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Key indicators

Single-crystal X-ray study T = 120 KMean σ (C–C) = 0.003 Å R factor = 0.048 wR factor = 0.125 Data-to-parameter ratio = 17.1

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

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Diethyl 2-(4-nitrophenyl)malonate forms a hydrogen-bonded chain of rings

Molecules of the title compound, $C_{13}H_{15}NO_6$, are linked by two C-H···O hydrogen bonds [H···O = 2.32 and 2.46 Å, C···O = 3.306 (2) and 3.393 (2) Å, and C-H···O = 171 and 156°] into a $C(4)C(7)[R_3^2(11)]$ chain of rings. Received 27 November 2003 Accepted 1 December 2003 Online 12 December 2003

Comment

Molecules of the title compound, (I) (Fig. 1), are linked by two C-H···O hydrogen bonds (Table 1) to form a chain of rings. The tertiary atom C7 lies between two carboxylate groups and thus carries one of the more acidic C-H bonds in the molecule; atom C7 in the molecule at (x, y, z) acts as hydrogenbond donor to carboxylate atom O92 in the molecule at (x, x) $\frac{3}{2} - y, \frac{1}{2} + z$), while C7 at $(x, \frac{3}{2} - y, \frac{1}{2} + z)$ in turn acts as donor to O92 at (x, y, 1+z). In this manner a C(4) chain is produced, running parallel to the [001] direction and generated by the *c*-glide plane at y = 0.75. This chain is reinforced by a second, weaker $C-H \cdots O$ hydrogen bond. Atom C81 in the molecule at (x, y, z) acts as hydrogen-bond donor, via H81B, to O92 in the molecule at (x, y, 1+z), so generating by translation a C(7) chain parallel to [001]. The combination of these two motifs thus generates a $C(4)C(7)[R_3^2(11)]$ chain of rings (Bernstein et al., 1995) running parallel to [001] (Fig. 2). There are two chains of this type, related to one another by inversion, running through each unit cell, but there are no direction-specific interactions between adjacent chains. In particular, there are no $C-H \cdots \pi$ (arene) hydrogen bonds nor any aromatic π - π stacking interactions in the structure.



It is noteworthy that a single atom, O92, acts as acceptor in both hydrogen bonds, whereas none of the other O atoms participates in the hydrogen bonding, and that both of the hydrogen-bond donors are aliphatic C atoms. Neither the aryl C-H bonds nor the nitro O atoms participate in any way in the supramolecular aggregation. The bond lengths and interbond angles in (I) show no unusual values.

Closely related to (I) is the diketone (II), which was found (Emsley *et al.*, 1989) to exist in the solid state solely as the enol (II*a*), stabilized by a short intramolecular $O-H\cdots O$ hydrogen bond. The intermolecular interactions were not





The molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.





Part of the crystal structure of (I), showing the formation of a chain of rings along [001]. For the sake of clarity, H atoms bonded to the C atoms which do not participate in the motifs shown have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions (x, x) $\frac{3}{2} - y, \frac{1}{2} + z$ and (x, y, 1 + z), respectively.

discussed in the original report, but scrutiny of the atomic coordinates retrieved from the Cambridge Structural Database (Allen, 2002), refcode SAPYEN, shows that the molecules are linked by a single C-H···O hydrogen bond into C(6) chains generated by the 2₁ screw axes in space group $P2_1/c$ (Fig. 3). We note here that the hydrogen-bond donor in (II) is one of the aryl C atoms, whereas in (I) the aryl C atoms play no role in the supramolecular aggregation. Likewise, the



Figure 3

Part of the crystal structure of (II), showing the formation of a C(6) chain along [010]; the atomic coordinates and the atom labels are as given in the original report (Emsley et al., 1989). For the sake of clarity, H atoms which do not participate in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions (1 - x, x) $y - \frac{1}{2}, \frac{1}{2} - z$ and $(1 - x, \frac{1}{2} + y, \frac{1}{2} - z)$, respectively.

hydrogen-bond acceptor in (II) is one of the nitro O atoms (Fig. 3), while the nitro group in (I) plays no role in the supramolecular aggregation.

Experimental

A sample of compound (I) was prepared by the CuBr-catalysed coupling reaction in dioxane solution between 4-nitrobromobenzene and the monosodium salt of diethylmalonate, Na[CH(COOEt)₂], following the general procedure described by Setsune et al. (1981). Crystals suitable for single-crystal X-ray diffraction were grown by slow evaporation of a solution in ethanol, and had a melting point of 330-332 K.

Crystal data

C ₁₃ H ₁₅ NO ₆	$D_x = 1.355 \text{ Mg m}^{-3}$
$M_r = 281.26$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 3150
a = 11.9275 (6) Å	reflections
$b = 13.8407 (8) \text{\AA}$	$\theta = 2.9-27.5^{\circ}$
c = 8.4595 (3) Å	$\mu = 0.11 \text{ mm}^{-1}$
$\beta = 99.192 \ (3)^{\circ}$	T = 120 (2) K
$V = 1378.60 (12) \text{ Å}^3$	Block, colourless
Z = 4	$0.35 \times 0.20 \times 0.08 \text{ mm}$

Data collection

Nonius KappaCCD diffractometer φ scans, and ω scans with κ offsets Absorption correction: multi-scan (SORTAV; Blessing, 1995, 1997) $T_{\min} = 0.967, \ T_{\max} = 0.991$ 14 747 measured reflections 3150 independent reflections

2007 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.067$ $\theta_{\rm max}=27.5^\circ$ $h = -12 \rightarrow 15$ $k = -17 \rightarrow 17$ $l = -10 \rightarrow 8$

Refinement

5	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0643P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.048$	+ 0.0884P]
$wR(F^2) = 0.125$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
3150 reflections	$\Delta \rho_{\rm max} = 0.30 \ {\rm e} \ {\rm \AA}^{-3}$
184 parameters	$\Delta \rho_{\rm min} = -0.32 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
-	Extinction coefficient: 0.023 (3)

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\overline{\begin{array}{c} C7 - H7 \cdots O92^{i} \\ C81 - H81B \cdots O92^{ii} \end{array}}$	1.00	2.32	3.306 (2)	171
	0.99	2.46	3.393 (2)	156

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

All H atoms were located in difference maps and subsequently treated as riding atoms, with C-H distances 0.95 (aromatic), 0.98 (CH₃), 0.99 (CH₂) or 1.00 Å (aliphatic CH), and with $U_{iso}(H) = 1.2U_{eq}(C)$ (CH and CH₂) or $1.5U_{eq}(C)$ (CH₃).

Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO–SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO–SMN*; program(s) used to solve structure: *OSCAIL* (McArdle, 2003) and *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *OSCAIL* and *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); soft-

ware used to prepare material for publication: *SHELXL*97 and *PRPKAPPA* (Ferguson, 1999).

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