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Key indicators

Single-crystal X-ray study

$T = 294\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$

R factor = 0.054

wR factor = 0.082

Data-to-parameter ratio = 19.9

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

N-[3-Acetamido-1,3-bis(4-ethylphenyl)butenyl]-acetamide

Both intra- and intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds are found in the structure of the title compound, $\text{C}_{24}\text{H}_{30}\text{N}_2\text{O}_2$. Molecules are assembled into chains, along the b axis, *via* intermolecular interactions.

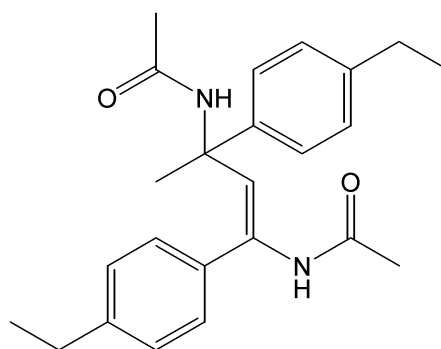
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Comment

The title compound, *N*-[3-acetamido-1,3-bis(4-ethylphenyl)butenyl]acetamide, (I), is one of the by-products obtained during the synthesis of *N*-[1-(4-ethylphenyl)ethenyl]acetamide (Burk *et al.*, 1998). In this study, the structure determination of (I) was conducted in order to obtain more knowledge about enamide compounds.



(I)

In the crystal structure (Fig. 1), the $\text{C13}-\text{C14}$ bond distance of $1.336(3)\text{ \AA}$ is indicative of double-bond character. The $\text{C13}-\text{C14}-\text{C17}-\text{C22}$, $\text{C15}-\text{N2}-\text{C14}-\text{C17}$ and $\text{C15}-\text{N2}-\text{C14}-\text{C13}$ torsion angles are $23.8(4)$, $63.2(3)$ and $-124.7(3)^\circ$, suggesting little conjugation in this residue, although the $\text{C3}-\text{C13}-\text{C14}-\text{C17}$ torsion angle is $177.0(2)^\circ$. Atom H2A of amine- N2 forms an intramolecular hydrogen bond with O1 (Table 1). Atom H1A of amine- N1 forms an intermolecular hydrogen bond with O2A of another molecule ($x, 1/2-y, z-1/2$). As illustrated in Fig. 2, these interactions link the molecules in chains along the b axis.

Experimental

The title compound was synthesized according to the literature method of Burk *et al.* (1998). A crystal suitable for X-ray analysis was slowly grown from a mixed solvent (ethyl acetate/hexane = 1:5) at room temperature. ^1H NMR (500 MHz, CDCl_3 , Varian): δ 1.18–1.22 (m , 6H), 1.35 (s , 3H), 1.63 (s , 3H), 2.09 (s , 3H), 2.59–2.63 (m , 4H), 5.37 (s , 1H), 6.02 (s , 1H), 7.12–7.20 (m , 6H), 7.35 (d , $J = 8.5\text{ Hz}$, 2H), 8.97 (s , 1H).

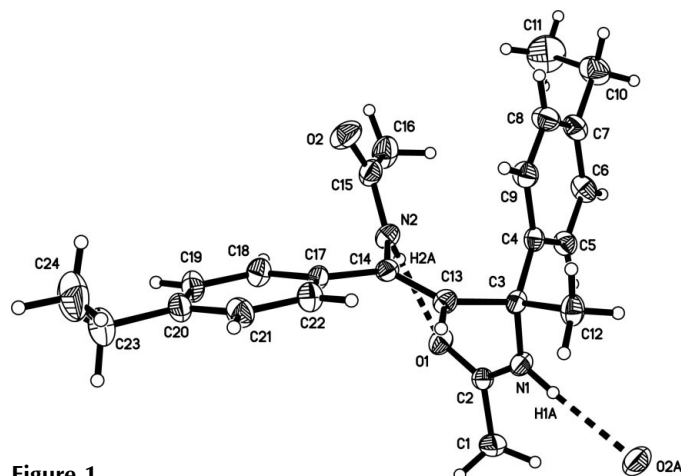


Figure 1
The molecular structure of (I), showing displacement ellipsoids at the 30% probability level (Siemens, 1995).

Crystal data

$C_{24}H_{30}N_2O_2$
 $M_r = 378.50$
 Monoclinic, $P2_1/c$
 $a = 20.773$ (4) Å
 $b = 6.5109$ (10) Å
 $c = 17.063$ (3) Å
 $\beta = 101.918$ (4)°
 $V = 2258.0$ (6) Å³
 $Z = 4$

$D_x = 1.113$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 1867 reflections
 $\theta = 1-27.5^\circ$
 $\mu = 0.07$ mm⁻¹
 $T = 294$ (2) K
 Prism, colorless
 $0.28 \times 0.23 \times 0.20$ mm

Data collection

Bruker CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.986$
 11299 measured reflections

5141 independent reflections
 1539 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.083$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -20 \rightarrow 26$
 $k = -7 \rightarrow 8$
 $l = -22 \rightarrow 16$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.054$
 $wR(F^2) = 0.082$
 $S = 0.91$
 5141 reflections
 258 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.005P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.17$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³

Table 1

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1A \cdots O2^i$	0.86	2.11	2.964 (3)	175
$N2-H2A \cdots O1$	0.86	2.01	2.753 (3)	144

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

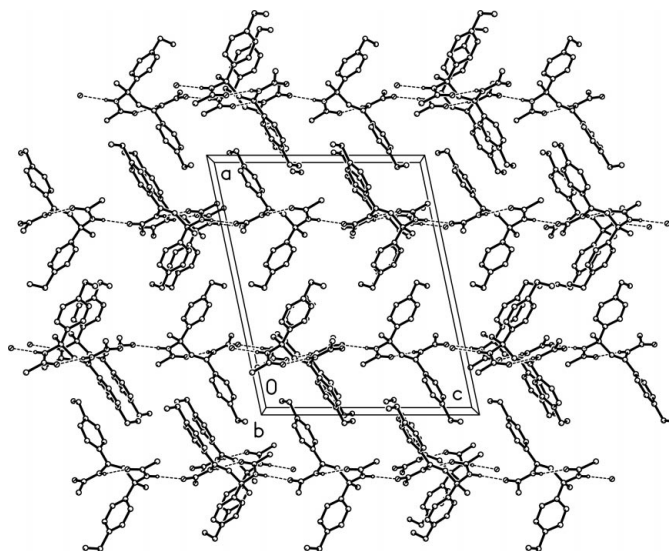


Figure 2

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

H atoms were included in the riding-model approximation, with U_{iso} equal to U_{eq} of the atom to which they were bound.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SHELXTL-NT* (Siemens, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL-NT*; software used to prepare material for publication: *SHELXTL-NT*.

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