

## 2,7-Dimethylocta-2,3,5,6-tetraene

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### Key indicators

Single-crystal X-ray study  
 $T = 178\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$   
 $R$  factor = 0.041  
 $wR$  factor = 0.121  
Data-to-parameter ratio = 14.0

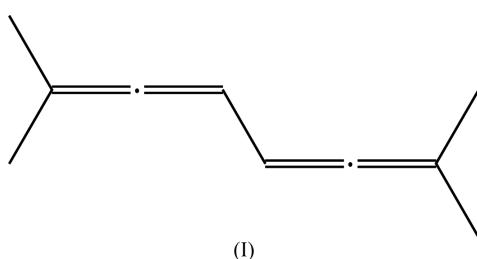
For details of how these key indicators were automatically derived from the article, see  
<http://journals.iucr.org/e>.

The molecule of the title compound,  $\text{C}_{10}\text{H}_{14}$ , displays inversion symmetry. The allenic bond lengths are 1.3067 (16) and 1.3126 (16)  $\text{\AA}$ .

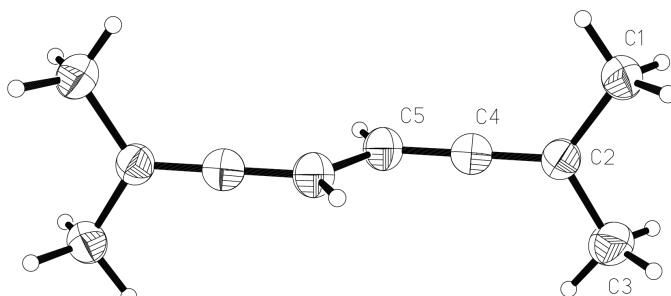
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### Comment

Conjugated bis-allenes are interesting starting materials in organic chemistry and have been used, in particular, as diene components in Diels–Alder addition (for a summary, see Hopf, 2000). In a study of the behaviour of bis-allenes in ionic reaction, we have prepared one of the oldest known bis-allenes, the title compound 2,7-dimethyl-octa-2,3,5,6-tetraene, (I) (Skattebøl & Solomon, 1965), and subjected it to structure determination.

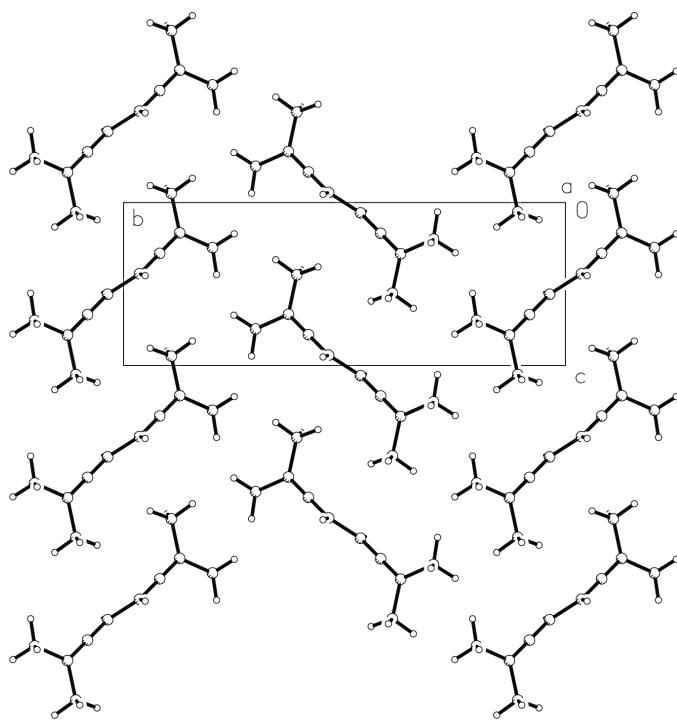


The molecule (Fig. 1) displays crystallographic inversion symmetry, because of which the standard IUPAC numbering was not used. Molecular dimensions may be regarded as normal [*cf. e.g.* the standard allenic  $\text{C}=\text{C}$  bond length of 1.307  $\text{\AA}$  (Allen *et al.*, 1987)]. The allenic angle  $\text{C}2-\text{C}4-\text{C}5$  is essentially linear at 179.89 (13) $^\circ$  and the angle  $\text{C}4-\text{C}5-\text{C}5^i$  124.04 (13) $^\circ$  somewhat wider than the standard  $sp^2$  value [symmetry code: (i)  $1-x, 1-y, -z$ ]. The shortest intermolecular contact is  $\text{H}1\text{A} \cdots \text{H}1\text{C}(-0.5+x, 0.5-y, -0.5+z)$ , 2.55 (2)  $\text{\AA}$ , and the molecules pack in the common herringbone pattern (Fig. 2).



**Figure 1**

The molecule of the title compound in the crystal. Ellipsoids represent 50% probability levels.

**Figure 2**

Packing diagram of the title compound, projected along the  $a$  axis. Radii are arbitrary.

## Experimental

Crystals were grown by sublimation.

### Crystal data

$C_{10}H_{14}$   
 $M_r = 134.21$   
 Monoclinic,  $P2_1/n$   
 $a = 4.644 (2) \text{ \AA}$   
 $b = 16.278 (6) \text{ \AA}$   
 $c = 6.037 (2) \text{ \AA}$   
 $\beta = 96.12 (3)^\circ$   
 $V = 453.8 (3) \text{ \AA}^3$   
 $Z = 2$

### Data collection

Nicolet R3 diffractometer  
 $\omega$  scans  
 Absorption correction: none  
 2589 measured reflections  
 1034 independent reflections  
 810 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.020$

$D_s = 0.982 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 Cell parameters from 50 reflections  
 $\theta = 10\text{--}12.5^\circ$   
 $\mu = 0.06 \text{ mm}^{-1}$   
 $T = 178 (2) \text{ K}$   
 Prism, colourless  
 $0.70 \times 0.25 \times 0.25 \text{ mm}$

$\theta_{\text{max}} = 27.5^\circ$   
 $h = -5 \rightarrow 6$   
 $k = -21 \rightarrow 21$   
 $l = -7 \rightarrow 7$   
 3 standard reflections  
 every 147 reflections  
 intensity decay: none

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.121$   
 $S = 1.06$   
 1034 reflections  
 74 parameters  
 All H-atom parameters refined

$$w = 1/[\sigma^2(F_o^2) + (0.0677P)^2 + 0.0479P]$$

where  $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\text{max}} < 0.001$ 
 $\Delta\rho_{\text{max}} = 0.14 \text{ e \AA}^{-3}$ 
 $\Delta\rho_{\text{min}} = -0.16 \text{ e \AA}^{-3}$

**Table 1**  
 Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

C1—C2	1.5076 (17)	C4—C5	1.3126 (16)
C2—C4	1.3067 (16)	C5—C5 <sup>i</sup>	1.470 (2)
C2—C3	1.5067 (17)		
C4—C2—C3	122.22 (11)	C2—C4—C5	179.89 (13)
C4—C2—C1	121.93 (10)	C4—C5—C5 <sup>i</sup>	124.04 (13)
C3—C2—C1	115.85 (10)		

Symmetry code: (i)  $1 - x, 1 - y, -z$ .

A rigid-body libration correction (Schomaker & Trueblood, 1968) was performed successfully ( $R_{\text{lib}} = 0.078$ ) and gave the following corrected bond lengths ( $\text{\AA}$ ): C1—C2 1.518, C2—C3 1.518, C5—C5<sup>i</sup> 1.477  $\text{\AA}$ . The corrections to the other bonds, C2—C4 and C4—C5, were calculated as zero, which is consistent with the expected libration pattern of the molecule.

Data collection:  $P3$  (Nicolet, 1987); cell refinement:  $P3$ ; data reduction:  $XDISK$  (Nicolet, 1987); 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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