

## SHORT COMMUNICATION

*Contributions intended for publication under this heading should be expressly so marked; they should not exceed about 1000 words; they should be forwarded in the usual way to the appropriate Co-editor; they will be published as speedily as possible.*

*Acta Cryst.* (1981). **B37**, 1161

**X-ray structure analysis and molecular conformation of *N*-propionylproline: erratum.** BY M. E. KAMWAYA, O. OSTER and H. BRADACZEK, *Institut für Kristallographie, Freie Universität Berlin, Takustrasse 6, D-1000 Berlin 33, Federal Republic of Germany*

(Received 16 March 1981)

### Abstract

In the paper by Kamwaya, Oster & Bradaczek [*Acta Cryst.* (1981), **B37**, 364–367] the first sentence on p. 367 should read: 'The  $N'C_4C_6C_7$  group is fairly planar.'

All the relevant information is contained in the *Abstract*.

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### Commission on Journals

#### Standards for the Publication of Powder Pattern Data

Standards for the publication of powder patterns, originally compiled by a subcommittee of the American Crystallographic Association and published in *National Bureau of Standards Special Publication 567* (1979), have been accepted by the Commission on Crystallographic Data and the Commission on Journals. Papers that present powder pattern data submitted for publication in IUCr journals are now required to follow these standards. Full details are given in *Acta Crystallographica*, Section A [*Acta Cryst.* (1981), **A37**, 443–444.]

*Acta Cryst.* (1981). **B37**, 1161–1162

### Commission on Journals

#### Deposition of Macromolecular Atomic Coordinates and Structure Factors with the Protein Data Bank

The policy that all structural papers, including low-resolution protein studies, should be equally subject to the requirement of deposition of atomic coordinates and lists of

structure factors was recently reaffirmed by the Commission on Journals. It was also decided that such depositions should be machine-readable, and an agreement has now been reached with the Protein Data Bank, Chemistry Department, Brookhaven National Laboratory, Upton, New York 11973, USA to incorporate these data in their system.

Authors of macromolecular structure determination manuscripts will, in future, be requested by their Co-editor to send their atomic coordinates and structure factors to the Protein Data Bank. A macromolecule is regarded as any substance containing a polypeptide, polynucleotide or polysaccharide chain longer than 25 residues. The manuscript should be accompanied by printed copies of the atomic coordinates which will be sent to the referees. The Protein Data Bank will acknowledge receipt of the machine-readable data both to the author and the Co-editor and will, at the same time, communicate to both the Reference number assigned to these data. As soon as the data have been processed into standard Data Bank format and the corresponding listings verified by the author, notification will be sent to the Co-editor. Such listings can usually be prepared in less than one month, with the cooperation of the author. Final acceptance of the manuscript requires that the atomic coordinates and structure factors have indeed been satisfactorily processed by the Data Bank. The data will also be deposited, as microfiche, with the British Library under the IUCr Supplementary Publication Scheme and may be requested from the Executive Secretary of the IUCr by quoting the Supplementary Publication Number in the usual way.

All information received by the Data Bank under this arrangement will be regarded as privileged and will not be released to others until the paper has been accepted or prior release has been authorized by the authors. Magnetic tape