



## Welcoming two new Co-editors

**Charles S. Bond, Elspeth F. Garman and Randy J. Read**

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*Acta Cryst. D – Structural Biology* continues to welcome articles covering all aspects of structural biology, especially structures of biological macromolecules or the methods used in their determination. Growth of new areas remains a focus, including neutron diffraction, application of artificial intelligence in structural biology, cryoEM, and electron crystallography. This month we are delighted to welcome two new Co-editors to *Acta Cryst. D* to help cover the breadth of structural biology we hope and aim to publish.



**Figure 1**  
Our new Co-editors, Dean Myles and Melanie Vollmar.

Dr Melanie Vollmar is currently an ARISE Marie Curie EMBL-EBI Fellow based at Hinxton in the UK. She is a structural biologist with particular expertise in the development of machine-learning and AI applications to automated data-analysis pipelines for macromolecular X-ray crystallography, and more recently in using AI to annotate PDB entries with relevant information. She has worked at Diamond Light Source, for the CCP4 Team, and has also interned at the IUCr Chester office as part of her ARISE Fellowship. She has wide experience of developing structural biology software tools, and we are delighted that she has joined the *Acta Cryst. D* team of Co-editors.

Dr Dean Myles is a member of the Distinguished Research Staff at Oak Ridge National Laboratory in Oak Ridge, Tennessee. His expertise is in neutron and X-ray diffraction of crystals of biological macromolecules, with an emphasis on technology development. He has a particular strength in hybrid approaches combining different diffraction and scattering methods with complementary biophysical techniques. With his expertise, he is an excellent addition to the *Acta Cryst. D* Co-editor team.

These appointments will enhance and renew the *Acta Cryst. D* Editorial Board while increasing the representation from different disciplines and different geographical regions around the world.