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A Crystal Structure Determination for Tyr-D-Nle-Gly-Phe-NleS [NleS = $MeCH_2CH_2CH_2CH(NH_2)SO_3H$]: an Active Synthetic Enkephalin Analogue

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A crystal structure determination for Tyr-D-NIe-GIy-Phe-NIeS [NIeS = MeCH₂CH₂CH₂CH(NH₂)SO₃H] provides two examples of a 2'- β -bend conformation for a biologically active, synthetic enkephalin analogue.

The synthetic pentapeptide analogue L-tyrosyl-D-norleucylglycyl-L-phenylalanyl-L- α -aminopentanesulphonic acid (Tyr-D-Nle-Gly-Phe-NleS) (1) has been found to display enkephalin-like activity.¹ An interesting consequence of its chemical structure is that the introduction of a chiral residue at position 2 of the sequence significantly reduces the conformational freedom of this pentapeptide relative to the natural enkephalins (*e.g.* Tyr-Gly-Gly-Phe-Leu, Leu⁵-enkephalin). We felt that a crystal structure determination for (1) would be particularly valuable since Leu⁵-enkephalin has been found to display very different conformations in two crystalline modifications.^{2.3} We report the crystal structure of (1) in which there are two molecules of the pentapeptide analogue per asymmetric unit and compare their conformations with eight examples available for Leu⁵-enkephalin.

Crystals of (1) were obtained by slow evaporation of an aqueous ethanol solution: monoclinic, space group $P2_1$, a = 8.588(7),b = 30.265(38),c = 15.024(18)Å, $\beta = 96.94(7)^{\circ}$ (at *ca.* 120 K); the composition of the asymmetric unit is: (C₃₁H₄₅N₅O₇)₂·5H₂O·C₂H₅OH. Diffraction intensities were measured from the cooled crystal to $2\theta_{max} = 50^{\circ}$ (monochromatized Mo- K_{α} radiation) with a Syntex P1 autodiffractometer (equipped with a Syntex LT-1 low temperature device) operating in the ω -scan mode with a scan rate of $0.5-24.0^{\circ}$ min⁻¹. Of 6944 unique reflections measured, 4307 had $I \ge 3\sigma(I)$. The initial structural model was determined by direct methods⁴ and developed⁵ by difference Fourier and least squares refinement techniques; 5834 reflections contributed to the refinement of 882 variables to give R = 0.092.[†]

The two peptide analogue molecules per asymmetric unit display similar 2'- β -bend conformations (Figure 1). Both molecules display an intramolecular hydrogen bond between the tyrosine carbonyl oxygen atom and the amide hydrogen atom of phenylalanine. In one molecule (1a) there is a second

intramolecular hydrogen bond between the protonated amino group of tyrosine and the carbonyl oxygen atom of phenylalanine that is not present in the second molecule; the



Figure 1. ORTEP-II⁷ plots illustrating the conformations for the two independent molecules of Tyr-D-Nle-Gly-Phe-NleS in the crystallographic unit. Hydrogen bonds are illustrated with dotted lines. The amino acid sequence is labelled for molecule (1a).

[†] The atomic co-ordinates for this work are available on request from the Director of the Cambridge Crystallographic Data Centre, University Chemical Laboratory, Lensfield Road, Cambridge CB2 1EW. Any request should be accompanied by the full literature citation for this communication.



Figure 2. A stereoscopic superposition diagram⁸ for the two molecules of Tyr-D-Nle-Gly-Phe-NleS, four molecules of Leu⁵-enkephalin reported by Blundell *et al.*,² and the four molecules of Leu⁵-enkephalin reported by Karle *et al.*³

respective $N \cdots O$ distances are 2.86(1) Å for (1a) and 3.23(1) Å for (1b). The conformations in this crystal structure are consistent with that proposed by Casey *et al.*,⁶ based on a 400 MHz ¹H n.m.r. study of (1) dissolved in Me₂SO.

Space limitations preclude a detailed comparison of the conformations of (1) with those reported for Leu⁵-enkephalin. However, a good qualitative impression of the similarities and differences can be achieved from examination of the stereoscopic superposition diagram⁸ presented in Figure 2. This figure also illustrates the marked differences in conformation adopted by Leu⁵-enkephalin in crystals obtained from aqueous alcohol² (β -bend) and from aqueous dimethylformamide (DMF)³ (extended backbone). A second crystal modification of (1) has been obtained from aqueous DMF and is presently under investigation.

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