

#### CATALYZING SCIENTIFIC ADVANCEMENTS CUTTING-EDGE WORKFLOWS AND COLLABORATIVE COMPUTING AT THE ADVANCED PHOTON SOURCE

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Argonne National Laboratory is a U.S. Department of Energy laboratory managed by UChicago Argonne, LLC.



# **X-Ray Light Sources**

#### Indispensable tools in the exploration of matter

- Wavelengths of the emitted photons span a range of dimensions from atoms to cells
- Scientific user facilities
- 5 funded by the US Department of Energy -Basic Energy Sciences
- Serves thousands of users per year from academia, government, and industry
- Diverse communities: materials research, biology, geosciences, life sciences, security, and many more



Advanced Photon Source (APS) at Argonne National Laboratory



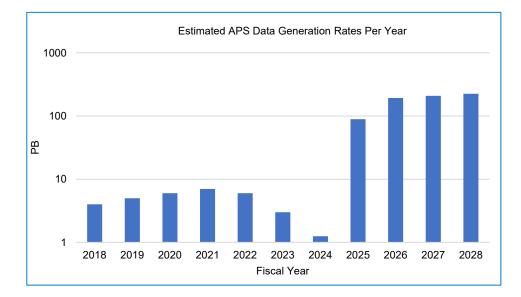


# THE UPGRADED ADVANCED PHOTON SOURCE

**Upgraded light source** up to 500x brighter than current APS for higher resolution Can image materials in less than a billionth of a second

#### Over the next decade the APS will

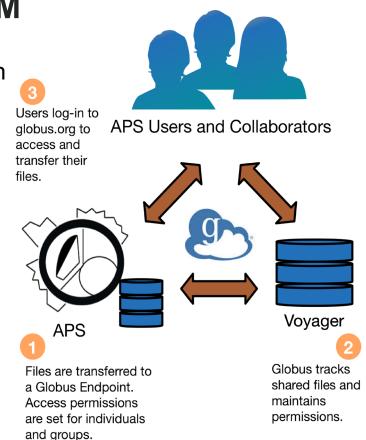
- Generate 100s of petabytes (PBs) of raw data per year
- Require 10s of petaflop/s of ondemand computing power for first pass data processing, reconstruction, and reduction





### **APS DATA MANAGEMENT SYSTEM**

- DM software is designed to help beamlines with various data management tasks
- Storage area management: movement of acquired data from local storage to a more permanent location, data archival, etc.
- Enabling users and applications to easily find and access data: metadata and replica catalog, remote data access tools
- Facilitating data processing and analysis with automated (in real-time) or user-initiated processing workflows







# **LEVERAGING GLOBUS SERVICES**

- Globus Connect Server 5 installed at central storage nodes allows secure access to data
- Globus Transfer provides data movement within flows and by users for post processing tasks
- Globus Compute offloads heavy computation to remote resources such as HPC
- Globus Flows executes complex data analysis pipelines
- Globus Groups secures access to experiment results in data portals
- Globus Auth executes flows as a beamline service account







## **BLUESKY INTEGRATION**



- Bluesky is an experiment specification and orchestration engine for control and collection of scientific data and metadata
- Bluesky interacts with hardware through a high-level abstraction
- Cross-facility, international collaboration
- Integration with the APS Data Management System means an end-to-end pipeline from detector to data portal



#### PARTNERING WITH THE ARGONNE LEADERSHIP COMPUTING FACILITY

- Service accounts for each beamline allow submitting jobs without each user maintaining their own compute allocations
- Trusted ALCF and APS Globus credentials to manage shared data access
- On-demand computing with preemption quickly executes APS jobs
- Dedicated log-in nodes for Polaris run Globus Compute Endpoints



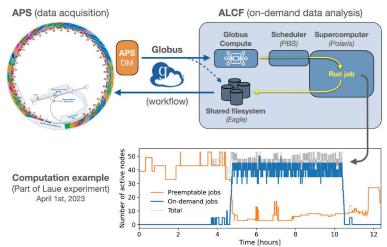
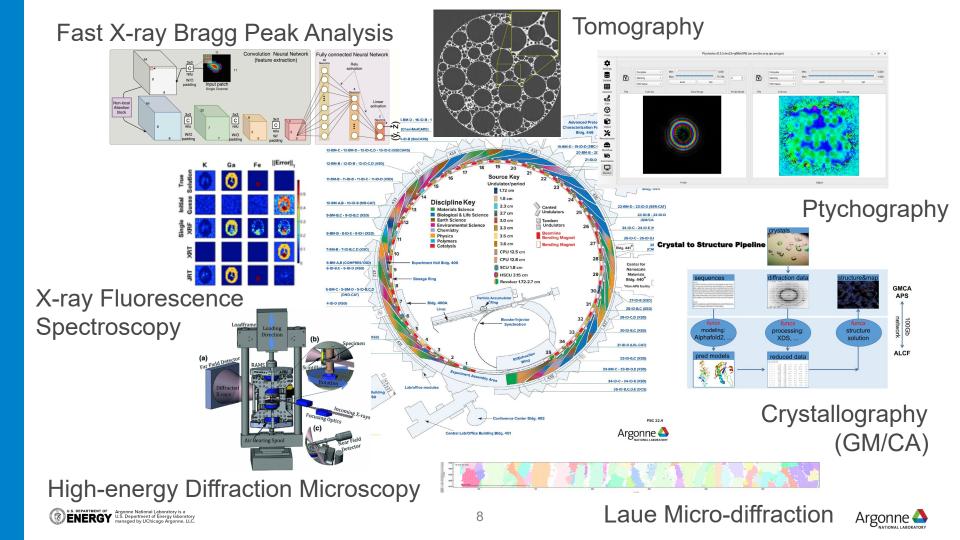


Diagram curtesy of Benoit Cote

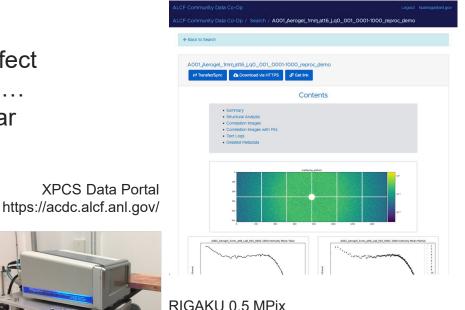






### X-RAY PHOTON CORRELATION SPECTROSCOPY AT BEAMLINE 8-ID-I

- Research outcomes include study of cartilage and joints, solar cells, flexible electronics, plastics, batteries, glass, effect of stress on the body, viruses and more...
- Hundreds of experiment users each year
- Over 400 TB data generated so far
- Hundreds of experiments processed



RIGAKU 0.5 MPix Ultra High Speed Sensor Q. Zhang *et al.,* J. Synchrotron Rad. (2021)





# **Adopting Digital Object Identifiers**

- A Digital Object Identifier (DOI) is a string of numbers, letters and symbols used to permanently identify an article or document and link to it on the web
- The Office of Science and Technology Policy's 2022 Public Access Memo states that federally funded research requires a unique digital persistent identifier for all scientific research and development awards
- DOIs created for 2023 beamtime awards



#### Structure determination of proteins from Sars-CoV-2

AWAF

https:/

RD	Award Description	
1 DDI: //doi.org/10.46936/APS-179063/6009021	Covid-19 Research Structural Analysis of proteins from Sars-Cov-2: Determine the crystal structures of listed proteins, in the complexes with small potential drug molecules, RNA, or other proteins from Sars-Cov-2. NSP3_PLP-LASCUDVS3 NSP9 NSP15 NSP16 NSP10 NSP8 NSP7 SGR_a SGR_b ADRP 3CLpro PLP Fabs mAbs	
	Investigator:	Gade, Priyanka ⑤ <sup>[1]</sup> ; Chang, Changsoo ⑤ <sup>[2]</sup> ; Finfrock, Yanhui ⑤ <sup>[2]</sup> ; Maltseva, Natalia ⑥ <sup>[2]</sup> ; Michalska, Karolina ⑥ <sup>[2]</sup> ; Joachimiak, Andrze <sup>[2]</sup> ; Kim, Youngchang <sup>[2]</sup> ; Lazarski, Krzysztof <sup>[2]</sup> ; Osipiuk, Jerzy <sup>[2]</sup> ; Tan, Kemin <sup>[2]</sup>
		1. The University of Texas at El Paso R 2. Argonne National Laboratory R
	Awarding Organization:	Advanced Photon Source (APS) R
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	Site Award Number:	179063
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- <u>https://github.com/AdvancedPhotonSource/XRF-Maps</u>



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