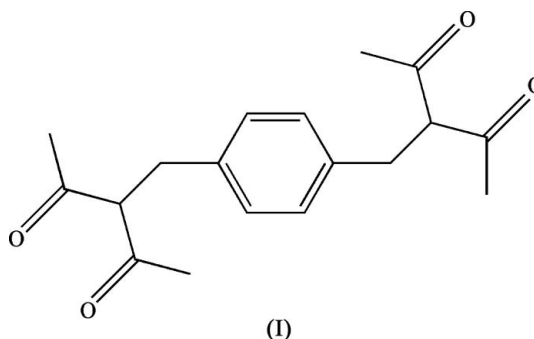
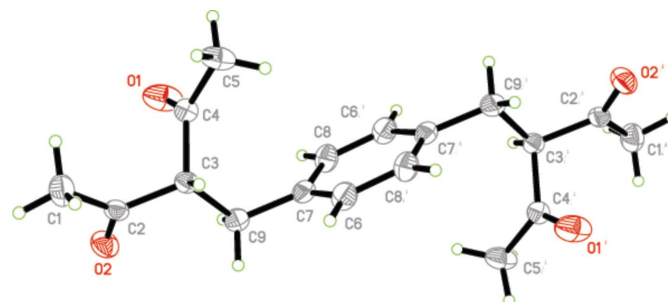


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shijingmin@beelink.com**Key indicators**Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
 R factor = 0.065
 wR factor = 0.157
Data-to-parameter ratio = 16.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**3,3'-(*p*-Phenylenedimethylene)bis(pentane-
2,4-dione)**The title molecule, $\text{C}_{18}\text{H}_{22}\text{O}_4$, lies on a crystallographic inversion center which is located at the center of the benzene ring. In the crystal structure, molecules are linked by weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming a two-dimensional network perpendicular to the a axis.

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CommentThe title compound, (I), was originally synthesized by Martin *et al.* (1959), but its crystal structure has not hitherto been reported. The presence of two β -diketone groups suggests that (I) may be a useful bridging ligand. The molecular structure of (I) is shown in Fig. 1. The molecule lies on a crystallographic inversion center located at the center of the benzene ring.In the crystal structure, weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds link molecules into a two-dimensional network perpendicular to the a axis (Table 1 and Fig. 2)**Experimental**The title compound was synthesized according to Martin *et al.* (1959). To a solution of potassium (3.9 g, 0.1 mol) in 2-methyl-2-propanol (100 ml), acetylacetone (10.0 g, 0.1 mol) was added dropwise and the**Figure 1**The molecular structure of (I), showing the atom-numbering scheme, with displacement ellipsoids drawn at the 30% probability level. [Symmetry code: (i) $-x, -y, -z + 1$.]

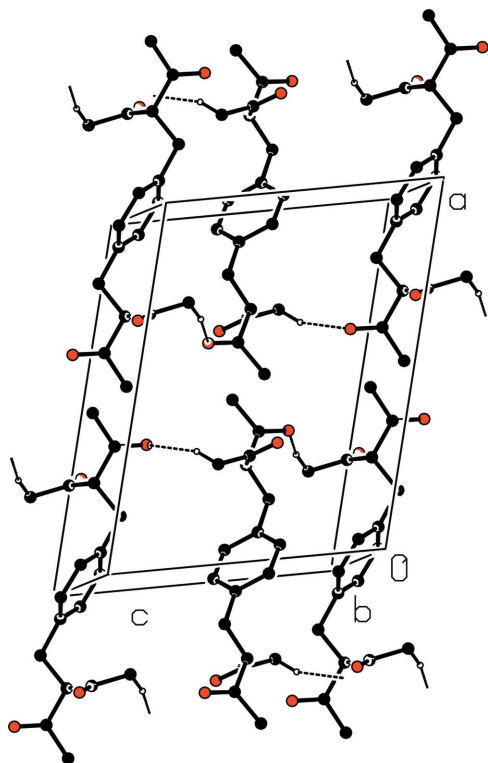


Figure 2
A packing plot (Spek, 2003), showing hydrogen bonds as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

mixture was refluxed for 30 min with stirring. 1,4-Dibromobenzene (13.2 g, 0.05 mol) was then added over a period of 40 min and KI (2.0 g, 0.012 mol) was then added. The mixture was stirred and refluxed for 4 h, and then 75% of the solvent was distilled and the solid residue was washed thoroughly with H₂O and dried. Single crystals were obtained by recrystallization from H₂O/acetonitrile in a 2:1 ratio. Elemental analysis found: C 71.72, H 7.15%; calculated for C₁₈H₂₂O₄: C 71.50, H 7.33%. The IR spectrum reveals a strong peak at 1724 cm⁻¹ for the vibration of carbonyl groups.

Crystal data

C₁₈H₂₂O₄
M_r = 302.36
 Monoclinic, *P*2₁/*c*
a = 13.959 (5) Å
b = 5.679 (2) Å
c = 10.860 (4) Å
 β = 107.386 (6)°
V = 821.6 (5) Å³

Z = 2
D_x = 1.222 Mg m⁻³
 Mo *K*α radiation
 μ = 0.09 mm⁻¹
T = 298 (2) K
 Prism, colorless
 0.30 × 0.21 × 0.12 mm

Data collection

Bruker SMART APEX CCD
 diffractometer
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
*T*_{min} = 0.975, *T*_{max} = 0.990

4365 measured reflections
 1669 independent reflections
 1189 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.038
 θ _{max} = 26.4°

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.065
wR(*F*²) = 0.157
S = 1.08
 1669 reflections
 102 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0555P)^2 + 0.239P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.22 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.13 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C5—H7C...O2 ⁱ	0.96	2.53	3.385 (4)	149

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

H atoms were placed in calculated positions and refined as riding, with benzene C—H = 0.93 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C), methyl C—H = 0.96 Å and *U*_{iso}(H) = 1.5*U*_{eq}(C), methylene C—H = 0.97 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C), methine C—H = 0.98 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C).

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

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