## MS27-O5 Improving rotation electron diffraction data quality by data merging

Yunchen Wang<sup>1</sup>, Xiaodong Zou<sup>1</sup>, Wei Wan<sup>1</sup>

 Berzelii Center EXSELENT on Porous Materials and Inorganic and Structural Chemistry, Department of Materials and Environmental Chemistry, Stockholm University, SE-106 91 Stockholm

### email: yunchen.wang@mmk.su.se

Automated three-dimensional electron diffraction methods, for example automated diffraction tomography (ADT)[1] and rotation electron diffraction (RED) [2-3], can be used to collect and process completed 3D electron diffraction data from nanometer sized crystals and have been very successful in application of solving complex structures. However, limited crystal tilting range and steps, electron beam damage and dynamical scattering of electrons degrade the RED data quality and result in high R values and inaccurate structural parameters from refinement. Often structure models obtained from RED data are refined against powder X-ray diffraction data for confirmation and better accuracy.

Here we propose an approach for improving RED data quality through merging multiple datasets collected from different crystals with different orientations. Individual datasets are first processed using RED following the standard procedure. The initial model for the refinement is obtained by direct methods using SHELXS. The HKL lists from the individual datasets are merged together for refinement with the initial model using SHELXL. To study the effects of merging datasets on data quality, we performed refinement by sequentially including the individual datasets from a test sample. We show that the number of observed reflections increases and the R values decrease while more datasets are merged. The structural parameters after refinement are improved by merging more datasets. The apparent improved data quality may be attributed partly to the increased data completeness and resolution, and reduced effedcts of dynamic effects through averaging. These factors will be examined carefully to evaluate their potential in improving RED data

- [1] U. Kolb et al, Ultramicroscopy 107 (2007) 507
- [2] D.L. Zhang, et al, Z. Kristallogr. 225 (2010) 94
- [3] W. Wan et al, J. Appl. Crystallogr. 46 (2013) 1863

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# MS28 New approaches in electron crystallography

Chairs: Louisa Meshi, Xiaodong Zou

### MS28-O1 When Precession Electron Diffraction Tomography goes dynamical

Philippe BOULLAY<sup>1</sup>, Gwladys Steciuk <sup>1</sup>, Olivier Pérez <sup>1</sup>, Sébastien Petit <sup>1</sup>, Moussa Zaarour <sup>2</sup>, Svetlana Mintova <sup>2</sup>, Karl Rickert <sup>3</sup>, Kenneth R. Poeppelmeier <sup>3</sup>, Wenrui Zhang <sup>4</sup>, Haiyan Wang <sup>4</sup>, Mariana Klementova <sup>5</sup>, Lukas Palatinus<sup>5</sup>

 Laboratoire de Cristallographie et de Sciences des Matériaux (CRISMAT) UMR CNRS 6508, ENSICAEN, 6 Boulevard Maréchal Juin, 14050 Caen, France

2. Laboratoire de Catalyse et de Spectroscopie (LCS), ENSICAEN, 6 Boulevard Maréchal Juin, 14050 Caen, France

3. Department of Chemistry, Northwestern University, Evanston, Illinois, 60208, USA

4. Department of Materials Science and Engineering, Texas A&M University, College Station, Texas 77843, USA

5. Institute of Physics of the AS CR, Na Slovance 2, Prague, Czech Republic

### email: philippe.boullay@ensicaen.fr

The use of electron diffraction (ED) data for the refinement of crystal structures as routinely as what is done in X-ray diffraction (XRD) has long appeared to be inaccessible. While the strong electron/mater interaction allows studying crystals of nanometer size, the kinematic approximation is not valid, which makes the analysis of the diffracted intensity more complex than for XRD. More complex, but not impossible, as the formalism to describe dynamic conditions in ED is known since long. However, its use by programs aiming structural refinements using ED patterns obtained from parallel beam met no success despite several attempts. [1]

In 2013 [2], L. Palatinus and co-authors have studied the feasibility of refining crystal structures from few zone axis patterns considering the dynamic conditions of diffraction, and shown, the importance of Precession Electron Diffraction (PED) [4] to get better convergence. Early in 2015, the Prague group reached a decisive step by implementing in JANA2006 a module dedicated to ED and such "dynamical refinements" [5]. Integrated in an interface familiar to most structural crystallographers, this module now allows to process PED data collected in a "tomography mode" [6] aiming to reach a high level of data completeness. Several groups across Europe have joined forces to test and validate this new approach and offer the community a reliable and functional tool. [7]

From structure solution to structure refinement (kinematical then dynamical), Precession Electron Diffraction Tomography (PEDT) is a mature technique readily to be used in any materials science laboratory. In the present communication, we will show what astonishing results can be achieved and what are the limitations using examples from incommensurate