Electron diffraction provides new insights into chemistry at the atomic level

Tim Grüne

Head Dr. Centre for X-ray Structure Analysis University of Vienna

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Facultad de Ciencias, Universidad de Zaragoza - CSIC C/ Pedro Cerbuna, 12. Zaragoza 50009. Spain





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Tim Gruene,ª

^aUniversity of Vienna

e-mail: tim.gruene@univie.ac.at

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X-ray diffraction is an important analytical method to visualise the atomic positions of many chemical compounds. It is applied in many disciplines in Chemistry, Materials Science, Pharmacology, and many others. Recently, electron diffraction has attracted great attention ¹, known as 3D ED², or microED³. 3D ED provides the same type of information as an X-ray structure, including the atom coordinates, lengths of bonded and non-bonded contacts, differentiation of elements, and determination of the absolute configuration of chiral compounds. Where X-ray diffraction requires crystals with at least a few µm in length, 3D ED determines the crystal structure from submicrometer sized crystals, even smaller than individual grains in powder. This is useful in cases where large crystals cannot be grown.

Instrumentation for 3D ED is based on transmission electron microscopes. Academic studies had to provide workarounds, because the control software lacks features essential for crystallographic data acquisition. There are now two commercial systems (ELDICO and RIGAKU), that change the situation, and provide a user interface optimised for crystallography.

This talk will focus on our realisation for an instrument, in particular the use of the JUNGFRAU

detector ⁴. Originally designed for X-ray Free Electron studies, this detector reveals an enhanced sensitivity, and provides us with great level of details. Our examples includes organic compounds, metal organic framework, and zeolites.

The zeolite A, and also the aluminosilicate albite acted as benchmark for the JUNGFRAU detector. Our data differentiated aluminium from silicon in two aluminosilicates, zeolite A and albite, a result which is difficult to achieve even with X-ray data.

Our latest study benefited from the superior quality of the JUNGFRAU detector. We localised coke in the pores of ZSM5, and we monitored the process of coking in the micropores.

Finally, I will also share my experience from sample preparation. Because samples are not visible with lightmicroscopes, one has to prepare them blindly, until the TEM reveals the quality of the grid. This might be a bottleneck, but is mainly a matter of experience.

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CV:

as of Feb. 2019 Head of the <u>X-ray Structure Analysis Centre</u> of the Faculty of Chemistry at the University of Vienna. The name "Centre for Chemical Structure Analysis" is closer to what we do, hence the abbreviation and url CCSA

Aug. 2015-Dec. 2018 Senior Staff Scientist at the the Paul Scherrer Institut

2007-2015 Staff Scientist with G. M. Sheldrick's group

2007 Sabbatical during the commissioning of the Macromolecular Beamlines at the Australian Synchrotron 2003-2007 post-doc position in George M. Sheldrick's group, University of Göttingen. Main research topic: automated tracing of nucleic acids

1999 - 2003 PhD at the EMBL Outstation Grenoble