

# On-demand Scientific Services: AlphaFold



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# funcX interface for on-demand HPC services

Leadership computing facilities:

- Extreme scale
- Specialized hardware
- Enormous datasets (simulation, reference, ML)

Also...

- Restrictive queueing models
- Strict authentication and authorization
- Auditing and reporting requirements
- Challenging environments



funcX can simplify access to data and compute for many communities

# AlphaFold as a Service at ALCF

Cutting edge ML technique to predict protein structure with applications in screening, therapeutics, light sources, crystallography, etc.

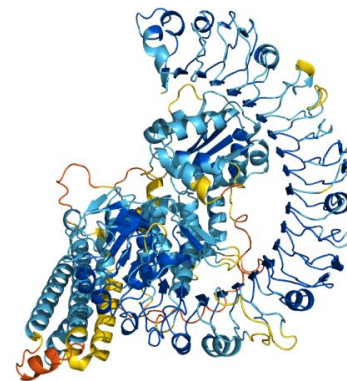
```
>GA98_DM.114 GA98 Deletion Mutation Sequence  
TTYKLILNLKQAKEEAIKELVDAGTAKYFKLIANAKTVEGVWTLKDE
```

Deployed funcX to run AlphaFold on-demand on ALCF resources



AlphaFold as a Service:

1. User provides FASTA input
2. Upload data to ALCF's Eagle storage system
3. funcX provisions GPU resources on Polaris
4. Task runs for ~1.5 hours
5. Results made available via Globus
6. Download link emailed to user



# funcX + AlphaFold + ALCF

funcX endpoint deployed on Polarislogin1

- Shared with private Globus group
  - Restricted to ALCF users
- Singularity-enabled
- Provisions GPU nodes
- Mounts necessary data
- Uses preemptable queue
- Runs in my account

Custom Singularity container

- [https://github.com/hyoo/alphafold\\_singularity](https://github.com/hyoo/alphafold_singularity)

Defined and registered a function to invoke alphafold in container

Also deployed a Globus flow to use the funcx function, runnable by the group

```
config = Config(  
    executors=[  
        HighThroughputExecutor(  
            max_workers_per_node=1,  
            strategy=SimpleStrategy(max_idletime=60),  
            address=address_by_hostname(),  
            scheduler_mode='soft',  
            worker_mode='singularity_reuse',  
            container_type='singularity',  
            container_cmd_options="--nv -H /home/ro  
            provider=PBSProvider(  
                cpus_per_node=32,  
                select_options="ngpus=4",  
                launcher=SingleNodeLauncher(),  
                account='APSDDataAnalysis',  
                queue='preemptable',  
                scheduler_options=user_opts['polaris']  
                worker_init=user_opts['polaris']['w  
                walltime='06:00:00',  
                nodes_per_block=1,  
                init_blocks=0,  
                min_blocks=0,  
                max_blocks=4,  
            ),  
        ],  
    ),  
)
```

# Using it

Run via funcX or Globus Flow

- CLI: <https://github.com/globus-labs/globus-alphafold-cli>

```
$ python cli.py run --fasta /path/to/file.fasta
```

- Or run the flow directly ->

Results emailed with download link:

## Globus AlphaFold flow completed.

You can collect the result here: [https://g-719d9.fd635.8443.data.globus.org/output/517e5fd2/GB98\\_DM\\_3.fasta.log](https://g-719d9.fd635.8443.data.globus.org/output/517e5fd2/GB98_DM_3.fasta.log)

```
flow_id = '7c277b80-2cca-42b7-a75d-a970841ee874'  
flow_scope = 'https://auth.globus.org/scopes/7c277b80-2c  
  
flow_input = {  
    "input": {  
        "fasta": fasta,  
        "email": email_address,  
    }  
}
```

Now start the flow. We create a `flows_client` and will then be prompted

```
flows_client = create_flows_client()  
  
flow_action = flows_client.run_flow(flow_id, flow_scope)  
flow_action_id = flow_action['action_id']  
print(f"flow started: https://app.globus.org/runs/{flow_action_id}")
```

Please log into Globus here:

```
-----  
https://auth.globus.org/v2/oauth2/authorize?client_id=0a100000-0000-0000-0000-000000000000  
Fauth.globus.org%2Fv2%2Fweb%2Fauth-code&scope=https%3A%2F%2Fauth.globus.org%2Fscopes%2F7c277b80-2c  
dfcfe%2Fflow_f616749e_ef70_4991_9429_dd803fdffcfe_user_credentials%2F7c277b80-2c  
fRa_KRhgGACrTYmyj16xASTXU8vvnw5c&code_challenge_method=S256  
mmand+Line+Interface+on+mbp.lan  
-----
```

Enter the resulting Authorization Code here: TYWzm5mQox  
flow started: <https://app.globus.org/runs/cadfa395-30b4>

Check the status of the flow. The AlphaFold step often takes roughly two

```
flow_action = flows_client.flow_action_status(flow_id,  
flow_status = flow_action['status']  
print(f'Flow status: {flow_status}')
```

Flow status: ACTIVE

# Production services with a funcX interface

Service account to operate endpoint and multiplex users

- Globus app credentials to own the endpoint

Project allocation to monitor and manage resource usage

Managing functions - vet, add, and remove functions

Manage user access - Globus group membership

Fine-grained auditing and reporting

Mechanism to stop bad tasks without impacting other users

Persist endpoints and restart as necessary (failures, maintenance)

- Service node to operate endpoint outside login nodes
- Considering podman for endpoint management

Ability to update execution environment and container



# Next steps

## ALCF

Define function vetting process

Work through service accounts

Persist endpoints on service nodes via systemd or similar

## funcX

Improve public auditing capabilities

Enhance function authorization/management for endpoints

Task cancellation and management

## AlphaFold service

Provide a web interface

Leverage in light source analysis loop with APS's GM/CA beamline

Integrate into ML training workflows

# Thanks!

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