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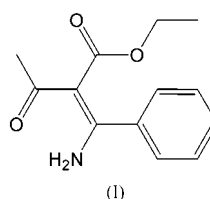
Key indicators

Single-crystal X-ray study
T = 294 K
Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$
R factor = 0.047
wR factor = 0.118
Data-to-parameter ratio = 18.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $\text{C}_{13}\text{H}_{15}\text{NO}_3$, is an *E* isomer and the phenyl ring does not conjugate with $\text{C}=\text{C}$. Both intra- and intermolecular $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds are found, and the infinite molecular chains stretch along the *b* axis.

Comment

The title compound, (I), is a by-product obtained in the synthesis of ethyl 3-acetamido-3-phenyl-propylenoate, a prochiral olefinic substrate for producing β -amino acids and derivatives by asymmetric hydrogenation (Hackler & Wickiser, 1985; Lubell *et al.*, 1991). The structure determination of (I) was conducted in order to obtain more stereochemical information about β -amino acids and their derivatives. In the structure of (I) (Fig. 1), the $\text{C}1-\text{C}2-\text{C}3-\text{C}8$ and $\text{C}1-\text{C}2-\text{C}3-\text{C}4$ torsion angles are $59.0(2)$ and $-124.07(18)^\circ$, respectively. This shows that the phenyl ring does not completely conjugate with the $\text{C}1=\text{C}2$ double bond in the solid state; the $\text{C}12-\text{C}1-\text{C}2-\text{N}1$, $\text{C}2-\text{C}1-\text{C}12-\text{O}3$ and $\text{C}2-\text{C}1-\text{C}12-\text{C}13$ torsion angles are $5.3(3)$, $-2.1(3)$ and $179.52(17)^\circ$, respectively. This illustrates that the atoms $\text{O}3-\text{C}12-\text{C}1-\text{C}2-\text{N}1-\text{H}$ are almost coplanar and the $\text{C}12=\text{O}3$ double bond and $\text{C}=\text{C}$ form a conjugated system (Table 1). As shown in Table 2 and the packing diagram (Fig. 2), the crystal structure of (I) is stabilized by both intra- and intermolecular hydrogen bonds, and infinite molecular chains stretch along the *b* axis.



Experimental

The title compound was synthesized according to Zhu *et al.* (1999). A crystal suitable for X-ray analysis was slowly grown in a mixed solvent of ethyl acetate and hexane at room temperature. ^1H NMR (400 MHz, acetone- d_6 , Bruker): δ 0.63–0.67 (*t*, $J = 7.1$ Hz, 3H), 2.24 (*s*, 3H), 3.64–3.69 (*q*, $J = 7.1$ Hz, 2H), 7.35–7.49 (*m*, 5H), 11.02 (*br*, 1H).

Crystal data

$\text{C}_{13}\text{H}_{15}\text{NO}_3$
 $M_r = 233.26$
Orthorhombic, *Pbca*
 $a = 17.158(3) \text{ \AA}$
 $b = 7.6070(12) \text{ \AA}$
 $c = 18.823(3) \text{ \AA}$
 $V = 2456.8(7) \text{ \AA}^3$
 $Z = 8$
 $D_x = 1.261 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation
Cell parameters from 4270 reflections
 $\theta = 1-27.5^\circ$
 $\mu = 0.09 \text{ mm}^{-1}$
 $T = 294(2) \text{ K}$
Block, colorless
 $0.50 \times 0.46 \times 0.30 \text{ mm}$

Data collection

Siemens SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.956$, $T_{\max} = 0.974$
 15490 measured reflections

2820 independent reflections
 1423 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.056$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -22 \rightarrow 22$
 $k = -9 \rightarrow 9$
 $l = -24 \rightarrow 21$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.118$
 $S = 1.02$
 2820 reflections
 157 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.05P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.16 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26 \text{ e } \text{\AA}^{-3}$
 Extinction correction: SHELXL97
 Extinction coefficient: 0.0249 (15)

Table 1

Selected geometric parameters (\AA , $^\circ$).

O1—C9	1.2098 (19)	C2—C3	1.490 (2)
O3—C12	1.244 (2)	C3—C4	1.380 (2)
C1—C2	1.401 (2)	C4—C5	1.381 (2)
C1—C12	1.451 (2)	C5—C6	1.365 (3)
C1—C9	1.466 (2)	C12—C13	1.504 (3)
C2—C1—C12	120.74 (15)	C3—C4—C5	120.11 (18)
C2—C1—C9	119.41 (14)	C6—C5—C4	120.61 (19)
C12—C1—C9	119.76 (14)	O1—C9—O2	121.87 (16)
N1—C2—C1	122.32 (14)	O1—C9—C1	126.39 (17)
C4—C3—C8	119.08 (15)	O3—C12—C1	121.76 (16)
C4—C3—C2	120.17 (15)		
C12—C1—C2—N1	5.3 (3)	C1—C2—C3—C8	59.0 (2)
C9—C1—C2—N1	-171.24 (16)	C5—C6—C7—C8	-0.3 (3)
C9—C1—C2—C3	11.4 (3)	C4—C3—C8—C7	-0.1 (3)
N1—C2—C3—C4	58.4 (2)	C2—C1—C12—O3	-2.1 (3)
C1—C2—C3—C4	-124.07 (18)	C2—C1—C12—C13	179.52 (17)

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O3	0.86	1.94	2.592 (2)	131
N1—H1B \cdots O1 ⁱ	0.86	2.23	3.048 (2)	158

Symmetry code: (i) $x, 1 + y, z$.

H atoms were positioned geometrically and refined in the riding-model approximation, with U_{iso} values equal to the U_{eq} value of the atom to which they are bound.

Data collection: SMART (Siemens, 1995); cell refinement: SMART; data reduction: SMART (Siemens, 1995) and SHELXTL (Siemens, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

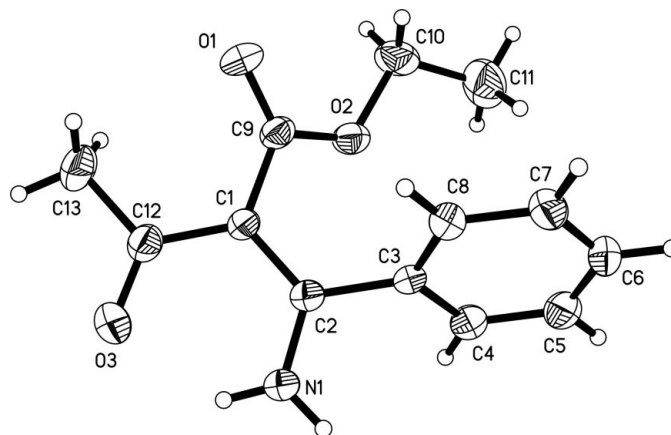


Figure 1

The molecular structure of (I), showing ellipsoids at the 50% probability level (Siemens, 1995).

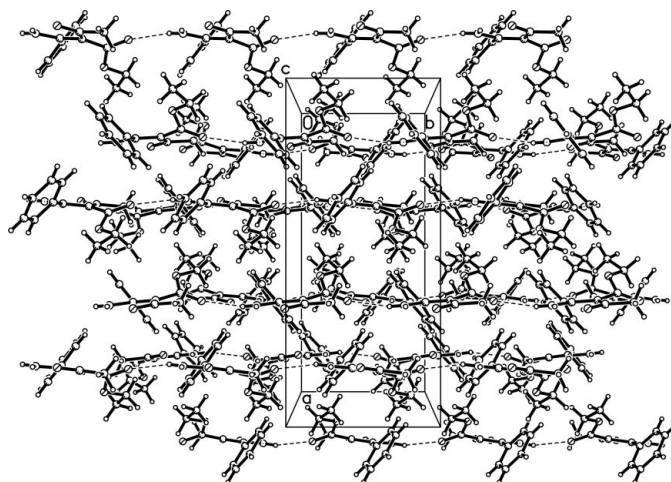


Figure 2

Packing diagram for (I). Hydrogen bonds are indicated by dashed lines.

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