

## Poster Presentation

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### *Displacement Parameter Restraints for Dealing with Limited Data*

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Standard crystallographic structure refinements employ anisotropic displacement parameters (ADPs) to represent the probability distribution of a scattering atom. Such distributions may be due to thermal motion of the atom and / or a spatial average of multiple discrete atomic positions. An anisotropic description of an atomic distribution requires six parameters, and - in cases where data is limited or poor quality - the optimal values of these parameters may be ill-defined. Application of restraints and constraints can impose some physical and chemical reality on the set of displacement parameters. Examples include those based on the Hirshfeld Rigid Bond Test [1], and more recently SHELXL's RIGU [2]. We have implemented these and other a.d.p. restraints in CRYSTALS [3], for introducing reasonable relationships amongst common arrangements of anisotropic atoms. Use of a priori information in the form of restraints must always be justified, and we present an assessment of the applicability of the new restraints against a large data set of high quality crystal structure determinations.

[1] Hirshfeld F. L. (1976) *Acta Cryst.* A32, 239–244, [2] Thorn A., Dittrich B. & Sheldrick G. M. (2012) *Acta Cryst.* A68, 448–451, [3] Betteridge P. W., Carruthers J. R., Cooper R. I., Prout K., & Watkin D. J. (2003) *J. Appl. Cryst.* 36, 1487

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