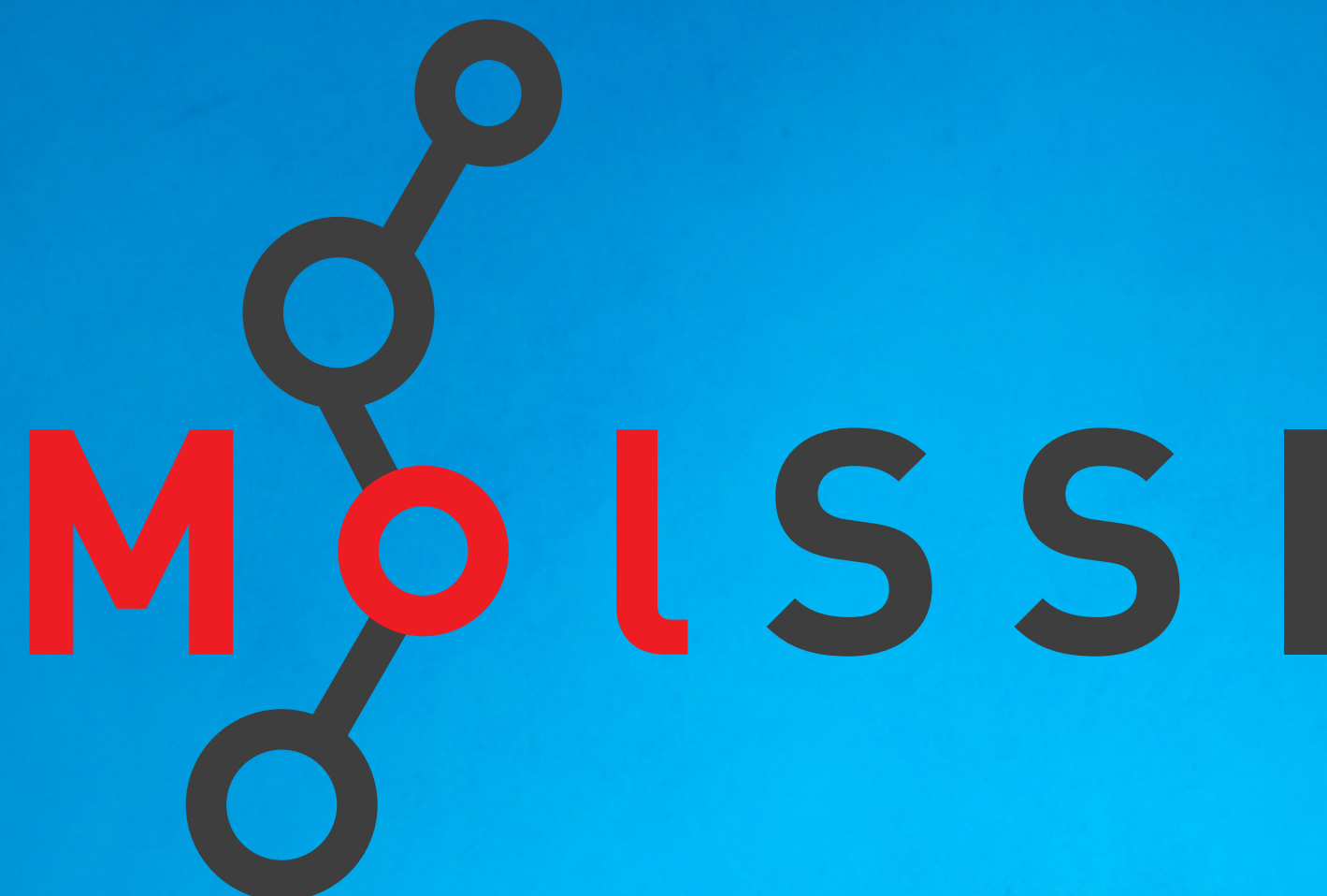


# The MolSSI Quantum Chemistry Archive Project



Daniel G. A. Smith, Levi N. Naden, Doaa Altarawy, and Matt Welborne

The Molecular Sciences Software Institute

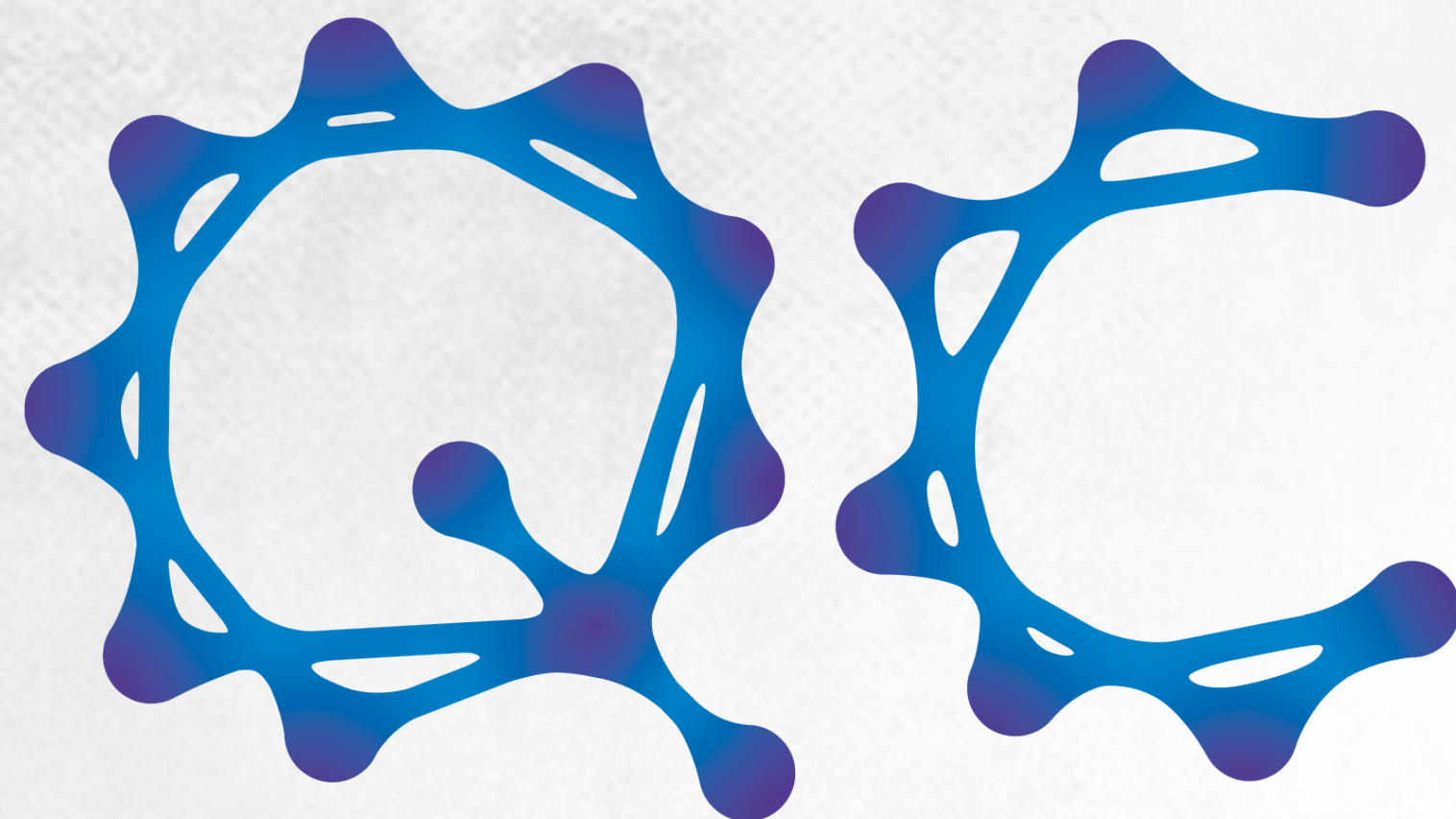
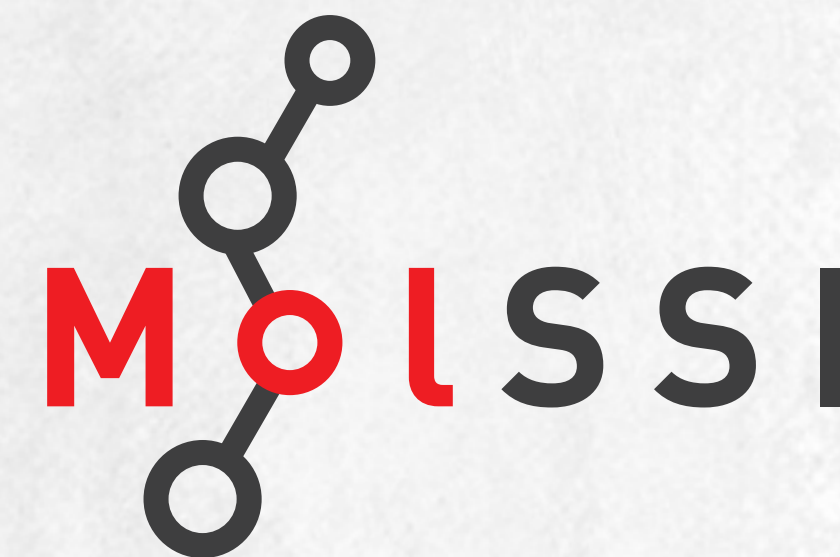
@dga\_smith

[qcarchive.molssi.org](http://qcarchive.molssi.org)



# QC Archive Overview

A central source to compile, aggregate, query, and share quantum chemistry data.



**QC Archive**

A MolSSI Project

## Design Goals

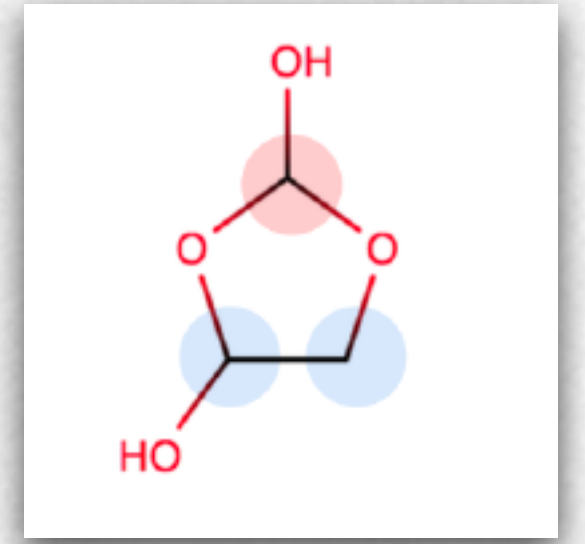
- Quantum chemistry data for all of the computational molecular sciences community.
- Analysis, visualization, and quick start data guides.
- Access data via Python, Jupyter notebook integration, REST API, and web apps.
- Evaluate on many community resources simultaneously.
- Store billions of quantum chemistry results.
- Prevent duplicate computation.
- Removing “the middle man”.

[qcarchive.molssi.org](http://qcarchive.molssi.org)



# Open Force Field

<https://openforcefield.org>

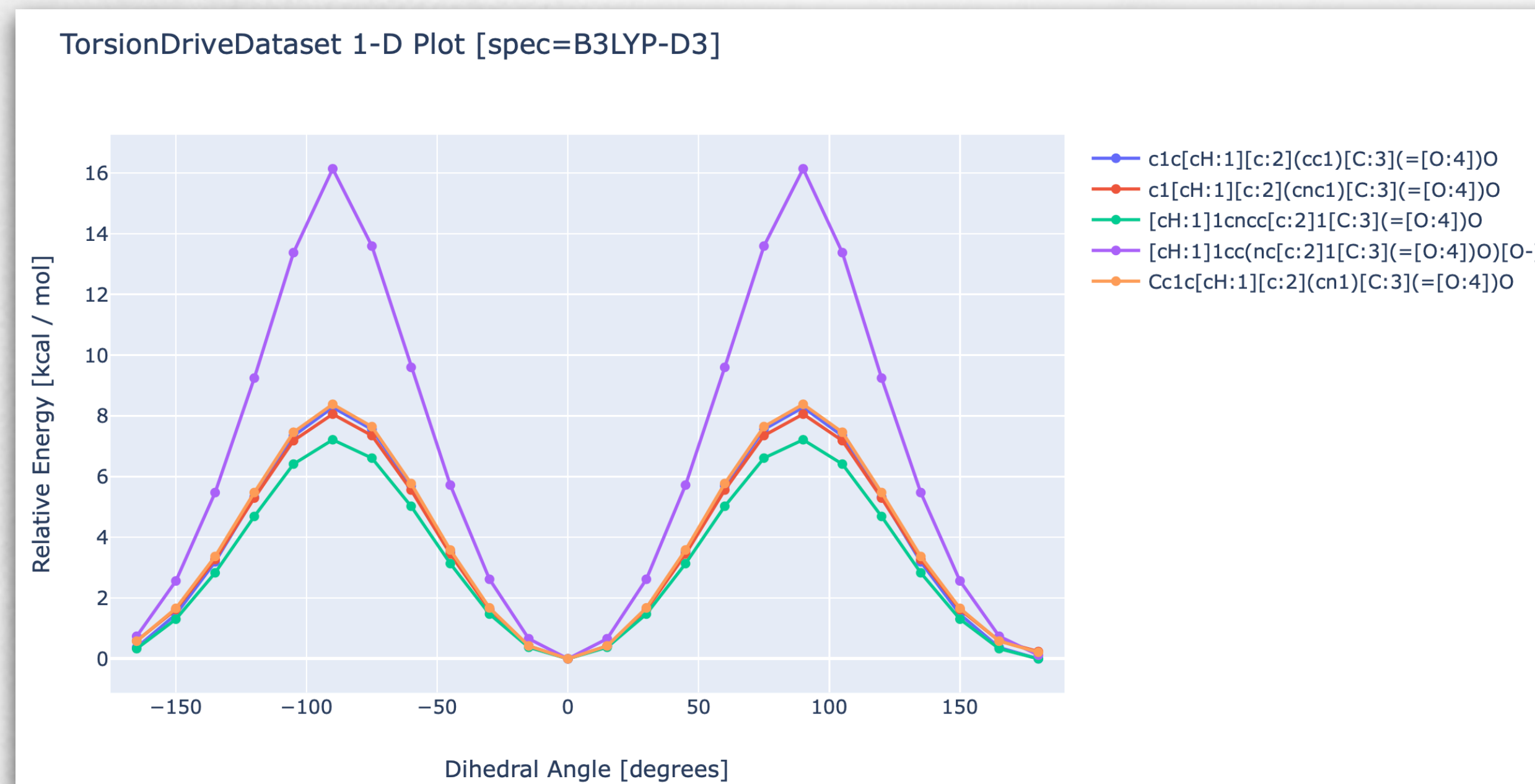


## Requirements

- Goal: Aggregate and compute open data for force field fitting, machine learning, and education.
- Store: Constrained geometry optimization, Torsion Drives, Hessians, partial charges, ESPs, and more!
- Search: SMILES, InChI, etc
- Compute: Multiple campus clusters, burst at XSEDE/DOE

## Computed (4 months)

- 2,400 torsion drives
- 170,000 geometry optimizations
- 50,000 Hessians
- ~60,000 geometry optimizations/week [limited by core time]



## QCArchive Sponsor

- Sponsoring features within QCArchive
- All open-source code, available to the community

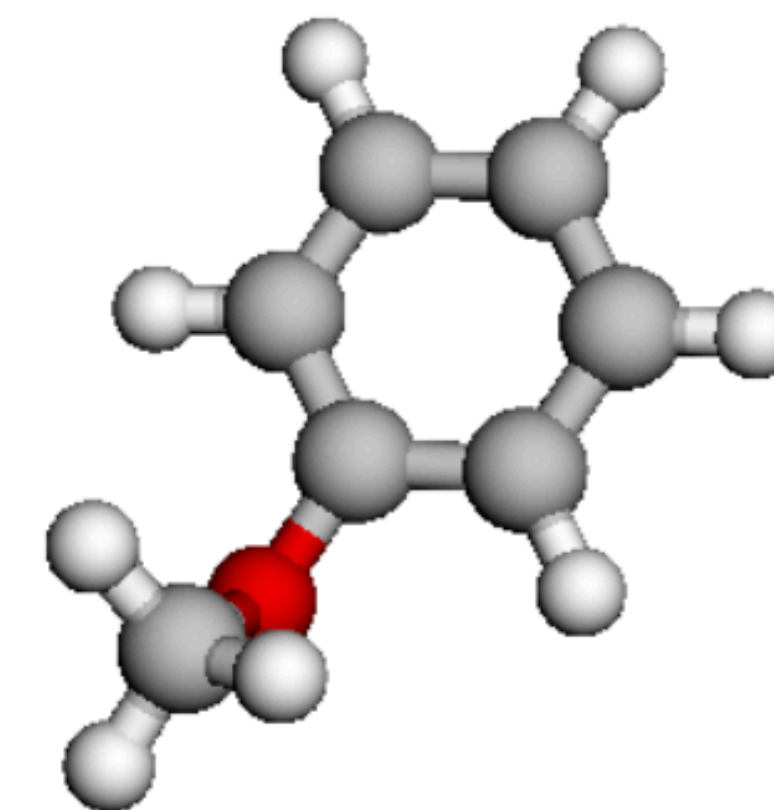


# Interactive and Gateway Sessions

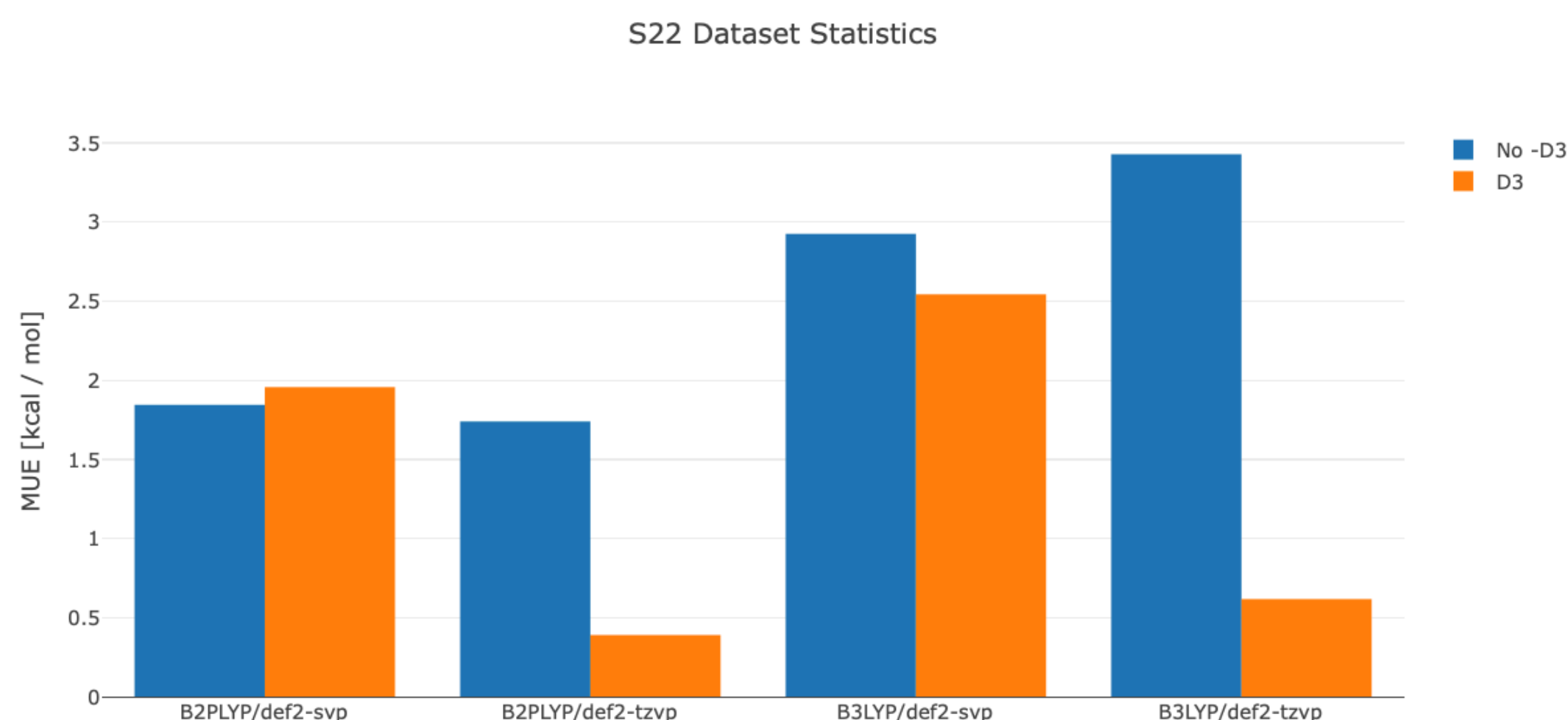
## Jupyter-Notebook Integration

- Molecular visualization, statistics, trajectories, etc
- Utilizing community-built and industry standard tools
- Interactive sessions to explore and compute new data
- Leveraging the greater Jupyter community of tools

```
td.get_final_molecules(90)
```



```
In [6]: ds.visualize(method=["B3LYP", "B3LYP-D3", "B2PLYP", "B2PLYP-D3"], basis=["def2-svp", "def2-tzvp"], groupby="D3")
```



## Web Apps

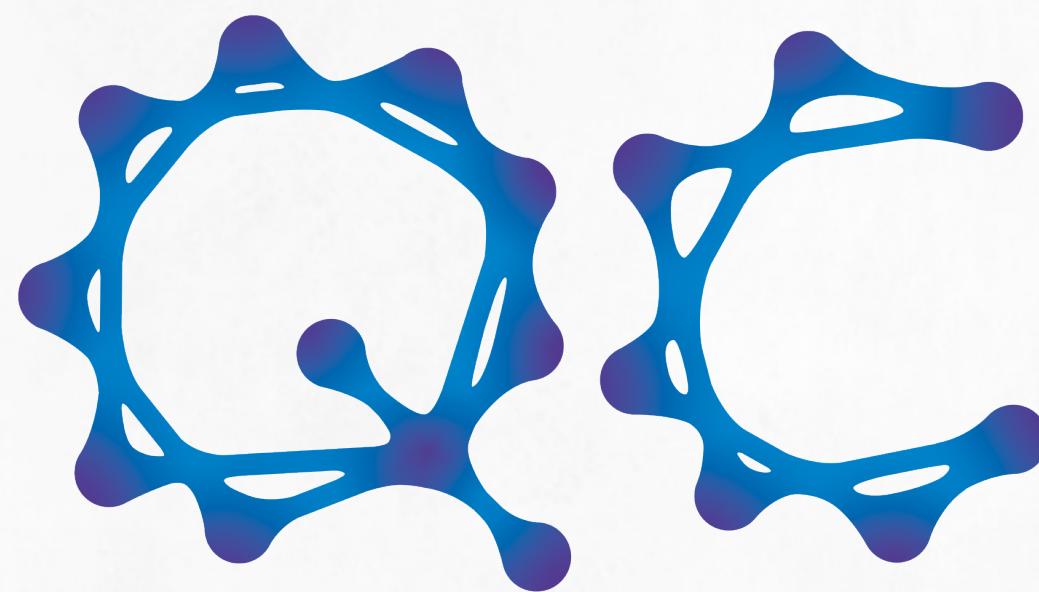
- Working in partnership with the Science Gateways Community Institute.
- Web-based statistics and visualization
- Targeting at CMS researchers and undergraduate educational initiatives
- Data-driven initiatives:
  - What is the best method for X
  - How long will X take?



# Engage with QCArchive

## View our data

- Browse our current 5.5M+ results and growing!
- Contribute “cookbook” use cases for interesting data applications.



**QC Archive**  
A MolSSI Project

## Extend our datasets

- Get in touch and help compute additional methods for our benchmark datasets.
- Add additional benchmark datasets to the Archive.

## Compute Open Data

- Work with us to compute additional open data.
- Expand our use cases for a web app framework.

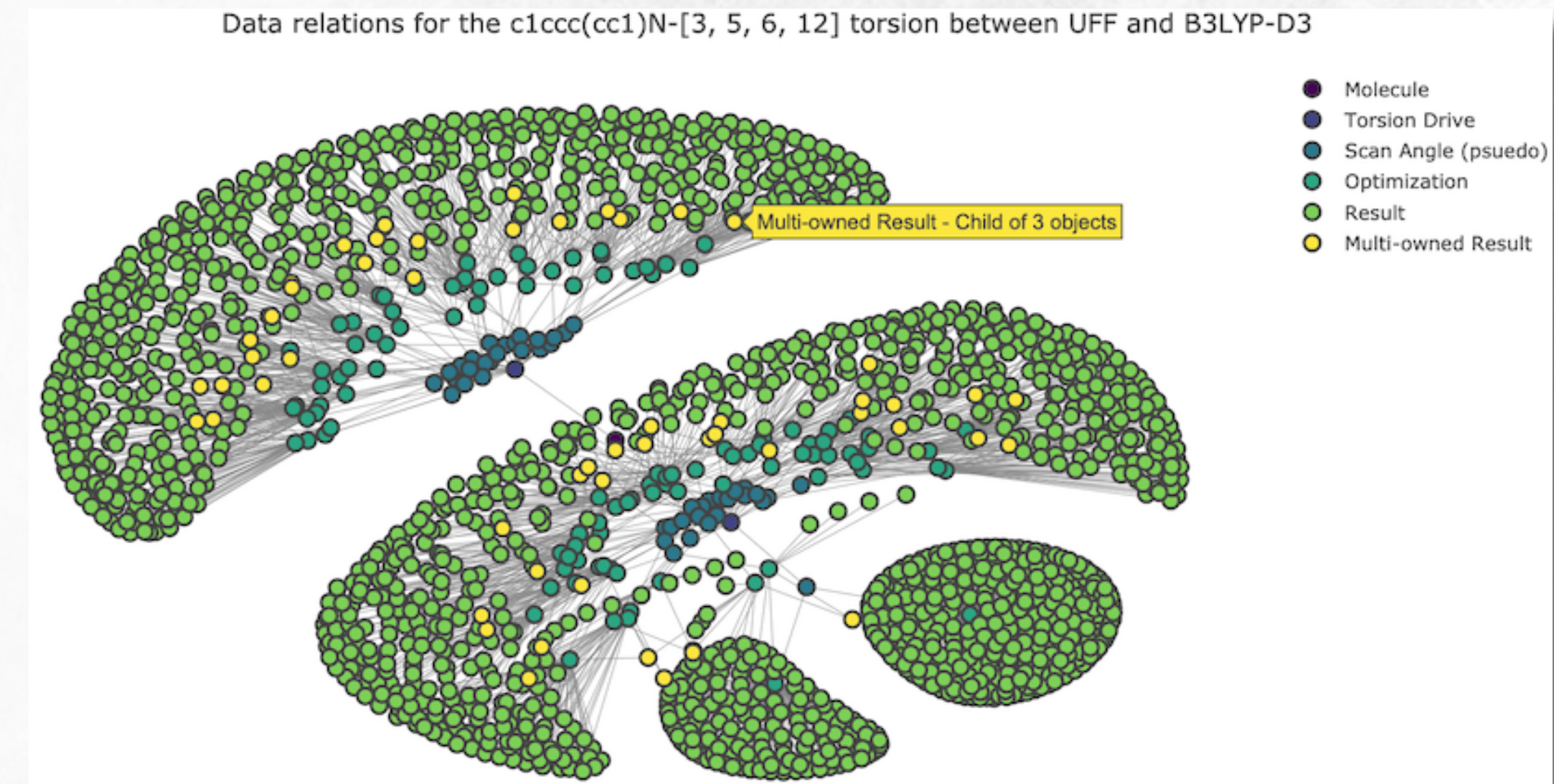
## Talk to us

- Tell us about science applications that we are missing.
- Chat about how to refine current presentations and ideas.



# QCArchive Infrastructure

- Quantum chemistry software projects for all CMS developers
- Composable building blocks
- Developed openly on GitHub ([github.com/MolSSI](https://github.com/MolSSI))
- Used by dozens of downstream programs



## QCSchema

- Standardized IO for quantum chemistry

## QCElemental

- Units
- QCSchema Models
- Molecule Parsing
- Visualization

## QCEngine

- Consume and produce QCSchema for many programs
- Not just quantum chemistry

## QCFractal

- High-throughput quantum chemistry
- Common pipelines
- Data Organization
- Visualization



# QCSchema

<https://github.com/MolSSI/QCSchema>

- Communication channel between all piece of the ecosystem.
- *Community* project useful for many aspects of quantum chemistry.
- Not only JSON, but any key/value/array language (BSON/HDF5/XML/YAML/msgpack/parquet)

```
{
  "molecule": {
    "geometry": [0, 0, 0, 0, 0, 1],
    "atoms": ["He", "He"]
  },
  "driver": "energy",
  "model": {
    "method": "SCF",
    "basis": "sto-3g",
  },
}
```



- Molecule
- QC Input/Output
- Optimization Structures
- Wavefunction Quantities

```
{
  ...Input
  "provenance": {
    "creator": "My QM Program",
    "version": "1.1rc1",
  },
  "properties": {
    "scf_n_iterations": 2.0,
    "scf_total_energy": -5.433191881443323,
    "nuclear_repulsion_energy": 2.11670883,
    "one_electron_energy": -11.67399006298,
    ..
  },
  "error": "",
  "success": true,
}
```



# QCEngine

<https://github.com/MolSSI/QCEngine>

- Quantum chemistry, semiempirical, AI energy evaluator, and force field agnostic backend to produce/consume Schema. Effectively our compute abstraction layer.
- Modular building block approach

```
>>> geometric_task = {
  "keywords": {
    "coordsys": "tric",
    "program": "rdkit"
  },
  "input_specification": {
    "driver": "gradient",
    "model": {"method": "UFF"},
  },
  "initial_molecule": qcengine.get_
```

```
>>> ret = qcengine.compute_procedure(
>>> ret.final_molecule.geometry
[0.0, 0.0, -0.1218741,
 0.0, -1.47972431, 1.02364509,
 0.0, 1.47972431, 1.02364509]
```

```
>>> geometric_task = {
  "keywords": {
    "coordsys": "tric",
    "program": "torchani"
  },
  "input_specification": {
    "driver": "gradient",
    "model": {"method": "ANI1"},
  },
  "initial_molecule": qcengine.get_
```

```
>>> ret = qcengine.compute_procedure(
>>> ret.final_molecule.geometry
[0.0, 0.0, -0.1123205,
 0.0, -1.4331881, 1.0188681,
 0.0, 1.4331881, 1.0188682]
```

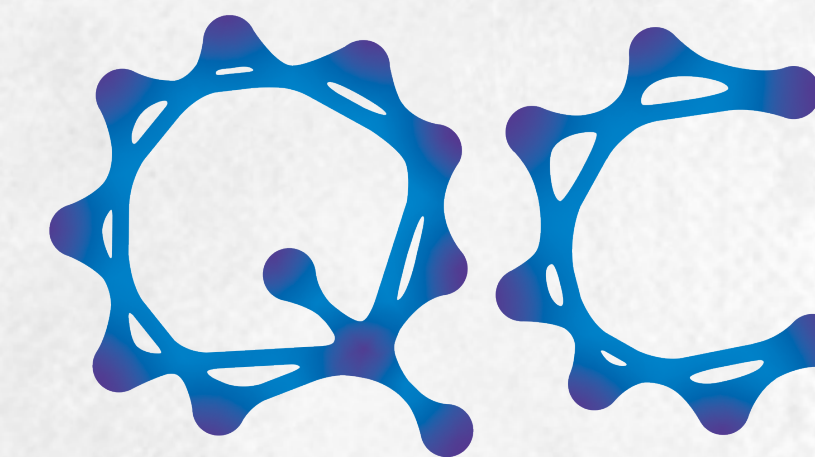
```
>>> geometric_task = {
  "keywords": {
    "coordsys": "tric",
    "program": "psi4"
  },
  "input_specification": {
    "driver": "gradient",
    "model": {"method": "WB97X-D"},
  },
  "initial_molecule": qcengine.get_
```

```
>>> ret = qcengine.compute_procedure(
>>> ret.final_molecule.geometry
[0.0, 0.0, -0.0690161,
 0.0, -1.49345852, 0.9901583,
 0.0, 1.49345852, 0.9901583]
```



# QCFractal

<https://github.com/MolSSI/QCFractal>

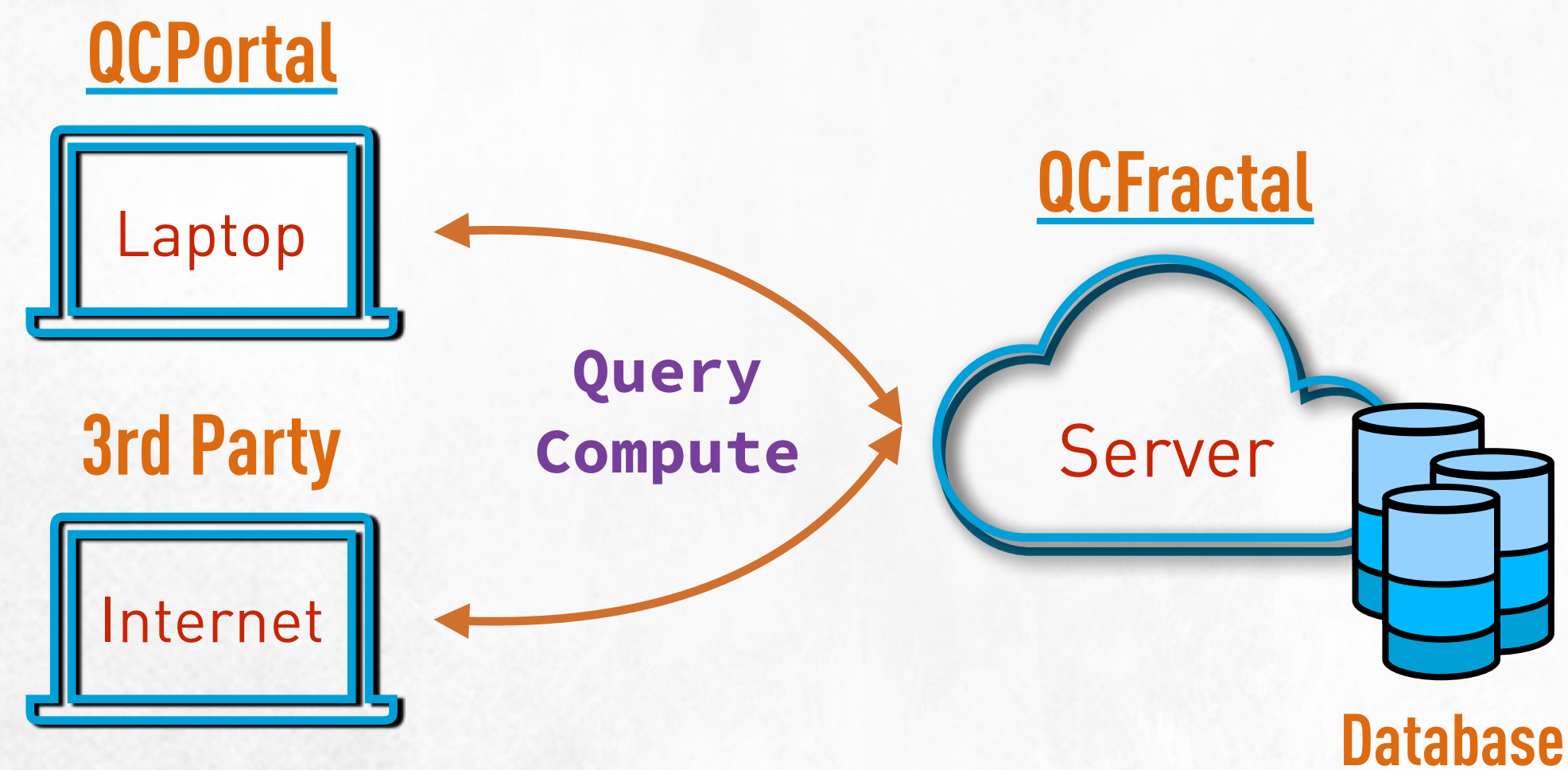


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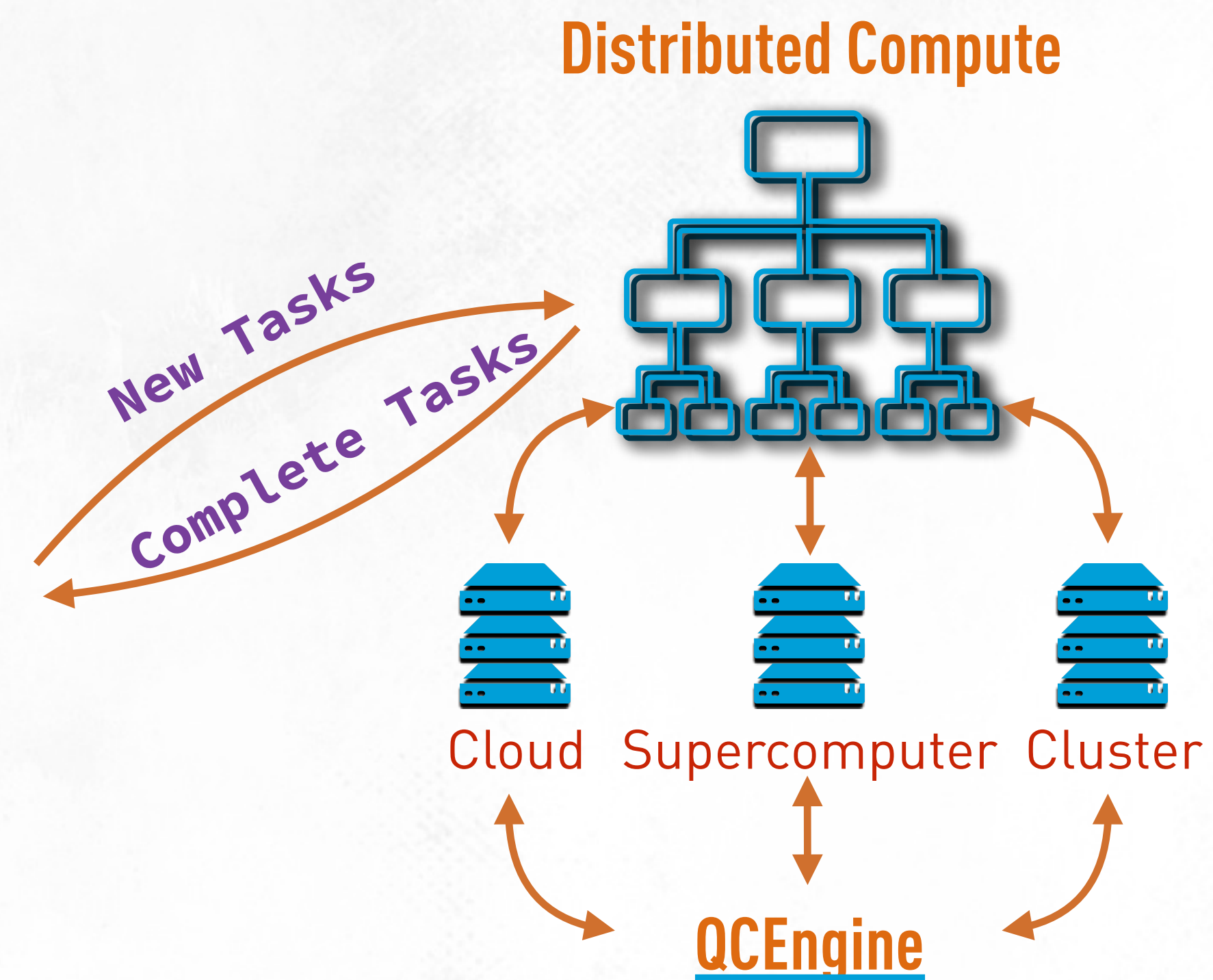
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## Goals:

- High-throughput quantum chemistry
- Laptop to campaign-scale compute orchestration
- Procedures run with a variety of different programs
- Organize data with common abstraction and collection layers
- Share and collaborate structured data
- Ease of use, less data parsing



[qcarchive.molssi.org](http://qcarchive.molssi.org)





# Reproducible Procedures and Workflows

## Procedures

- Procedures = small reproducible series of computations
- Exact input of pipeline and version data available
- Geometry optimizations, torsion evaluations, finite difference computations, spectral computations, etc

```
optimization = client.query_procedures(procedure="optimization", id=1724500)[0]
```

```
optimization
```

```
<OptimizationRecord(id='1724500' status='COMPLETE')>
```

```
optimization.keywords
```

```
{'coordsys': 'tric',  
 'enforce': 0.1,  
 'reset': True,  
 'qccnv': True,  
 'epsilon': 0,  
 'constraints': {'set': [{'type': 'dihedral',  
   'indices': [1, 0, 4, 2],  
   'value': -45}]},  
 'program': 'psi4'}
```

```
ds.get_history(method="B3LYP-D3M")  
ds.df.head()
```

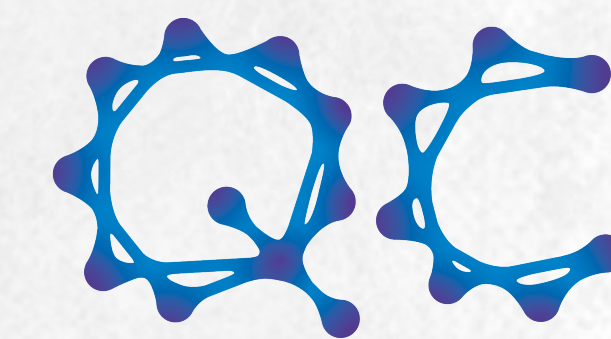
|                          | S220   | S22a   | S22b    | B3LYP-D3M/def2-svp | B3LYP-D3M/def2-tzvp |
|--------------------------|--------|--------|---------|--------------------|---------------------|
| <b>Ammonia Dimer</b>     | -3.17  | -3.15  | -3.133  | -6.248386          | -4.049052           |
| <b>Water Dimer</b>       | -5.02  | -5.07  | -4.989  | -9.002674          | -6.427460           |
| <b>Formic Acid Dimer</b> | -18.61 | -18.81 | -18.753 | -25.933297         | -20.668411          |
| <b>Formamide Dimer</b>   | -15.96 | -16.11 | -16.062 | -21.689185         | -17.436781          |
| <b>Uracil Dimer HB</b>   | -20.65 | -20.69 | -20.641 | -25.623412         | -21.922461          |

## Collections

- After 100+ interviews there seems to be very little common ground on data organization.
- Many single computations or procedure grouped together known as *Collections*
- Reproducible, recomputable, and tweakable
- Data organization for ML, methodology assessment, forcefield creation, etc

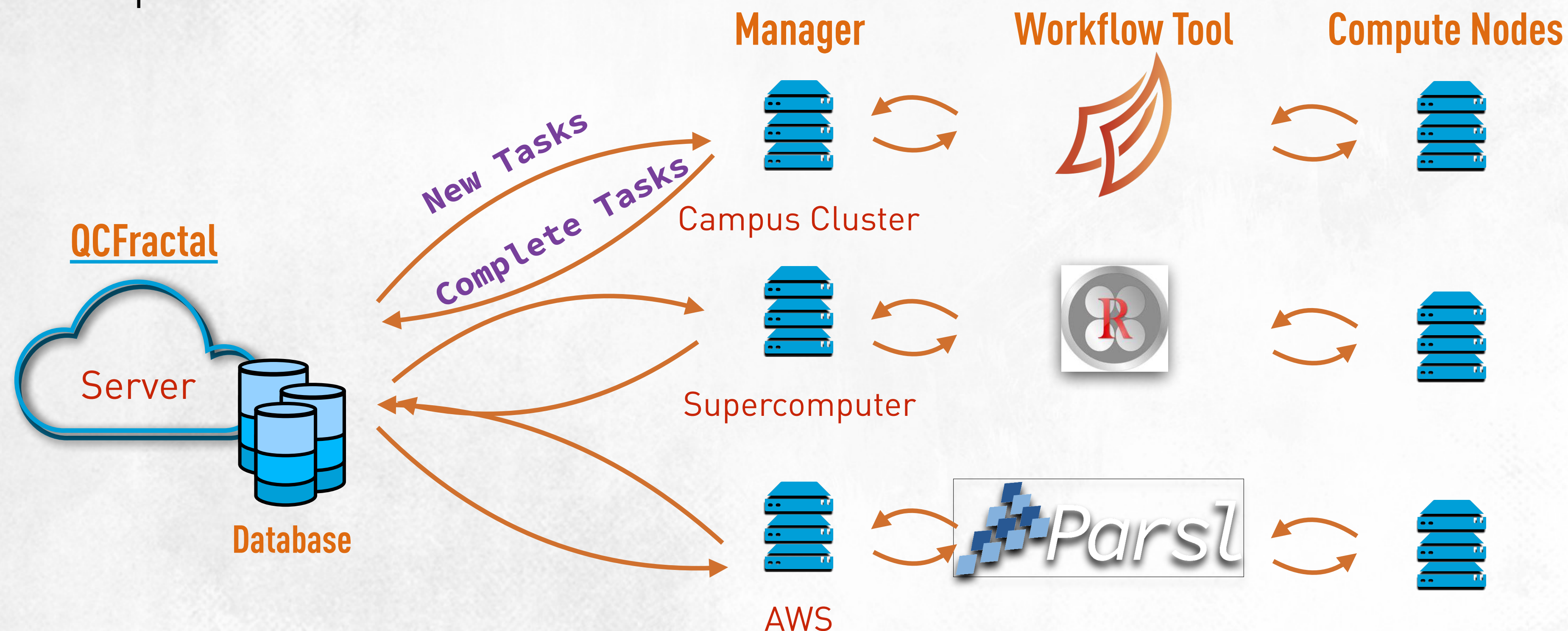


# Distributed compute



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- Multi-physical-site compute or a single laptop
- Scale up to 500 tasks/second, 300,000 concurrent tasks @ 10 minutes each
- Setup once and walk away
- Managers:
  - Runs on head node or local compute
  - Smart task acquisition
  - ~20 ms per-task overhead



Additional Tools



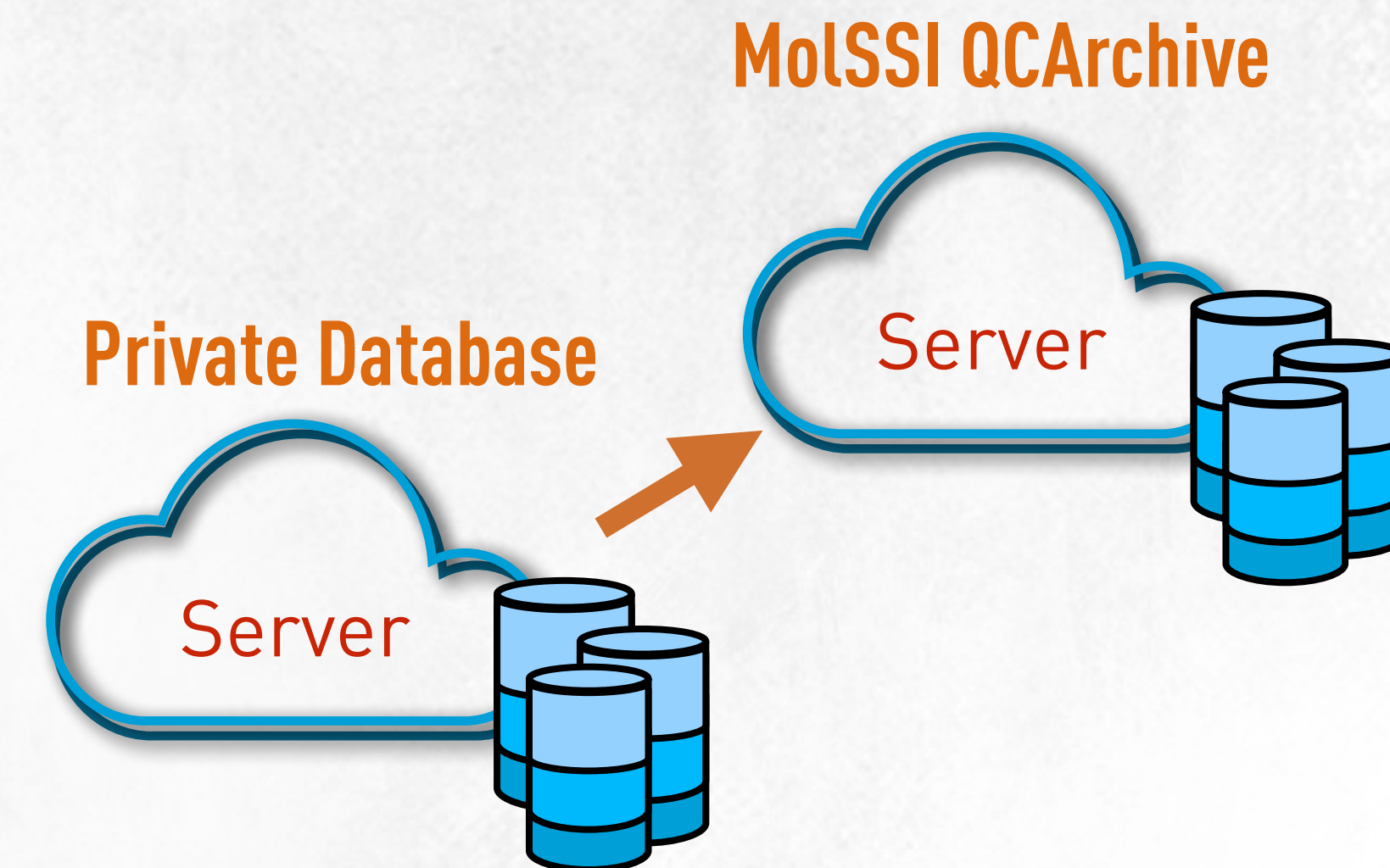


# MolSSI and Self-Hosted Databases

- A domain specific SQL database layer
- Generation and computation of new quantum chemistry tasks
- Central MolSSI-hosted server for community data accessed via REST or Python API
- Open-software (QCFractal) used at scale at MolSSI, research groups, and individuals

## Self-Hosted

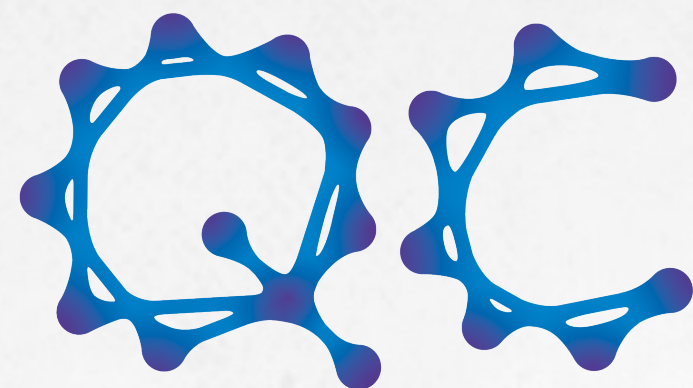
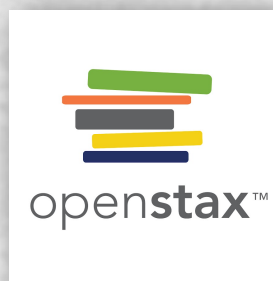
- Long-term private data with access controls
- (or) Quick testing and evaluation environments
- Can migrate data to central MolSSI server after publication
- Identical infrastructure and technology as MolSSI central repository



## MolSSI QCArchive

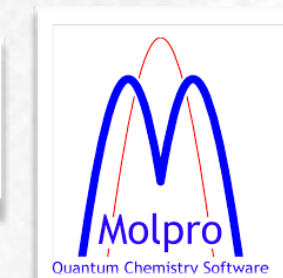
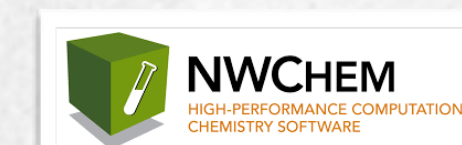
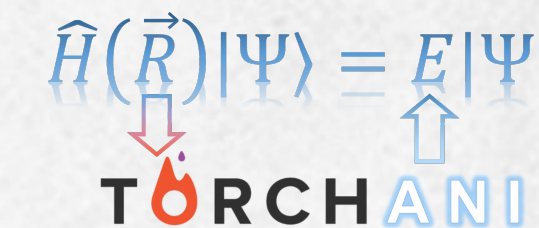
- Open community data
- FAIR Data standards
- ~5.5M current results
- ~60 community datasets
- Can host ~1B results with current hardware, looking to expand!





QC Archive  
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# CMS and Community Software

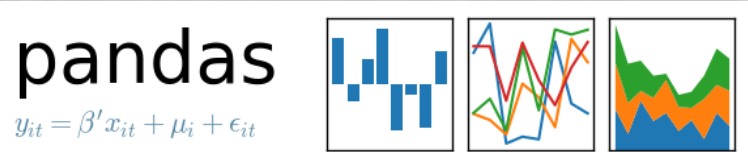


## Gateways Portal

- Reach non-CMS community
- Research-focused web portals
- Educational initiatives



3dMolJs

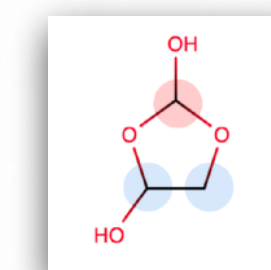


Beta as of August 26th  
 Monthly substantial releases  
 Use cases from 100+ research groups  
 60,000+ downloads  
 4.5M computations  
 Rapidly expanding

## Software Developers



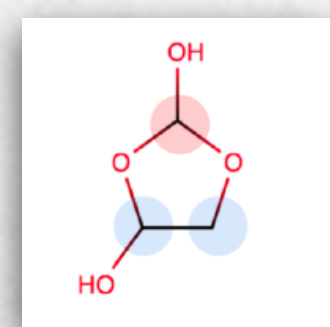
OptKing



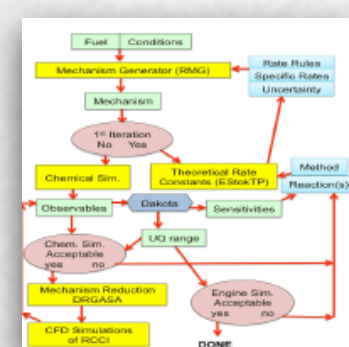
## View and Analyze data

- Large-scale analytics
- Valuable community insights

## The MolSSI Community Database



The Open Force Field Consortium



PACChem



## Private Databases

- Custom ML Datasets
- Methodology assessment
- Modern access to QC computations

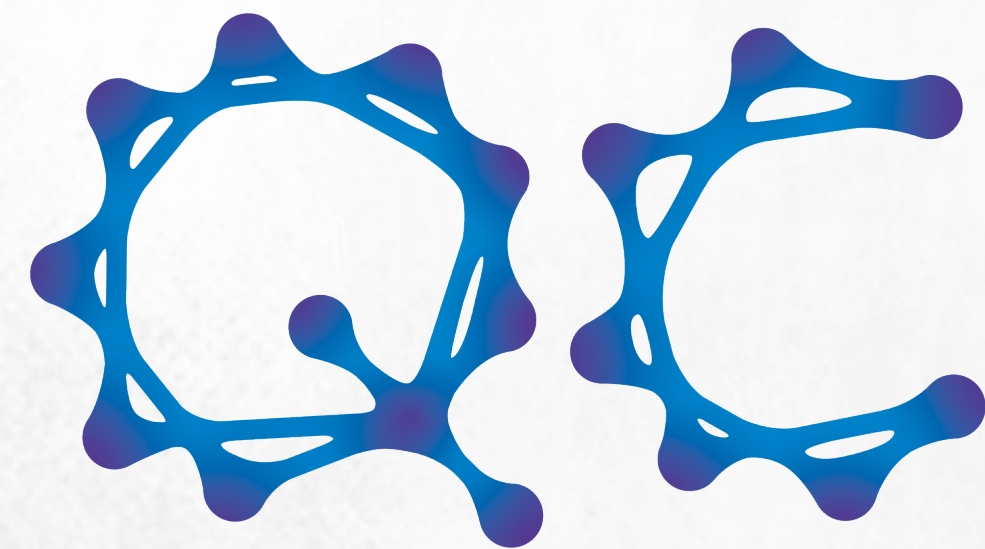
## Cyberinfrastructure





# Thank You!

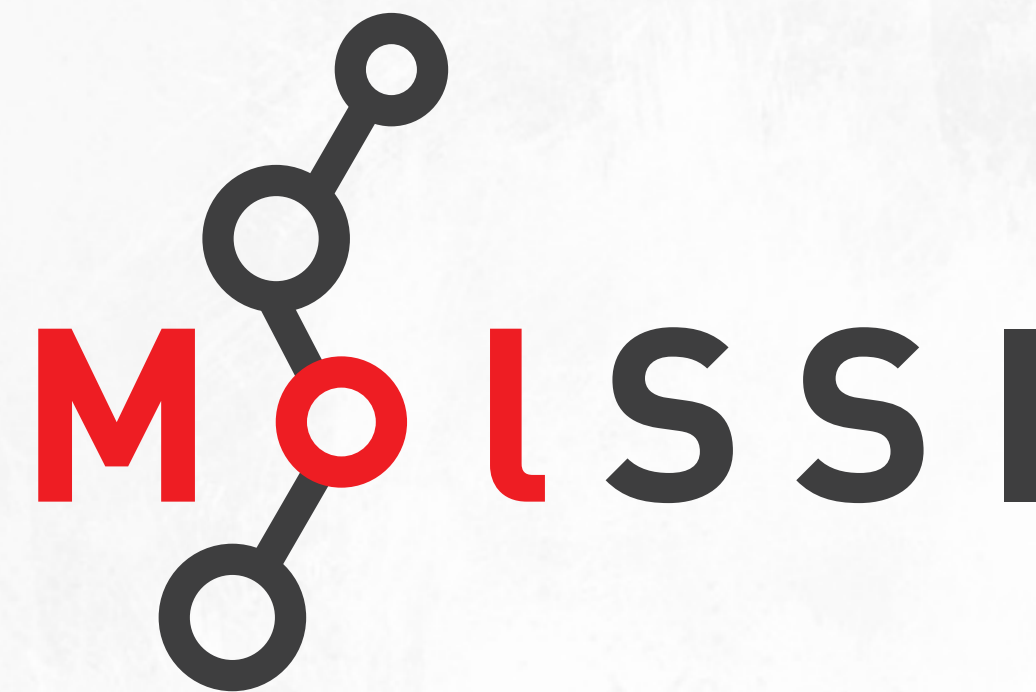
- Doaa Altarawy (MolSSI)
- Levi Naden (MolSSI)
- Matt Wellborne (MolSSI)
- Lori Burns (Georgia Tech)
- Sam Ellis (MolSSI)
- Jessica Nash (MolSSI)
- Ben Pritchard (MolSSI)
- Chaya Sten (MSKCC)
- Yudong Qiu (UC Davis)
- Fang Liu (MIT)
- Sebastian Lee (Cal Tech)
- David Sherrill (Georgia Tech)
- Daniel Crawford (MolSSI)
- Lee-Ping Wang (UC Davis)
- Jeff Wagner (UCI)
- John D. Chodera (MSKCC)
- Dom Sirianni (Georgia Tech)
- Daniel Nascimento (PNNL)
- Nick Petosa (Microsoft)
- Justin Turney (UGA)
- Bert de Jong (LBNL)
- Theresa Windus (Iowa State)
- Aaron Virshup (Arzeda)
- Marcus Hanwell (Kitware)
- Shantenu Jha (Rutgers)
- Matteo Turilli (Rutgers)
- Kyle Chard (U. Chicago)



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[molssi.org](http://molssi.org)

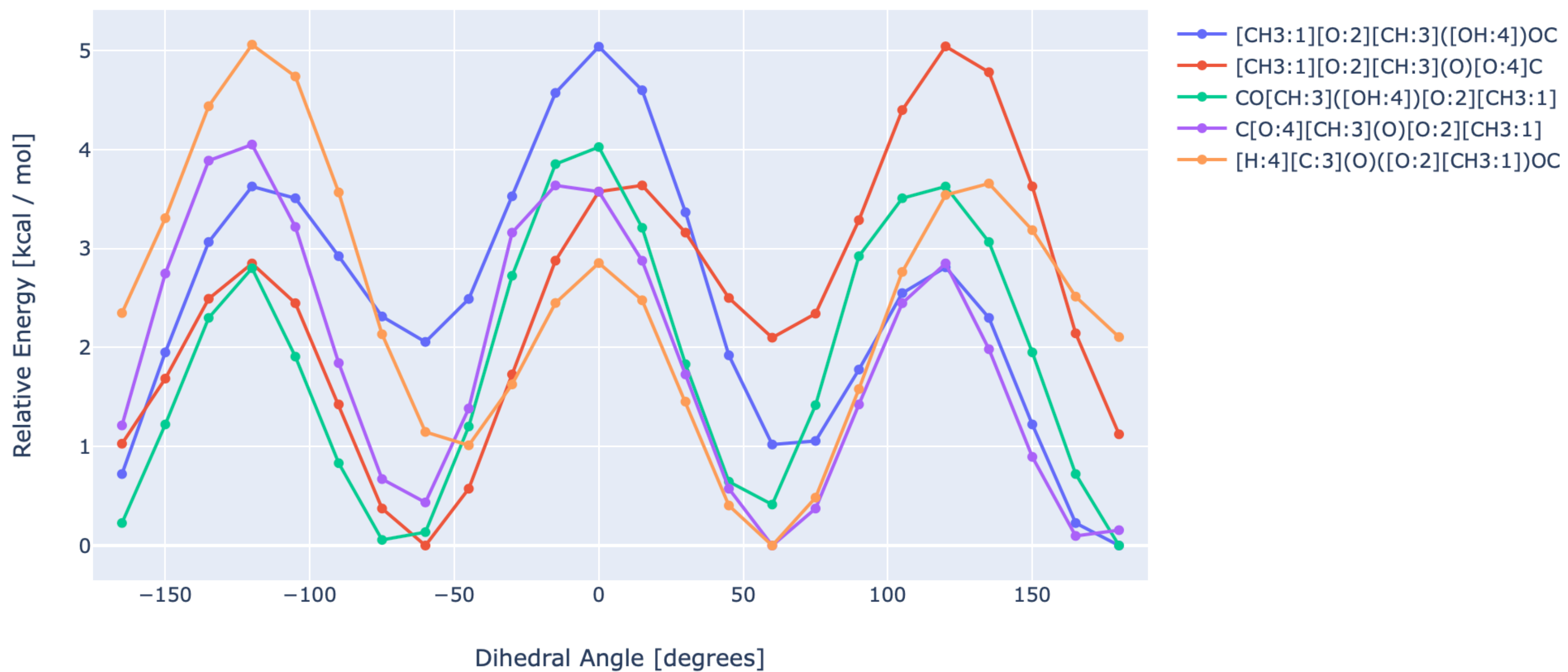


```
In [13]: import qcportal as ptl
```

```
In [14]: client = ptl.FractalClient()
ds = client.get_collection("TorsiondriveDataset", "SMIRNOFF Coverage Torsion Set 1")
```

```
In [15]: ds.visualize(['[CH3:1][O:2][CH:3]([OH:4])OC',
                      '[CH3:1][O:2][CH:3](O)[O:4]C',
                      'CO[CH:3]([OH:4])[O:2][CH3:1]',
                      'C[O:4][CH:3](O)[O:2][CH3:1]',
                      '[H:4][C:3](O)([O:2][CH3:1])OC'],
                  "default")
```

TorsionDriveDataset 1-D Plot [spec=default]





# Notebook Demonstration

[https://docs.qcarchive.molssi.org/en/latest/basic\\_examples/torsiondrive\\_datasets.html](https://docs.qcarchive.molssi.org/en/latest/basic_examples/torsiondrive_datasets.html)