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Processing incommensurately modulated protein diffraction data with *Eval15*. Corrigendum

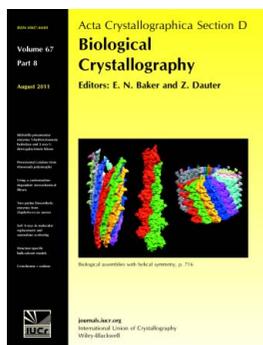
Jason Porta, Jeffrey J. Lovelace, Antoine M. M. Schreurs, Loes M. J. Kroon-Batenburg and Gloria E. O. Borgstahl

Acta Cryst. (2011). **D67**, 745

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Acta Crystallographica Section D: Biological Crystallography welcomes the submission of papers covering any aspect of structural biology, with a particular emphasis on the structures of biological macromolecules and the methods used to determine them. Reports on new protein structures are particularly encouraged, as are structure–function papers that could include crystallographic binding studies, or structural analysis of mutants or other modified forms of a known protein structure. The key criterion is that such papers should present new insights into biology, chemistry or structure. Papers on crystallographic methods should be oriented towards biological crystallography, and may include new approaches to any aspect of structure determination or analysis. Papers on the crystallization of biological molecules will be accepted providing that these focus on new methods or other features that are of general importance or applicability.

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addenda and errata

Processing incommensurately modulated
protein diffraction data with *Eva/15*.
CorrigendumJason Porta,^{a,b} Jeffrey J. Lovelace,^a Antoine M. M.
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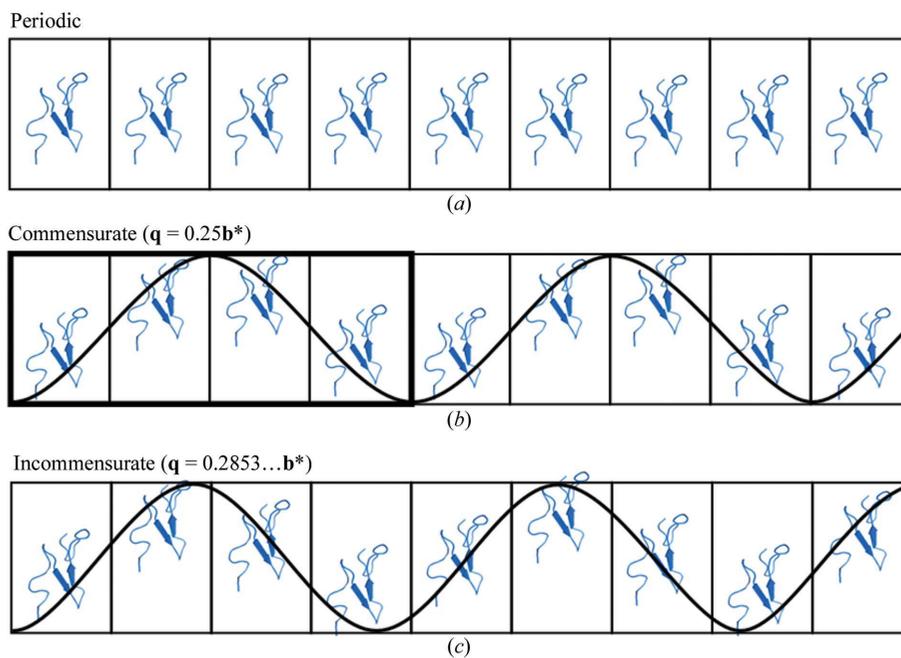
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A correction is made to a figure in the article by Porta *et al.* [(2011). *Acta Cryst. D67*, 628–638].

In the article by Porta *et al.* (2011) Fig. 2 was incorrect and the sine/cosine waves superimposed on panels (b) and (c) did not illustrate the correct frequencies. A corrected version of Fig. 2 is given here.

References

Porta, J., Lovelace, J. J., Schreurs, A. M. M., Kroon-Batenburg, L. M. J. & Borgstahl, G. E. O. (2011). *Acta Cryst. D67*, 628–638.

**Figure 2**

Three categories of crystals. (a) Periodic case with identical unit cells. (b) Commensurate modulation. (c) Incommensurate modulation with harmonic modulation wave. Protein structure from PDB entry 2rro.