

CCP-EM

A multi-platform suite of computational tools covering all aspects of cryoEM data processing, from image manipulation to building atomic models.



The Collaborative Computational Project for Electron cryo-Microscopy (CCP-EM) was initiated in 2012 to support the computational needs of the electron cryo-microscopy (cryoEM) community. CCP-EM is mandated to provide user training and developer support and to establish a coherent community for the exchange of best practices and novel ideas.

The CCP-EM software suite is a multi-platform suite of tools that aims to cover all aspects of cryoEM data processing from image manipulation to the building of atomic models, and to cover multiple techniques such as single-particle reconstruction (SPR), tomography and diffraction.

The software suite was conceived as a generic framework that could support a wide variety of functionalities, and has a modular organisation which can be divided into three layers. The top-level GUI layer is written using the PyQt toolkit. This provides a simple graphical interface to the associated programs. Distinct from this is the mid-level management layer, which is written in pure Python. This provides a bridge between the GUI layer and the third layer: the set of functional programs. These programs originate from collaborating developers and are written in a wide variety of languages (including C, C++, Fortran and Python) with distinct control methods and input conventions.

References

1. Tom Burnley, Colin M. Palmer and Martyn Winn (June 2017), <https://journals.iucr.org/d/issues/2017/06/00/rr5147/index.html>, Acta Cryst., 73, 469-477
2. Chris Wood, Tom Burnley, Ardan Patwardhan, Sjors Scheres, Maya Topf, Alan Roseman and Martyn Winn (January 2015), <https://journals.iucr.org/d/issues/2015/01/00/gm5035/index.html>, Acta Cryst, 71, 123-126

Category

Software/CCP-EM

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