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## Key indicators

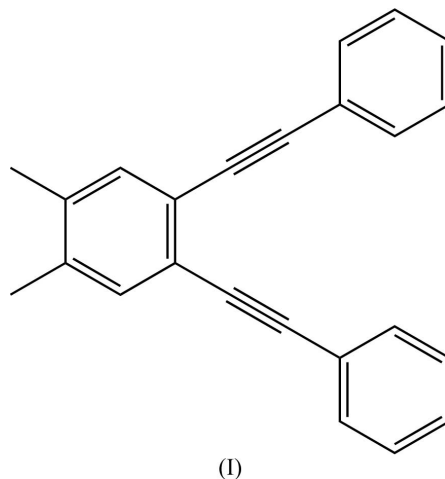
Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.058  
 $wR$  factor = 0.159  
Data-to-parameter ratio = 13.6For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

## 1,2-Dimethyl-4,5-bis(phenylethynyl)benzene

The title compound,  $\text{C}_{24}\text{H}_{18}$ , possesses a twofold rotation axis, which bisects the 1,2-dimethylbenzene ring. In the crystal structure, all three benzene rings are essentially coplanar, the angle between the central and terminal rings being  $3.8$  ( $1^\circ$ ).

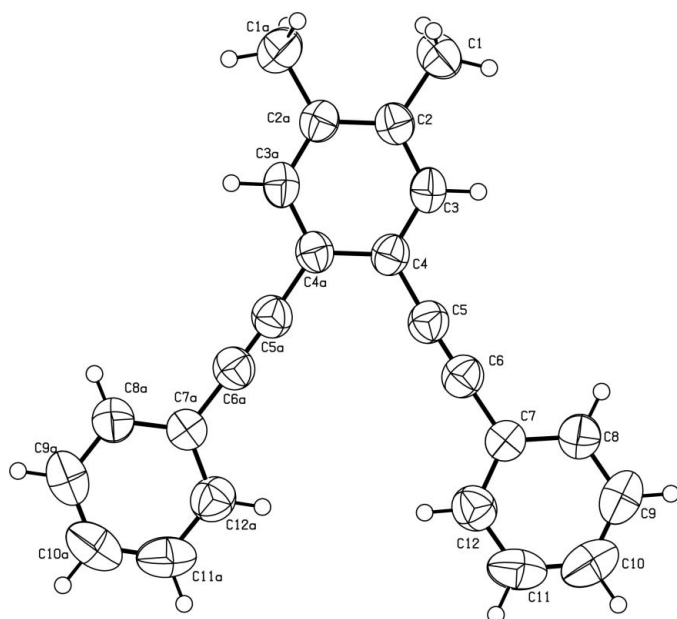
## Comment

A study of the thermal cycloaromatization of enediynes led to the suggestion of a benzene 1,4-biradical intermediate (Lockhart *et al.*, 1981). 1,2-Dimethyl-4,5-bis(phenylethynyl)benzene is an important 1,4-biradical intermediate for substituted polyphenylenes. The latter have been shown to exhibit excellent thermal and chemical resilience, interesting semiconducting properties upon doping, and applications in light-emitting diodes (Kovacic & Jones, 1987; Grem *et al.*, 1992). In this paper, we present the X-ray crystallographic analysis of 1,2-dimethyl-4,5-bis(phenylethynyl)benzene, (I). The crystal structure of the isomeric compound, 1,4-bis(*p*-tolylethynyl)benzene, has been reported by Filatov & Petrukhina (2005).

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The molecule of (I) possesses a twofold rotation axis, which passes through the mid-points of bonds  $\text{C}2-\text{C}2a$  and  $\text{C}4-\text{C}4a$  (Fig. 1). All three benzene rings are essentially coplanar, the angle between the central and terminal rings being  $3.8$  ( $1^\circ$ ). Selected bond distances and angles are given in Table 1. In the crystal structure of (I), molecules are aligned in a herring-bone fashion (Fig. 2).

It has been shown previously that methyl groups can function as hydrogen-bond donors towards aromatic  $\pi$  systems (Desiraju, 2002). Weak  $\text{C}-\text{H}\cdots\pi$  interactions are observed in the crystal structure (Fig. 3 and Table 2). Additionally, weak  $\pi-\pi$  stacking can be observed in the crystal structure.



**Figure 1**  
View of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary size. The suffix *a* in atom labels indicates the symmetry position  $(-x, y, \frac{3}{2} - z)$ .

## Experimental

The title compound was synthesized according to a literature procedure (Kabalka *et al.*, 2001). Crystals appropriate for data collection were obtained by slow evaporation of a dichloromethane solution at 283 K.

### Crystal data

$C_{24}H_{18}$   
 $M_r = 306.38$   
 Orthorhombic, *Pbcn*  
 $a = 11.735$  (2) Å  
 $b = 12.964$  (3) Å  
 $c = 11.820$  (2) Å  
 $V = 1798.3$  (6) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.132$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation  
 Cell parameters from 1966 reflections  
 $\theta = 1.0$ – $27.0^\circ$   
 $\mu = 0.06$  mm<sup>-1</sup>  
 $T = 292$  (2) K  
 Block, colourless  
 $0.30 \times 0.20 \times 0.10$  mm

### Data collection

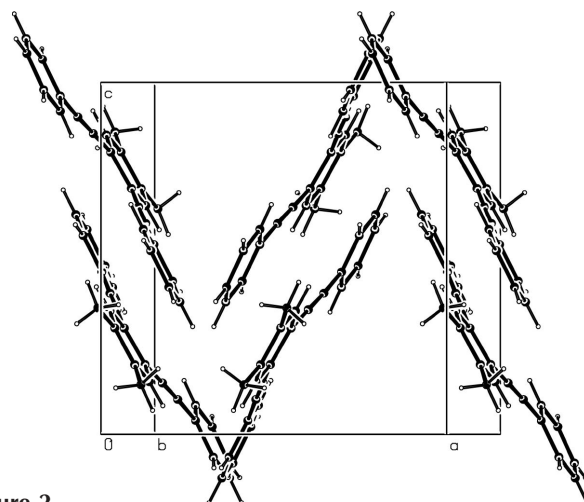
Bruker SMART 4K CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: none  
 9915 measured reflections  
 1966 independent reflections

1207 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.043$   
 $\theta_{max} = 27.0^\circ$   
 $h = -14 \rightarrow 14$   
 $k = -15 \rightarrow 16$   
 $l = -15 \rightarrow 8$

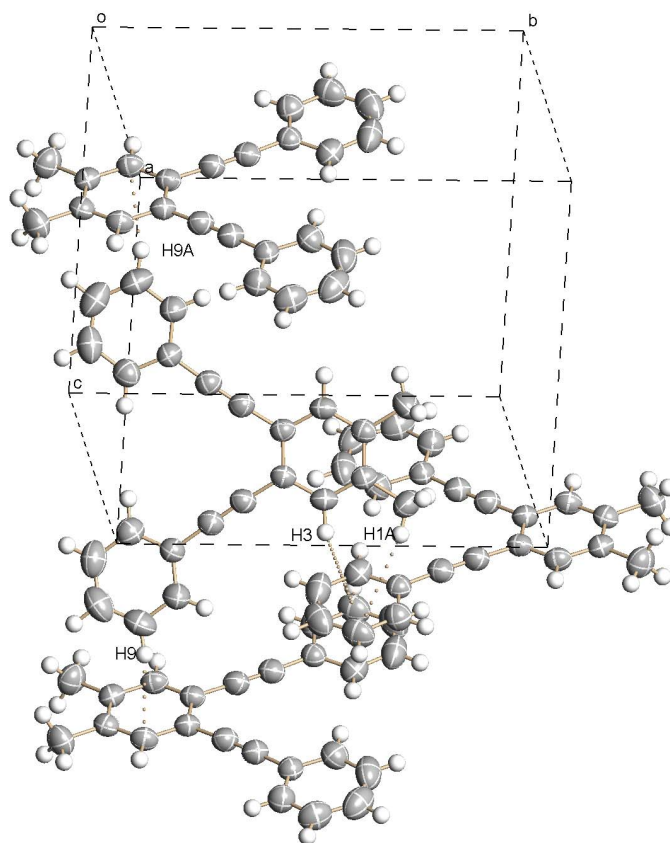
### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.058$   
 $wR(F^2) = 0.159$   
 $S = 1.03$   
 1966 reflections  
 145 parameters  
 All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0646P)^2 + 0.2012P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} < 0.001$   
 $\Delta\rho_{max} = 0.11$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.13$  e Å<sup>-3</sup>



**Figure 2**  
The packing of (I), viewed along the *b* axis.



**Figure 3**  
The C–H... $\pi$ (arene) interactions, indicated by dotted lines.

**Table 1**  
Selected geometric parameters (Å, °).

C1–C2	1.506 (3)	C6–C7	1.434 (3)
C5–C6	1.191 (2)		
C4 <sup>i</sup> –C4–C5	121.48 (10)	C5–C6–C7	177.8 (2)
C6–C5–C4	177.2 (2)	C8–C7–C6	120.32 (18)

Symmetry code: (i)  $-x, y, \frac{3}{2} - z$ .

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$C1-H1A \cdots Cg2^{ii}$	0.94 (3)	3.35 (2)	4.061 (3)	135 (2)
$C3-H3 \cdots Cg2^{ii}$	0.990 (19)	3.035 (18)	3.849 (2)	140 (1)
$C9-H9 \cdots Cg1^{iii}$	0.95 (3)	2.92 (2)	3.575 (3)	128 (2)
$C9-H9 \cdots Cg1^{iv}$	0.95 (3)	2.92 (2)	3.575 (3)	128 (2)

Symmetry codes: (ii)  $x, -y, z - \frac{1}{2}$ ; (iii)  $\frac{1}{2} - x, \frac{1}{2} - y, \frac{1}{2} + z$ ; (iv)  $-x, y, \frac{5}{2} - z$ .  $Cg1$  and  $Cg2$  are the centroids of rings C2–C4/C4a–C2a and C7–C12, respectively.

All H atoms were freely refined; C–H = 0.91 (3)–0.99 (2) Å for aromatic H atoms and 0.94 (3)–1.00 (3) Å for methyl H atoms.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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