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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.048$
$w R$ factor $=0.087$
Data-to-parameter ratio $=18.0$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Methyl (E)-2-acetyl-3-amino-2-pentenoate

The structure of the title compound, $\mathrm{C}_{8} \mathrm{H}_{13} \mathrm{NO}_{3}$, refined in space group $\mathrm{Pna2}_{1}$, which was chosen over Pnam as the space group because of the lack of mirror or inversion symmetry. An $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ intermolecular hydrogen bond is observed.

## Comment

The title compound, (I), is one of the by-products obtained during the synthesis of methyl 3-acetamido-2-pentenoate from the reaction of methyl 3-amino-2-pentenoate refluxed with acetic anhydride in THF for 24 h (Hackler \& Wickiser, 1985; Lubell et al., 1991; Yasutake et al., 2001). The structure determination of (I) was conducted in order to obtain more stereochemical information.


In the crystal structure of (I) (Fig. 1), the $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5$, $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5-\mathrm{O} 3$ and $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5-\mathrm{C} 6$ torsion angles are -2.0 (4), -1.1 (4) and 179.4 (3) ${ }^{\circ}$, respectively. This shows that atoms $\mathrm{N} 1 / \mathrm{C} 1 / \mathrm{C} 2 / \mathrm{C} 5 / \mathrm{O} 3 / \mathrm{C} 6$ are almost coplanar and the $\mathrm{C} 1=\mathrm{C} 2$ and $\mathrm{C} 5=\mathrm{O} 3$ double bonds form a conjugated system (Table 1). The molecules in the crystal structure are interconnected by $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding (Table 2). As shown in the packing diagram (Fig. 2), the hydrogen bonds links the molecules along the $c$ axis.

## Experimental

The title compound, (I), was synthesized according a literature procedure (Zhu et al., 1999). A crystal suitable for X-ray analysis was grown slowly from a mixed solution in ethyl acetate and hexane (1:5) at room temperature.

## Crystal data

| $\mathrm{C}_{8} \mathrm{H}_{13} \mathrm{NO}_{3}$ | Mo $K \alpha$ radiation |
| :--- | :--- |
| $M_{r}=171.19$ | Cell parameters from 1585 |
| Orthorhombic, Pna $_{1}$ | reflections |
| $a=23.998(7) \AA$ | $\theta=1-27.5^{\circ}$ |
| $b=4.2785(13) \AA$ | $\mu=0.10 \mathrm{~mm}^{-1}$ |
| $c=8.944(3) \AA$ | $T=294(2) \mathrm{K}$ |
| $V=918.3(5) \AA^{3}$ | Block, colorless |
| $Z=4$ | $0.50 \times 0.40 \times 0.38 \mathrm{~mm}$ |
| $D=1.238 \mathrm{Mg} \mathrm{m}^{-3}$ |  |

$D_{x}=1.238 \mathrm{Mg} \mathrm{m}^{-3}$

## Data collection

Siemens SMART CCD area-
detector diffractometer

## $\varphi$ and $\omega$ scans

Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\min }=0.954, T_{\max }=0.965$
5793 measured reflections

> Mo $K \alpha$ radiation Cell parameters from 1585 $\quad$ reflections $\theta=1-27.5^{\circ}$ $\mu=0.10 \mathrm{~mm}^{-1}$ $T=294(2) \mathrm{K}$ Block, colorless $0.50 \times 0.40 \times 0.38 \mathrm{~mm}$

2037 independent reflections
903 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.069$
$\theta_{\text {max }}=27.6^{\circ}$
$h=-27 \rightarrow 31$
$k=-5 \rightarrow 5$
$l=-11 \rightarrow 10$

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Figure 1
The molecular structure of (I), showing ellipsoids at the $50 \%$ probability level (Siemens, 1995).

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& (\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.15 \mathrm{e}^{-3} \AA^{-3} \\
& \Delta \rho_{\min }=-0.14 \mathrm{e} \AA^{-3} \\
& \text { Extinction correction: } \text { SHELXXL97 } \\
& \text { Extinction coefficient: } 0.0098(18) \\
& \text { Absolute structure: } \text { Flack (1983) } \\
& \text { Flack parameter }=-0.5(16)
\end{aligned}
$$

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.048$
$w R\left(F^{2}\right)=0.088$
$S=1.03$
2037 reflections
113 parameters
H -atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.015 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3$

Table 1
Selected geometric parameters ( $\mathrm{A},{ }^{\circ}$ ).

| $\mathrm{O} 1-\mathrm{C} 3$ | $1.196(4)$ | $\mathrm{C} 1-\mathrm{C} 2$ | $1.393(4)$ |
| :--- | :---: | :--- | ---: |
| $\mathrm{O} 2-\mathrm{C} 3$ | $1.339(3)$ | $\mathrm{C} 2-\mathrm{C} 5$ | $1.440(4)$ |
| $\mathrm{O} 3-\mathrm{C} 5$ | $1.247(3)$ | $\mathrm{C} 5-\mathrm{C} 6$ | $1.499(4)$ |
| $\mathrm{N} 1-\mathrm{C} 1$ | $1.311(3)$ | $\mathrm{C} 7-\mathrm{C} 8$ | $1.521(4)$ |
|  |  |  |  |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2$ | $122.5(3)$ | $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5$ | $121.4(3)$ |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 7$ | $114.4(3)$ | $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | $117.1(2)$ |
|  |  |  |  |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5$ | $-2.0(4)$ | $\mathrm{C} 7-\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | $-11.7(4)$ |
| $\mathrm{C} 7-\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5$ | $172.4(3)$ | $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5-\mathrm{O} 3$ | $-1.1(4)$ |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | $173.9(3)$ | $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 5-\mathrm{C} 6$ | $179.4(3)$ |

Table 2
Hydrogen-bonding geometry $\left(\AA{ }^{\circ}{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 A \cdots \mathrm{O} 3$ | 0.86 | 1.95 | $2.597(3)$ | 131 |
| $\mathrm{~N} 1-\mathrm{H} 1 B \cdots \mathrm{O} 3$ | 0.86 | 2.04 | $2.890(3)$ | 173 |

Symmetry code: (i) $\frac{1}{2}-x, \frac{1}{2}+y, z-\frac{1}{2}$.


Figure 2
Packing diagram for (I). The hydrogen bonds are indicated by dashed lines.

H atoms were included in the riding-model approximation, with $U_{\text {iso }}$ values equal to $U_{\text {eq }}$ of the atom to which they are bonded. The Flack (1983) parameter is indeterminate in the absence of significant anomalous scattering.

Data collection: SMART (Siemens, 1995); cell refinement: SMART; data reduction: SAINT 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.

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