Xport Collaboratory for X-ray Crystallography

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Macromolecular Crystallography

Mission critical tool in biochemistry
 Structure of large biological molecules
 Function from structure

- Crystallographic experiment
 - Crystals diffract X-ray wavelength light
 - Diffraction pattern can be used to determine size of the unit cell of the crystal and location of atoms
 - For large molecules diffraction pattern is spatially compact requiring high image resolution
 - Requires high-brilliance X-ray source
 - Advanced Light Source (LBL)
 - Advanced Photon Source (ANL)

The Crystallographic Experiment



Problems

- Travel/presence
 - Beam time is hard to come by and expen\$ive
 Researcher must be at beam line for effective use of beam time
- Data management
 - Typical experiment generates 200-400 images at 5-20MB per image
 - Different researchers use different packages for analysis and visualization
 - File formats

Xport Architectural Goals

Framework for

- experiment setup and monitoring
 Data storage and management
 Data reduction and analysis
 Collaboration
- Leverage existing middleware and services
- Use existing hardware and infrastructure
- "Minimally invasive"
- Allow remote users to use the tools they are familiar with

Xport Architecture Goals (2)

- Leverage existing middleware and services
 - Globus for authentication and job execution
 - Common Component Architecture Toolkit (CCAT) for modularization and reuse
 - CCAT Active Notebook as focal point for data and process management
 - GARA for bandwidth management
 - HPSS for storage

Xport Architecture Goals (3)

Use existing hardware and infrastructure Add "Proxy-cache" to beamline data collection system – consistent interface Grid computing facilities ESNet and Abilene/I2 QoS testbeds Support multiple sites and users with different data collection systems and analysis software

Xport Architecture

Xport Crystallography Collaboratory



Globus

Grid resources (computers, storage, instruments)

The SC2000 Xport Testbed



Xport Testbed Components



Demonstration

 With crystallographers Dr. John Bollinger at the IU-MSC and Kia Huffman
 Source is diffractometer at IU
 Remote user at sc2000

