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

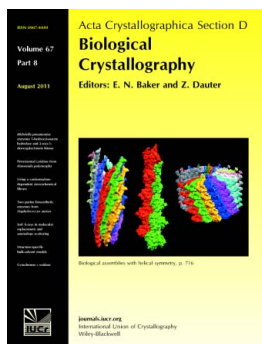
**Jason Porta, Jeffrey J. Lovelace, Antoine M. M. Schreurs, Loes M. J. Kroon-Batenburg and Gloria E. O. Borgstahl**

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

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## addenda and errata

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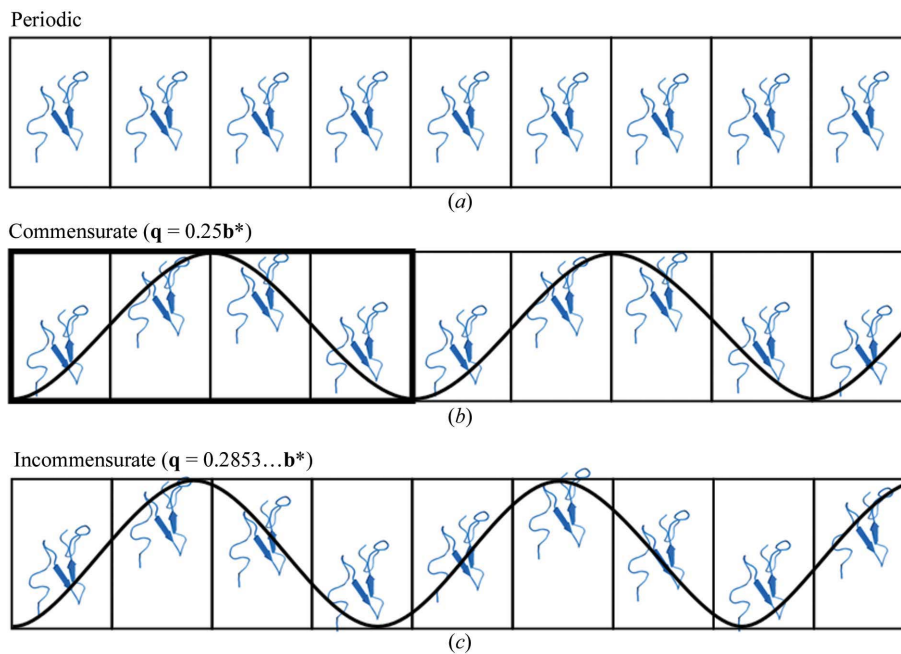
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A correction is made to a figure in the article by Porta *et al.* [(2011). *Acta Cryst. D67*, 628–638].

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.