I), was synthesized and its structure is reported here.



Compound (I) crystallizes with two (essentially equivalent) independent centrosymmetric molecules in the asymmetric unit, one of which is shown in Fig. 1. Each molecule has a center of symmetry at the midpoint of the central C–C bond, namely C35–C35<sup>i</sup> and C45–C45<sup>ii</sup> [symmetry codes: (i) -x, 1 - y, -z; (ii) 1 - x, 2 - y, 2 - z]. The torsion angles in the aliphatic sequence [S–(CH<sub>2</sub>)<sub>10</sub>–S] are all trans, indicating that the molecules are in the fully extended conformation (Table 1). The dihedral angles formed between the aromatic

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### Figure 1

The molecular structure and atomic numbering of one independent molecule of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms have been omitted. The unlabeled part of the molecule is related by the symmetry transformation (-x, 1 - y, -z).



### Figure 2

A crystal packing diagram of (I). H atoms have been omitted.

rings and the  $[S-(CH_2)_{10}-S]$  sequences are 21.8 (2) and 23.1 (2)° for molecules 1 and 2, respectively. The phenyl groups of neighboring molecules form a dihedral angle of 62.8 (1)°. The bond distances and angles in (I) are in the normal ranges (Table 1) (*International Tables for Crystallography*, 1995, Vol. C).

In the crystal structure of (I) (Fig. 2), there are no significant  $\pi$ - $\pi$  stacking interactions but, in addition to normal van der Waals interactions, there are some weak C-H··· $\pi$ (arene) interactions (Table 2).

## Experimental

1,10-Bis(phenylthio)decane,  $L^{10}$ , (I), was synthesized according to the published procedure of Hartley *et al.* (1979). Compound (I) was obtained as a crystalline powder from which a platelet crystal suitable for X-ray analysis was isolated (yield 91%). Analysis, found: C 73.47, H 8.59%; calculated for C<sub>22</sub>H<sub>30</sub>S<sub>2</sub>: C 73.68, H 8.43%. <sup>1</sup>H NMR (acetone- $d_6$ ,  $\delta$ , p.p.m.): 1.28 [t, 4H, -S-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH<sub>2</sub>-CH

Crystal data

C22H30S2 Z = 2 $M_r = 358.58$  $D_r = 1.186 \text{ Mg m}^{-3}$ Triclinic, P1 Cu  $K\alpha$  radiation a = 5.570 (2) Å Cell parameters from 25 b = 7.943 (3) Å reflections c = 22.953 (7) Å  $\theta = 20.0-25.0^{\circ}$  $\mu = 2.38 \text{ mm}^{-1}$  $\alpha = 82.82(3)^{\circ}$  $\beta = 85.05(3)^{\circ}$ T = 298 (2) K  $\gamma = 89.55 (3)^{\circ}$ Platelet, colorless  $0.25 \times 0.11 \times 0.05 \ \text{mm}$ V = 1003.8 (6) Å<sup>3</sup>

2520 reflections with  $I > 2\sigma(I)$ 

 $R_{\rm int} = 0.038$  $\theta_{\rm max} = 69.9^{\circ}$ 

 $h = -6 \rightarrow 6$ 

 $k = -9 \rightarrow 9$ 

 $l = -28 \rightarrow 28$ 4 standard reflections

frequency: 60 min

intensity decay: none

## Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega$  scans Absorption correction: Gaussian, from crystal shape (*ABSORP* in *NRCVAX*; Gabe *et al.*1989)  $T_{min} = 0.528$ ,  $T_{max} = 0.868$ 24003 measured reflections 3799 independent reflections

#### Refinement

Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.050$ $wR(F^2) = 0.147$	H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0908P)^2]$ where $P = (F_o^2 + 2F_o^2)/3$
S = 0.91	$(\Delta/\sigma)_{\rm max} < 0.001$
3799 reflections	$\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$
217 parameters	$\Delta \rho_{\rm min} = -0.47 \ {\rm e} \ {\rm \AA}^{-3}$

#### Table 1

Selected geometric parameters (Å, °).

S1-C11	1.764 (2)	\$2-C41	1.790 (3)
S1-C31	1.796 (3)	C35-C35 <sup>i</sup>	1.513 (4)
S2-C21	1.759 (2)	C45-C45 <sup>ii</sup>	1.511 (4)
C11-S1-C31	104.59 (11)	C21-S2-C41	105.20 (12)
S1-C31-C32-C33	179.33 (19)	S2-C41-C42-C43	178.6 (2)
C31-C32-C33-C34	179.8 (2)	C41-C42-C43-C44	178.7 (2)
C32-C33-C34-C35	-179.9(2)	C42-C43-C44-C45	178.9 (2)
C33-C34-C35-C35 <sup>i</sup>	179.8 (3)	C43-C44-C45-C45 <sup>ii</sup>	179.4 (3)

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

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

## Cg2 is the centroid of the C21–C26 ring.

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C12-H12\cdots Cg2^{iii}$ $C15-H15\cdots Cg2^{iv}$	0.93 0.93	2.88 2.91	3.621 (3) 3.638 (3)	137 136
		(1)		

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

H atoms were positioned geometrically (C–H distances of 0.93–0.97 Å) and were included in the refinement in the riding-model approximation, with  $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$ . A final verification of possible voids was performed using the VOID routine of the *PLATON* program (Spek, 2003).

Data collection: *CAD-4 Software* (Enraf–Nonius 1989); cell refinement: *CAD-4 Software*; data reduction: local program; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *UdMX* (Maris, 2004).

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