

## 1,2-Bis(diphenylthioarsinoyl)ethane

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## Key indicators

Single-crystal X-ray study

T = 293 K

Mean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$ 

R factor = 0.043

wR factor = 0.097

Data-to-parameter ratio = 24.1

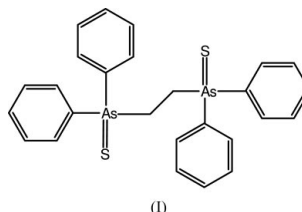
For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.The structure of the title compound,  $[\text{As}_2\text{S}_2(\text{C}_2\text{H}_4)(\text{C}_6\text{H}_5)_4]$ , which has twofold symmetry, features an  $\text{As}=\text{S}$  bond distance of 2.0674 (13)  $\text{Å}$ .

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## Comment

The title compound, (I), was prepared for use as a ligand in novel nickel complexes (Smith, 2002) as part of a wider study to prepare synthetic compounds with features similar to those of the active sites of the nickel-containing enzymes: hydrogenase, carbon monoxide dehydrogenase and acetyl-CoA synthase (Smith *et al.*, 2003; Evans & Pickett, 2003).

The structure of (I) (Fig. 1 and Table 1) lies about a twofold rotation axis which bisects the ethane bond. The As atom is tetrahedrally coordinated, with  $\text{S}-\text{As}-\text{C}$  angles lying in the range  $111.51(13)$ – $114.04(12)^\circ$  and  $\text{C}-\text{As}-\text{C}$  angles lying in the range  $105.60(17)$ – $106.92(15)^\circ$ . Bond lengths within the molecule are as expected, with  $\text{As}-\text{C}$  lengths lying in the range  $1.924(4)$ – $1.946(3) \text{ \AA}$  and  $\text{As}-\text{S}$  being  $2.0674(13) \text{ \AA}$ . The torsion angle for the ethane bridge [ $\text{As}-\text{C}-\text{C}^i-\text{As}^i$ ; symmetry code (i)  $1-x, y, \frac{1}{2}-z$ ] is  $156.4(2)^\circ$ .

The molecules, separated by normal van der Waals contacts, are arranged so that circular channels run parallel to the crystallographic  $a$  axis (bounded by four molecules) and rectangular channels run parallel to the  $c$  axis (bounded by eight molecules), as highlighted in the two views of Fig. 2.

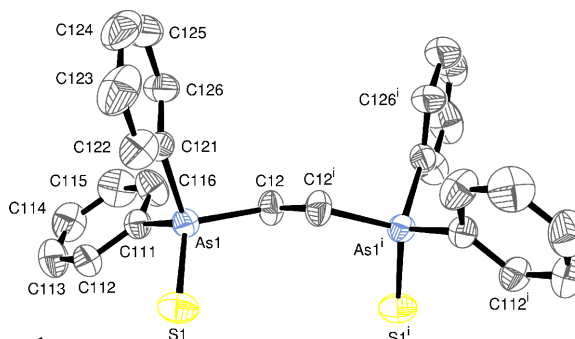


Figure 1

A view of (I). Displacement ellipsoids are drawn at the 50% probability level. Symmetry code (i)  $1-x, y, \frac{1}{2}-z$

## Experimental

Under an N<sub>2</sub> atmosphere, solid elemental S (0.153 g, 4.77 mmol) was added to a slurry of [(Ph)<sub>2</sub>AsCH<sub>2</sub>CH<sub>2</sub>As(Ph)<sub>2</sub>] (1.16 g, 2.39 mmol; Aldrich) in ethanol (50 ml). The mixture was refluxed for 5 h, giving a light-coloured orange–brown solution. Upon cooling and standing overnight, large colourless needles formed that were collected by filtration and dried *in vacuo* (0.21 g, 16%). Expected for C<sub>26</sub>H<sub>24</sub>As<sub>2</sub>S<sub>2</sub>: C 56.7, H 4.4, S 11.6%; found: C 56.8, H 4.3, S 12.8%.

### Crystal data

[As <sub>2</sub> S <sub>2</sub> (C <sub>2</sub> H <sub>4</sub> )(C <sub>6</sub> H <sub>5</sub> ) <sub>4</sub> ]	$D_x = 1.482 \text{ Mg m}^{-3}$
$M_r = 550.43$	Mo–K $\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 25 reflections
$a = 15.976(3) \text{ \AA}$	$\theta = 10\text{--}11^\circ$
$b = 9.168(4) \text{ \AA}$	$\mu = 2.89 \text{ mm}^{-1}$
$c = 17.635(3) \text{ \AA}$	$T = 293(2) \text{ K}$
$\beta = 107.213(13)^\circ$	Needle, colourless
$V = 2467.3(13) \text{ \AA}^3$	$0.52 \times 0.12 \times 0.06 \text{ mm}$
$Z = 4$	

### Data collection

Enraf–Nonius CAD-4 diffractometer	$R_{\text{int}} = 0.021$
$\omega/\theta$ scans	$\theta_{\text{max}} = 30.0^\circ$
Absorption correction: $\psi$ scan (EMPABS; Sheldrick <i>et al.</i> , 1977)	$h = -22 \rightarrow 21$
$T_{\text{min}} = 0.713$ , $T_{\text{max}} = 0.841$	$k = -1 \rightarrow 12$
3937 measured reflections	$l = -1 \rightarrow 24$
3573 independent reflections	3 standard reflections
1815 reflections with $I > 2\sigma(I)$	frequency: 167 min
	intensity decay: 13.2%

### Refinement

Refinement on $F^2$	Only H-atom $U$ 's refined
$R[F^2 > 2\sigma(F^2)] = 0.043$	$w = 1/[\sigma^2(F_o^2) + (0.0218P)^2]$
$wR(F^2) = 0.097$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.01$	$(\Delta/\sigma)_{\text{max}} < 0.001$
3573 reflections	$\Delta\rho_{\text{max}} = 0.53 \text{ e \AA}^{-3}$
148 parameters	$\Delta\rho_{\text{min}} = -0.49 \text{ e \AA}^{-3}$

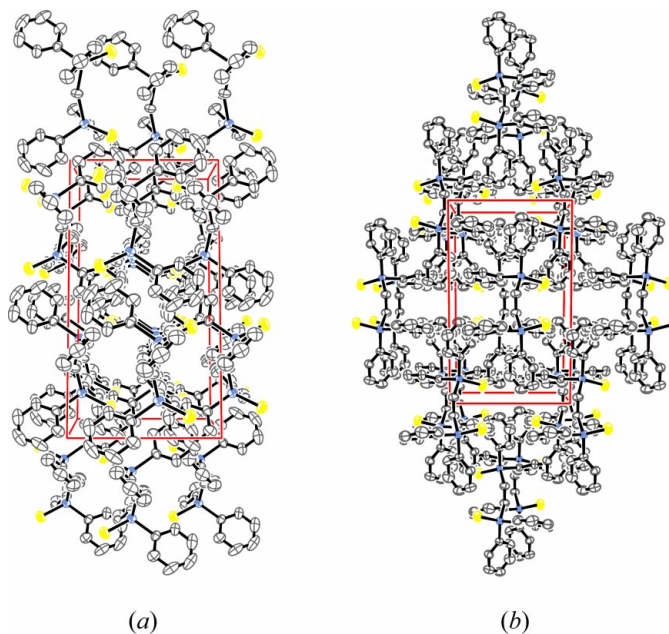
**Table 1**

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

As1–S1	2.0674 (13)	As1–C121	1.924 (4)
As1–C12	1.946 (3)	C12–C12 <sup>i</sup>	1.511 (7)
As1–C111	1.928 (3)		
C12–As1–S1	111.51 (13)	C121–As1–C12	105.60 (17)
C111–As1–S1	114.04 (12)	C121–As1–C111	106.92 (15)
C121–As1–S1	112.53 (12)	C12 <sup>i</sup> –C12–As1	110.4 (3)
C111–As1–C12	105.64 (14)		
As1–C12–C12 <sup>i</sup> –As1 <sup>i</sup>	156.4 (2)		

Symmetry code: (i)  $1 - x, y, \frac{1}{2} - z$ .

All H atoms were positioned geometrically and allowed to ride on the parent atoms, with C–H distances of 0.93  $\text{\AA}$  for phenyl H atoms and 0.97  $\text{\AA}$  for methyl H atoms; isotropic displacement parameters were refined freely.



**Figure 2**

Packing diagrams for (I) showing (a) a view in the direction of the crystallographic [100] vector and (b) a view in the direction of the crystallographic [001] vector.

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1992); cell refinement: *CAD-4 EXPRESS*; data reduction: *CAD-4* (Hursthouse, 1976) and *BAYES* (French & Wilson, 1978); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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