

The Crystal Structure of t-Butyloxycarbonyltetra-L-proline Benzyl Ester*

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Crystals of t-butyloxycarbonyltetra-L-proline benzyl ester monohydrate, $C_{32}H_{44}N_4O_7 \cdot H_2O$, are monoclinic with cell dimensions $a = 21.426 \pm 0.003$, $b = 8.738 \pm 0.002$, $c = 17.469 \pm 0.003 \text{ \AA}$, $\beta = 91.11 \pm 0.02^\circ$. The space group is $C2$, $Z = 4$. The structure was refined by a block-matrix least-squares method including anisotropic thermal parameters. The final R value for 2530 reflexions, which were measured on a diffractometer, is 0.080. The four proline residues in the molecule form one turn of poly-L-proline II-like helix. The torsion angles of the main chain are $\omega_0 = -11^\circ$, $\varphi_1 = -67.7^\circ$, $\psi_1 = 151.8^\circ$, $\omega_1 = 168.8^\circ$, $\varphi_2 = -69.9^\circ$, $\psi_2 = 164.9^\circ$, $\omega_2 = 176.4^\circ$, $\varphi_3 = -61.1^\circ$, $\psi_3 = 152.3^\circ$, $\omega_3 = 179.8^\circ$, $\varphi_4 = -57.8^\circ$, $\psi_4 = -38.5^\circ$. The pyrrolidine rings in Pro(2) and Pro(4) residues take the puckered form in which the β -carbon atom deviates from the plane of the remaining four atoms in the ring. Several short intramolecular distances were found, the shortest C-C and C-O distances of which were 2.921 and 2.618 \AA , respectively.

Introduction

The conformations of a series of L-proline oligomer derivatives [t-butyloxycarbonyl-(L-proline)_n] benzyl ester; $n = 2, 3, 4, 5, 6$ and t-amyloxycarbonyl-(L-proline)_n; $n = 2, 3, 4, 5, 6, 8$ were investigated, with regard to the degree of polymerization, by infrared spectroscopy (Isemura, Okabayashi & Sakakibara, 1968), ultraviolet absorption spectroscopy (Okabayashi & Isemura, 1968, 1970a), Raman spectroscopy (Rippon, Koenig & Walton, 1970), nuclear magnetic resonance (Deber, Bovey, Carvar & Blout, 1970; Okabayashi & Isemura, 1970b) and calculation of the intramolecular energies (Tonelli, 1970). The results of these investigations agree well in the case of the oligomers containing five or more residues, which were assumed to have a left-handed helical structure identical with poly-L-proline II except for the terminal residues. For the conformation of the tetramer, however, Deber *et al.* (1970) indicated from their 220-MHz n.m.r. study that out of 16 possible peptide bond conformational isomers of the tetramer, eight are apparently present, while the other investigators reported that the tetramer assumes a helical structure similar to that of poly-L-proline II, though not conclusively as in the case of the pentamer. Thus, the tetramer seems to be on a boundary where the poly-L-proline II-like helical conformation begins to appear. The present paper describes the crystal structure of t-butyloxycarbonyltetra-L-proline benzyl ester (BOC-Pro₄-Bz), which crystallizes with a water molecule. The serial numbering

of the proline residues and the atoms in the molecule is shown in Fig. 1.

Experimental

The material was synthesized according to the method described by Miyoshi, Kimura & Sakakibara (1970). The crystals recrystallized from ethyl acetate solution were colourless transparent plates with well developed (001) faces. The density was measured by flotation in a carbon tetrachloride-ether mixture. The unit-cell dimensions were measured on a single-crystal diffractometer using $Cu K\alpha$ ($\lambda = 1.5405 \text{ \AA}$) radiation monochromated by a graphite crystal and refined by a least-squares method.

Crystal data

t-Butyloxycarbonyl-tetra-L-proline benzyl ester monohydrate, $C_{32}H_{44}N_4O_7 \cdot H_2O$
Monoclinic
 $C2$
 $Z = 4$
 $a = 21.426 \pm 0.003 \text{ \AA}$
 $b = 8.738 \pm 0.002$
 $c = 17.469 \pm 0.003$
 $\beta = 91.11 \pm 0.02^\circ$
 $U = 3269.9 \text{ \AA}^3$
 $D_m = 1.23 \text{ g cm}^{-3}$
 $D_x = 1.24 \text{ g cm}^{-3}$
 $\mu(Cu K\alpha) = 8.6 \text{ cm}^{-1}$
 $F(000) = 1320$

The intensity data were collected on the diffractometer by an $\omega-2\theta$ continuous scan. 3165 independent reflexions with $\sin \theta/\lambda$ less than 0.61 were measured, of which 5230 reflexions had $|F|$ greater than $2\sigma(|F|)$ and were used in the structure determination. Since the size of the crystal was $0.2 \times 0.2 \times 0.2 \text{ mm}$, no absorption

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correction was applied. After the Lorentz and polarization corrections, the value of an overall temperature factor B was determined to be 5.75 \AA^2 by a Wilson plot.

Determination of the structure

The structure was solved by the symbolic addition method and the application of the tangent formula (Karle & Karle, 1966). The procedure was similar to that described before (Matsuzaki & Iitaka, 1971). Fifteen atoms of the BOC-Pro₄-Bz molecule were recognized in an E map calculated with 833 phases having $|E| \geq 1.1$. The phases were refined and extended by applying the tangent formula to the structure factors which were calculated with the parameters of the 15 atoms. The second E map based on 874 phases revealed 40 atoms of the molecule and three remaining atoms were found in a subsequent difference Fourier map. The parameters of the 43 atoms were then refined by a block-matrix least-squares method to yield an R index of 0.146, where the function minimized was $\sum(|F_o| - |F_c|)^2$. At this stage a new difference Fourier map was calculated. The map showed two additional peaks (A and B) 1.127 \AA apart and near the diad axis at $x=0$ and $z=\frac{1}{2}$. The interatomic distances between these two peaks and carbonyl oxygen atoms of the tetramers, O(1) and O(2), are shown in Fig. 2, where the values are based on the final parameters. It seems reasonable to assume that the two peaks should be assigned to two independent water molecules and that one molecule occupies the A peak and the other the B' peak, which is related to B by the diad axis. In this case the water molecule at A is hydrogen-bonded to O(1)' and O(2)' atoms while another water molecule at B' is hydrogen-bonded to O(2)' and the oxygen atom of the water molecule at A . The values of the angles, 108.1° for O(1)'-A-O(2) and 96.4° for O(2)'-B'-A, together with the distances, support this interpretation. Should this be the case, the true space group would be $P2_1$. But there were no extra reflexions of $h+k=2n+1$ type even in the heavily exposed Weissenberg photographs. A disordered structure was therefore assumed for the water molecules and a halved occupancy factor was assigned to the A and B oxygen atoms.

Refinement of the structure

The parameters of 45 non-hydrogen atoms were refined by eight cycles of block-matrix least-squares calculations with anisotropic temperature factors. In the last cycle of the refinement every parameter shift was within one half of its standard deviation. The final R value was 0.080 for 2530 reflexions. The final difference Fourier map did not show any peaks or troughs with magnitudes of electron density greater than 0.2 e \AA^{-3} and no attempt was made to locate the hydrogen atoms.

The atomic scattering factors were taken from *International Tables for X-ray Crystallography* (1962).

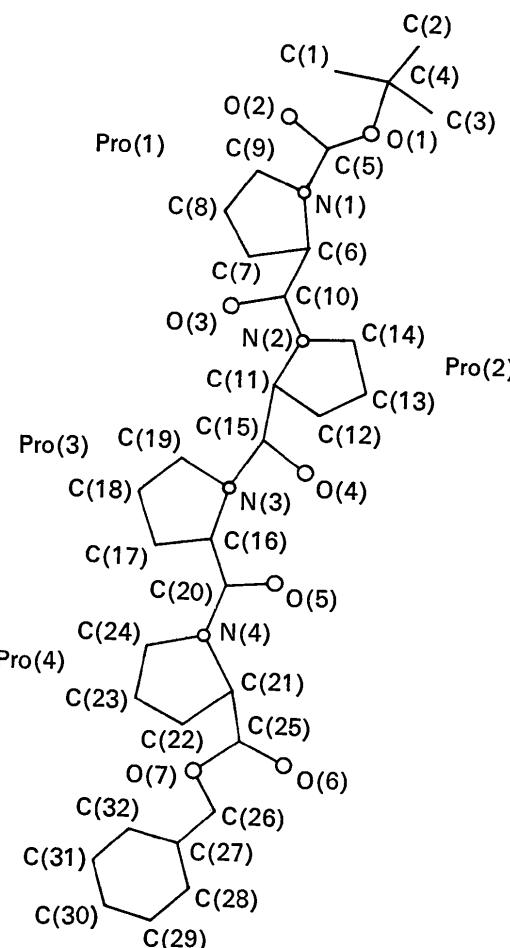


Fig. 1. Serial numbering of the proline residues and atoms in a BOC-Pro₄-Bz molecule.

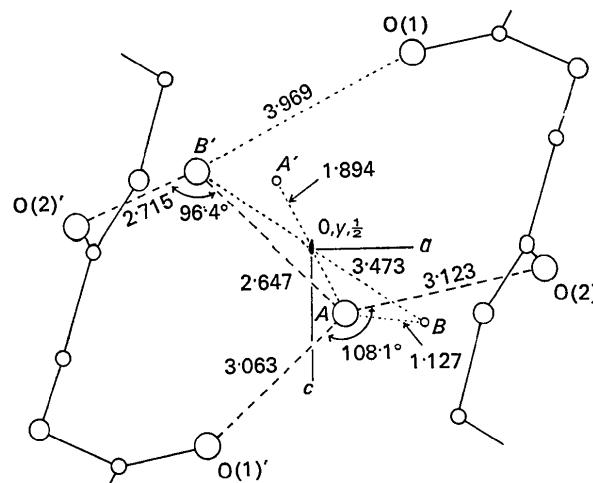


Fig. 2. The arrangement of atoms around the diad axis at $x=0, z=\frac{1}{2}$. A' , B' , $O(1)'$ and $O(2)'$ are symmetry-equivalent atoms to A , B , $O(1)$ and $O(2)$.

Table 1. Observed and calculated structure factors ($\times 10$)
	θ₁	θ₂	θ₃	θ₄	θ₅	θ₆	θ₇	θ₈	θ₉	θ₁₀	θ₁₁	θ₁₂	θ₁₃	θ₁₄	θ₁₅	θ₁₆	θ₁₇	θ₁₈	θ₁₉	θ₂₀	θ₂₁	θ₂₂	θ₂₃	θ₂₄	θ₂₅	θ₂₆	θ₂₇	θ₂₈	θ₂₉	θ₃₀	θ₃₁	θ₃₂	θ₃₃	θ₃₄	θ₃₅	θ₃₆	θ₃₇	θ₃₈	θ₃₉	θ₄₀	θ₄₁	θ₄₂	θ₄₃	θ₄₄	θ₄₅	θ₄₆	θ₄₇	θ₄₈	θ₄₉	θ₅₀	θ₅₁	θ₅₂	θ₅₃	θ₅₄	θ₅₅	θ₅₆	θ₅₇	θ₅₈	θ₅₉	θ₆₀	θ₆₁	θ₆₂	θ₆₃	θ₆₄	θ₆₅	θ₆₆	θ₆₇	θ₆₈	θ₆₉	θ₇₀	θ₇₁	θ₇₂	θ₇₃	θ₇₄	θ₇₅	θ₇₆	θ₇₇	θ₇₈	θ₇₉	θ₈₀	θ₈₁	θ₈₂	θ₈₃	θ₈₄	θ₈₅	θ₈₆	θ₈₇	θ₈₈	θ₈₉	θ₉₀	θ₉₁	θ₉₂	θ₉₃	θ₉₄	θ₉₅	θ₉₆	θ₉₇	θ₉₈	θ₉₉	θ₁₀₀	θ₁₀₁	θ₁₀₂	θ₁₀₃	θ₁₀₄	θ₁₀₅	θ₁₀₆	θ₁₀₇	θ₁₀₈	θ₁₀₉	θ₁₁₀	θ₁₁₁	θ₁₁₂	θ₁₁₃	θ₁₁₄	θ₁₁₅	θ₁₁₆	θ₁₁₇	θ₁₁₈	θ₁₁₉	θ₁₂₀	θ₁₂₁	θ₁₂₂	θ₁₂₃	θ₁₂₄	θ₁₂₅	θ₁₂₆	θ₁₂₇	θ₁₂₈	θ₁₂₉	θ₁₃₀	θ₁₃₁	θ₁₃₂	θ₁₃₃	θ₁₃₄	θ₁₃₅	θ₁₃₆	θ₁₃₇	θ₁₃₈	θ₁₃₉	θ₁₄₀	θ₁₄₁	θ₁₄₂	θ₁₄₃	θ₁₄₄	θ₁₄₅	θ₁₄₆	θ₁₄₇	θ₁₄₈	θ₁₄₉	θ₁₅₀	θ₁₅₁	θ₁₅₂	θ₁₅₃	θ₁₅₄	θ₁₅₅	θ₁₅₆	θ₁₅₇	θ₁₅₈	θ₁₅₉	θ₁₆₀	θ₁₆₁	θ₁₆₂	θ₁₆₃	θ₁₆₄	θ₁₆₅	θ₁₆₆	θ₁₆₇	θ₁₆₈	θ₁₆₉	θ₁₇₀	θ₁₇₁	θ₁₇₂	θ₁₇₃	θ₁₇₄	θ₁₇₅	θ₁₇₆	θ₁₇₇	θ₁₇₈	θ₁₇₉	θ₁₈₀	θ₁₈₁	θ₁₈₂	θ₁₈₃	θ₁₈₄	θ₁₈₅	θ₁₈₆	θ₁₈₇	θ₁₈₈	θ₁₈₉	θ₁₉₀	θ₁₉₁	θ₁₉₂	θ₁₉₃	θ₁₉₄	θ₁₉₅	θ₁₉₆	θ₁₉₇	θ₁₉₈	θ₁₉₉	θ₂₀₀	θ₂₀₁	θ₂₀₂	θ₂₀₃	θ₂₀₄	θ₂₀₅	θ₂₀₆	θ₂₀₇	θ₂₀₈	θ₂₀₉	θ₂₁₀	θ₂₁₁	θ₂₁₂	θ₂₁₃	θ₂₁₄	θ₂₁₅	θ₂₁₆	θ₂₁₇	θ₂₁₈	θ₂₁₉	θ₂₂₀	θ₂₂₁	θ₂₂₂	θ₂₂₃	θ₂₂₄	θ₂₂₅	θ₂₂₆	θ₂₂₇	θ₂₂₈	θ₂₂₉	θ₂₃₀	θ₂₃₁	θ₂₃₂	θ₂₃₃	θ₂₃₄	θ₂₃₅	θ₂₃₆	θ₂₃₇	θ₂₃₈	θ₂₃₉	θ₂₄₀	θ₂₄₁	θ₂₄₂	θ₂₄₃	θ₂₄₄	θ₂₄₅	θ₂₄₆	θ₂₄₇	θ₂₄₈	θ₂₄₉	θ₂₅₀	θ₂₅₁	θ₂₅₂	θ₂₅₃	θ₂₅₄	θ₂₅₅	θ₂₅₆	θ₂₅₇	θ₂₅₈	θ₂₅₉	θ₂₆₀	θ₂₆₁	θ₂₆₂	θ₂₆₃	θ₂₆₄	θ₂₆₅	θ₂₆₆	θ₂₆₇	θ₂₆₈	θ₂₆₉	θ₂₇₀	θ₂₇₁	θ₂₇₂	θ₂₇₃	θ₂₇₄	θ₂₇₅	θ₂₇₆	θ₂₇₇	θ₂₇₈	θ₂₇₉	θ₂₈₀	θ₂₈₁	θ₂₈₂	θ₂₈₃	θ₂₈₄	θ₂₈₅	θ₂₈₆	θ₂₈₇	θ₂₈₈	θ₂₈₉	θ₂₉₀	θ₂₉₁	θ₂₉₂	θ₂₉₃	θ₂₉₄	θ₂₉₅	θ₂₉₆	θ₂₉₇	θ₂₉₈	θ₂₉₉	θ₃₀₀	θ₃₀₁	θ₃₀₂	θ₃₀₃	θ₃₀₄	θ₃₀₅	θ₃₀₆	θ₃₀₇	θ₃₀₈	θ₃₀₉	θ₃₁₀	θ₃₁₁	θ₃₁₂	θ₃₁₃	θ₃₁₄	θ₃₁₅	θ₃₁₆	θ₃₁₇	θ₃₁₈	θ₃₁₉	θ₃₂₀	θ₃₂₁	θ₃₂₂	θ₃₂₃	θ₃₂₄	θ₃₂₅	θ₃₂₆	θ₃₂₇	θ₃₂₈	θ₃₂₉	θ₃₃₀	θ₃₃₁	θ₃₃₂	θ₃₃₃	θ₃₃₄	θ₃₃₅	θ₃₃₆	θ₃₃₇	θ₃₃₈	θ₃₃₉	θ₃₄₀	θ₃₄₁	θ₃₄₂	θ₃₄₃	θ₃₄₄	θ₃₄₅	θ₃₄₆	θ₃₄₇	θ₃₄₈	θ₃₄₉	θ₃₅₀	θ₃₅₁	θ₃₅₂	θ₃₅₃	θ₃₅₄	θ₃₅₅	θ₃₅₆	θ₃₅₇	θ₃₅₈	θ₃₅₉	θ₃₆₀	θ₃₆₁	θ₃₆₂	θ₃₆₃	θ₃₆₄	θ₃₆₅	θ₃₆₆	θ₃₆₇	θ₃₆₈	θ₃₆₉	θ₃₇₀	θ₃₇₁	θ₃₇₂	θ₃₇₃	θ₃₇₄	θ₃₇₅	θ₃₇₆	θ₃₇₇	θ₃₇₈	θ₃₇₉	θ₃₈₀	θ₃₈₁	θ₃₈₂	θ₃₈₃	θ₃₈₄	θ₃₈₅	θ₃₈₆	θ₃₈₇	θ₃₈₈	θ₃₈₉	θ₃₉₀	θ₃₉₁	θ₃₉₂	θ₃₉₃	θ₃₉₄	θ₃₉₅	θ₃₉₆	θ₃₉₇	θ₃₉₈	θ₃₉₉	θ₄₀₀	θ₄₀₁	θ₄₀₂	θ₄₀₃	θ₄₀₄	θ₄₀₅	θ₄₀₆	θ₄₀₇	θ₄₀₈	θ₄₀₉	θ₄₁₀	θ₄₁₁	θ₄₁₂	θ₄₁₃	θ₄₁₄	θ₄₁₅	θ₄₁₆	θ₄₁₇	θ₄₁₈	θ₄₁₉	θ₄₂₀	θ₄₂₁	θ₄₂₂	θ₄₂₃	θ₄₂₄	θ₄₂₅	θ₄₂₆	θ₄₂₇	θ₄₂₈	θ₄₂₉	θ₄₃₀	θ₄₃₁	θ₄₃₂	θ₄₃₃	θ₄₃₄	θ₄₃₅	θ₄₃₆	θ₄₃₇	θ₄₃₈	θ₄₃₉	θ₄₄₀	θ₄₄₁	θ₄₄₂	θ₄₄₃	θ₄₄₄	θ₄₄₅	θ₄₄₆	θ₄₄₇	θ₄₄₈	θ₄₄₉	θ₄₅₀	θ₄₅₁	θ₄₅₂	θ₄₅₃	θ₄₅₄	θ₄₅₅	θ₄₅₆	θ₄₅₇	θ₄₅₈	θ₄₅₉	θ₄₆₀	θ₄₆₁	θ₄₆₂	θ₄₆₃	θ₄₆₄	θ₄₆₅	θ₄₆₆	θ₄₆₇	θ₄₆₈	θ₄₆₉	θ₄₇₀	θ₄₇₁	θ₄₇₂	θ₄₇₃	θ₄₇₄	θ₄₇₅	θ₄₇₆	θ₄₇₇	θ₄₇₈	θ₄₇₉	θ₄₈₀	θ₄₈₁	θ₄₈₂	θ₄₈₃	θ₄₈₄	θ₄₈₅	θ₄₈₆	θ₄₈₇	θ₄₈₈	θ₄₈₉	θ₄₉₀	θ₄₉₁	θ₄₉₂	θ₄₉₃	θ₄₉₄	θ₄₉₅	θ₄₉₆	θ₄₉₇	θ₄₉₈	θ₄₉₉	θ₅₀₀	θ₅₀₁	θ₅₀₂	θ₅₀₃	θ₅₀₄	θ₅₀₅	θ₅₀₆	θ₅₀₇	θ₅₀₈	θ₅₀₉	θ₅₁₀	θ₅₁₁	θ₅₁₂	θ₅₁₃	θ₅₁₄	θ₅₁₅	θ₅₁₆	θ₅₁₇	θ₅₁₈	θ₅₁₉	θ₅₂₀	θ₅₂₁	θ₅₂₂	θ₅₂₃	θ₅₂₄	θ₅₂₅	θ₅₂₆	θ₅₂₇	θ₅₂₈	θ₅₂₉	θ₅₃₀	θ₅₃₁	θ₅₃₂	θ₅₃₃	θ₅₃₄	θ₅₃₅	θ₅₃₆	θ₅₃₇	θ₅₃₈	θ₅₃₉	θ₅₄₀	θ₅₄₁	θ₅₄₂	θ₅₄₃	θ₅₄₄	θ₅₄₅	θ₅₄₆	θ₅₄₇	θ₅₄₈	θ₅₄₉	θ₅₅₀	θ₅₅₁	θ₅₅₂	θ₅₅₃	θ₅₅₄	θ₅₅₅	θ₅₅₆	θ₅₅₇	θ₅₅₈	θ₅₅₉	θ₅₆₀	θ₅₆₁	θ₅₆₂	θ₅₆₃	θ₅₆₄	θ₅₆₅	θ₅₆₆	θ₅₆₇	θ₅₆₈	θ₅₆₉	θ₅₇₀	θ₅₇₁	θ₅₇₂	θ₅₇₃	θ₅₇₄	θ₅₇₅	θ₅₇₆	θ₅₇₇	θ₅₇₈	θ₅₇₉	θ₅₈₀	θ₅₈₁	θ₅₈₂	θ₅₈₃	θ₅₈₄	θ₅₈₅	θ₅₈₆	θ₅₈₇	θ₅₈₈	θ₅₈₉	θ₅₉₀	θ₅₉₁	θ₅₉₂	θ₅₉₃	θ₅₉₄	θ₅₉₅	θ₅₉₆	θ₅₉₇	θ₅₉₈	θ₅₉₉	θ₆₀₀	θ₆₀₁	θ₆₀₂	θ₆₀₃	θ₆₀₄	θ₆₀₅	θ₆₀₆	θ₆₀₇	θ₆₀₈	θ₆₀₉	θ₆₁₀	θ₆₁₁	θ₆₁₂	θ₆₁₃	θ₆₁₄	θ₆₁₅	θ₆₁₆	θ₆₁₇	θ₆₁₈	θ₆₁₉	θ₆₂₀	θ₆₂₁	θ₆₂₂	θ₆₂₃	θ₆₂₄	θ₆₂₅	θ₆₂₆	θ₆₂₇	θ₆₂₈	θ₆₂₉	θ₆₃₀	θ₆₃₁	θ₆₃₂	θ₆₃₃	θ₆₃₄	θ₆₃₅	θ₆₃₆	θ₆₃₇	θ₆₃₈	θ₆₃₉	θ₆₄₀	θ₆₄₁	θ₆₄₂	θ₆₄₃	θ₆₄₄	θ₆₄₅	θ₆₄₆	θ₆₄₇	θ₆₄₈	θ₆₄₉	θ₆₅₀	θ₆₅₁	θ₆₅₂	θ₆₅₃	θ₆₅₄	θ₆₅₅	θ₆₅₆	θ₆₅₇	θ₆₅₈	θ₆₅₉	θ₆₆₀	θ₆₆₁	θ₆₆₂	θ₆₆₃	θ₆₆₄	θ₆₆₅	θ₆₆₆	θ₆₆₇	θ₆₆₈	θ₆₆₉	θ₆₇₀	θ₆₇₁	θ₆₇₂	θ₆₇₃	θ₆₇₄	θ₆₇₅	θ₆₇₆	θ₆₇₇	θ₆₇₈	θ₆₇₉	θ₆₈₀	θ₆₈₁	θ₆₈₂	θ₆₈₃	θ₆₈₄	θ₆₈₅	θ₆₈₆	θ₆₈₇	θ₆₈₈	θ₆₈₉	θ₆₉₀	θ₆₉₁	θ₆₉₂	θ₆₉₃	θ₆₉₄	θ₆₉₅	θ₆₉₆	θ₆₉₇	θ₆₉₈	θ₆₉₉	θ₇₀₀	θ₇₀₁	θ₇₀₂	θ₇₀₃	θ₇₀₄	θ₇₀₅	θ₇₀₆	θ₇₀₇	θ₇₀₈	θ₇₀₉	θ₇₁₀	θ₇₁₁	θ₇₁₂	θ₇₁₃	θ₇₁₄	θ₇₁₅	θ₇₁₆	θ₇₁₇	θ₇₁₈	θ₇₁₉	θ₇₂₀	θ₇₂₁	θ₇₂₂	θ₇₂₃	θ₇₂₄	θ₇₂₅	θ₇₂₆	θ₇₂₇	θ₇₂₈	θ₇₂₉	θ₇₃₀	θ₇₃₁	θ₇₃₂	θ₇₃₃	θ₇₃₄	θ₇₃₅	θ₇₃₆	θ₇₃₇	θ₇₃₈	θ₇₃₉	θ₇₄₀	θ₇₄₁	θ₇₄₂	θ₇₄₃	θ₇₄₄	θ₇₄₅	θ₇₄₆	θ₇₄₇	θ₇₄₈	θ₇₄₉	θ₇₅₀	θ₇₅₁	θ₇₅₂	θ₇₅₃	θ₇₅₄	θ₇₅₅	θ₇₅₆	θ₇₅₇	θ₇₅₈	θ₇₅₉	θ₇₆₀	θ₇₆₁	θ₇₆₂	θ₇₆₃	θ₇₆₄	θ₇₆₅	θ₇₆₆	θ₇₆₇	θ₇₆₈	θ₇₆₉	θ₇₇₀	θ₇₇₁	θ₇₇₂	θ₇₇₃	θ₇₇₄	θ₇₇₅	θ₇₇₆	θ₇₇₇	θ₇₇₈	θ₇₇₉	θ₇₈₀	θ₇₈₁	θ₇₈₂	θ₇₈₃	θ₇₈₄	θ₇₈₅	θ₇₈₆	θ₇₈₇	θ₇₈₈	θ₇₈₉	θ₇₉₀	θ₇₉₁	θ₇₉₂	θ₇₉₃	θ₇₉₄	θ₇₉₅	θ₇₉₆	θ₇₉₇	θ₇₉₈	θ₇₉₉	θ₈₀₀	θ₈₀₁	θ₈₀₂	θ₈₀₃	θ₈₀₄	θ₈₀₅	θ₈₀₆	θ₈₀₇	θ₈₀₈	θ₈₀₉	θ₈₁₀	θ₈₁₁	θ₈₁₂	θ₈₁₃	θ₈₁₄	θ₈₁₅	θ₈₁₆	θ₈₁₇	θ₈₁₈	θ₈₁₉	θ₈₂₀	θ₈₂₁	θ₈₂₂	θ₈₂₃	θ₈₂₄	θ₈₂₅	θ₈₂₆	θ₈₂₇	θ₈₂₈	θ₈₂₉	θ₈₃₀	θ₈₃₁	θ₈₃₂	θ₈₃₃	θ₈₃₄	θ₈₃₅	θ₈₃₆	θ₈₃₇</

Table 1 (*cont.*)

In the last three cycles of the least-squares refinement, the following weighting system was adopted:

$$\begin{aligned}\sqrt{\omega} &= 59/|F_o| \quad \text{when } |F_o| > 59 \\ \sqrt{\omega} &= 1.0 \quad \text{when } 59 \geq |F_o| \geq 5.9 \\ \sqrt{\omega} &= 0.7 \quad \text{when } 5.9 > |F_o|.\end{aligned}$$

A comparison of the observed and calculated structure factors is given in Table 1. The final atomic parameters and their estimated standard deviations are listed in Table 2. In the next section the atoms are designated as listed in the last column of Table 2, in accordance with the IUPAC-IUB Commission (1970) rules.

Discussion of the structure

Molecular structure

A stereoscopic drawing of the molecule is shown in Fig. 3. The bond lengths and angles are shown in Fig. 4. The standard deviations are 0.012 Å for lengths and 0.7° for angles. The values are reasonable except for two lengths; 1.581 Å for Pro(2)C α -Pro(2)C β and 1.402 Å for Pro(4)C β -Pro(4)C γ , which are probably result from the large temperature factors. The four

Table 2. Final atomic parameters and their standard deviations ($\times 10^4$)

Temperature factors are of the form: $T = \exp [-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + 2\beta_{12}hk + 2\beta_{13}hl + 2\beta_{23}kl)]$. Standard deviations are listed in parentheses denoting the least significant digits.

x	y	z	β_{11}	β_{22}	β_{33}	β_{12}	β_{13}	β_{23}		
C(1)	396 (5)	2498 (15)	1339 (7)	51 (4)	242 (22)	119 (7)	-43 (8)	24 (4)	-48 (12)	BOC-C(1)
C(2)	1150 (5)	3796 (14)	413 (5)	68 (4)	270 (22)	41 (4)	-4 (9)	15 (3)	-20 (8)	BOC-C(2)
C(3)	1582 (6)	2044 (13)	1460 (7)	59 (4)	186 (20)	96 (6)	45 (8)	-2 (4)	-43 (10)	BOC-C(3)
C(4)	1055 (4)	3169 (10)	1234 (5)	41 (3)	167 (14)	50 (3)	-4 (5)	3 (2)	-24 (6)	BOC-C(4)
C(5)	782 (3)	5629 (9)	1822 (4)	32 (2)	132 (12)	31 (2)	0 (4)	0 (2)	8 (5)	BOC-C(5)
C(6)	1386 (4)	6024 (10)	3024 (4)	36 (2)	165 (14)	32 (3)	-5 (5)	-9 (2)	-4 (5)	pro(1)C α
C(7)	1384 (4)	7391 (12)	3590 (5)	48 (3)	188 (16)	60 (4)	-14 (7)	-8 (3)	-31 (8)	pro(1)C β
C(8)	998 (6)	8603 (16)	3237 (9)	69 (5)	316 (30)	145 (9)	62 (10)	-37 (6)	-143 (14)	pro(1)C γ
C(9)	611 (4)	7936 (11)	2570 (5)	37 (3)	201 (18)	69 (5)	18 (6)	-5 (3)	-48 (8)	pro(1)C δ
C(10)	1199 (3)	4561 (10)	3449 (4)	27 (2)	186 (14)	30 (2)	-11 (4)	0 (2)	-8 (5)	pro(1)C'
C(11)	1529 (3)	2463 (10)	4242 (4)	31 (2)	147 (12)	29 (2)	-1 (4)	1 (2)	1 (5)	pro(2)C α
C(12)	2169 (4)	1573 (12)	4323 (5)	35 (2)	205 (17)	64 (4)	10 (6)	6 (3)	11 (8)	pro(2)C β
C(13)	2646 (4)	2842 (13)	4175 (7)	32 (3)	251 (22)	95 (6)	-2 (6)	-4 (3)	35 (10)	pro(2)C γ
C(14)	2344 (3)	3946 (12)	3624 (5)	25 (2)	231 (18)	56 (4)	-3 (5)	-2 (2)	22 (7)	pro(2)C δ
C(15)	1329 (3)	3101 (10)	5007 (4)	28 (2)	174 (14)	35 (3)	0 (4)	-3 (2)	-8 (5)	pro(2)C'
C(16)	876 (3)	2688 (12)	6243 (4)	26 (2)	260 (18)	36 (3)	3 (5)	2 (2)	-14 (6)	pro(3)C α
C(17)	553 (4)	1368 (15)	6597 (6)	41 (3)	320 (24)	66 (5)	-27 (8)	19 (3)	-19 (10)	pro(3)C β
C(18)	760 (7)	19 (17)	6178 (7)	87 (6)	293 (28)	83 (6)	-21 (11)	38 (5)	32 (11)	pro(3)C γ
C(19)	919 (4)	519 (12)	5366 (5)	44 (3)	188 (17)	54 (4)	-29 (6)	9 (3)	7 (7)	pro(3)C δ
C(20)	1478 (3)	3155 (10)	6704 (4)	28 (2)	198 (15)	33 (3)	10 (5)	4 (2)	-2 (6)	pro(3)C'
C(21)	1955 (4)	4665 (11)	7693 (4)	33 (2)	197 (16)	44 (3)	-4 (5)	-4 (2)	-3 (6)	pro(4)C α
C(22)	1670 (5)	5789 (12)	8287 (6)	46 (3)	200 (18)	64 (4)	8 (6)	0 (3)	-48 (8)	pro(4)C β
C(23)	1062 (6)	6114 (31)	8023 (13)	36 (4)	1101 (85)	251 (17)	29 (15)	-5 (6)	-424 (34)	pro(4)C γ
C(24)	824 (4)	5042 (14)	7446 (6)	35 (3)	308 (24)	73 (5)	13 (7)	7 (3)	-79 (9)	pro(4)C δ
C(25)	2276 (3)	3343 (11)	8102 (4)	28 (2)	203 (16)	39 (3)	-7 (5)	2 (2)	-9 (6)	pro(4)C'
C(26)	2195 (4)	1054 (13)	8779 (6)	32 (3)	271 (21)	80 (5)	-2 (6)	3 (3)	69 (9)	Bz-C(7)
C(27)	1690 (3)	99 (9)	9135 (4)	28 (2)	154 (13)	42 (3)	5 (4)	3 (2)	2 (5)	Bz-C(1)
C(28)	1860 (4)	-751 (12)	9781 (5)	34 (2)	236 (18)	50 (3)	13 (6)	-8 (2)	-2 (7)	Bz-C(2)
C(29)	1428 (4)	-1689 (12)	10137 (5)	39 (3)	217 (17)	43 (3)	3 (6)	3 (2)	18 (7)	Bz-C(3)
C(30)	822 (4)	-1775 (11)	9857 (5)	37 (2)	205 (17)	50 (4)	18 (6)	2 (2)	14 (7)	Bz-C(4)
C(31)	654 (4)	-890 (13)	9223 (5)	34 (2)	237 (19)	69 (4)	1 (6)	-9 (3)	22 (9)	Bz-C(5)
C(32)	1082 (4)	24 (11)	8863 (5)	30 (2)	208 (17)	55 (4)	-12 (5)	-2 (2)	24 (7)	Bz-C(6)
N(1)	907 (3)	6448 (7)	2451 (3)	32 (2)	133 (10)	38 (2)	9 (4)	-3 (2)	-7 (4)	pro(1)N
N(2)	1673 (2)	3679 (7)	3726 (3)	25 (1)	158 (10)	32 (2)	-2 (3)	-2 (1)	4 (4)	pro(2)N
N(3)	1048 (3)	2192 (8)	5488 (3)	29 (2)	190 (11)	31 (2)	-1 (4)	2 (1)	6 (5)	pro(3)N
N(4)	1408 (3)	4189 (8)	7242 (3)	30 (2)	203 (13)	33 (2)	16 (4)	4 (1)	-6 (5)	pro(4)N
O(1)	1162 (2)	4400 (7)	1790 (3)	34 (1)	148 (9)	44 (2)	18 (3)	-3 (1)	-14 (4)	BOC-O(1)
O(2)	400 (3)	5992 (7)	1330 (3)	36 (2)	207 (11)	45 (2)	14 (4)	-10 (1)	-4 (4)	BOC-O(2)
O(3)	659 (2)	4260 (9)	3579 (3)	24 (1)	285 (13)	55 (2)	-3 (4)	2 (1)	17 (5)	pro(1)O
O(4)	1425 (3)	4448 (7)	5175 (3)	54 (2)	173 (10)	44 (2)	-3 (4)	2 (2)	-16 (4)	pro(2)O
O(5)	1983 (2)	2536 (8)	6586 (3)	27 (1)	276 (13)	52 (2)	26 (4)	-1 (1)	-31 (5)	pro(3)O
O(6)	2839 (2)	3305 (9)	8186 (4)	27 (1)	287 (14)	69 (3)	-19 (4)	-6 (2)	11 (6)	pro(4)O
O(7)	1892 (2)	2325 (7)	8377 (3)	25 (1)	205 (10)	58 (2)	-1 (3)	-1 (1)	24 (5)	Bz-O
O(8)	206 (7)	6291 (21)	5484 (13)	47 (6)	237 (30)	201 (19)	2 (11)	12 (7)	-33 (19)	water A
O(9)	-681 (10)	6835 (25)	4444 (12)	95 (9)	321 (46)	128 (12)	-48 (17)	3 (8)	49 (20)	water B'

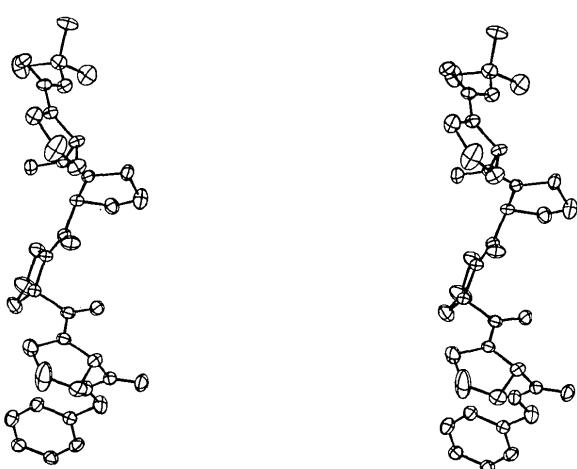


Fig. 3. Stereoscopic drawing of BOC-Pro₄-Bz molecule. Each ellipsoid encloses the region in which the atom is found with probability 20%.

proline residues in the BOC-Pro₄-Bz molecule form one turn of poly-L-proline II-like helix as shown in Fig. 3.

The internal rotation angles [designated on the basis of the IUPAC-IUB (1970) rules] are shown in Fig. 5. The values of φ , ψ , ω and the average translation per residue along the helix axis, h , are listed in Table 3 together with those of poly-L-proline II and collagen 1-bonded and 2-bonded models. In Fig. 6 these structures are plotted on a φ - ψ chart. The values of Pro(1) and Pro(3) residues lie between those of collagen 1-bonded and 2-bonded models and Pro(2) is located near the collagen 2-bonded model. The chart indicates that the three successive proline residues from the N-terminal do not assume the exact threefold screw-axis symmetry as poly-L-proline II does and that their structures are close to those of collagen models rather than that of poly-L-proline II. As can be seen in Table 3, the unit height of BOC-Pro₄-Bz is also consistent with those of collagen models. The C-terminal Pro(4)

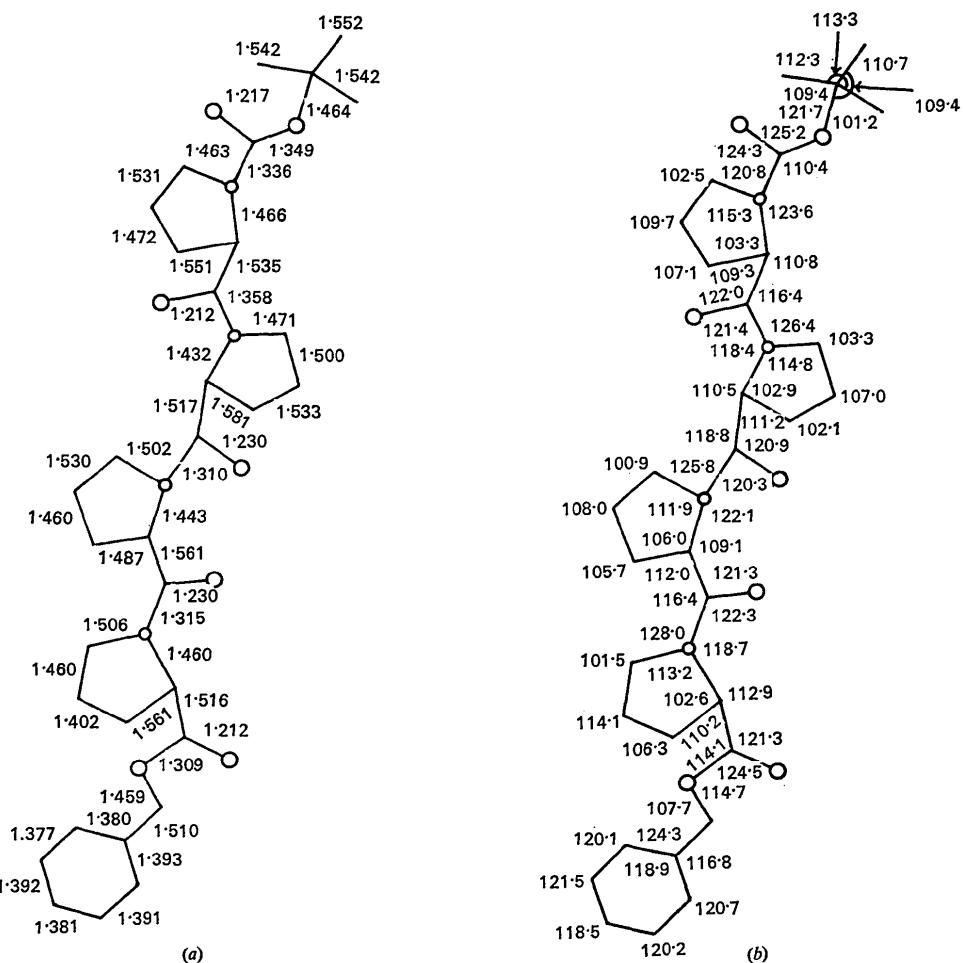


Fig. 4. (a) Bond lengths (\AA). E.s.d. is $\pm 0.012 \text{\AA}$. (b) Bond angles ($^\circ$). E.s.d. is $\pm 0.7^\circ$.

Table 3. Torsion angles and unit heights

	Pro(0)	Pro(1)	Pro(2)	Pro(3)	Pro(4)	Poly-L-proline II	1-bonded model	2-bonded model
ϕ (°)		-67.7	-69.9	-61.1	-57.8	-75	-52.5	-77
ψ (°)		151.8	164.9	152.3	-38.5	145	148.5	157
ω (°)	-1.1	168.8	176.4	179.8		180	180	180
h (Å)				2.87		3.12	2.86	2.91
Reference	(1)					(2)	(3)	(4)

(1) Present study. (2) Sasisekharan (1959). (3) Rich & Crick (1961). (4) Ramachandran & Sasisekharan (1965).

residue takes the α -helix type conformation which was found in a simple peptide, acetyl-L-proline-N-methylamide (Matsuzaki & Iitaka, 1970) and seems to be another stable conformation of proline residues. The N-terminal peptide bond takes the *cis* conformation, the presence of which was suggested by Deber *et al.* (1970) and Tonelli (1970). As for the planarity of the peptide bonds, the bond between Pro(1) and Pro(2) deviates from a plane at an angle ω of 168.8°, while the other three are nearly planar.

The equations of the least-squares planes of the pyrrolidine rings and the distances of the individual atoms from the planes are given in Table 4. The puckered form in which the β -carbon atom deviates from the least-squares plane is observed in Pro(2) and Pro(4) residues. In Pro(1) and Pro(3) residues, γ -carbon atoms are shifted from the planes as is usually found in proline residues. The directions of the displacements of the γ -carbon atoms from the planes with respect to the carbonyl carbon atoms are *trans*, *cis*, *cis* and *trans* for Pro(1), Pro(2), Pro(3) and Pro(4), respectively. The intramolecular distances that are shorter than the allowed values proposed by Ramachandran, Rama-

Table 4. Least-squares planes through the pyrrolidine rings and the benzene ring

The equations of the planes are of the form

$$AX + BY + CZ = D$$

where X , Y and Z are coordinates (in Å) referred to the orthogonal axis, $X \parallel a^*$, $Y \parallel b$ and $Z \parallel c$ respectively, and D is the origin-to-plane distance.

Pro(1) residue

$$0.715X + 0.447Y - 0.551Z = 6.371$$

Deviations of atoms from the plane

N	-0.008 Å	$C\gamma^*$	0.218 Å
$C\alpha$	0.008	$C\delta$	0.005
$C\beta$	-0.004		

Pro(2) residue

$$-0.047X + 0.673Y + 0.739Z = 0.330$$

Deviations of atoms from the plane

N	0.021 Å	$C\gamma$	0.012 Å
$C\alpha$	-0.013	$C\delta$	-0.020
$C\beta^*$	-0.500		

Pro(3) residue

$$0.875X - 0.264Y + 0.389Z = 1.828$$

Deviations of atoms from the plane

N	-0.038 Å	$C\gamma^*$	0.392 Å
$C\alpha$	0.038	$C\delta$	0.023
$C\beta$	-0.022		

Table 4 (cont.)

Pro(4) residue	$0.220X + 0.671Y - 0.713Z = 0.312$
	Deviations of atoms from the plane
N	0.015 Å
$C\alpha$	-0.010
$C\beta^*$	-0.225
$C\gamma$	0.009 Å
$C\delta$	-0.015

Benzene ring	$-0.262X + 0.761Y + 0.599Z = 3.450$
	Deviations of atoms from the plane
Bz-C(1)	-0.008 Å
Bz-C(2)	0.008
Bz-C(3)	0.000
Bz-C(4)	-0.009 Å
Bz-C(5)	0.010
Bz-C(6)	-0.001

*Not included in the least-squares calculation.

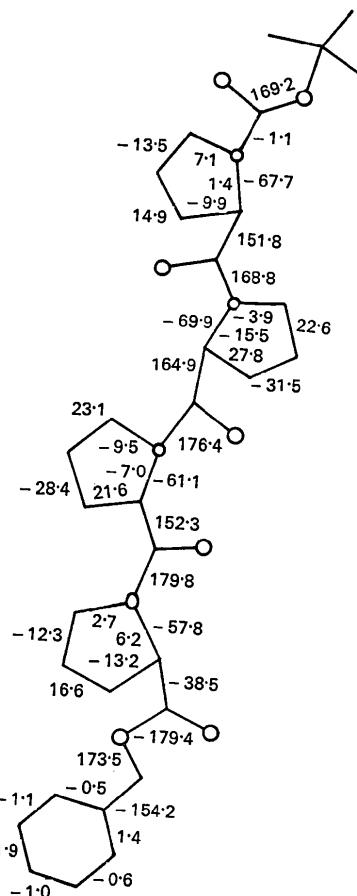


Fig. 5. Internal rotation angles (°).

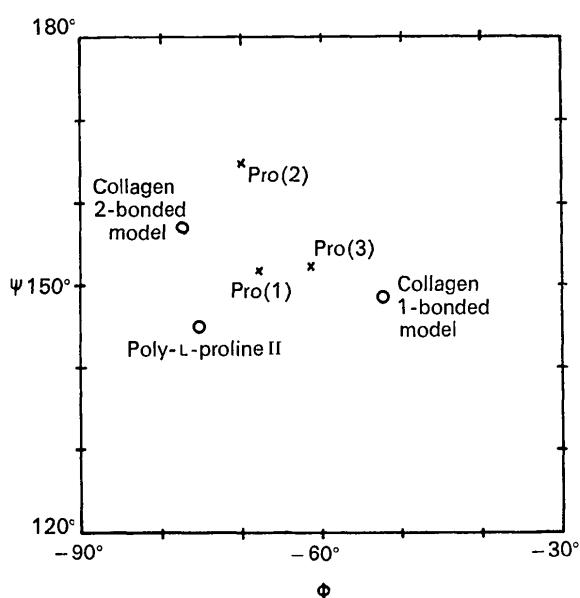


Fig. 6. ϕ - ψ chart. Pro(1), Pro(2) and Pro(3) are the proline residues in the BOC-Pro₄-Bz molecule.

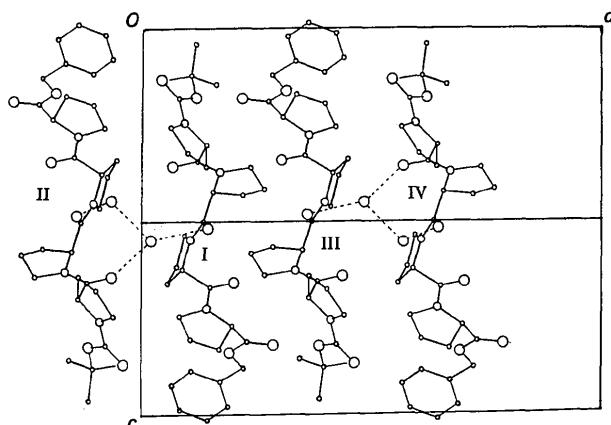


Fig. 7. Projection of the crystal structure along the b axis. The coordinates of the molecules I, II, III and IV are shown in Table 6.

krishnan & Sasisekharan (1963) are listed in Table 5. The distances between the α -carbon and δ -carbon atom of the next residue, between the carbonyl carbon and carbonyl carbon atom of the next residue, and between the α -carbon and carbonyl oxygen atom of the preceding residue are all included in this table. Some are shorter than the outer limit by about 0.08 Å.

Crystal structure

The projections of the crystal structure along the b axis and c axis are shown in Figs. 7 and 8, respectively [ORTEP, Johnson (1965)]. Two antiparallel chains of the tetramers I and II are linked together by the hydrogen bonds through the water molecules. Two of these hydrogen-bonded pairs of BOC-Pro₄-Bz molecules constitute a unit cell but there is no hydrogen bond between the pairs.

The short intermolecular contacts within 3.8 Å are listed in Table 6. The shortest, with a length of 2.991 Å, is between B' and pro(3) $C\gamma$ (010). But this is not the C-H...O type hydrogen bond suggested for the structures of poly-L-proline II (Sasisekharan, 1959) and the collagen 2-bonded model (Ramachandran & Sasisekharan, 1965), since the distance between the oxygen atom of the water and the hydrogen atom attached to the $C\gamma$ atom is 2.40 Å, where the coordinates of the hydrogen atom were calculated to be $x=0.0423$, $y=0.0774$ and $z=0.6151$.

Table 6. Packing distances shorter than 3.8 Å

From atom of I(000)	To atom	Distance
BOC-O(2)	Bz-C(3)	3.672 Å
BOC-O(2)	Bz-C(4)	3.366
Pro(2) $C\gamma$	Pro(2)O	3.736
Pro(3) $C\gamma$	water A	3.665
water B'	Pro(3) $C\gamma$	2.991
Pro(3)O	Pro(1) $C\alpha$	3.787
Pro(3)O	Pro(1) $C\beta$	3.521
Pro(3)O	Pro(2) $C\delta$	3.475
Pro(4)O	BOC-C(5)	3.769
Pro(4)O	Pro(1) $C\alpha$	3.369
Pro(4)O	Pro(1) $C\beta$	3.639
Pro(4)O	Pro(1) $C\delta$	3.613

Table 5. Intramolecular distances between non-bonded atoms

C-C distances		
Allowed value	3.20 Å	
Outer limit	3.00	
Pro(1) $C\alpha$ -Pro(2) $C\delta$	2.921 Å	
Pro(2) $C\alpha$ -Pro(3) $C\delta$	2.923	
Pro(3) $C\alpha$ -Pro(4) $C\delta$	2.945	
BOC-C(5)-Pro(1) C'	3.106	
Pro(1) C' -Pro(2) C'	3.015	
Pro(2) C' -Pro(3) C'	2.976	
Pro(3) C' -Pro(4) C'	2.959	
BOC-C(1)-BOC-C(5)	2.975	
BOC-C(2)-BOC-C(5)	3.052	
N-O distance		
Allowed value	2.70 Å	
Outer limit	2.60	
Pro(2)N-Pro(2)O	2.683 Å	
C-O distances		
Allowed value	2.80 Å	
Outer limit	2.70	
Pro(1) $C\alpha$ -BOC-O(1)	2.618 Å	
Pro(2) $C\alpha$ -Pro(1)O	2.683	
Pro(3) $C\alpha$ -Pro(2)O	2.704	
Pro(4) $C\alpha$ -Pro(3)O	2.685	
Pro(1) $C\delta$ -BOC-O(2)	2.783	
Pro(4) C' -Pro(3)O	2.800	
Bz-C(7)-Pro(4)O	2.626	
Bz-C(6)-Bz-O	2.799	

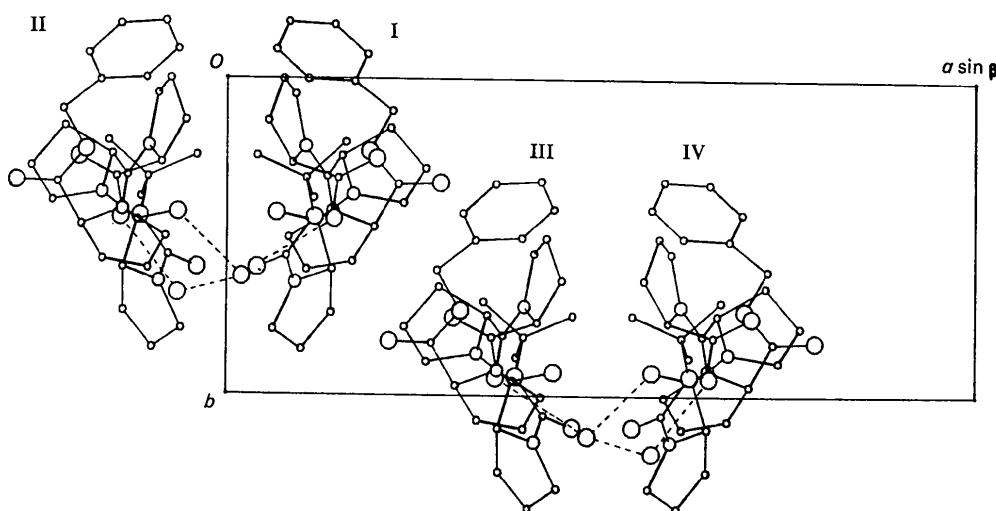
Fig. 8. Projection of the crystal structure along the *c* axis.

Table 6 (cont.)

Pro(4)O	Pro(1)N	III(0̄10)	3.348
BOC-C(5)	Pro(4)Cδ	II(000)	3.729
Pro(1)Cγ	Pro(4)O	III(000)	3.564
Pro(4)Cγ	BOC-O(2)	II(000)	3.353
BOC-O(2)	Pro(4)Cδ	II(000)	3.513
Pro(1)O	Pro(3)Cx	II(000)	3.583
Pro(1)O	Pro(3)Cβ	II(000)	3.632
Pro(1)O	Pro(4)Cδ	II(000)	3.680
Pro(2)O	Pro(2)Cβ	III(000)	3.631
Pro(4)O	BOC-C(3)	III(000)	3.545
water B'	Pro(2)C'	II(000)	3.680
water B'	Pro(4)Cδ	II(000)	3.663
Pro(4)Cγ	Bz-C(4)	I(010)	3.742
Pro(4)Cγ	Bz-C(5)	I(010)	3.477
Pro(4)Cγ	Bz-C(6)	I(010)	3.718
BOC-O(2)	Bz-C(5)	II(010)	3.657
water B'	Pro(3)Cδ	II(010)	3.277
Bz-C(4)	Bz-C(4)	II(001)	3.566
Bz-C(4)	Bz-C(5)	II(001)	3.657
Pro(4)O	Bz-C(2)	III(001)	3.690
Pro(4)O	Bz-C(3)	III(001)	3.296
	e.s.d.	(0.014)	

The coordinates of the equivalent positions are

I	x	y	z
II	-x	y	1-z
III	$\frac{1}{2}$ -x	$\frac{1}{2}$ +y	1-z
IV	$\frac{1}{2}$ +x	$\frac{1}{2}$ +y	z

followed by the components of translation vectors in parentheses.

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