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Key indicators

Single-crystal X-ray study T = 122 KMean $\sigma(\text{C}-\text{C}) = 0.001 \text{ Å}$ R factor = 0.061 wR factor = 0.149 Data-to-parameter ratio = 48.6

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

Diethyl 4,7-dimethoxyindan-2,2-dicarboxylate

The title compound, $C_{17}H_{22}O_6$, crystallizes with two independent molecules in the asymmetric unit. The bond lengths and angles are generally within the normal ranges. The crystal packing is stabilized by weak intermolecular $C-H\cdots O$ hydrogen bonds and weak $C-H\cdots \pi$ (arene) interactions.

Comment

The title compound, (I), was prepared for use as a building block in the syntheses of single-walled carbon nanotube (SWNT) interacting compounds. It crystallizes in the monoclinic space group $P2_I/c$ with two independent molecules in the asymmetric unit (Fig. 1). The corresponding bond lengths and angles of the two molecules agree with each other, but the molecules differ in the orientation of an ester group (Table 1). In both molecules, the methoxy groups are almost coplanar with the benzene ring and the five-membered rings of the indane ring system adopt envelope conformations. The crystal packing of (I) is stabilized mainly by weak intermolecular C– $H \cdots O$ hydrogen bonds and weak C– $H \cdots \pi$ (arene) interactions (Table 2 and Fig. 2). In Table 2, Cg1 and Cg2 denote the centroids of rings C1A–C6A and C1B–C6B, respectively.



Experimental

Diethyl 5-bromo-4,7-dimethoxyindan-2,2-dicarboxylate (10 g, 24.9 mmol), prepared as described by Hammershøj & Christensen (2005), was dissolved in methanol (300 ml) and Pd/C 10% (cat. amount) was added. The mixture was hydrogenated (5.8 p.s.i.) at room temperature for 20 h. The reaction mixture was filtered and evaporated to dryness *in vacuo*, yielding an off-white solid. Purification by crystallization from ethanol yielded compound (I) as a white powder (yield 63%, m.p. 362–364 K). Single crystals were obtained by slow evaporation from ethanol ¹H NMR (300 MHz, CDCl₃): δ 6.7 (2H, *s*), 4.2 (4H, *q*, *J* = 7.3 Hz), 3.41 (4H, *s*), 1.25 (6H, *t*, *J* = 6.23 Hz); ¹³C NMR (100 MHz, CDCl₃): δ 171.5, 149.7, 129.5, 109.2, 61.5, 59.7, 55.5, 37.9, 13.9; *m/e*: 322 (62), 248 (83.5), 175 (100) and 161

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Figure 1

The asymmetric unit of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radii.

(40%); analysis calculated for C₁₇H₂₂O₆: C 63.3, H 6.8%; found: C 63.1, H 6.8%.

Crystal data

a	\mathbf{D} (accord -3
$C_{17}H_{22}O_6$	$D_x = 1.306 \text{ Mg m}^{-3}$
$M_r = 322.35$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 37980
a = 16.569 (1)Å	reflections
b = 8.033 (1) Å	$\theta = 1.240.0^{\circ}$
c = 24.643 (3) Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 90.232 \ (7)^{\circ}$	T = 122 (2) K
V = 3279.9 (6) Å ³	Prism, white
Z = 8	$0.4 \times 0.31 \times 0.25 \text{ mm}$

Data collection

Nonius KappaCCD area-detector	20169 independent reflections
diffractometer	12472 reflections with $I > 2\sigma(I)$
ω and φ scans	$R_{\rm int} = 0.079$
Absorption correction: Gaussian	$\theta_{\rm max} = 40.0^{\circ}$
integration (Coppens, 1970)	$h = -29 \rightarrow 29$
$T_{\min} = 0.977, \ T_{\max} = 0.988$	$k = -14 \rightarrow 14$
141916 measured reflections	$l = -43 \rightarrow 44$

Refinement

$w = 1/[\sigma^2(F_o^2) + (0.0458P)^2]$
+ 1.6487 <i>P</i>]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} = 0.001$
$\Delta \rho_{\rm max} = 0.65 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.32 \text{ e } \text{\AA}^{-3}$

Table 1

Selected torsion angles (°).

13.47 (14)
2.88 (15)
81.15 (11)
-1.98(15)
-12.14 (14)
-163.66 (9)





Table 2

 $I > 2\sigma(I)$

	H٧	vdrogen-bond	geometry	(Å, °).
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$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C7A - H7AB \cdots O3A$	0.99	2.48	2.8639 (13)	102
$C9A - H9AB \cdots O5A$	0.99	2.48	2.8501 (12)	102
$C11A - H11AC \cdot \cdot \cdot O3B^{i}$	0.98	2.55	3.3757 (15)	142
$C7B - H7BA \cdots O5B$	0.99	2.54	2.8778 (13)	100
$C10B - H10BA \cdots O5A$	0.98	2.55	3.4685 (15)	156
$C13A - H13AB \cdots Cg1^{ii}$	0.99	2.65	3.615 (1)	164
$C13B-H13BA\cdots Cg2^{iii}$	0.99	2.67	3.633 (1)	165

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

H atoms were placed in idealized positions and allowed to ride on their parent atoms, with C-H = 0.95-0.99 Å and $U_{iso}(H) = 1.2$ - $1.5U_{eq}(C).$

Data collection: COLLECT (Nonius, 1999); cell refinement: DIRAX (Duisenberg, 1992); data reduction: EVALCCD (Duisenberg et al., 2003); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett & Johnson, 1996); software used to prepare material for publication: SHELXL97.

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