

Sheldrick, G. M. (1993). *SHELXL93. Program for the Refinement of Crystal Structures*. University of Göttingen, Germany.  
 Sivaraman, J., Subramaniam, K., Velmurugan, D., Subramanian, E. & Seetharaman, J. (1996). *J. Mol. Struct.* **385**, 123–128.  
 Starikov, E. B. & Steiner, T. (1998). *Acta Cryst.* **B54**, 94–96.  
 Zsolnai, L. (1997). *ZORTEP. An Interactive ORTEP Program for Structure Illustration*. University of Heidelberg, Germany.

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### *tert*-Butoxycarbonyl-L-leucyl-L-threoninamide

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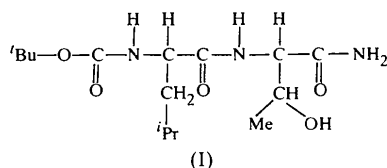
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#### Abstract

The peptide chain in C<sub>15</sub>H<sub>29</sub>N<sub>3</sub>O<sub>5</sub> adopts an extended conformation. The peptide unit is *trans* and shows significant deviations from planarity. The crystal packing enables neighbouring molecules to interact through hydrogen bonding in an anti-parallel fashion.

#### Comment

The conformation of the *tert*-butoxycarbonyl group in the title peptide, (I), is characterized by the torsion angles  $\theta_0 = 175.6(3)^\circ$  (C1—O1—C0'—N1) and  $\omega_0 = -173.7(3)^\circ$  (O1—C0'—N1—C1A) as *trans-trans* (Benedetti *et al.*, 1980). The peptide unit is *trans* and shows significant deviation from planarity ( $\Delta\omega = 10.7^\circ$ ). The carboxyl group makes a dihedral angle of  $37.5^\circ$  with the adjacent peptide unit. The peptide chain backbone is folded with the conformation  $\varphi_1 = -119.8(3)$ ,  $\psi_1 = 0.9(4)$ ,  $\omega_1 = 169.3(3)$ ,  $\varphi_2 = -139.6(3)$  and  $\psi_2 = 161.1(3)^\circ$ .



The side chain of the leucyl residue has torsion angles  $\chi_1 = -58.9(4)$ ,  $\chi_{21} = 174.4(3)$  and  $\chi_{22} = -60.5(5)^\circ$ , corresponding to the commonly observed

*g*<sup>-</sup> (*tg*<sup>-</sup>) conformation (Benedetti *et al.*, 1983). The side-chain conformation of the threonyl residue is described by  $\chi_{11}$  (N2—C2A—C2B—C2G) =  $-51.5(3)^\circ$  and  $\chi_{12}$  (N2—C2A—C2B—O2G) =  $73.2(3)^\circ$ . For comparison, in Gly—Thr—H<sub>2</sub>O,  $\chi_{11} = -62$  and  $\chi_{12} = 61^\circ$  (Yadava & Padmanabhan, 1973) and in Boc—Phe—D—Leu—Thr—OMe,  $\chi_{11} = -69.4(5)$  and  $\chi_{12} = 49.9(4)^\circ$  (Doi *et al.*, 1993).

The peptide chain folding introduces an intramolecular interaction [2.629(3) Å] between N2 and O2'. Such an interaction and an accompanying large deviation from planarity of the peptide unit have been observed in other peptides, such as Leu—Leu (Mitra & Subramanian, 1994) and Leu—Ala (Mitra *et al.*, 1996).

The packing gives rise to layers of molecules perpendicular to the *c* axis, with three intermolecular N—H...O hydrogen bonds holding the molecules together in each layer, while the layers are stabilized through hydrophobic and van der Waals interactions.

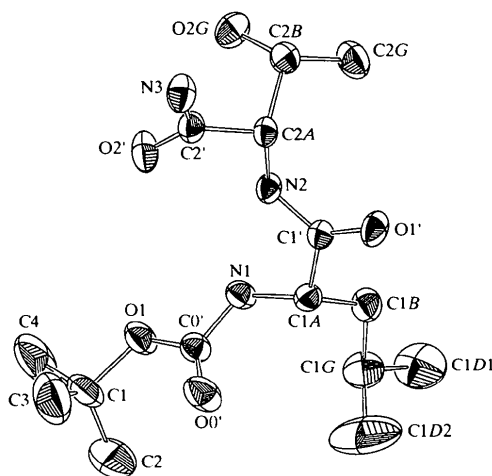


Fig. 1. Perspective view of the molecule, with displacement ellipsoids shown at the 50% probability level. H atoms have been omitted for clarity.

#### Experimental

The crystals were obtained by slow evaporation of a methanol/water solution at room temperature.

#### Crystal data

C<sub>15</sub>H<sub>29</sub>N<sub>3</sub>O<sub>5</sub>  
 $M_r = 331.41$   
 Orthorhombic  
*P*2<sub>1</sub>2<sub>1</sub>2<sub>1</sub>  
 $a = 7.713(2) \text{ \AA}$   
 $b = 8.674(1) \text{ \AA}$   
 $c = 28.325(6) \text{ \AA}$   
 $V = 1895.0(7) \text{ \AA}^3$   
 $Z = 4$   
 $D_x = 1.162 \text{ Mg m}^{-3}$   
 $D_m$  not measured

Cu *K*α radiation  
 $\lambda = 1.5418 \text{ \AA}$   
 Cell parameters from 25 reflections  
 $\theta = 15\text{--}22^\circ$   
 $\mu = 0.719 \text{ mm}^{-1}$   
 $T = 293(2) \text{ K}$   
 Parallelepiped  
 $0.40 \times 0.28 \times 0.20 \text{ mm}$   
 Colourless

**Data collection**

Enraf–Nonius CAD-4 diffractometer	$\theta_{\max} = 73^\circ$
$\omega/2\theta$ scans	$h = 0 \rightarrow 9$
Absorption correction: none	$k = 0 \rightarrow 10$
2140 measured reflections	$l = 0 \rightarrow 35$
2140 independent reflections	3 standard reflections
2030 reflections with $I > 2\sigma(I)$	every 100 reflections
	intensity decay: <1%

**Refinement**

Refinement on $F^2$	Extinction correction:
$R[F^2 > 2\sigma(F^2)] = 0.059$	<i>SHELXL93</i>
$wR(F^2) = 0.176$	Extinction coefficient:
$S = 1.008$	0.0040 (12)
2137 reflections	Scattering factors from
215 parameters	<i>International Tables for Crystallography</i> (Vol. C)
H atoms: see below	Absolute structure:
$w = 1/[\sigma^2(F_o^2) + (0.0708P)^2]$	Flack (1983)
where $P = (F_o^2 + 2F_c^2)/3$	Flack parameter = 0.1 (4)
$(\Delta/\sigma)_{\max} = 0.003$	
$\Delta\rho_{\max} = 0.357 \text{ e } \text{\AA}^{-3}$	
$\Delta\rho_{\min} = -0.249 \text{ e } \text{\AA}^{-3}$	

Table 1. Selected torsion angles ( $^\circ$ )

C1—O1—C0'—N1	175.6 (3)
O1—C0'—N1—C1A	−173.7 (3)
C0'—N1—C1A—C1'	−119.8 (3)
N1—C1A—C1B—C1G	−58.9 (4)
C1A—C1B—C1G—C1D2	−60.5 (6)
C1A—C1B—C1G—C1D1	174.4 (3)
N1—C1A—C1'—N2	0.9 (4)
C1A—C1'—N2—C2A	169.3 (3)
C1'—N2—C2A—C2'	−139.6 (3)
N2—C2A—C2B—O2G	73.2 (3)
N2—C2A—C2B—C2G	−51.5 (3)

Table 2. Selected hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ )

D—H...A	D—H	H...A	D...A	D—H...A
N1—H1N1...O2G'	0.86	2.40	3.056 (3)	134
N3—H2N3...O1''	0.86	2.17	2.941 (4)	163
O2G...O2'	—	—	2.750 (4)	—

Symmetry codes: (i)  $\frac{1}{2} + x, \frac{1}{2} - y, -z$ ; (ii)  $x - \frac{1}{2}, \frac{3}{2} - y, -z$ .

All H-atom positions, except that of O2G, were geometrically fixed and allowed to ride on the corresponding non-H atoms.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989). Cell refinement: *SDP* (Frenz, 1978). Data reduction: *CAD-4 Software*. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *ZORTEP* (Zsolnai, 1997). Software used to prepare material for publication: *SHELXL* and *PARST* (Nardelli, 1995).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: VJ1089). Services for accessing these data are described at the back of the journal.

**References**

- Benedetti, E., Morelli, G., Nemethy, G. & Scheraga, H. A. (1983). *Int. J. Pept. Protein Res.* **22**, 1–15.

- Benedetti, E., Pedone, C., Toniolo, C., Nemethy, G., Pottle, M. S. & Scheraga, H. A. (1980). *Int. J. Pept. Protein Res.* **16**, 156–172.
- Doi, M., In, Y., Tanaka, M., Inoue, M. & Ishida, T. (1993). *Acta Cryst.* **C49**, 1530–1532.
- Enraf–Nonius (1989). *CAD-4 Software*. Version 5.0. Enraf–Nonius, Delft, The Netherlands.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Frenz, B. A. (1978). *The Enraf–Nonius CAD-4 SDP – A Real-Time System for Concurrent X-ray Data Collection and Crystal Structure Solution. Computing in Crystallography*, edited by H. Schenk, R. Olthof-Hazekamp, H. van Koningsveld & G. C. Bassi, pp. 64–71. Delft University Press.
- Mitra, S. N., Govindasamy, L. & Subramanian, E. (1996). *Acta Cryst.* **C52**, 2871–2873.
- Mitra, S. N. & Subramanian, E. (1994). *Biopolymers*, **34**, 1139–1143.
- Nardelli, M. (1995). *J. Appl. Cryst.* **28**, 659.
- Sheldrick, G. M. (1990). *Acta Cryst.* **A46**, 467–473.
- Sheldrick, G. M. (1993). *SHELXL93. Program for the Refinement of Crystal Structures*. University of Göttingen, Germany.
- Yadava, V. S. & Padmanabhan, V. M. (1973). *Acta Cryst.* **B29**, 854–857.
- Zsolnai, L. (1997). *ZORTEP. An Interactive ORTEP Program for Structure Illustration*. University of Heidelberg, Germany.

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## Dineopentylprehnitene (1,2,3,4-tetramethyl-5,6-dineopentylbenzene) at 296 and 223 K

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**Abstract**

The title substance,  $C_{20}H_{34}$ , crystallized in the centrosymmetric space group  $C2/c$  with a single molecule as the asymmetric unit. The molecule is quite crowded intramolecularly, and has a notably non-planar benzene ring. There is, however, virtually no intermolecular crowding.

**Comment**

Interest in dineopentylprehnitene, (I), is associated with its potential for substantial crowding in the crystalline state. It crystallized in the centrosymmetric space group

