Biostruct-X Integration: DIALS Synchrotron and Free-Electron Laser Diffraction Integration

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Diamond Light Source, CCP4

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Overview

- Background the Biostruct-X project
- What are we doing?
- Why are we doing this?
- How is this different?



Acknowledgements

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- Thinking: Andrew Leslie, Phil Evans, Gwyndaf Evans, David Waterman, Gleb Bourenkov, Garib Murshudov

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- Collaborating: Nick Sauter, Ralf Grosse-Kunstleve
- Writing: developer community at large

Biostruct-X

- EU framework 7 contract no. 283570
- Main focus of project on trans-national access to facilities for MX, SAXS etc.
- Workpackage 6: data integration and analysis for synchrotron and FEL crystallography



WP6 objectives

Develop data analysis suite which is forward compatible¹ to:

- correctly handle pixel array detectors
- deliver high speed
- handle high mosaicity, overlapping reflections
- cope with weak diffraction
- To attempt to unify where possible analysis of synchrotron and FEL data: unified tools for structural biologists, handling of coherence effects.
- Develop tools to manage massive data sets via
 pre-classification²





WP6 tasks

- 6.1 Development of common frameworks for data management, integration and results inspection (DLS, XFEL, CFEL.)
- 6.2 Development of key algorithm modules for integration program optimised for PAD and challenging cases (DLS, CFEL, XFEL, Dectris, PSI.)
- 6.3 Development of data simulation software and visualisation for raw data and results. Systematic count rate optimisation studies and new hybrid detector technology tests (EMBL-GR, DLS, HZB, CFEL, XFEL, DECTRIS, PSI.)
- 6.4 Development of integration pipeline to handle communication with up and down stream analysis software (DLS.)

 6.5 Release of integration software to all partner sites for testing (DLS, all.)

What are we doing? WP 6.1 and 6.2

Physics-based modelling of experiment

- Take maximum likelihood approach
- Work with smart people for algorithm design
- Emphasise global refinement
- Enable proper modelling of errors from 1st principles
- Extensible framework
- Build on existing tools
- Speed and smart algorithms





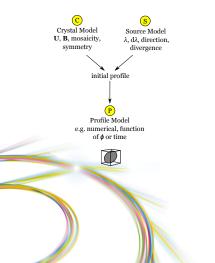




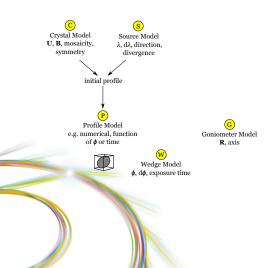




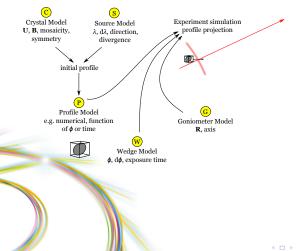




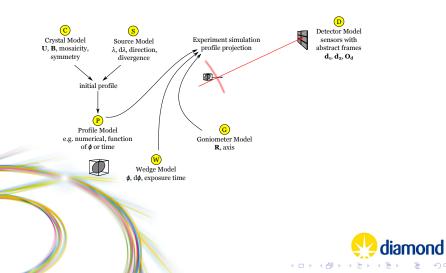


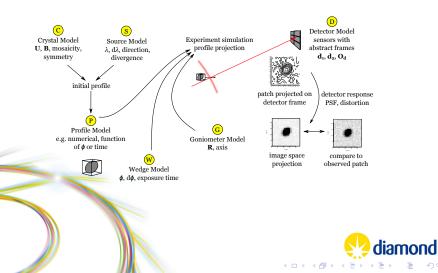












- Illustration had integration on detector surface, on Ewald sphere should also be designed in
- Correctly following equations will be critical for parameter refinement via ML - going back from the predicted and observed profiles to model adjustments



Extensible Framework

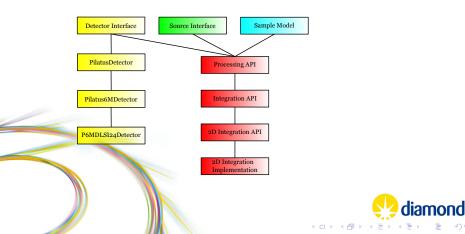
- Separate out things with proper abstraction layers should map to real world
- Design system to allow new detector types (and technologies) to be added dynamically c.f. dxtbx
- Also new models for other components e.g. profile modelling
- Options for 2D / 3D / hybrid profiles within same framework
- Synchrotron and free electron laser sources primarily, option
 for lab sources



Abstraction Layers

An illustration - how the API framework can be organised - which will evolve as the project progresses

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Aside - dxtbx

- Framework to allow *dynamic* detector support extension in xia2
- Wrote from the ground up as a toolbox for cctbx will import into iotbx
- If enough time later will describe this showing code, else can discuss this week



Everything in Library

Not a single user interface - allow operation through GUI, expert system, scripts.

EDNA / PyQT GUI / xia2 /	
Python Bindings	Python "wrapper" for C++ application
Library	C++ "main" application (binary)
User-defined extensions	(ondig)
Dependencies (crystallographic libraries, openCL, etc.)	



Existing Tools

- Work within and extend CCTBX framework open source by design, from the start, can piggy-back on their infrastructure e.g. subversion, testing framework etc.
- Collaborating with CCTBX developers, including Nick Sauter (NIH funded project)



Why are we doing this?

- Open source
- Future proof
- Framework for novel algorithm development
- Enable use of modern computing infrastructure

None of these goals are really special - they were desirable in MADNES in the 1980's - 1990's - however the computing hardware was not available at the time to allow this. Now, it is.



Next Steps / ongoing activity

- Define series of use cases (with Nick Sauter):
 - straightforward integration (summation) using masks defined by nearby peaks
 - xds-style "kabsch" transformation
- Defining model components, integrating into use case code
- Starting with well behaved Pilatus data from MX and small molecule crystallography
- Aim at this stage is to determine the nature of the problem,
 to inform design

