Ultra-high Throughput Macromolecular Crystallography **Data Collection Using the Bluesky Framework** D. Perl, N. Frisina, D. Oram, N. Paterson, Diamond Light Source, Didcot, UK

MX data collection



•Align a protein crystal with the X-ray beam. Capture diffraction images as the crystal is rotated.

Bluesky¹

 Modern Python library for experiment control. • Developed at NSLS-II, in collaboration with other facilities including DLS.

•Used for fragment-based drug discovery. •Completely automated at some beamlines.



A

В

Resolution: 1.43Å

 Good separation of harware interaction and data management. • Easy integration with other scientific Python libraries.







Application for unattended data collection, built with Bluesky. • Uses DODAL, a library to contain and organise all hardware at DLS. • Part of the new DLS Athena² platform to support Diamond-II. • Step-by-step replacement of routines in the legacy data acquisition software. Modular design follows naturally from the way Bluesky works. • Well tested against beamline simulations. • Currently in transition from prototype to full version.

Hyperion



References

Faster data acquision

- A fragment screening experiment takes hundreds or thousands of datasets.
- Hyperion will support new kinds of experiments allowing

[1] D. Allan, T. Caswell, S. Campbell, and M. Rakitin, 'Bluesky's Ahead: A Multi-Facility Collaboration for an a la Carte Software Project for Data Acquisition and Management', Synchrotron Radiat. News, vol. 32, no. 3, pp. 19–22, May 2019, doi: 10.1080/08940886.2019.1608121.

[2] J. Shannon, C. Forrester, and K. Ralphs, 'Diamond Light Source Athena Platform', presented at the ICALEPCS'23, Cape Town, South Africa. Paper TH1BCO05

thousands of data collections per day.

• Initial prototype for centring has already increased throughput from ~380 to ~500 samples per day.

