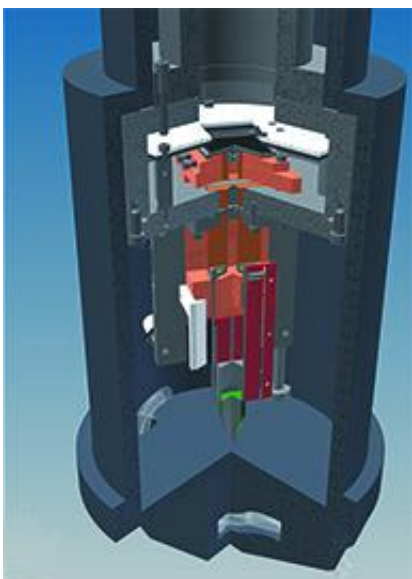


Determining the structures of nanocrystalline pharmaceuticals by electron diffraction

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90° cut-off schematic of the camera as designed for the CM200 (Technische Universiteit Delft, The Netherlands). Credit: van Genderen et al

Reliable information about the structure of pharmaceutical compounds is important for patient safety, for the development of related drugs and for patenting purposes. However, working out the structures of pharmaceuticals can be tough. The individual molecules can pack together in the solid in different ways to form different polymorphs, and pertinent properties such as stability, bioavailability or how fast they dissolve in the stomach can vary from one polymorph to another. Single crystals (as used in standard X-ray diffraction experiments) therefore

might not be representative of the bulk sample, or indeed might not even be available.

Moreover, the compounds themselves can be damaged by the high energy of the X-radiation used. As electrons are less damaging than X-rays by several orders of magnitude, using electron diffraction should be an attractive alternative, particularly when only nanometre-sized crystals are available. Cooling the sample to liquid-nitrogen temperatures ('cryo-cooling') can also help to minimize [radiation damage](#), but the compound might change structure on cooling, so the structure that is obtained is not actually that of the material as taken by the patient at [room temperature](#).

A group of scientists from a number of European countries have tackled all aspects of these problems by using low-dose electron diffraction, rotating the sample so that individual nanocrystals are not in the electron beam long enough to be damaged and collecting the [diffraction data](#) using a new type of detector developed by CERN [van Genderen et al. (2016), Acta Cryst. A72, [DOI: 10.1107/S2053273315022500](https://doi.org/10.1107/S2053273315022500)]. This new detector combines a [high dynamic range](#) with a very high signal-to-noise ratio and sensitivity to single electrons. Radiation damage was reduced so much that cooling the sample was not found to be necessary, allowing the team to study the anticonvulsant drug carbamazepine and nicotinic acid (vitamin B3) at room temperature. The data they collected were high enough quality that they could solve the structures of the two compounds using direct methods and software developed for X-ray crystallography.

Based on their experience with these case studies, the authors are planning to improve the design of their experimental setup further, and will also be developing programs specifically designed for handling electron-diffraction data.

More information: E. van Genderen et al. structure determination of

nanocrystals of organic pharmaceutical compounds by electron diffraction at room temperature using a Timepix quantum area direct electron detector , *Acta Crystallographica Section A Foundations and Advances* (2016). [DOI: 10.1107/S2053273315022500](https://doi.org/10.1107/S2053273315022500)

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