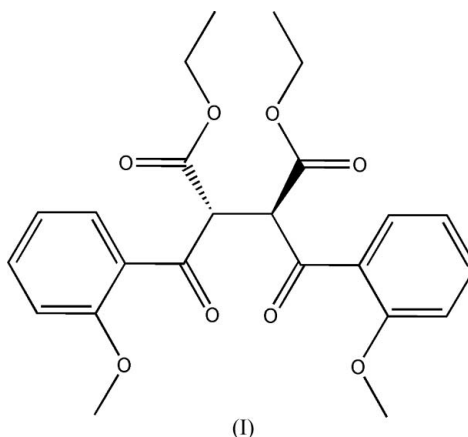
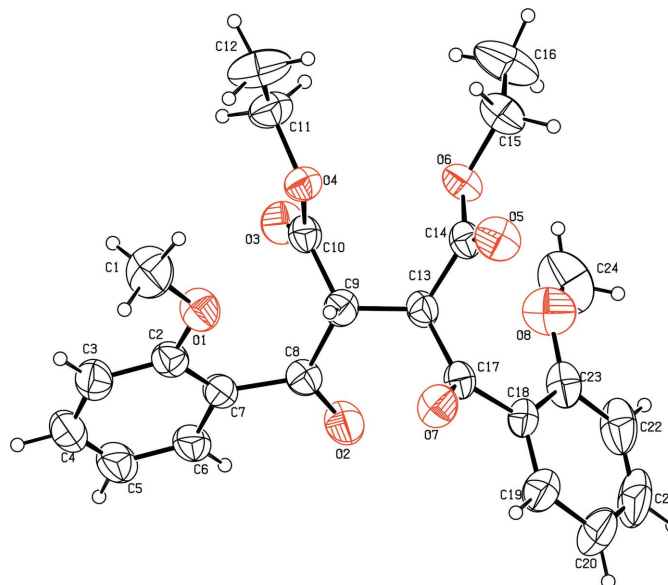
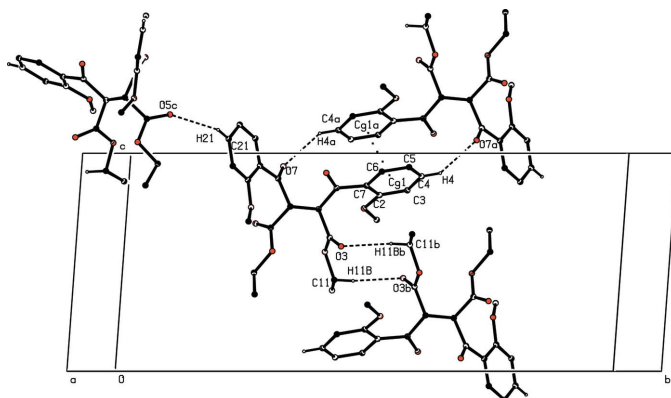


**rac-Diethyl 2,3-bis(2-methoxybenzoyl)succinate****Xiang-Gao Meng\*** and  
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of ChinaCorrespondence e-mail:  
mengxianggao@mail.ccnu.edu.cn,  
chwax@mail.ccnu.edu.cn**Key indicators**Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å  
 $R$  factor = 0.065  
 $wR$  factor = 0.177  
Data-to-parameter ratio = 13.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.In the *rac* isomer mixture of the title compound,  $\text{C}_{24}\text{H}_{26}\text{O}_8$ , the molecules are weakly linked *via*  $\text{C}-\text{H}\cdots\text{O}$  and  $\pi-\pi$  interactions, forming a three-dimensional network.Received 30 August 2005  
Accepted 6 September 2005  
Online 14 September 2005**Comment**We synthesized the title compound, a 2,3-disubstituted succinate derivative, (I), according to the literature method of Wu *et al.* (1998), and report its crystal structure here.

The molecular structure of (I) is illustrated in Fig. 1, and selected bond lengths and torsion angles are given in Table 1.

**Figure 1**  
The molecular structure of the title compound, showing 30% probability displacement ellipsoids.


**Figure 2**

Plot of the crystal packing, showing the linking of the molecules by C—H...O hydrogen bonds and a  $\pi$ – $\pi$  interaction (dashed lines). H atoms not involved in the hydrogen bonds have been omitted for clarity. [Symmetry codes: (ii)  $1 - x, 1 - y, 2 - z$ ; (iii)  $1 - x, 1 - y, 1 - z$ ; (iv)  $1 + x, \frac{1}{2} - y, \frac{1}{2} + z$ ]. Cg1 is the centroid of ring C2–C7. Labels having the suffix a indicate atoms at  $(-x, 1 - y, 2 - z)$ .

Chiral atoms C9 and C13 possess the same configuration. The central C8—C9—C13—C17 chain is in a *cis* configuration with a torsion angle of  $53.0(4)^\circ$ . Analysis with *PLATON* (Spek, 2003) shows that no classical hydrogen bonds exist in the molecular structure, although C—H...O hydrogen bonds and a  $\pi$ – $\pi$  interaction are present, forming a three-dimensional network in the crystal structure (Fig. 2 and Table 2). A  $\pi$ – $\pi$  interaction is observed between the benzene rings [C2–C7 and C2<sup>v</sup>–C7<sup>v</sup>; symmetry code: (v)  $-x, 1 - y, 2 - z$ ] with a centroid–centroid distance of  $3.696(2) \text{ \AA}$ .

## Experimental

Compound (I) was synthesized according to the literature procedure of Wu *et al.* (1998). Single crystals suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution at room temperature.

### Crystal data

C<sub>24</sub>H<sub>26</sub>O<sub>8</sub>  
*M<sub>r</sub>* = 442.45  
 Monoclinic, *P*2<sub>1</sub>/*c*  
*a* = 7.6510 (10) Å  
*b* = 28.146 (4) Å  
*c* = 10.9770 (14) Å  
 $\beta$  = 103.214 (2)°  
*V* = 2301.2 (5) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.277 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 4946 reflections  
 $\theta$  = 2.4–21.7°  
 $\mu$  = 0.10 mm<sup>-1</sup>  
*T* = 292 (2) K  
 Block, colorless  
 0.30 × 0.30 × 0.20 mm

### Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: none  
 16206 measured reflections  
 4040 independent reflections

2830 reflections with  $I > 2\sigma(I)$   
*R*<sub>int</sub> = 0.030  
 $\theta_{\text{max}}$  = 25.0°  
*h* = -8 → 9  
*k* = -33 → 33  
*l* = -12 → 13

### Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.065  
*wR*(*F*<sup>2</sup>) = 0.177  
*S* = 1.09  
 4040 reflections  
 293 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0669P)^2 + 1.0101P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.18 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

C7–C8	1.493 (4)	C13–C14	1.508 (4)
C8–C9	1.517 (4)	C13–C17	1.513 (4)
C9–C13	1.429 (4)	C17–C18	1.484 (4)
C9–C10	1.503 (4)		
C7–C8–C9–C13	-175.5 (3)	C8–C9–C13–C17	53.0 (4)
C7–C8–C9–C10	50.8 (4)	C9–C13–C17–C18	-136.3 (3)
C8–C9–C13–C14	-175.0 (3)	C13–C17–C18–C23	-14.3 (4)

**Table 2**

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>
C21–H21...O5 <sup>i</sup>	0.93	2.56	3.366 (5)	146
C4–H4...O7 <sup>ii</sup>	0.93	2.50	3.323 (4)	147
C11–H11B...O3 <sup>iii</sup>	0.97	2.49	3.448 (5)	169

Symmetry codes: (i)  $x + 1, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (ii)  $-x + 1, -y + 1, -z + 2$ ; (iii)  $-x + 1, -y + 1, -z + 1$ .

All H atoms were located at idealized positions (methyl C–H = 0.96 Å, methylene C–H = 0.97 Å, methine C–H = 0.98 Å and aromatic C–H = 0.93 Å) and included in the refinement using the riding-model approximation, with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$  for methyl H atoms and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for the other H atoms.

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: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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