Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.034 wR factor = 0.072 Data-to-parameter ratio = 10.9

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

N-Acetyl-L-valine

The title compound, $C_7H_{13}NO_3$, was prepared by acetylation of L-valine. The C(carbonyl)—N bond length is 1.324 (2) Å, which corresponds to the C–N bond length of a typical acylamine group.

Received 7 April 2004 Accepted 29 April 2004 Online 8 May 2004

Comment

N-Acetyl-L-valine, (I), is a versatile synthon in the synthesis of several pharmaceuticals and their key intermediates (Reddy *et al.*, 1999; Heavner, 1997). In a continuation of our work on the structure–activity relationship of the avermectin B1 derivatives, we have obtained a colourless crystalline compound that was the product of acetylation of L-valine. The structural identity of our product, (I), was resolved using single-crystal X-ray diffraction to determine the molecular structure.



The molecular structure of (I) is illustrated in Fig. 1. Selected bond lengths, angles and torsion angles are listed in Table 1. In (I), the C(carbonyl)—N bond length is 1.324 (2) Å, which corresponds to the C—N bond length of a typical acylamine group (Ganis *et al.*, 1971). Atoms C2, C3, N1 and C1 are coplanar [deviations within 0.0136 (9) Å], and atoms N1, C1, C4 and O3 are also coplanar [deviations within 0.0290 (8) Å]. The dihedral angle between the two planes is $49.75 (13)^{\circ}$.

Experimental

Compound (I) was obtained by acetylating L-valine with acetic anhydride and was recrystallized from ethanol to give colourless prisms (m.p. 437–438 K), $[\alpha]_{25}^{D}$ +4.1 (H₂O, 1%) (Cadogan *et al.*, 1996). L-Valine was purchased from the Shanghai Chemical Reagents Co.

Crystal data

$C_7H_{13}NO_3$		
$M_r = 159.18$	Mo $K\alpha$ radiation	
Orthorhombic, $P2_12_12_1$	Cell parameters from 25	
a = 6.654 (1) Å	reflections	
b = 9.444(2) Å	$\theta = 3.1-27.5^{\circ}$	
c = 13.182(3) Å	$\mu = 0.10 \text{ mm}^{-1}$	
V = 828.4 (3) Å ³	T = 293 (2) K	
Z = 4	Prism, colourless	
$D_x = 1.276 \text{ Mg m}^{-3}$	$0.20 \times 0.20 \times 0.18 \text{ mm}$	

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organic papers

Data collection

Rigaku R-AXIS RAPID diffractometer $\omega/2\theta$ scans Absorption correction: none 1841 measured reflections 1099 independent reflections *Refinement*

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.072$ S = 0.911099 reflections 101 parameters

H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

O1-C3	1.245 (2)	O3-C4	1.326 (2)
N1-C3	1.324 (2)	O2-C4	1.191 (2)
N1-C1	1.452 (2)		
C3-N1-C1	124.54 (14)	N1-C3-C2	116.12 (16)
O1-C3-N1	122.83 (17)	N1-C1-C4	107.37 (15)
O1-C3-C2	121.05 (17)	N1-C1-C5	112.91 (15)
C1-N1-C3-C2	177.95 (17)	C3-N1-C1-C5	104.6 (2)
C3-N1-C1-C4	-129.47 (18)	O3-C4-C1-N1	-175.47 (15)

830 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_o^2) + (0.0340P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

Extinction correction: SHELXL97

Extinction coefficient: 0.052 (4)

 $R_{\rm int} = 0.021$

 $\theta_{\rm max}=27.5^\circ$

 $h = -8 \rightarrow 8$ $k = -11 \rightarrow 12$

 $l = -16 \rightarrow 17$

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.12 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.13 \ {\rm e} \ {\rm \AA}^{-3}$

H atoms were included at calculated positions and refined using a riding model, with $U_{\rm iso} = 1.2$ (or 1.5 for methyl H atoms) times $U_{\rm eq}$ (parent atom). C–H distances were constrained to 0.96 Å for methyl H atoms and 0.98 Å for the remainder. The O–H distance was constrained to 0.82 Å and the N–H distance to 0.86 Å. In the absence of significant anomalous scattering effects, the Friedel pairs were merged. The absolute configuration was assumed to be that of the starting material.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/



Figure 1

The structure of (I), drawn with 30% probability displacement ellipsoids.

MSC, 2003); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

We are very grateful to the National Basic Science Research and Development Grants (973) (No. 2003CB114402).

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