

Building the Model Research Data Portal

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with Kyle Chard, Eli Dart, Steve
Tuecke, Jason Williams



The Modern Research Data Portal: A Design Pattern for Networked, Data-Intensive Science

<https://docs.globus.org/mrdp>

The Modern Research Data Portal is a new design pattern for providing secure, scalable, and high performance access to research data.



GitHub Repo

provides code for the simple data portal that you can experiment with online

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Example Data Portal

allows you to experiment with an example implementation of the design pattern

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Code Walkthrough

provides a narrative description of the simple data portal code

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Jupyter Notebook

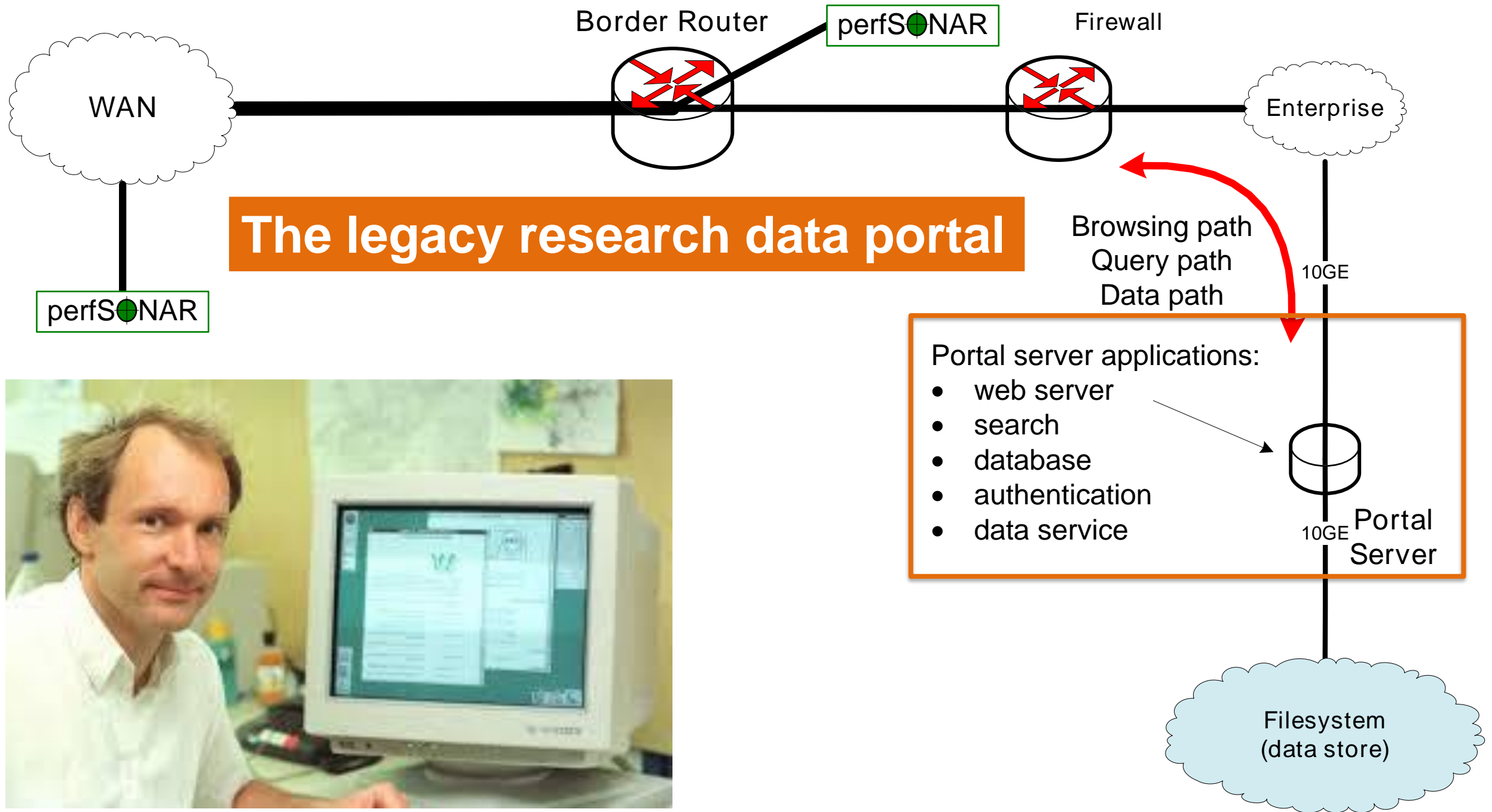
demonstrates some Globus features described in the technical article

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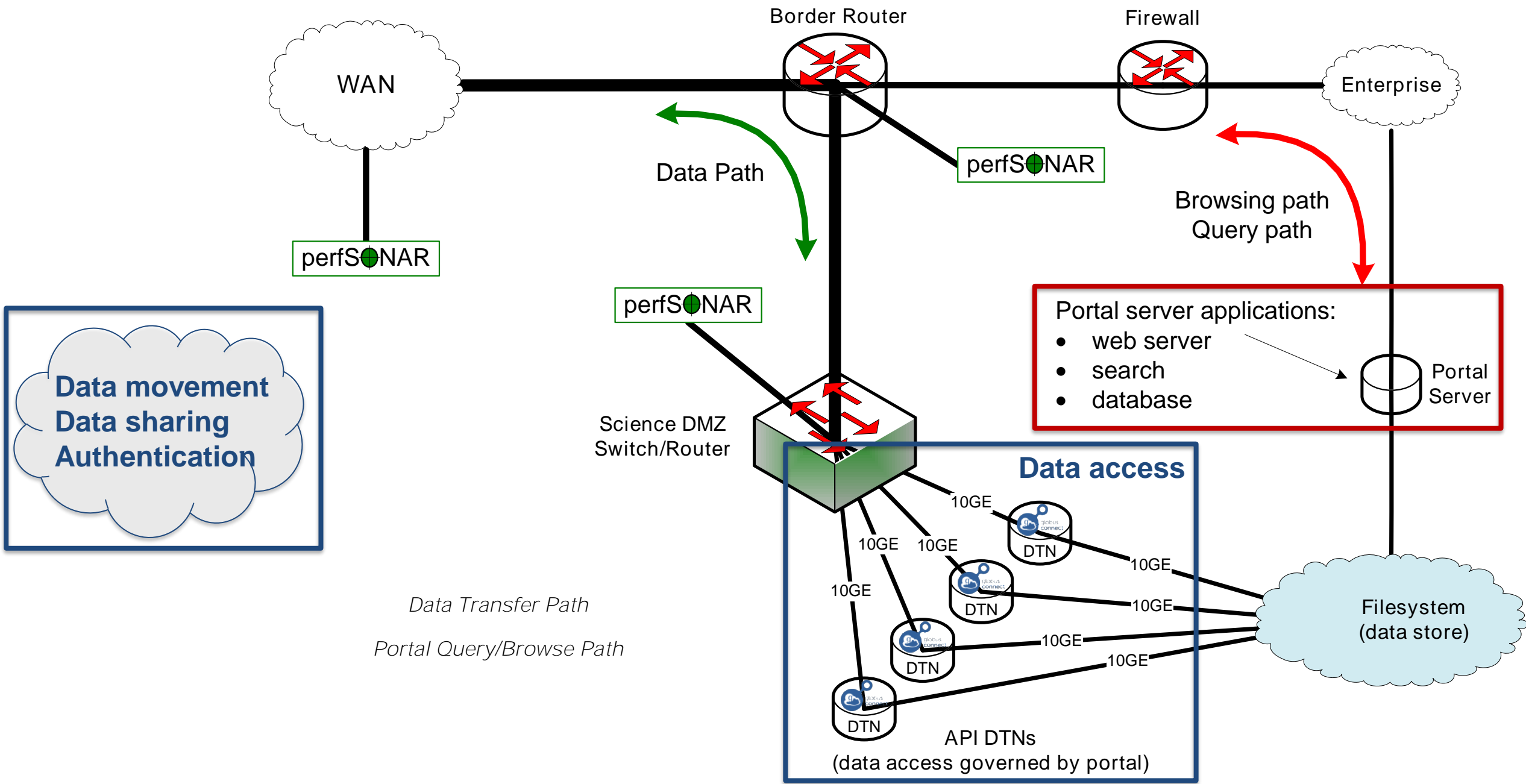
A technical article describes this design pattern, reviews representative examples at research laboratories and universities (see below), and uses coding examples to demonstrate how Globus APIs can be used to implement a range of research data portal capabilities.

Chard K, Dart E, Foster I, Shifflett D, Tuecke S, Williams J. (2017) The Modern Research Data Portal: A design pattern for networked, data-intensive science. PeerJ Preprints5:e3194v1
<https://doi.org/10.7287/peerj.preprints.3194v1>

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The modern research data portal

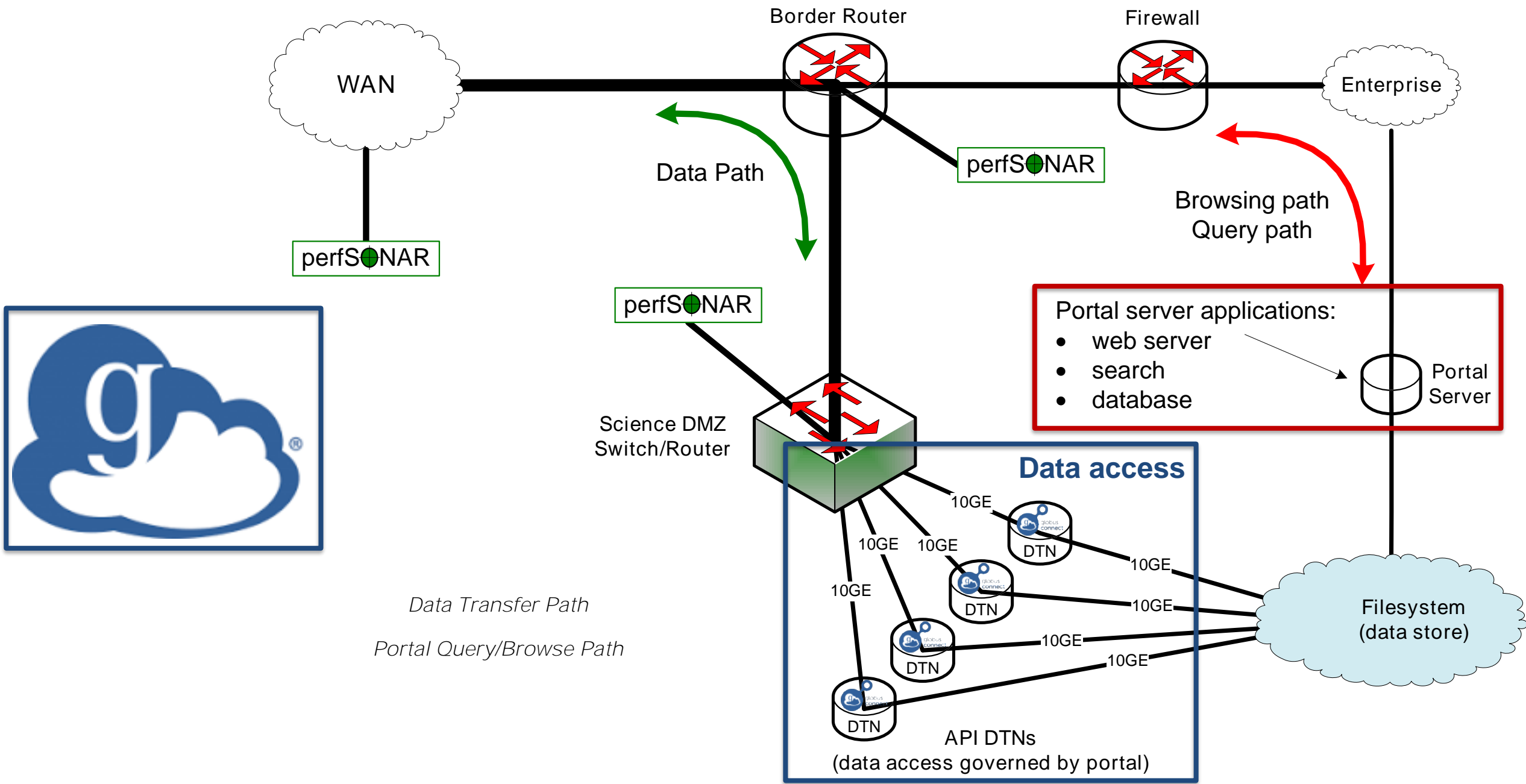


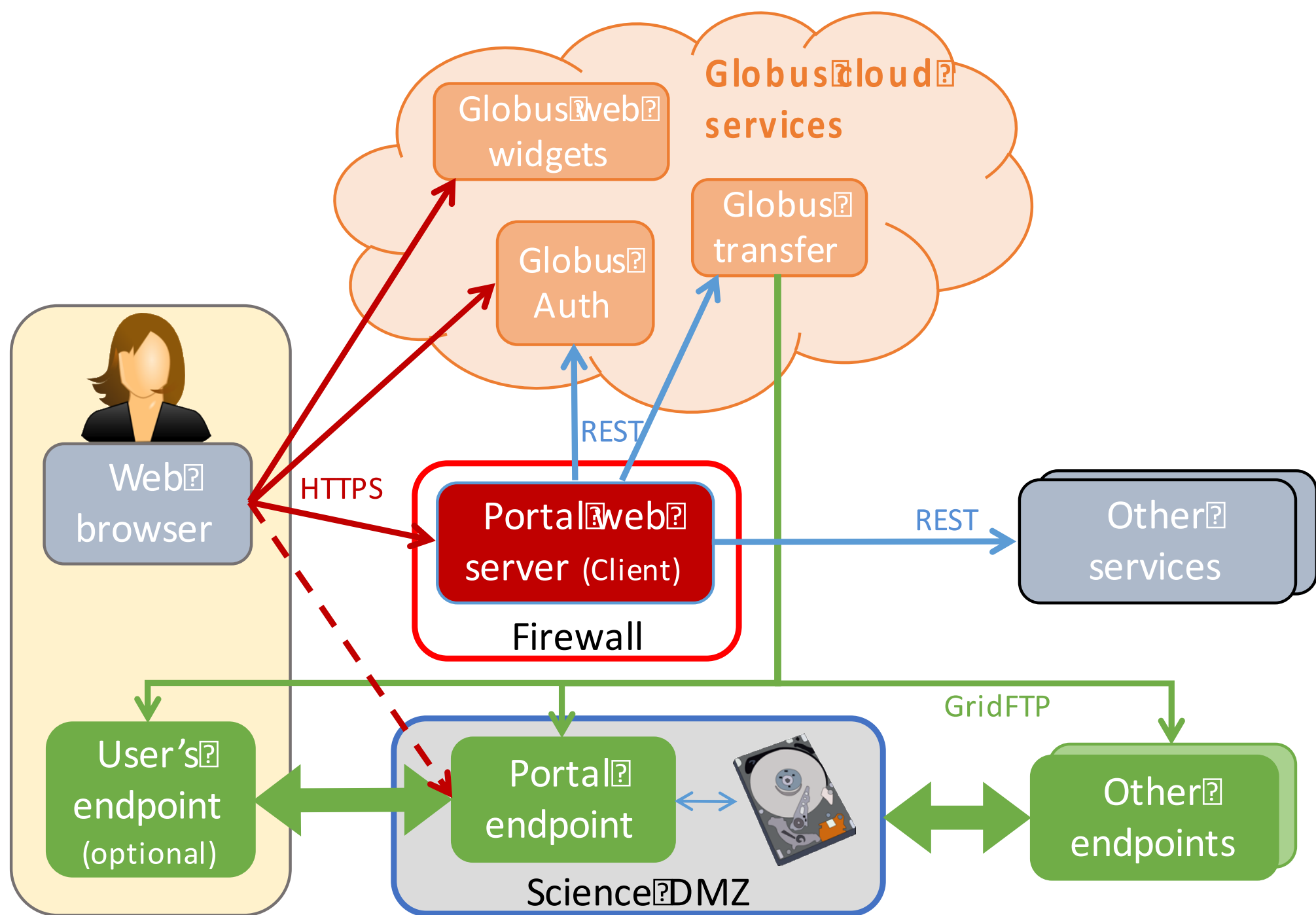


A key message: Outsource all that you can

- Outsource responsibility for determining user identities
- Outsource control over who can access different data and services within the portal
- Outsource responsibility for managing data uploads and downloads between various locations and storage systems
- Leverage standard web user interfaces for common user actions

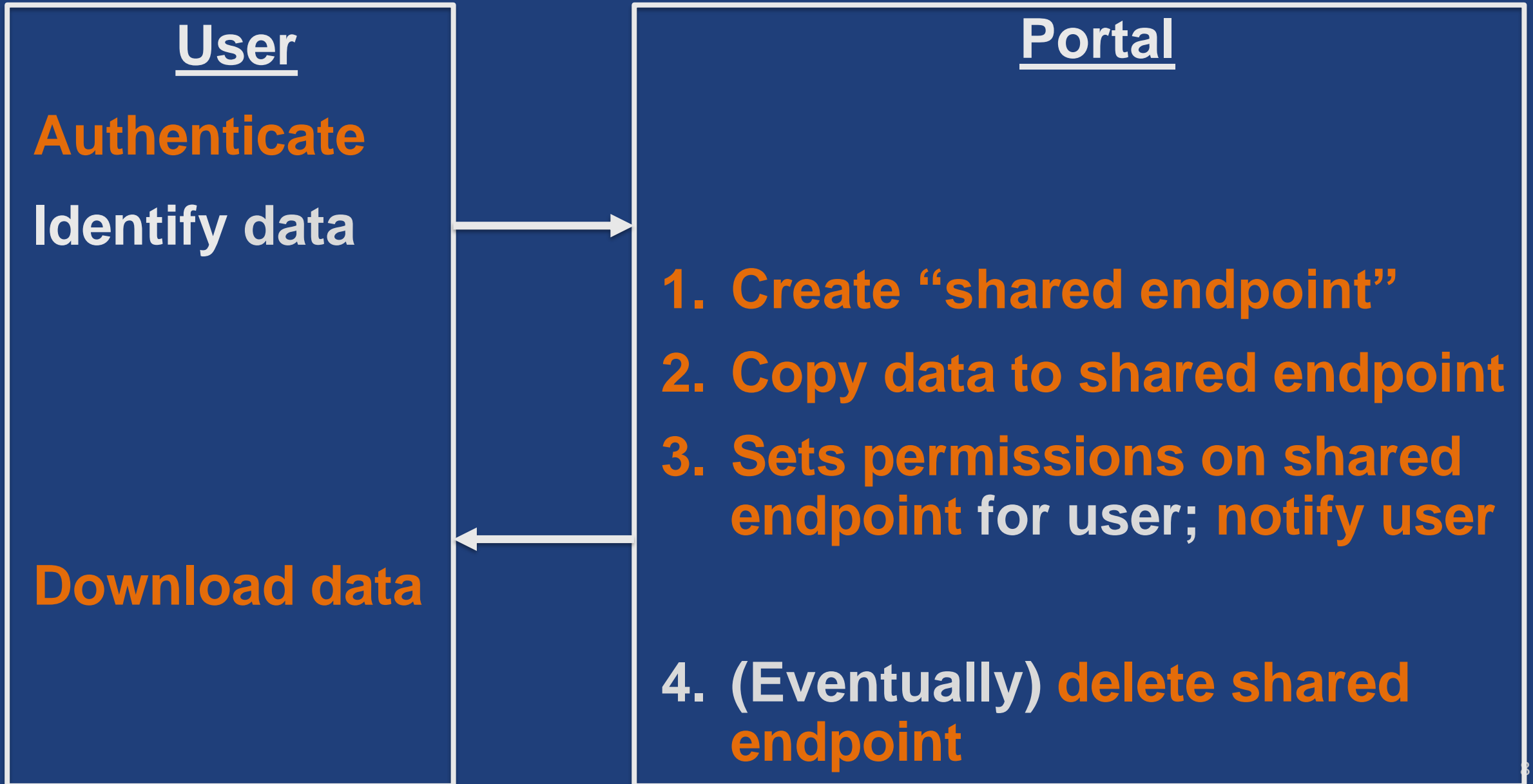
The modern research data portal







A simple example of MDRDP logic





1. Create “shared endpoint”
2. Copy data to shared endpoint
3. Set permissions on shared endpoint for user; notify user
- ...
4. (Eventually) delete shared endpoint

```
from globus_sdk import TransferClient, TransferData
from globus_sdk import AuthClient
import sys, random, uuid
```

```
def rdp(host_id,          # Endpoint for shared endpoint
        source_path,     # Directory to copy data from
        email):          # Email address to share with
```

```
    tc = TransferClient()
```

```
    ac = AuthClient()
```

```
    tc.endpoint_autoactivate(host_id)
```

Connect
to storage
system

```
    # (1) Create shared endpoint:
```

```
    # (a) Create directory to be shared
```

```
    share_path = '/~/ ' + str(uuid.uuid4()) + '/'
```

```
    tc.operation_mkdir(host_id, path=share_path)
```

```
    # (b) Create shared endpoint on directory
```

```
    shared_ep_data = {
```

```
        'DATA_TYPE': 'shared_endpoint',
```

```
        'host_endpoint': host_id,
```

```
        'host_path': share_path,
```

```
        'display_name': 'RDP shared endpoint',
```

```
        'description': 'RDP shared endpoint'
```

```
    }
```

```
    r = tc.create_shared_endpoint(shared_ep_data)
```

```
    share_id = r['id']
```



1. Create “shared endpoint”
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```
# (2) Copy data into the shared endpoint
tc.endpoint_autoactivate(share_id)
tdata = TransferData(tc, host_id, share_id,
                    label='RDP copy', sync_level='checksum')
tdata.add_item(source_path, '/', recursive=True)
r = tc.submit_transfer(tdata)
tc.task_wait(r['task_id'], timeout=1000,
            polling_interval=10)
```

```
# (3) Enable access by user
r = ac.get_identities( usernames=email)
user_id = r['identities'][0]['id']
rule_data = {
    'DATA_TYPE': 'access',
    'principal_type': 'identity', # Grantee is
    'principal': user_id,        # a user.
    'path': '/',                 # Path is /
    'permissions': 'r',          # Read-only
    'notify_email': email,       # Email invite
    'notify_message':            # Invite msg
        'Requested data are available.'
}
tc.add_endpoint_acl_rule(share_id, rule_data)

# (4) Ultimately, delete the shared endpoint
tc.delete_endpoint(share_id)
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An example MRDP

<https://docs.globus.org/mrdp>

Modern Research Data Portal

TRANSFER

| GRAPH

| PROFILE

| LOGOUT

| IAN@GLOBUSID.ORG

Repository

Select some dataset(s) to transfer **or** click on a dataset name to browse its files.

Dataset Name	Select
 Atlanta International Airport Climate Data	<input type="checkbox"/>
 Boston Logan International Airport Climate Data	<input type="checkbox"/>

Many variants possible

- Manage access to data at multiple locations
- Manage access to data on cloud
- Upload data for analysis
- Data download from scientific instruments
- Data publication
- Transfer data to computer for analysis

Sanger Imputation Service **Beta** Home About Instructions Resources Status

Sanger Imputation Service

This is a free genotype **imputation** and **phasing** service provided by the [Wellcome Trust Sanger Institute](#). You can upload GWAS data in VCF or 23andMe format and receive imputed and phased genomes back. Click [here](#) to learn more and [follow us on Twitter](#).

Before you start

Be sure to [read through the instructions](#).

You will need to set up a free account with [Globus](#) and have [Globus Connect](#) running at your institute or on your computer to transfer files to and from the service.

Ready to start?

If you are ready to upload your data, please fill in the details below to **register an imputation and/or phasing job**. If you need more information, see the [about](#) page.

Full name

Organisation

Email address

What is this? [?](#)

Globus user identity

[Next](#)

News



[@sangerimpute](#)

30/1/2017
Support for chromosome X has been added to all pipelines. PBWT has been updated to increase imputation accuracy of dosages and fix some bugs. See [ChangeLog](#).

31/10/2016
New African Genome Resources panel with 9,912 haplotypes (6,230 African) is [now available](#).

11/04/2016
Thanks to [EAGLE2](#), we can now return **phased data**. The HRC panel has been updated to r1.1 to fix a [known issue](#). See [ChangeLog](#) for more details.

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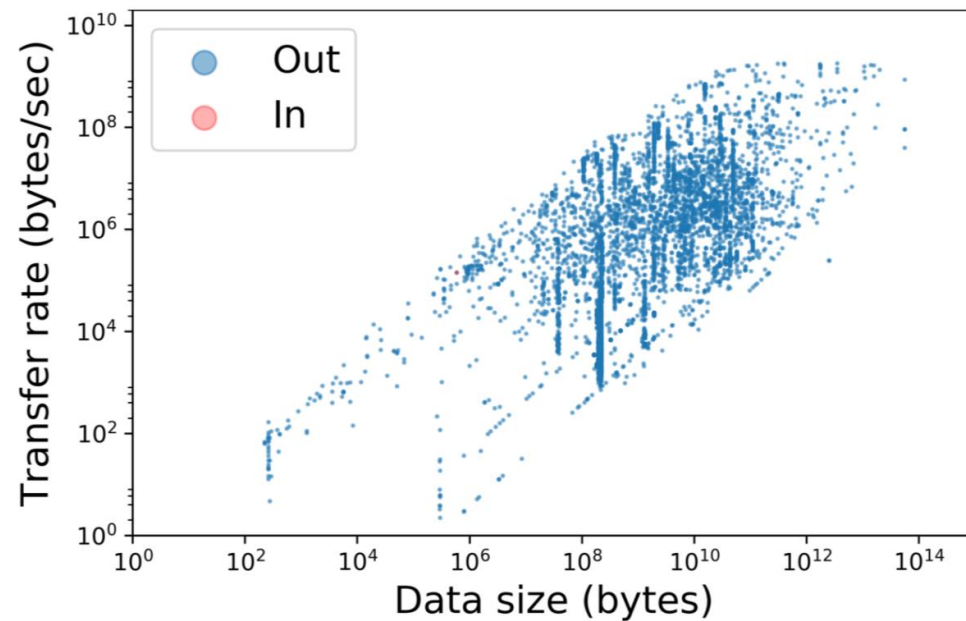
Publication of large datasets
MDF offers researchers access to petabytes (PB) of reliable and high performance data storage via NCSA



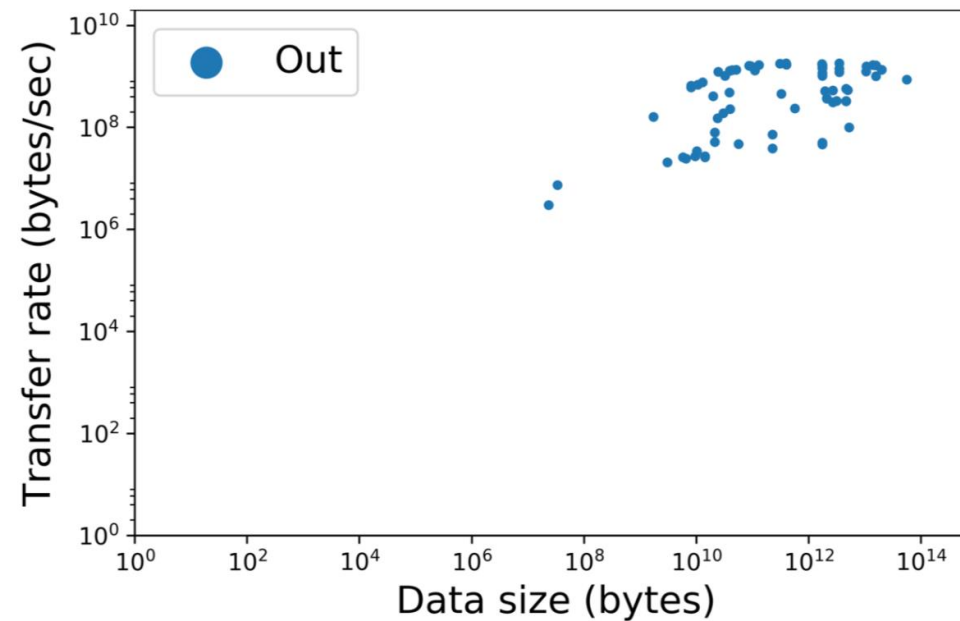
Customizable metadata descriptions
MDF collection owners can define and use their own materials-specific metadata schemas to describe their published



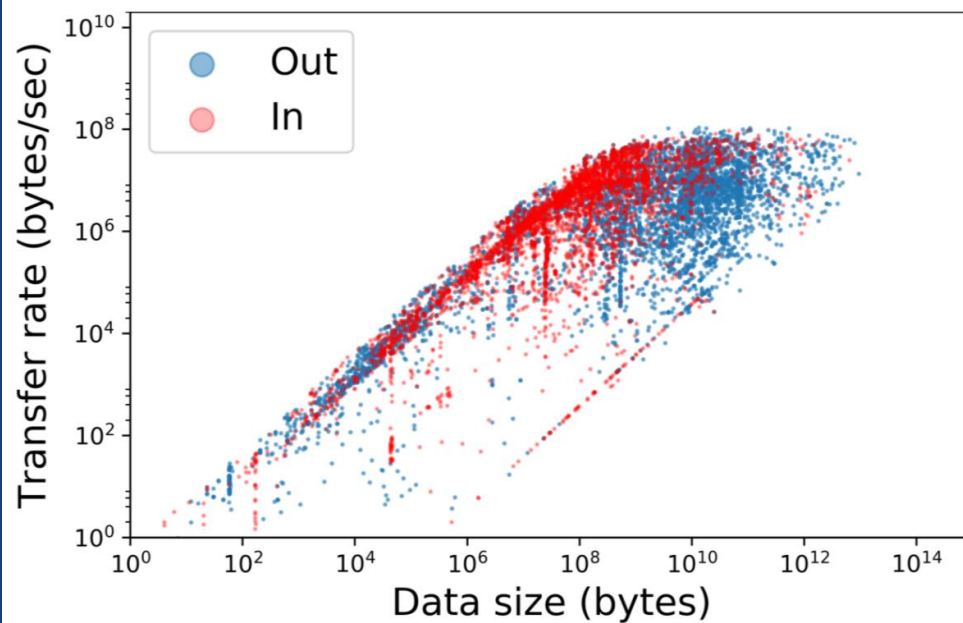
Flexible access control
Published datasets may be private, shared with a particular group of users, or shared publicly



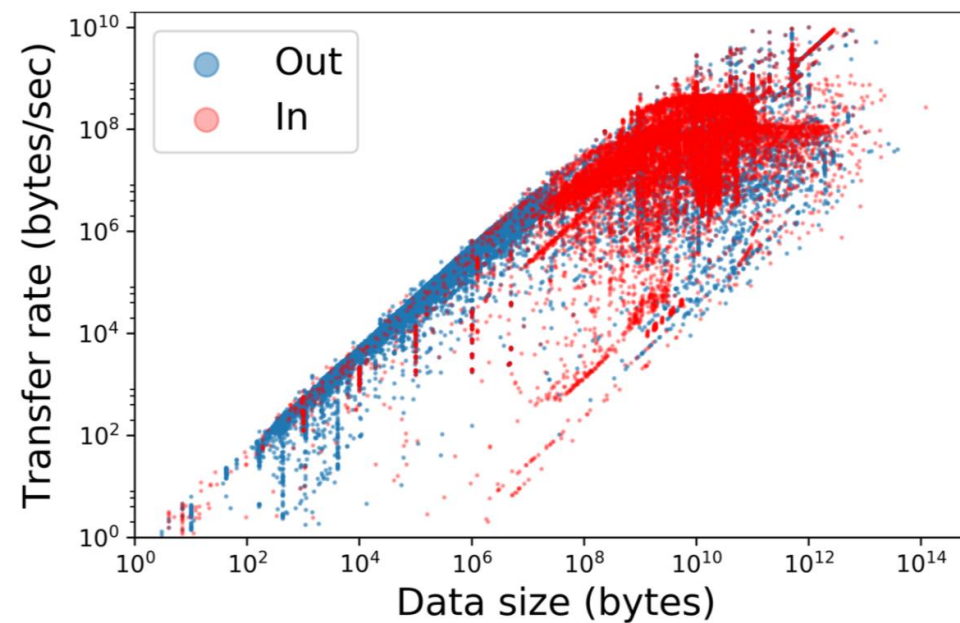
(a) RDA



(b) RDA to NERSC subset



(c) Sanger



(d) Petrel

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