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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.045
 wR factor = 0.119
Data-to-parameter ratio = 13.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

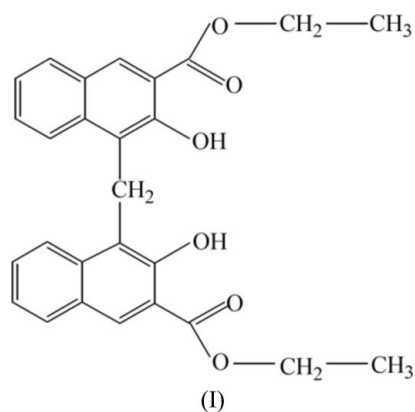
Ethyl pamoate

In the title compound, $\text{C}_{27}\text{H}_{24}\text{O}_6$, the dihedral angle between the two naphthalene ring systems is $84.79(6)^\circ$. The unit-cell packing features weak π - π interactions along the a axis between adjacent molecules.

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Comment

Organotin compounds have been widely used in homogeneous catalysis, and they have shown increased selectivity in a variety of chemical transformations (Edelman *et al.*, 1990; Orita *et al.*, 1999). In this work, the title compound, (I), was synthesized under the catalysis of an organotin compound. Although pamoic acid is a well known industrial precursor (Fei *et al.*, 2001), to the best of our knowledge, the crystal structure of its ethyl ester has never been reported.



The two naphthalene ring systems of (I) are almost perpendicular, with a dihedral angle of $84.79(6)^\circ$ (Fig. 1). The bond distances and bond angles are all normal. Two intramolecular $\text{O}-\text{H}\cdots\text{O}$ interactions (Table 1) are observed. The

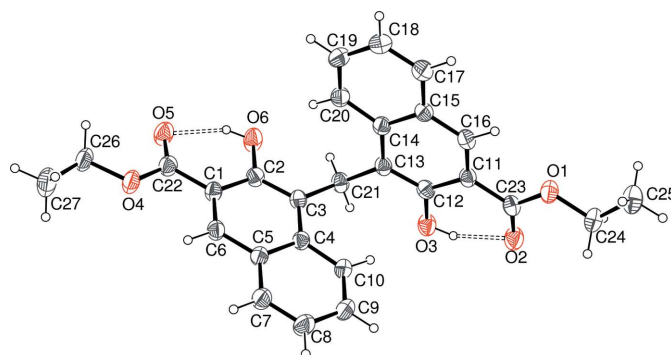


Figure 1
A view of (I), showing 30% displacement ellipsoids (arbitrary spheres for the H atoms). Hydrogen bonds are indicated by dashed lines.

naphthalene rings of adjacent molecules show centroid-to-centroid distances of 3.97 Å along the *a* axis (Fig. 2), indicating the presence of weak π - π interactions (Dong *et al.*, 2005).

Experimental

A mixture of pamoic acid (0.194 g, 0.5 mmol), NaOH (0.040 g, 1 mmol) and Bu₂SnCl₂ (0.152 g, 0.5 mmol) in ethanol (8 ml) was sealed in a 15 ml Teflon-lined stainless steel bomb and heated to 413 K for 3 days. Yellow crystals of (I) were obtained after slowly cooling to room temperature at the rate of 10 K h⁻¹.

Crystal data

C ₂₇ H ₂₄ O ₆	Z = 8
<i>M_r</i> = 444.46	<i>D_x</i> = 1.319 Mg m ⁻³
Monoclinic, <i>C2/c</i>	Mo <i>K</i> α radiation
<i>a</i> = 14.556 (5) Å	μ = 0.09 mm ⁻¹
<i>b</i> = 11.153 (5) Å	<i>T</i> = 293 (2) K
<i>c</i> = 27.617 (5) Å	Block, yellow
β = 93.514 (5)°	0.32 × 0.25 × 0.23 mm
<i>V</i> = 4475 (3) Å ³	

Data collection

Bruker APEX CCD diffractometer	10534 measured reflections
ω scans	4127 independent reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996)	2992 reflections with <i>I</i> > 2σ(<i>I</i>)
<i>T_{min}</i> = 0.941, <i>T_{max}</i> = 0.965	<i>R_{int}</i> = 0.021
	θ_{\max} = 25.6°

Refinement

Refinement on <i>F</i> ²	$w = 1/[\sigma^2(F_o^2) + (0.057P)^2 + 1.3961P]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.120$	(Δ/σ) _{max} < 0.001
<i>S</i> = 1.01	$\Delta\rho_{\max} = 0.17 \text{ e \AA}^{-3}$
4127 reflections	$\Delta\rho_{\min} = -0.18 \text{ e \AA}^{-3}$
298 parameters	
H-atom parameters constrained	

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>
O3—H3···O2	0.82	1.86	2.589 (2)	148
O6—H4···O5	0.82	1.86	2.597 (2)	148

All C-bound H atoms were positioned geometrically and refined as riding, with C—H = 0.93–0.97 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C) or 1.5*U*_{eq}(methyl C). The hydroxyl H atoms were located in difference maps, repositioned in idealized locations and refined as riding, with *U*_{iso}(H) = 1.5*U*_{eq}(O).

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve

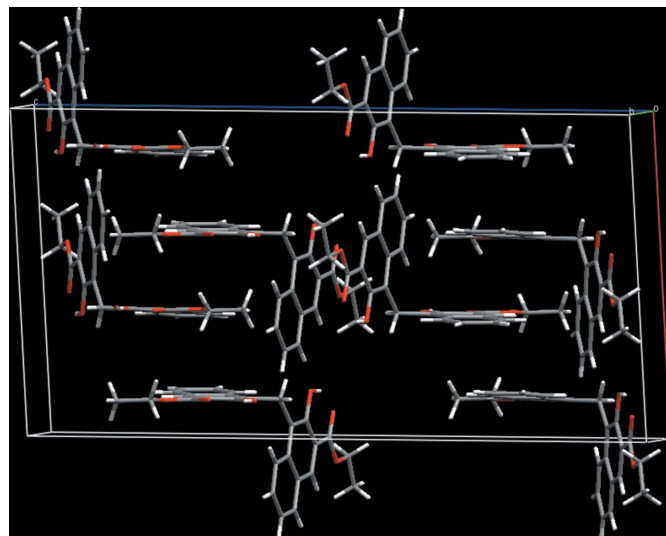


Figure 2
Packing diagram of (I).

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL-Plus* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL97*.

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