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Key indicators

Single-crystal X-ray study
 $T = 173\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.009\text{ \AA}$
 R factor = 0.070
 wR factor = 0.182
Data-to-parameter ratio = 9.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.A short depsipeptide: *tert*-butoxycarbonyl-
L-alanyl-L-lactic acid benzyl ester
(Boc-L-Ala-L-Lac-OBzl)The title compound, $\text{C}_{18}\text{H}_{25}\text{N}_1\text{O}_6$, crystallizes in an extended sheet-like conformation. In the crystal structure, independent molecules are linked by an $\text{N}-\text{H}\cdots\text{O}=\text{C}$ hydrogen bond, yielding an infinite arrangement.Received 18 March 2004
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Comment

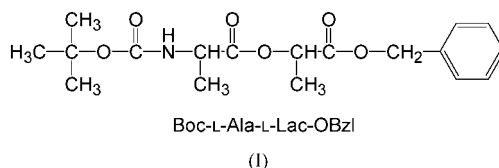
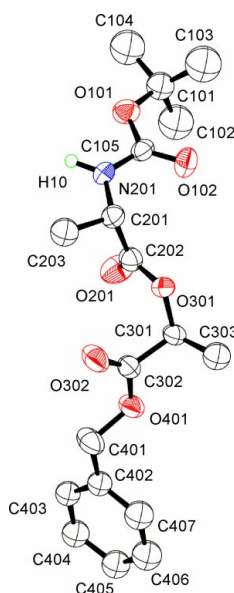
The title compound, (I), is a key starting material in the production of bioabsorbable polymers, such as $-(\text{Ala}-\text{Ala}-\text{Lac})_n-$ (Katakai & Goodman, 1982). Similar compounds, containing polydepsipeptide and oligodepsipeptide sequences, have been extensively studied in our laboratory (Katakai & Goodman, 1982; Katakai, 1988; Katakai *et al.*, 1995, 2004; Ohyama *et al.*, 2000, 2001; Yasuno *et al.*, 2001; Yoshida *et al.*, 1990).The molecular structure of (I) is shown in Fig. 1; the bond lengths and angles are unexceptional. The molecule has an extended sheet-like conformation. As shown in Fig. 2, independent molecules are linked by an $\text{NH}\cdots\text{O}=\text{C}$ interaction, yielding an infinite arrangement.

Figure 1

The structure of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. All H atoms have been omitted, except for those involved in hydrogen bonding.

The melting point of (I) is 356–359 K. Relatively low values are often observed for short sequences, especially those containing a hydroxyacid residue (Oku *et al.*, 2003, 2003a,b). In these cases, the number of hydrogen bonds definitely influences the melting temperature.

Experimental

The title compound, (I), was isolated from a reaction mixture of a liquid-phase peptide synthesis (Ohya *et al.*, 2000, 2001). Crystals of the title compound were successfully grown by the slow evaporation of an ethyl acetate solution. Analytical data (melting point, $^1\text{H NMR}$, ESI-MS, and $[\alpha]_D^{20}$) are in accordance with the expected structure; m.p. 356–359 K, $[\alpha]_D^{20} = -44.8^\circ$ (*c* 0.1, chloroform).

Crystal data

$\text{C}_{18}\text{H}_{25}\text{NO}_6$	$D_x = 1.200 \text{ Mg m}^{-3}$
$M_r = 351.40$	Cu $K\alpha$ radiation
Monoclinic, $P2_1$	Cell parameters from 4037 reflections
$a = 10.365$ (6) Å	$\theta = 4.2\text{--}67.5^\circ$
$b = 8.827$ (5) Å	$\mu = 0.75 \text{ mm}^{-1}$
$c = 11.001$ (6) Å	$T = 173.1 \text{ K}$
$\beta = 104.93$ (2)°	Prism, colorless
$V = 972.4$ (9) Å ³	$0.40 \times 0.30 \times 0.05 \text{ mm}$
$Z = 2$	

Data collection

Rigaku R-Axis RAPID diffractometer	1747 independent reflections
ω scans	1624 reflections with $F^2 > 2\sigma(F^2)$
Absorption correction: refined from ΔF (DIFABS; Walker & Stuart, 1983)	$R_{\text{int}} = 0.029$
$T_{\text{min}} = 0.789$, $T_{\text{max}} = 0.963$	$\theta_{\text{max}} = 68.2^\circ$
15424 measured reflections	$h = -12 \rightarrow 12$
	$k = -10 \rightarrow 10$
	$l = -13 \rightarrow 12$

Refinement

Refinement on F^2	All H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.070$	$w = 1/[0.005F_o^2 + 20\sigma(F_o^2) + 0.5]/(4F_o^2)$
$wR(F^2) = 0.182$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 0.94$	$\Delta\rho_{\text{max}} = 0.37 \text{ e \AA}^{-3}$
1747 reflections	$\Delta\rho_{\text{min}} = -0.33 \text{ e \AA}^{-3}$
191 parameters	

Table 1

Selected torsion angles (°).

O101–C105–N201–C201	177.4 (4)	C202–O301–C301–C302	–69.3 (5)
C105–N201–C201–C202	–66.9 (6)	O301–C301–C302–O401	162.7 (5)
N201–C201–C202–O301	145.9 (5)	C301–C302–O401–C401	174.9 (4)
C201–C202–O301–C301	166.4 (4)		

Table 2

Hydrogen-bonding geometry (Å, °).

$D\text{--}H\cdots A$	$D\text{--}H$	$H\cdots A$	$D\cdots A$	$D\text{--}H\cdots A$
N201–H10 \cdots O302 ⁱ	0.95	2.07	2.946 (6)	153

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

The *tert*-butyl C atoms, those of the benzene ring, and C203 and C303 were refined isotropically; all other non-H atoms were refined anisotropically. This procedure was adopted because of the limited

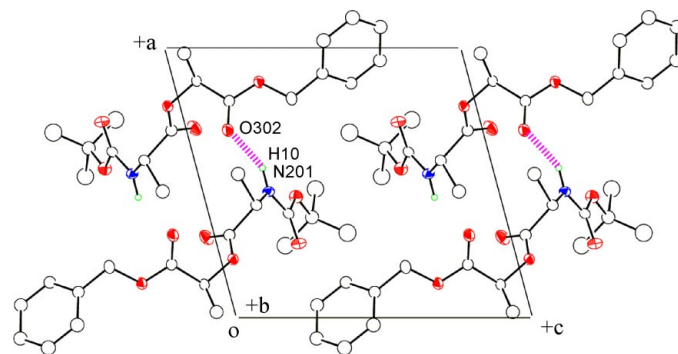


Figure 2

A packing diagram of (I), projected down the *b* axis. Hydrogen bonds are indicated as dashed lines.

number of observed reflections. H atoms were positioned geometrically and refined using a riding model, with C–H and N–H distances of 0.95 Å. $U_{\text{iso}}(\text{H})$ values were constrained to be $1.2U_{\text{eq}}$ of the carrier atom. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged. The absolute configuration was confirmed from the analytical data, α_D , of the compound.

Data collection: *RAPID-AUTO* (Rigaku/MS and Rigaku, 2003); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS and Rigaku, 2003); program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *CRYSTALS* (Watkin *et al.*, 1996); molecular graphics: *ORTEP* (Johnson, 1965); software used to prepare material for publication: *CrystalStructure*.

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