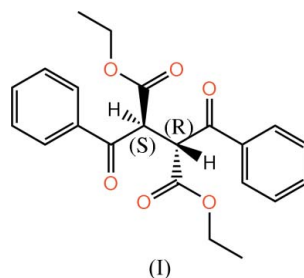


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Key indicators

Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
Disorder in main residue
 R factor = 0.068
 wR factor = 0.219
Data-to-parameter ratio = 13.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.(2*R*,3*S*)-Diethyl 2,3-dibenzoylsuccinateIn the crystal structure of the title compound, $\text{C}_{22}\text{H}_{22}\text{O}_6$, the asymmetric unit consists of one half-molecule with the other half generated by a centre of inversion. The crystal packing is stabilized by $\text{C}-\text{H}\cdots\text{O}$ intermolecular interactions.Received 14 March 2006
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Comment

1,4-Diketones are important intermediates for the synthesis of some natural products consisting of cyclopentanone and furan ring systems (McMurry & Meiton, 1971; Ito *et al.*, 1975, 1977). As part of a study of these compounds, we report the crystal structure of (2*R*,3*S*)-2,3-dibenzoylsuccinic acid diethyl ester, (I). The two aromatic rings are parallel (Fig. 1), the ethyl groups have a disordered methyl group and the crystal packing is stabilized by $\text{C}-\text{H}\cdots\text{O}$ intermolecular interactions as detailed in Table 1.

Experimental

The title compound was synthesized according to the literature procedure of Wu *et al.* (1997). Crystals appropriate for data collection were obtained by slow evaporation of a methanol–ethyl acetate (1:1 *v/v*) solution of (I).

Crystal data

$\text{C}_{22}\text{H}_{22}\text{O}_6$	$Z = 4$
$M_r = 382.40$	$D_x = 1.251$ Mg m ⁻³
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 11.328$ (4) Å	$\mu = 0.09$ mm ⁻¹
$b = 11.228$ (4) Å	$T = 292$ (2) K
$c = 15.986$ (6) Å	Block, colourless
$\beta = 92.868$ (6)°	$0.30 \times 0.20 \times 0.20$ mm
$V = 2030.7$ (12) Å ³	

Data collection

Bruker SMART CCD area-detector diffractometer	1785 independent reflections
φ and ω scans	1095 reflections with $I > 2\sigma(I)$
Absorption correction: none	$R_{\text{int}} = 0.067$
5177 measured reflections	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.068$ $wR(F^2) = 0.219$ $S = 1.06$

1785 reflections

133 parameters

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.116P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.28 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.17 \text{ e } \text{\AA}^{-3}$ **Table 1**Hydrogen-bond geometry (\AA , $^\circ$).

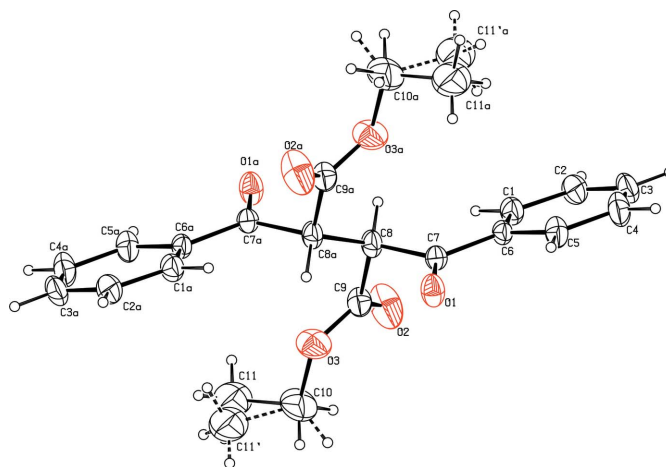
$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C3-H3\cdots O2^{ii}$	0.93	2.59	3.388 (4)	145

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

The methyl group was found to be disordered over two orientations. The occupancies of the disordered positions C11/C11' (and C11a/C11'a) refined to 0.540 (12)/0.460 (12). Suitable restraints were applied to the C–C distances involving the disordered atoms. The methyl H atoms were constrained to an ideal geometry with C–H distances of 0.96 \AA and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$, but each group was allowed to rotate freely about its C–C bond. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C–H distances of 0.93–0.98 \AA and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2003); software used to prepare material for publication: *SHELXTL*.

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**Figure 1**

View of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. Atoms labelled with the suffix a are related by $(-x + 1, -y + 2, -z + 1)$. Both disordered methyl components of the ethyl group are shown.

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