

# First steps to data portals: Globus, data catalog, and You

Andy Boughton  
([abought@globus.org](mailto:abought@globus.org))  
APECx Demos – March 2025



THE UNIVERSITY OF  
CHICAGO



Argonne  
NATIONAL LABORATORY





# Globus Transfer: Part of an integrated platform



## Learn more:

<https://www.globus.org/platform>

<https://www.go-fair.org/fair-principles/>



## Today's notes:

<https://github.com/globus/apecx-demos/>



## Try it yourself

Get an endpoint: <https://www.globus.org/globus-connect-personal>

Manage files: <https://app.globus.org/>

Example static search portal: <https://abought.github.io/apecx-demo-static-search-portal>

-> Real version *not yet publicly available*

**F**indable

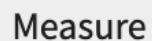
**A**ccessible

**I**nteroperable

**R**eusable



The SPHERICAL data repository contains inputs from many teams. Explore example datasets below:



### Experimental data

## Protein structure examples



## Computational models

## Demo render files



Curated insights from across the community

## Publications

## Literature chatbot

10MW



**DOI:** <https://doi.org/10.2210/pdb1OMW/pdb>

**Description:** Crystal Structure of the complex between G Protein-Coupled Receptor Kinase 2 and Heterotrimeric G Protein beta 1 and gamma 2 subunits

**Organism:** bos taurus

**Deposited:** 2003-02-26T00:00:00+00:00

Released: 2003-06-03T00:00:00+00:00

**Authors:** Lodowski, D.T.; Pitcher, J.A.; Capel, W.D.; Lefkowitz, R.J.; Tesmer, J.J.G.

**Funders:** (none provided)

protein , gprotein , gpcr , xrd , experimental

Publication Type

<input type="checkbox"/> JOUR	(24)
<input type="checkbox"/> CHAP	(1)
<input type="checkbox"/> COMP	(1)
<input type="checkbox"/> CONF	(1)
<input type="checkbox"/> THES	(1)

## Top Tags

<input type="checkbox"/> chemistry	(14)
<input type="checkbox"/> spectroscopy	(14)
<input type="checkbox"/> genetics	(10)
<input type="checkbox"/> protein	(8)
<input type="checkbox"/> software	(8)
<input type="checkbox"/> visualization	(5)
<input type="checkbox"/> crowdsourcing	(4)
<input type="checkbox"/> cell imaging	(3)
<input type="checkbox"/> covid	(3)
<input type="checkbox"/> machine learning	(3)

## Results

19 datasets found

Big image

**Description:** A big background picture



## JSON blob

**Description:** A json blob with odd formatting, to test code highlighting of rendered JSON

```
{ "key": "value", "items": [1,2,3], "nothing": null, "nested": {} }
```

Demo of portal-in-progress

**(not public yet)**



# Core Technologies





# Technology highlights

- **Globus provides:**
  - One **auth** system across services
  - **Groups** to control access
  - Full text search features
- **Reusable tools**
  - Python and JavaScript SDKs
  - “Search portal” helper tools
  - Prebuilt UI for workflows and file transfer



# Django Globus Portal Framework

DOI 10.5281/zenodo.14919979 Tests passing pypi v0.4.13 wheel yes License Apache 2.0

The Django Globus Portal Framework is a collection of tools that enable you to rapidly create an easily accessible portal for your data. Globus provides robust Auth and Search services, both powerful tools to help manage who has access to your data and metadata. Tailoring your portal to your data can be done simply by modifying existing built-in Bootstrap templates, allowing many levels of customization to suit the required needs.

Framework components. You can copy-paste the individual settings below, or use our [Settings Example](#) for a complete `settings.py` file reference.

`settings.py` file to enable user auth and Globus Portal

SEARCH

ABOUT

Q

Type

- ☐ protein (2)
- ☐ xrd (2)
- ☐ Experimental (1)
- ☐ experimental (1)
- ☐ gpcr (1)
- ☐ gprotein (1)
- ☐ rift valley fever (1)
- ☐ viral protein (1)

Results

2 datasets found

```
# Your portal credentials for a Globus Auth Flow
SOCIAL_AUTH_GLOBUS_KEY = 'Put your Client ID here'
SOCIAL_AUTH_GLOBUS_SECRET = 'Put your Client Secret Here'

# This is a general Django setting if views need to redirect to login
# https://docs.djangoproject.com/en/3.2/ref/settings/#login-url
LOGIN_URL = '/login/globus'

# This dictates which scopes will be requested on each user login
SOCIAL_AUTH_GLOBUS_SCOPE = [
    'urn:globus:auth:scope:search.api.globus.org:search',
]

# Installed apps tells Django which packages to load on startup
INSTALLED_APPS = [
    ...
    'globus_portal_framework',
    'social_django',
]
```





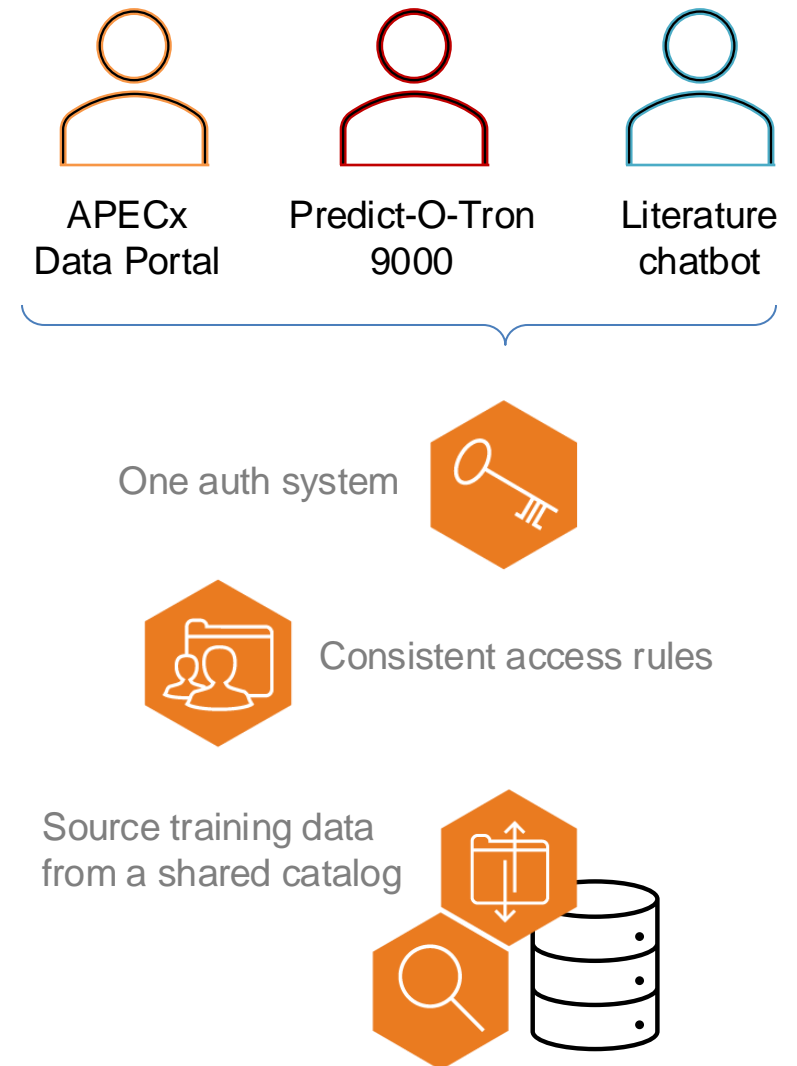
# Build your own adventure

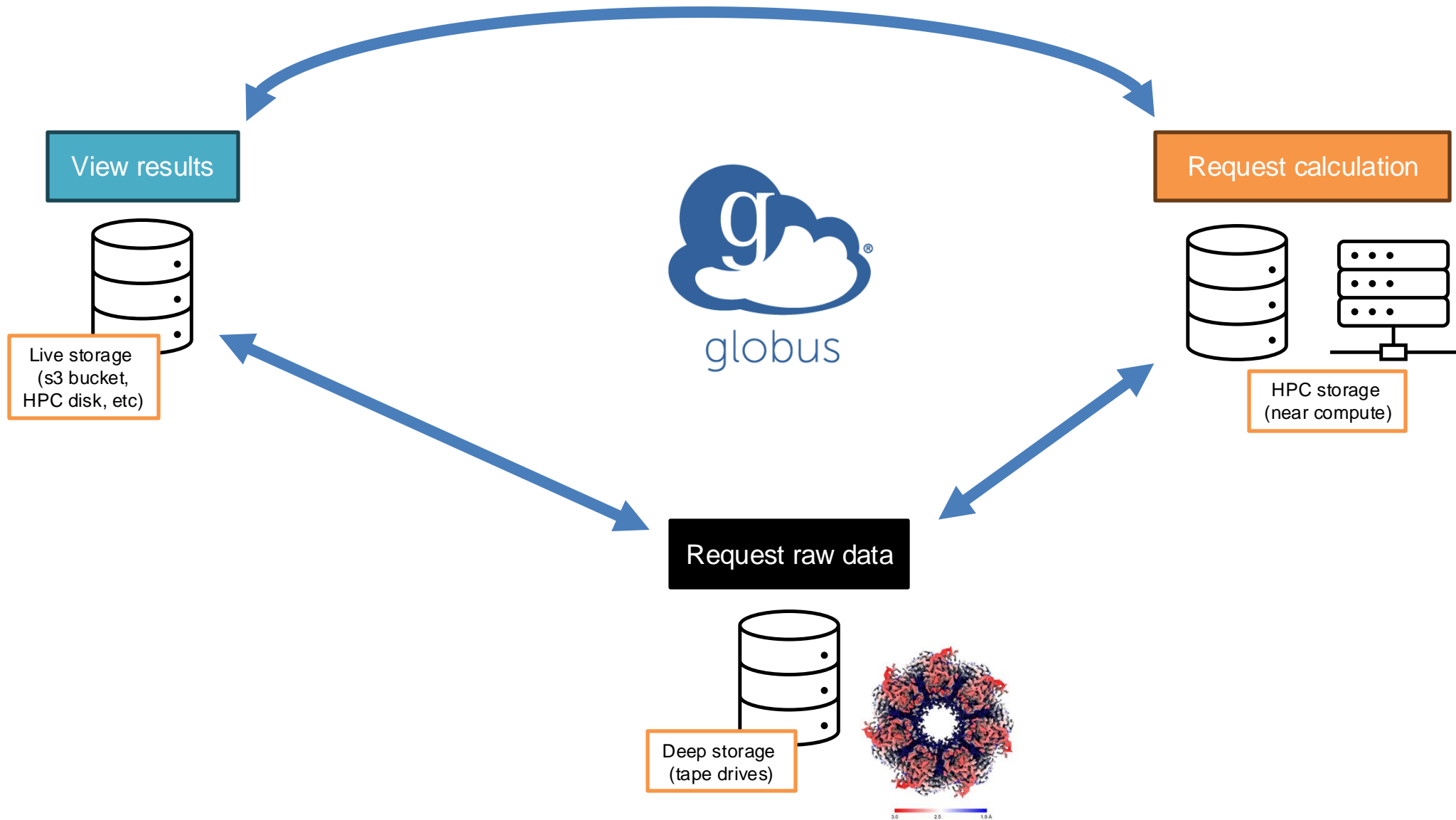
- **SDKs control every aspect of Globus**
  - Python: <https://globus-sdk-python.readthedocs.io/en/stable/>
  - JS: <https://github.com/globus/globus-sdk-javascript>
- **Higher-order helpers build websites fast**
  - Django: <https://github.com/globus/django-globus-portal-framework>
- **Reusable libraries let you adopt just one feature**
  - Python Social auth: [https://github.com/python-social-auth/social-core/blob/master/social\\_core/backends/globus.py](https://github.com/python-social-auth/social-core/blob/master/social_core/backends/globus.py)



# Findable and reusable

- **Multiple websites can build on the shared data catalog / portal**
- **Shared foundation allows reuse of data**
- **We do the fiddly stuff (“members of APECx, only”)**

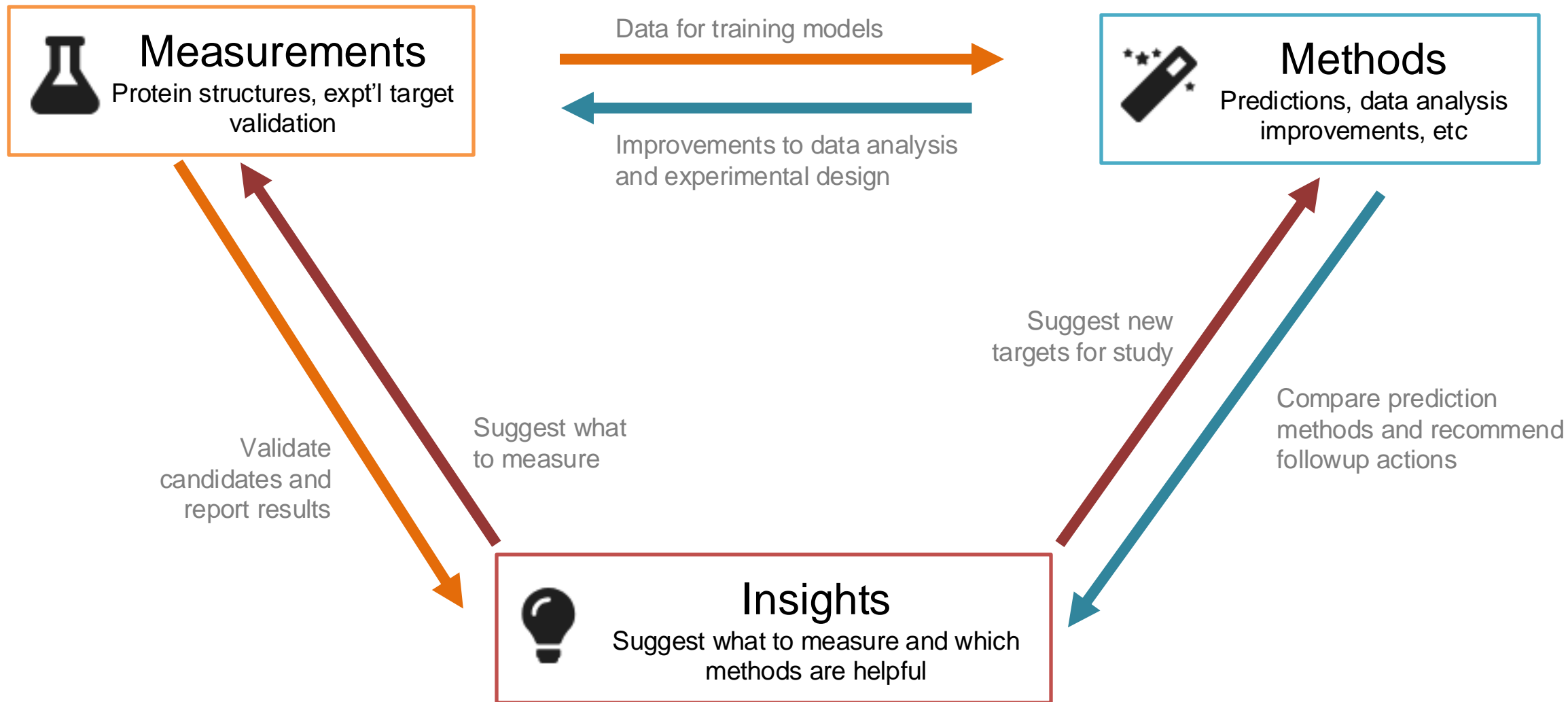




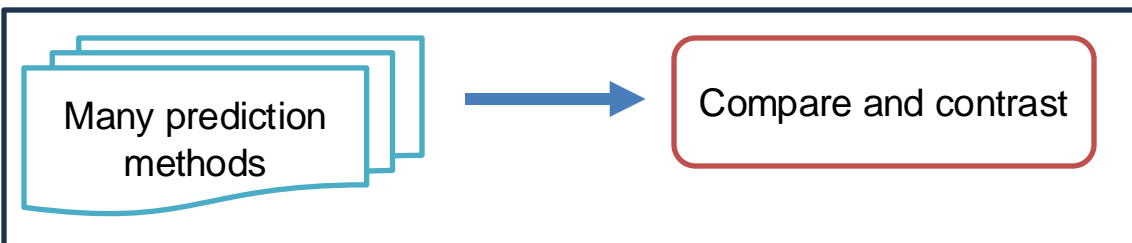
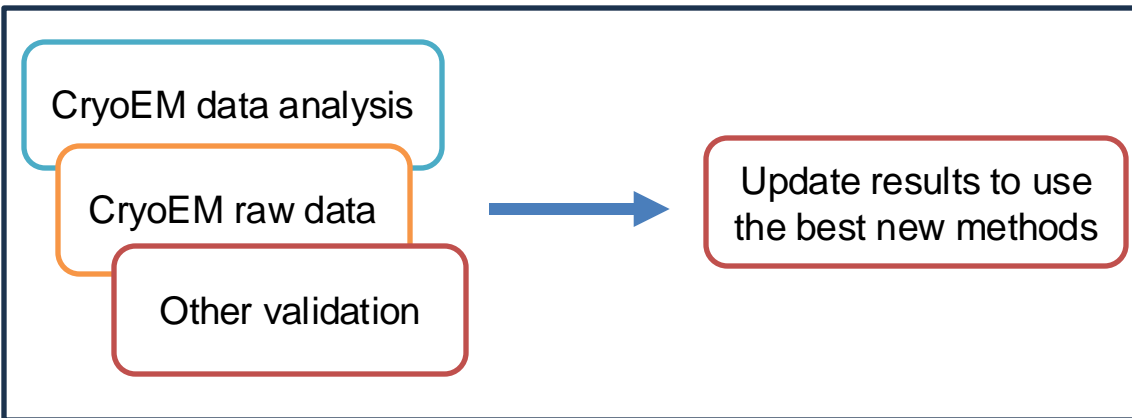
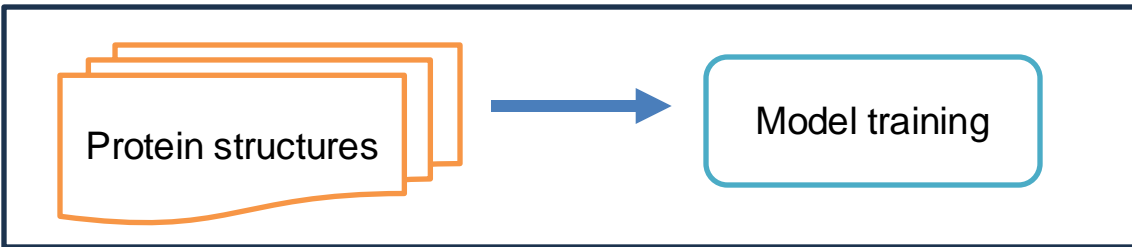
All the fancy stuff depends on data

**Where do we start?**





# Points of collaboration and connection



- **Create categories based on value-add capabilities**
  - Identify “AI/ML ready” models
  - Run a method that requires inputs of a certain type
  - Incorporate QC for things that might feed into methods
- **Search features that allow discovering reusable items**



# Finding a Taxonomy

- Encourage **discovery** across research groups
  - Identify what datasets have in common
- Share a **common language**
  - Eg all submissions should use the same molecule/virus name/ID symbol
- Unrealistic to build custom apps for every individual dataset- what are the **big wins**?

Some example data types  
(maybe)

Protein structures (measured and predicted)

Predicted druggable targets

*In silico* estimates of antibody efficacy

Clinical validation

Available squishy wet lab things

...Many others

# Discussion questions







# Audience Questions: Data Catalog

- What are some **patterns** in the data types we have catalogued to date?
- Is there a **standard** ID format / language for entities?
- What general categories would encourage **interoperable** re-use?

Some example data types  
(maybe)

Protein structures (measured and predicted)

Predicted druggable targets

*In silico* estimates of antibody efficacy

Clinical validation

Available squishy wet lab things

...Many others



**Measurements**  
Protein structures, expt'l  
target validation



**Methods**  
Predictions, data analysis  
improvements, etc



**Insights**  
Suggest what to measure and  
which methods are helpful



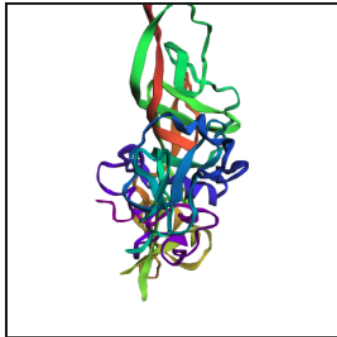
# Audience Questions: First goal

- **What is a highly reusable piece of data that multiple groups want to access?**
  - And what is needed to make that data “AI/ML ready”
- **What is a useful presentation of that data?**
  - How does a user decide which dataset will meet their needs?
  - Schema + interactive features

6F8P

DOWNLOAD VIA HTTPS

RUN CALCULATION



**PDB ID:** 6F8P

**DOI:** <https://doi.org/10.2210/pdb6F8P/pdb>

**Description:** Crystal structure of Gn from Rift Valley fever virus

**Organism:** Rift Valley fever virus

**Deposited:** 2017-12-13T00:00:00+00:00

**Released:** 2018-01-31T00:00:00+00:00

**Authors:** Halldorsson, S.; Bowden, T.A.; Harlos, K.

**Funders:** Medical Research Council (United Kingdom)

protein , viral protein , rift valley fever , xrd , Experimental



# Audience questions: General

- **What topics would be useful for a future talk?**
  - Globus technologies (workflows/ compute, auth, etc)
  - Deep dive on data
  - Portal features
- **What do we want the portal to *do*?**
  - We have the technology
  - You have the science





# Future questions

- **Where does the website live?**
- **Who has access?**