# organic papers

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#### Key indicators

Single-crystal X-ray study T = 295 KMean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ Å}$  R factor = 0.040 wR factor = 0.104 Data-to-parameter ratio = 10.2

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

# 1,4-Di-9-anthrylbutane

In the title compound,  $C_{32}H_{26}$ , the molecule has an inversion centre at the mid-point of the central C–C bond. Weak intermolecular C–H··· $\pi$  interactions help to stabilize the crystal structure.

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## Comment

Bisanthrylalkanes are extensively used in studies aimed at gaining information on the photophysical properties and electron donor-acceptor complexations of the related photoconductive polymers (Masnovi *et al.*, 1985; Becker & Andersson, 1987; Becker *et al.*, 1992; Rettig *et al.*, 1999). The spacing and orientation of the chromophore groups are determining factors in the photophysical and photochemical features of these dimers. For the complexation studies of a series of bis-9-anthrylalkanes with a number of electron acceptors, we have prepared the title compound, (I), and carried out a single-crystal X-ray analysis to establish its structure.



Compound (I) (Fig. 1) crystallizes in the monoclinic space group  $P2_1/n$ . The molecule has an inversion centre at the midpoint of the central C-C bond. Bond lengths and angles in the anthracene unit are in agreement with those in related compounds (Becker *et al.*, 1984; Becker *et al.*, 1992). The 14 atoms of the anthracene skeleton are coplanar to within 0.019 (2) Å. The methylene chain connecting the two ring systems adopts an *anti-anti-anti* conformation. The two anthracene ring systems are parallel to each other.

The crystal packing of (I) is mainly determined by weak intermolecular C-H··· $\pi$  interactions (Table 1), together with van der Waals forces. No intermolecular  $\pi$ - $\pi$  interactions are observed.

#### **Experimental**

© 2007 International Union of Crystallography All rights reserved The title compound was prepared according to the literature procedure of Dunand *et al.* (1980) *via* the formation of a di-Grignard





The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 40% probability level. Unlabelled atoms are related to labelled atoms by the symmetry operator (-x, -y + 1, -z + 1).



#### Figure 2

A packing diagram for (I), viewed down the a axis. Dashed lines represent  $C-H\cdots\pi$  contacts.

reagent from 1,4-dibromobutane followed by its reaction with anthrone. The di-Grignard reagent was prepared by stirring magnesium (0.50 g, 20 mmol) (washed twice with acetone and dried) and 1,4-dibromobutane (1.0 ml, 8.2 mmol) in anhydrous diethyl ether (20 ml) with a small chunk of iodine. The mixture was stirred under a nitrogen atmosphere for 24 h under ambient conditions. A hot solution of anthrone (0.01 mol) in anhydrous benzene (40 ml) was then added. The reaction mixture was stirred and refluxed under a nitrogen atmosphere for 5 h, until the colour of the mixture turned green. The green mixture was cooled and decomposed with ice and dilute hydrochloric acid; the organic solvents were extracted and evaporated. The residue was washed eight times with hot 20% NaOH solution to remove unreacted anthrone. Column chromatography of the crude product on basic alumina using hexane-dichloromethane (9:1 v/v) as the eluting solvents gave 1.19 g (2.90 mmol, 29% yield) of the title compound as pale-yellow needles [m.p. 527-528 K; literature value 527 K (Dunand et al., 1980)]. Single crystals of (I) suitable for

X-ray diffraction analysis were grown from a chloroform-hexane (1:1 v/v) solvent mixture using the slow evaporation technique.

## Crystal data

CaeHar	V - 11
$M_{10}$ 52	7 - 11
$M_r = 410.55$	Z = Z
Monoclinic, $P2_1/n$	Mo Ka
a = 11.3964 (8) Å	$\mu = 0.0$
b = 7.9000 (10)  Å	T = 29
c = 12.7887 (6) Å	$0.5 \times 0$
$\beta = 94.747 \ (5)^{\circ}$	

#### Data collection

Enraf-Nonius CAD-4 diffractometer Absorption correction: none 2015 measured reflections 2015 independent reflections

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.040$  $wR(F^2) = 0.104$ S = 1.032015 reflections

47.44 (17) Å<sup>3</sup> x radiation 07 mm<sup>-</sup> 5 (2) K  $0.4 \times 0.3 \text{ mm}$ 

1396 reflections with  $I > 2\sigma(I)$ 3 standard reflections frequency: 120 min intensity decay: 0.1%

197 parameters All H-atom parameters refined  $\Delta \rho_{\rm max} = 0.13 \ {\rm e} \ {\rm \AA}^{-2}$  $\Delta \rho_{\rm min} = -0.1 \text{ e } \text{\AA}^{-3}$ 

## Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C4a/C9a/C9/C8a/C10a/C10 ring and Cg2 is the centroid of the C5/C6/C7/C8/C8a/C10a ring.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C5-H5\cdots Cg1^i$	0.98 (2)	2.57 (2)	3.496 (2)	160
$C11 - H11A \cdots Cg2^{ii}$	1.02(2)	2.85 (2)	3.563 (2)	128
$C11 - H11B \cdots Cg1^{ii}$	1.01 (2)	2.82 (2)	3.550 (2)	130

Symmetry codes: (i)  $-x + \frac{3}{2}$ ,  $y + \frac{1}{2}$ ,  $-z + \frac{1}{2}$ ; (ii) -x + 1, -y + 1, -z.

All H atoms were located in difference Fourier maps and refined freely. The range of refined C-H distances is 0.96 (2)-1.02 (2) Å and the range of  $U_{iso}(H)$  values is 0.054 (4)–0.097 (7) Å<sup>2</sup>.

Data collection: CAD-4-PC Software (Enraf-Nonius, 1993); cell refinement: CAD-4-PC Software; data reduction: DATRD2 in NRCVAX (Gabe et al., 1989); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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