

Cyberinfrastructure for Autonomous Science

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Data Science and Learning

09 December 2019

Outline

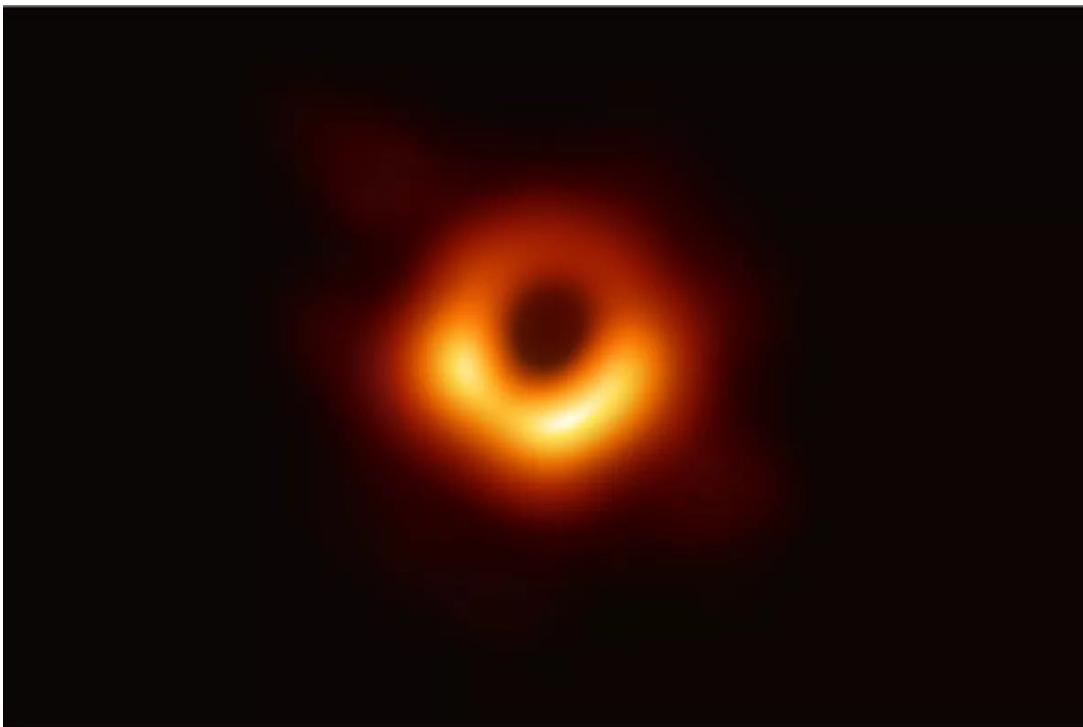
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- Scientific data management challenges
- Automation goals
- Cyberinfrastructure for automation
- Simplifying remote execution
- ML in the loop



Data management challenges as volumes increase

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Event Horizon Telescope

- 12 telescopes
- Telescopes generate roughly 900TB per 5 day run
- Data written to ~1000 HDDs
- Transported to MIT & Max Planck via airplane
- Aggregated and analyzed

Global resources, long timescales

Too much data for manual processing

Data loss due to HDD failure

Data management challenges as volumes increase

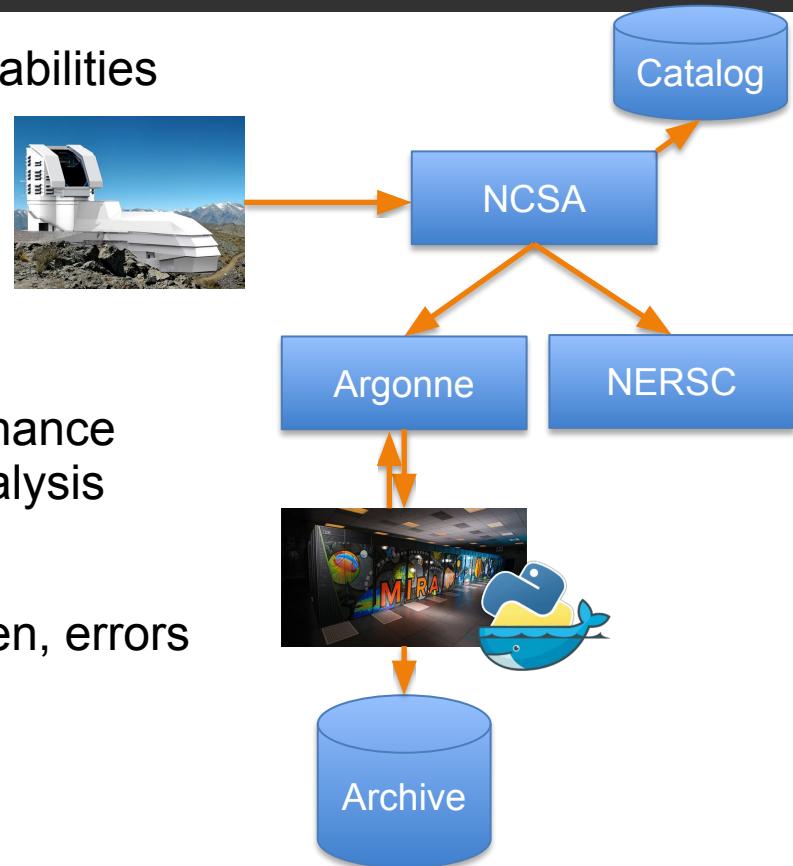
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Scientific data are overwhelming finite human capabilities

Scientific results are dependent on

- Data acquired at various locations/times
- Analysis processes executed on distributed resources
- Catalogs of descriptive metadata and provenance
- Dynamic collaborations around data and analysis

Best practices are overlooked, useful data forgotten, errors propagate through pipelines, ...



LSST data distribution and analysis pipeline

Experimental Science

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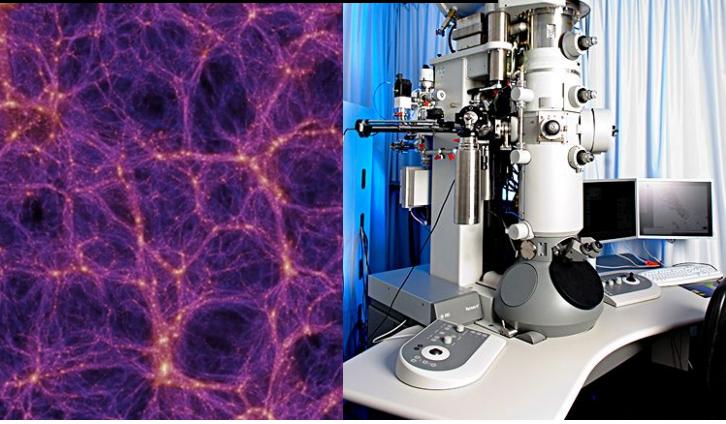
Data issues are particularly evident in large scale experimental science

Researchers are allocated short periods of instrument time

- Must maximize experiment efficiency and output data quality/accuracy
- Analysis requirements exceed local resource capabilities

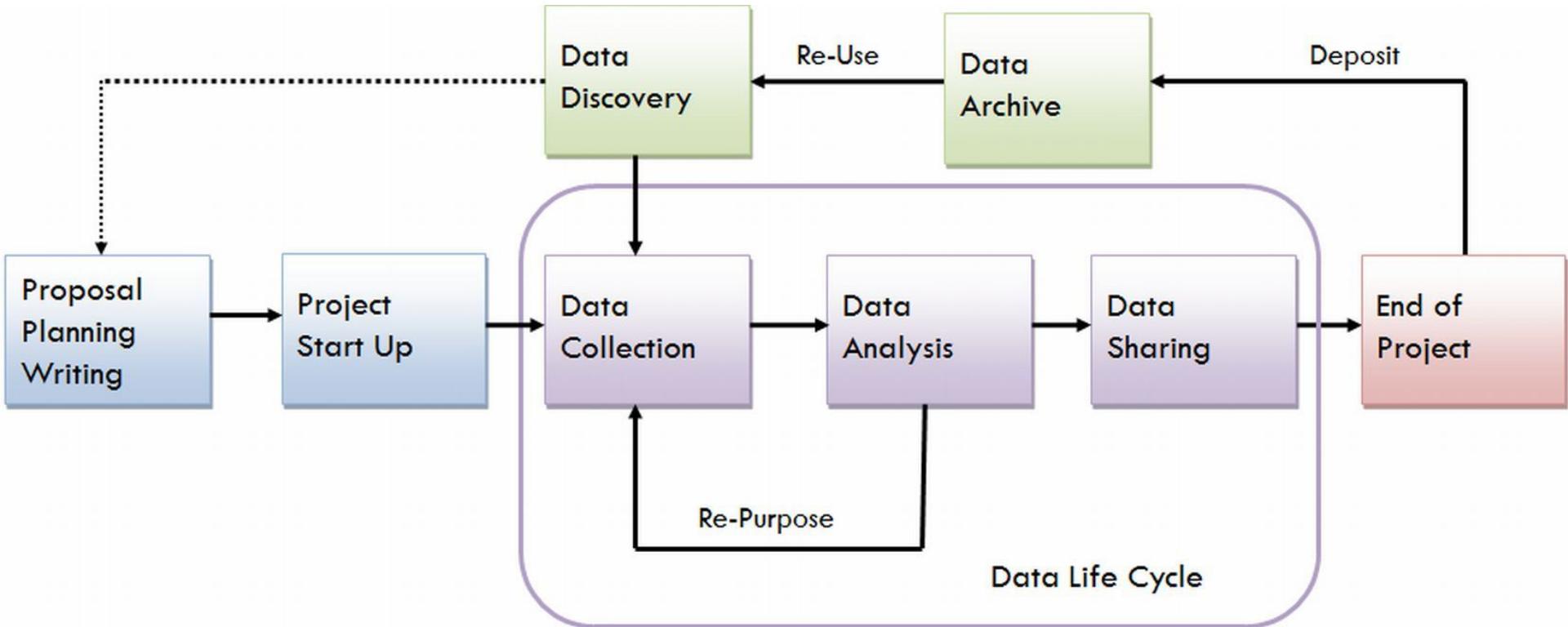
Inefficiencies mean less science is performed

Automating data lifecycle is key



Scientific Data Lifecycle

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<https://data.library.virginia.edu/data-management/lifecycle/>

Automation Use Cases

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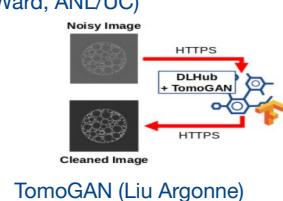
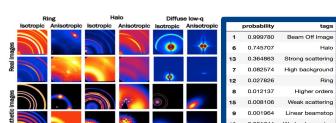
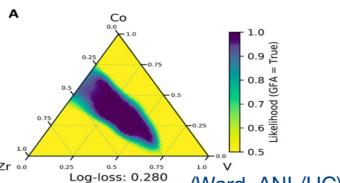
Best practices



Publication & archiving



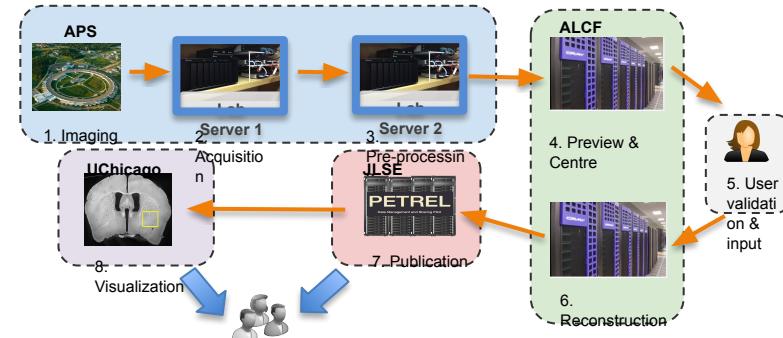
Model-in-the-Loop



XRD image tagging (Yager, BNL)

TomoGAN (Liu Argonne)

Analysis



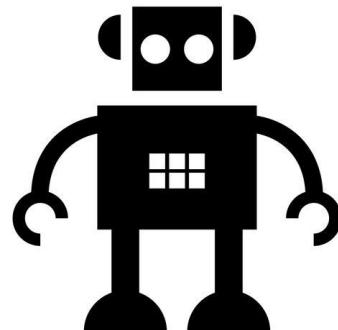
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Orchestrating End-to-End Automations

Requirements

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- Data driven: automation in response to data events
- Accessible to any scientist
- Scalable, secure, and fault tolerant
- Applicable to laptops and supercomputers
- Reliable end-to-end pipelines
- Supports human-in-the-loop timescales



Globus Automate

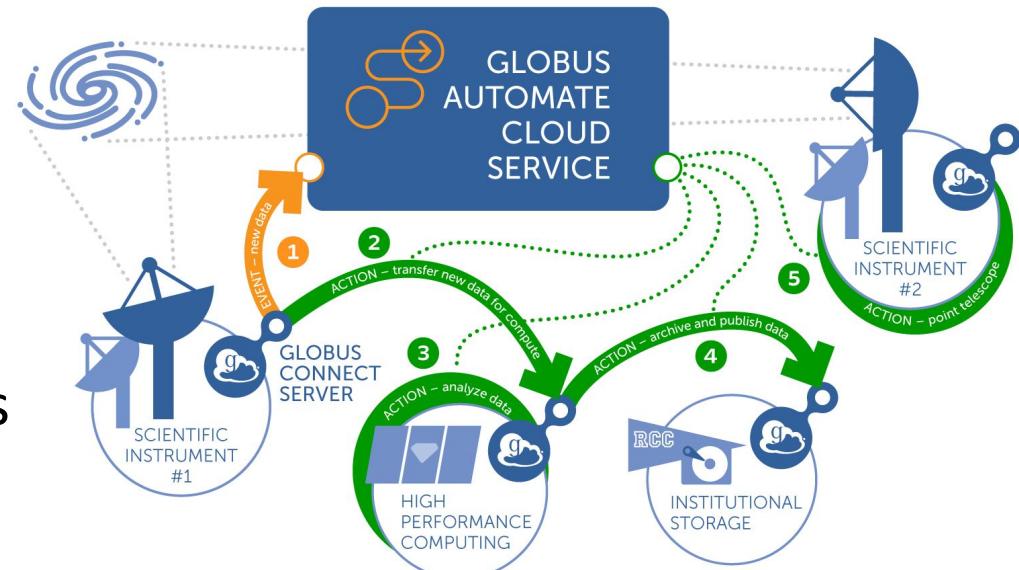
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A distributed automation PaaS

Compose pipelines that *step* between Web services

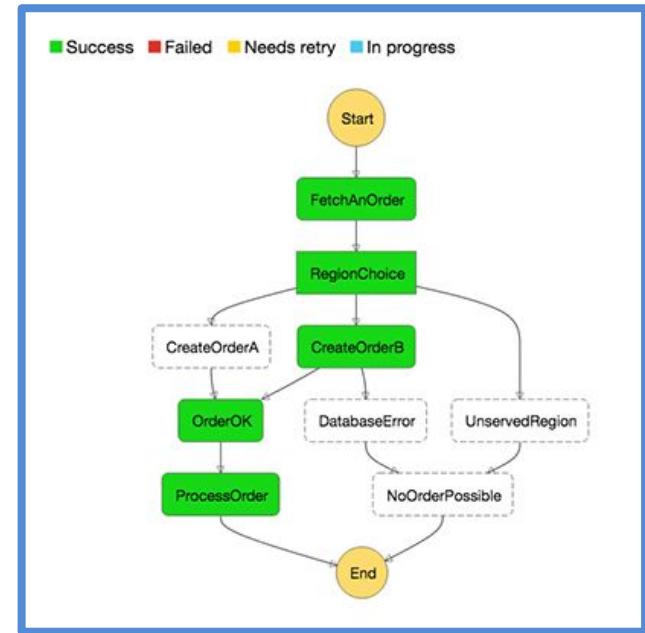
Supports user-defined actions

Integrates Globus Auth at every step



Globus automation capabilities

- Built on AWS Step Functions
 - Simple JSON-based state machine language
 - Conditions, loops, fault tolerance, etc.
 - Propagates state through the flow
- Standardized API for integrating custom event and action services
 - Actions: synchronous or asynchronous
 - Custom Web forms prompt for user input
- Actions secured with Globus Auth



Automate Actions

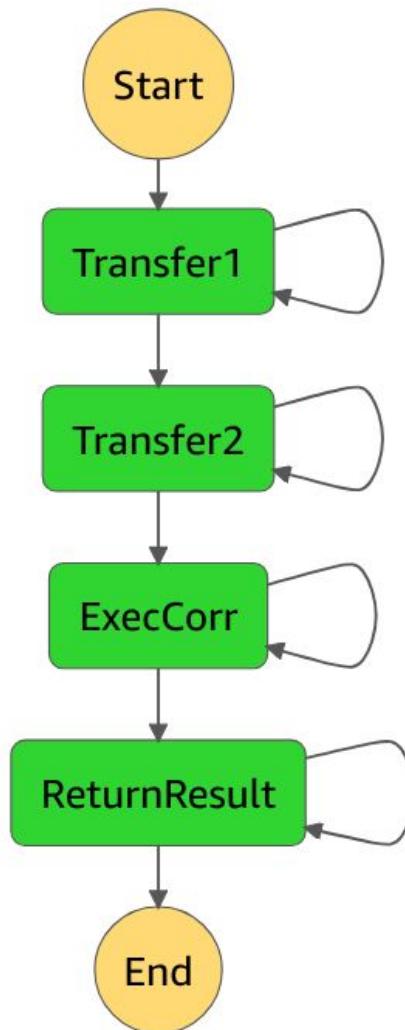
Any Web service can expose the Action API

- /automate/v1/<action>/run, status, cancel, introspect,
- .../status used to enable polling

Framework to create actions

Actions are asynchronous, timeout after 1 year

- Supports human-in-the-loop tasks



Automate Flows

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Definition extends SFN State
Machine Language

Definitions are transformed into
flows, then into state machines

- Adds auth, async, etc.
- Returns uuid

Accepts JSON blob as input

```
flow_definition = {
    "Comment": "Two step transfer",
    "StartAt": "Transfer1",
    "States": {
        "Transfer1": {
            "Comment": "Initial Transfer from Cam",
            "Type": "Action",
            "ActionUrl": "https://actions.automat",
            "ActionScope": "https://auth.globus.co",
            "InputPath": "$.Transfer1Input",
            "ResultPath": "$.Transfer1Result",
            "Next": "Transfer2"
        },
        "Transfer2": {
            "Comment": "Transfer from DMZ to data",
            "Type": "Action",
            "ActionUrl": "https://actions.automat",
            "ActionScope": "https://auth.globus.co",
            "InputPath": "$.Transfer2Input",
            "ResultPath": "$.Transfer2Result",
            "End": True
        }
    }
}
```

Running Flows

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```
flow_input = {
    "Transfer1Input": {
        "source_endpoint_id": "go#ep1",
        "destination_endpoint_id": "go#ep2",
        "transfer_items": [
            {
                "source_path": "/~/globus_publish.auto",
                "destination_path": "/~/globus_publish_2.auto",
                "recursive": False
            }
        ]
    },
    "Transfer2Input": {
        "source_endpoint_id": "go#ep2",
        "destination_endpoint_id": "go#ep1",
        "transfer_items": [
            {
                "source_path": "/~/globus_publish_2.auto",
                "destination_path": "/~/globus_publish_3.auto",
                "recursive": False
            }
        ]
    }
}
```

Authenticate

Specify input: \$.Transfer1Input

```
fc.run_flow(flow_id,
flow_scope, flow_input)
```

Monitor

Automate Actions

Auth



User login

Secure service interactions

App identity and interactions

Identifier



Manage namespace

Mint DOI

Search



Catalog

Faceted search

Search query

Transfer



File operations

Transfer data

Set permission

funcX



Remote execution

Secure connections

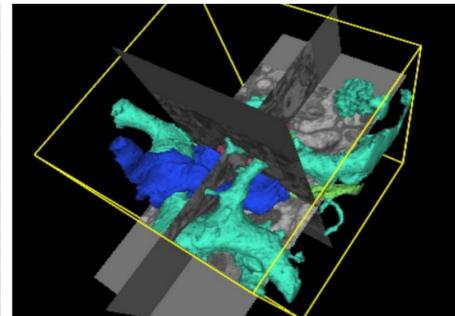
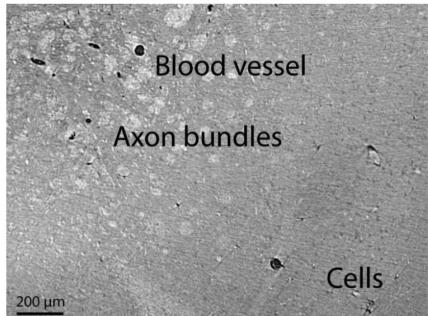
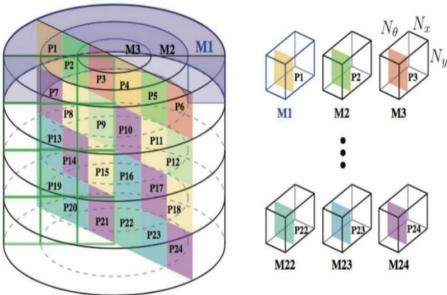
Isolated sandboxes

End-to-end Analysis Pipeline

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UChicago's Kasthuri Lab study brain aging and disease

- Construct connectomes -- mapping of neuron connections
- Use synchrotron (APS) to rapidly image brains (and other things)
- Beam time once every few months
- Generate segmented datasets/visualizations for the community
- ~20GB/minute for large (cm) unsectioned brains
- Perform semi-standard reconstruction on all data across HPC resources



Neuroanatomy automation

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APS



1. Imaging



2. Acquisition



3. Pre-processing

ALCF



4. Preview & Centre

UChicago



8. Visualization

JLSE



7. Publication



5. User validation & input

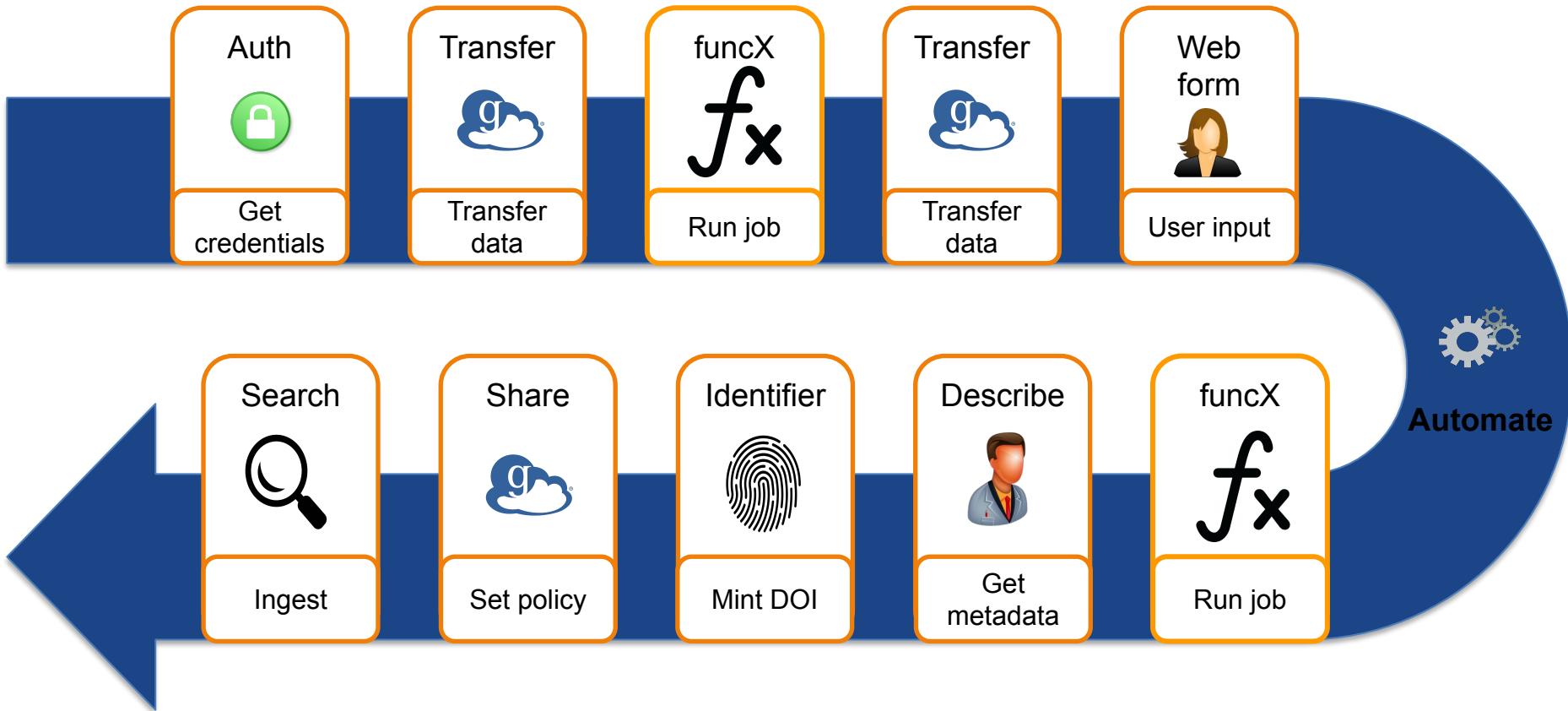


6. Reconstruction



Neuroanatomy automation

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Experimental Control

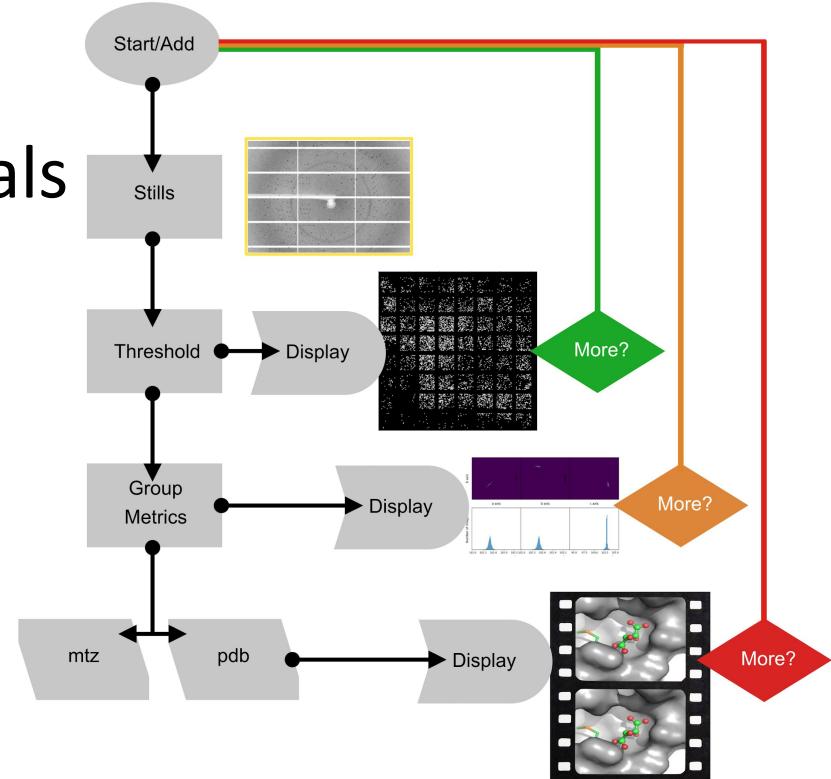
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Serial crystallography

- Serially image 26000 crystals

Quality control first 1000

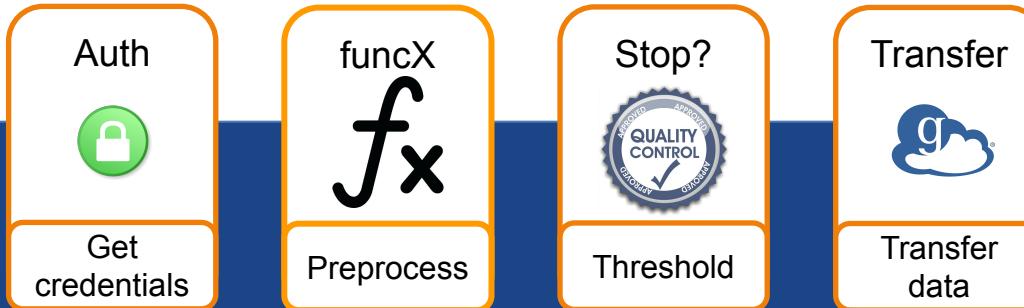
Analyze full 26,000



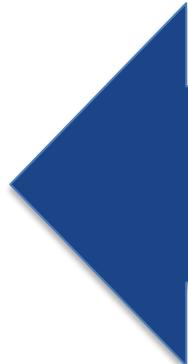
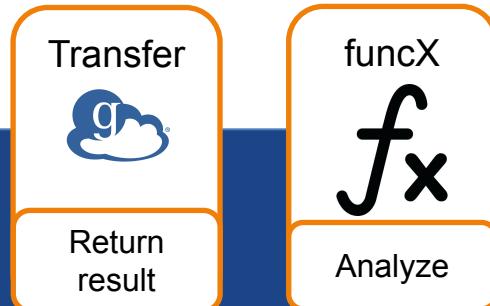
Return crystal structure to scientist

SSX Automation

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Automate





Compute Abstractions: From Laptops to Supercomputers

Laptops to Supercomputers

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Automation tasks need to be applied to laptops, desktops, local clusters, cloud, and supercomputers

Heterogeneous resources, permissions, environments, ...

Need a common way to interact with arbitrary resources and perform tasks

Serverless computing

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Serverless computing is revolutionizing enterprise IT

Function as a Service (FaaS)

- Pick a runtime (python/JS/R etc.)
- Write function code
- Run at any scale

Low latency, on-demand

Abstracts computing infrastructure



funcX: Serverless Supercomputing

Turn any machine into a function serving platform

Remove barriers to HPC

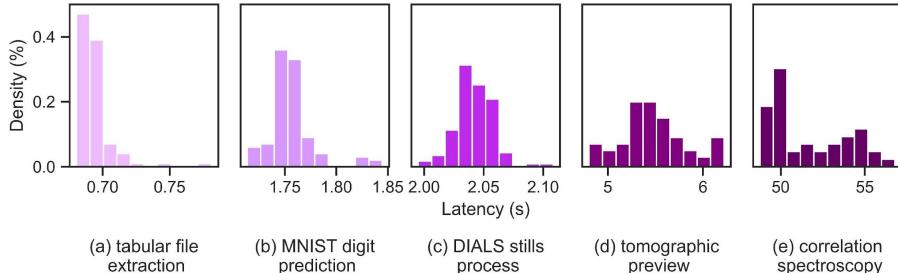
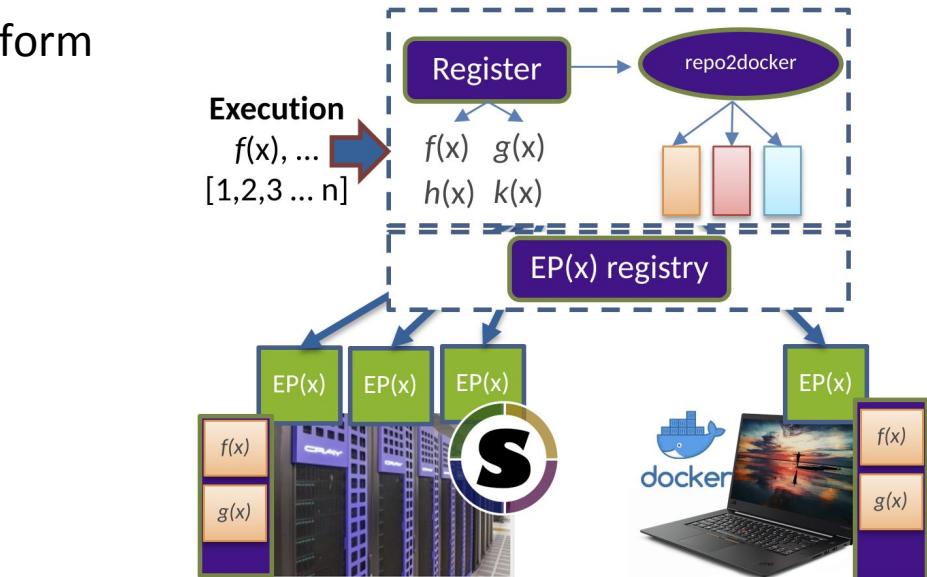
Backfill queues enable elastic HPC

Endpoints:

- Lightweight agent with Parsl config
- Abstracts underlying resource

Functions:

- Register once, run anywhere
- Container encapsulation
- Globus Auth



Parsl's resource configuration
abstracts compute resource

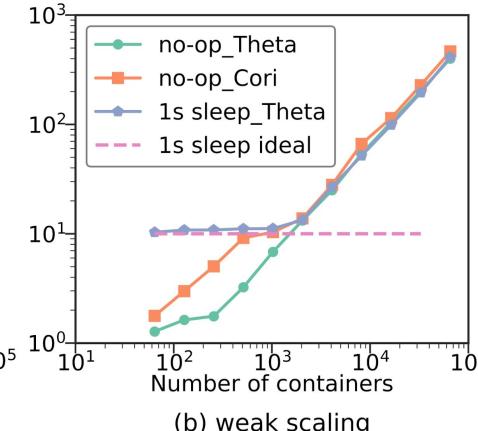
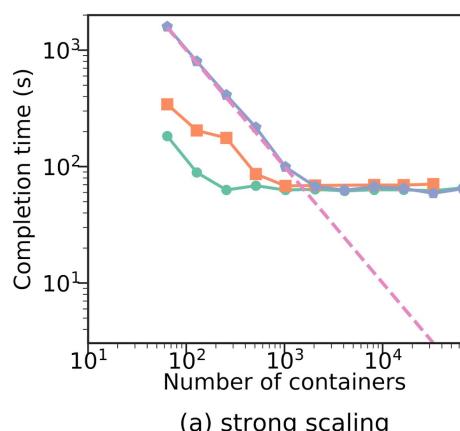
Portable containers encapsulate
dependencies

Run >65,000 containers on
Theta and Cori

funcX simplifies using HPC



```
config = Config(  
    executors=[  
        HighThroughputExecutor(  
            label="stampede2_htex",  
            address=address_by_hostname(),  
            provider=SlurmProvider(  
                channel=LocalChannel(),  
                nodes_per_block=128,  
                init_blocks=1,  
                partition="skx-normal",  
                walltime="12:00:00"  
            )  
        ]  
    )  
)
```

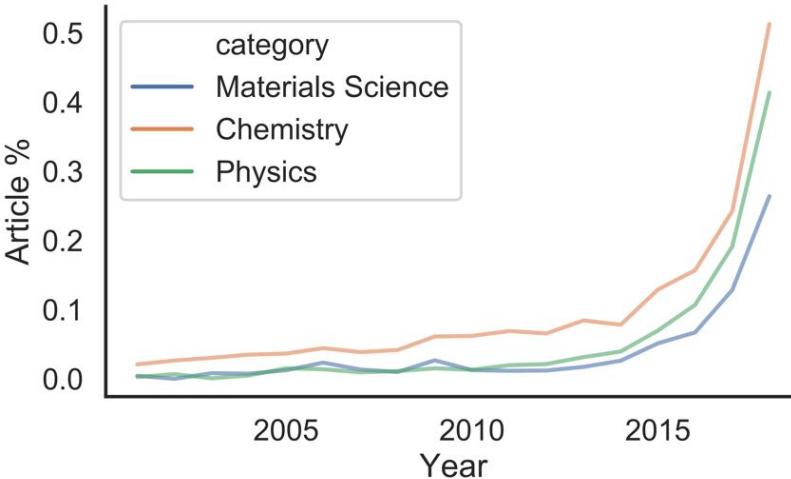




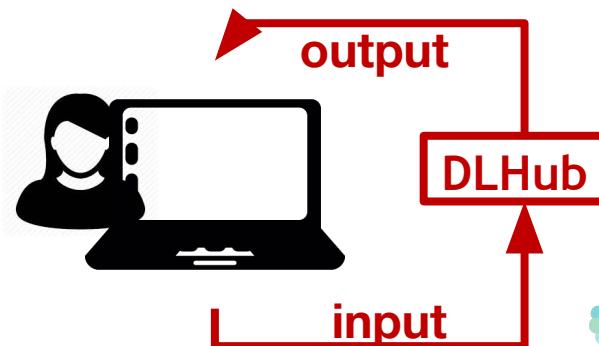
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ML in the Loop

Scientific ML



- Where are the model and trained weights?
- How do I run the model on my data?
- Should I run the model on my data?
- How do I share my model with the community?
- How can I build on this work?
- How do I cite it?



Model /
transform
containers

ARGONNE LEADERSHIP
COMPUTING FACILITY

parsl



MATERIALS
DATA
FACILITY

PETREL

Data and Learning Hub for Science (DLHub)



- Fits into the MDF/Globus ecosystem
- Collect, publish, categorize models from many disciplines (materials science, physics, chemistry, genomics, etc.)
- Serve model inference on-demand via API to simplify sharing, consumption, and access
- Enable new science through reuse, real-time model-in-the-loop integration, and synthesis & ensembling of existing models

Using DLHub is Easy!

Python SDK

```
$ pip install dlhub_sdk
```

Command Line Interface

```
$ pip install dlhub_cli
```

①

Describe

```
model = KerasModel.create_model("p1bl-example.h5")
model.set_title("CANDLE Pilot 1 - Benchmark 1")
model.set_name("candle_p1bl")
model.set_domains(["genomics", "biology", "HPC"])
```

②

Publish

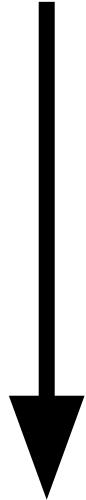
```
from dlhub_sdk.client import DLHubClient
dl = DLHubClient.login()
dl.publish_servable(model)
```

③

Run

```
data = np.load(pilot1.npy)
pred = dl.run("blaiszik_uchicago/candle_p1bl", data,
              type="python")
```

Marking up a Model – Python SDK



Existing Model

User Mark Up with
SDK

SDK Extracts
Metadata for Known
Model Types

Send to DLHub
(via Globus or HTTPS)

DLHub
Containerization

Populate Search
Index / Mint
Identifiers

```
from dlhub_toolbox.models.servables.keras import KerasModel
import pickle as pkl
import json

# Describe the keras model
model = KerasModel('model.hd5', list(map(str, range(10)))))

# Describe the model
model.set_title("MNIST Digit Classifier")
model.set_name("mnist_tiny_example")
model.set_domains(["general","digit recognition"])

# Add link to paper describing the dataset
model.add_related_identifier("10.1109/CVPR.2007.383157", "DOI",
| | | | | | | | "IsDescribedBy")

model.set_authors(["Lecunn, Yann", "Cortes, Corinna"])

# Describe the outputs in more detail
model.output['description'] = 'Probabilities of being 0-9'
model.input['description'] = 'Image of a digit'
```

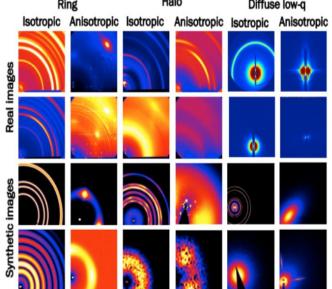
Model-in-the-Loop

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Enable automated use of models in analysis pipelines

Accessible as Automate Action

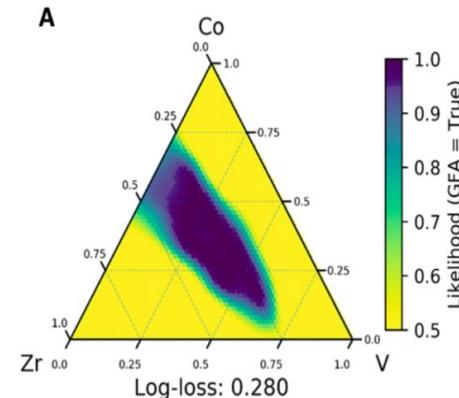
Tag beamline images



	probability	tags
1	0.999780	Beam Off Image
6	0.745707	Halo
13	0.364863	Strong scattering
7	0.082574	High background
12	0.027826	Ring
8	0.012137	Higher orders
15	0.008106	Weak scattering
9	0.001964	Linear beamstop
16	0.001744	Wedge beamstop
2	0.000982	Circular Beamstop

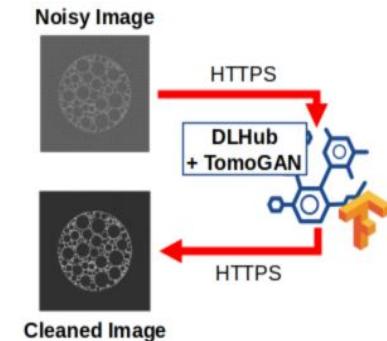
XRD image tagging (Yager, BNL)

Guide experiments



(Ward, ANL/UC)

Improve quality



TomoGAN (Liu Argonne)



Thanks!

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Questions?

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