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#### **Key indicators**

Single-crystal X-ray study T = 110 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.033 wR factor = 0.093 Data-to-parameter ratio = 8.8

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

# 1,3-Dipropionyl-2,4,6-trimethylbenzene

In the molecule of the title compound,  $C_{15}H_{20}O_2$ , the two propionyl groups are in unusual positions with the two carbonyl groups in *syn* orientation relative to each other and bent at angles of 94.3 (3) and 93.6 (2)° away from the central methyl group. Received 5 December 2006 Accepted 8 December 2006

## Comment

In a previous study, involving orientation of carbonyl groups by adjacent methyls (Pinkus et al., 2006), dipropiodurene [systematic name: 1,1'-(2,3,5,6-tetramethyl-p-phenylene)dipropan-1-one] was found to have a trans orientation relative to the central ring with the planes of the carbonyl group and attached methylene C atom at a dihedral angle of 86.50 (11)° relative to the central ring. In a related crystallographic study of 1,4-bis(4-chlorobenzoyl)-2,3,5,6-tetramethylbenzene by Ferguson et al. (1993), it was reported that the two p-chlorobenzovl groups were also in a trans orientation, with a corresponding angle of 85.9 (1)°. Casarini & Lunazzi (1996) separated orthogonal syn/anti isomers of 1,4-bis(2,2-dimethyl propanoyl)durene by preparative TLC. In a more recent study Coluccini et al. (2003) observed more complicated rotamers in both syn and anti forms of 1,4-bis(mesitoyl)-2,3,5,6-tetramethylbenzene in low temperature NMR spectra. It was of interest to examine the related angles in the molecule of the title compound, (I), for which syn/anti orientations of keto groups are also possible.



In the molecule of (I) (Fig. 1) the bond lengths and angles (Table 1) are within expected ranges. The two propionyl groups are in unusual positions with the two carbonyl groups in *syn* orientation relative to each other and bent at angles of 94.3 (3) and 93.6 (2)° away from the central C7 methyl group.

#### **Experimental**

Dipropionylmesitylene was prepared (yield 79%; m.p. 373–374 K, literature 374–375 K) according to a previously reported procedure (Weil, 1897). Crystals suitable for X-ray analysis were obtained from absolute ethanol.

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#### Crystal data

 $C_{15}H_{20}O_2$   $M_r = 232.31$ Monoclinic, *Cc a* = 7.9058 (10) Å *b* = 21.191 (3) Å *c* = 8.8305 (13) Å *β* = 111.989 (4)° *V* = 1371.8 (3) Å<sup>3</sup>

#### Data collection

Bruker X8 APEX area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\min} = 0.977, T_{\max} = 0.985$

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0613P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.033$	+ 0.257P]
$wR(F^2) = 0.093$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} < 0.001$
1399 reflections	$\Delta \rho_{\rm max} = 0.20 \text{ e} \text{ Å}^{-3}$
159 parameters	$\Delta \rho_{\rm min} = -0.16 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	

Z = 4

 $D_x = 1.125 \text{ Mg m}^{-3}$ 

Mo  $K\alpha$  radiation

 $\mu = 0.07 \text{ mm}^{-1}$ 

T = 110 (2) K

 $R_{\rm int} = 0.030$  $\theta_{\rm max} = 26.4^{\circ}$ 

Block, colorless

 $0.32 \times 0.25 \times 0.21 \text{ mm}$ 

6948 measured reflections 1399 independent reflections 1336 reflections with  $I > 2\sigma(I)$ 

### Table 1

Selected geometric parameters (Å, °).

O1-C10	1.210 (2)	C6-C13	1.503 (3)
O2-C13	1.207 (2)	C10-C11	1.502 (3)
C2-C10	1.505 (3)	C14-C13	1.504 (3)
O1-C10-C11	121.89 (18)	O2-C13-C6	121.42 (17)
O1-C10-C2	120.94 (18)	O2-C13-C14	122.52 (18)
C11-C10-C2	117.16 (16)	C6-C13-C14	116.06 (16)

H atoms were positioned geometrically, with C-H = 0.95, 0.99 and 0.98 Å for aromatic, methylene and methyl H atoms, respectively, and constrained to ride on their parent atoms, with  $U_{iso}(H) = xU_{eq}(C)$ , where x = 1.5 for methyl and x = 1.2 for all other H atoms. In the absence of significant anomalous scattering effects, Friedel pairs were merged.

Data collection: APEX2 (Bruker, 2003); cell refinement: APEX2; data reduction: SAINT-Plus (Bruker, 2003); program(s) used to solve



#### Figure 1

A view of the molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

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