

Publication**Reducing dynamical electron scattering reveals hydrogen atoms****JournalArticle (Originalarbeit in einer wissenschaftlichen Zeitschrift)****ID** 4530684**Author(s)** Clabbers, Max T. B.; Gruene, Tim; van Genderen, Eric; Abrahams, Jan Pieter**Author(s) at UniBasel** [Abrahams, Jan Pieter](#) ;**Year** 2019**Title** Reducing dynamical electron scattering reveals hydrogen atoms**Journal** Acta crystallographica. Section A, Foundations and advances**Volume** 75**Number** Pt 1**Pages / Article-Number** 82-93**Keywords** dynamical scattering; electron diffraction; hybrid pixel detector; hydrogen atoms; nanocrystals**Mesh terms** Crystallography, methods; Electrons; Hydrogen, chemistry; Likelihood Functions; Models, Molecular; Molecular Structure; Nanoparticles, chemistry; Proteins, chemistry; Scattering, Radiation

Compared with X-rays, electron diffraction faces a crucial challenge: dynamical electron scattering compromises structure solution and its effects can only be modelled in specific cases. Dynamical scattering can be reduced experimentally by decreasing crystal size but not without a penalty, as it also reduces the overall diffracted intensity. In this article it is shown that nanometre-sized crystals from organic pharmaceuticals allow positional refinement of the hydrogen atoms, even whilst ignoring the effects of dynamical scattering during refinement. To boost the very weak diffraction data, a highly sensitive hybrid pixel detector was employed. A general likelihood-based computational approach was also introduced for further reducing the adverse effects of dynamic scattering, which significantly improved model accuracy, even for protein crystal data at substantially lower resolution.

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