

MANAGING COMPLEXITY: LESSONS FROM GROWING AND MAINTAINING THE Psi4 Ecosystem

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REUSEABLE LIBRARIES FOR QUANTUM CHEMISTRY, MAJVIK, FINLAND

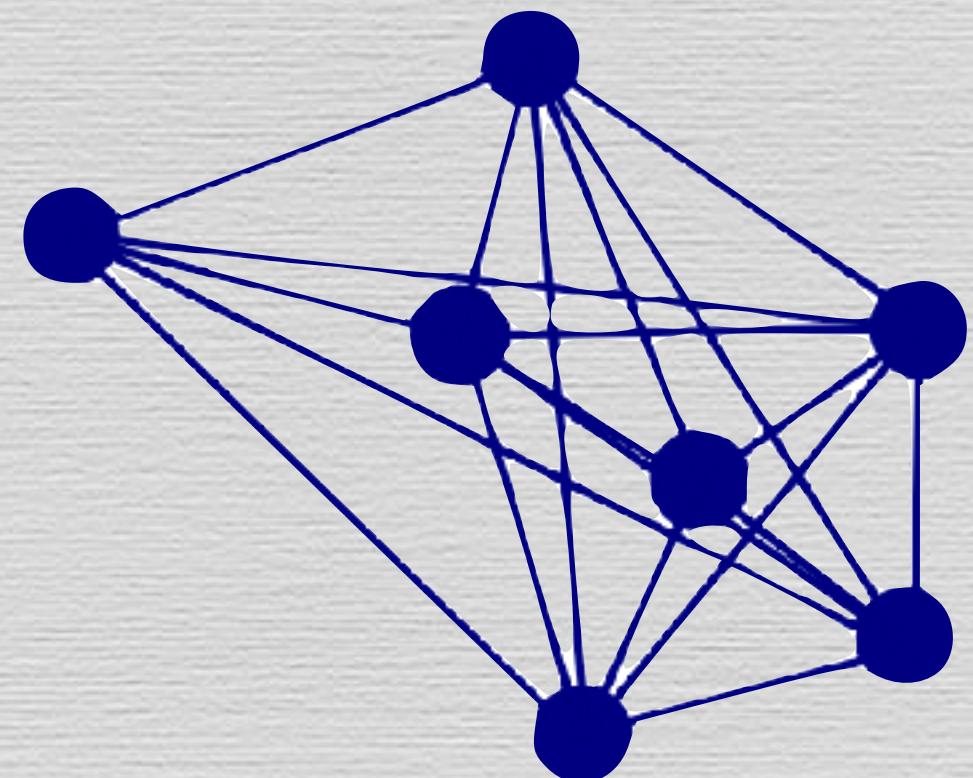
1 JULY 2025



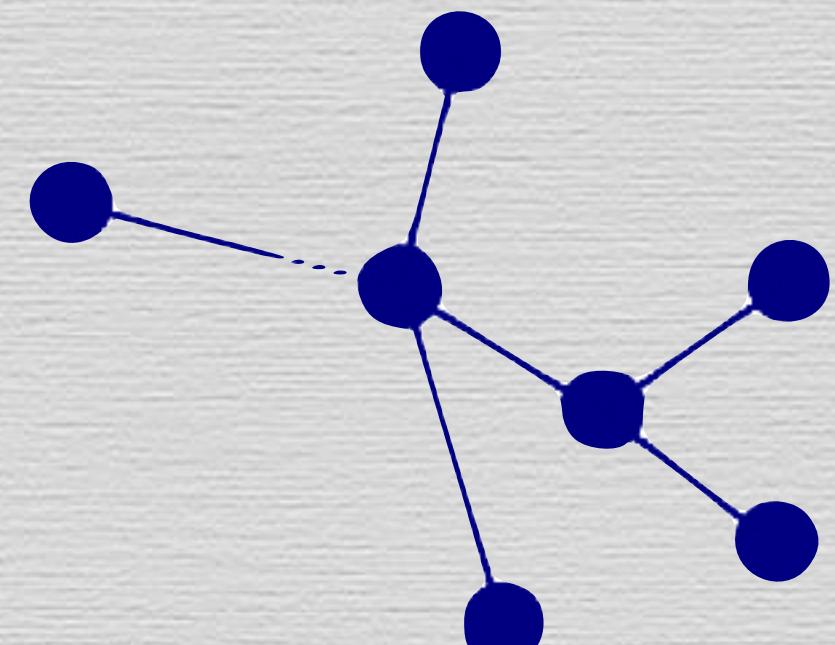
REFACTORING AN ECOSYSTEM

prefer loose coupling and high cohesion

STRONG COUPLING

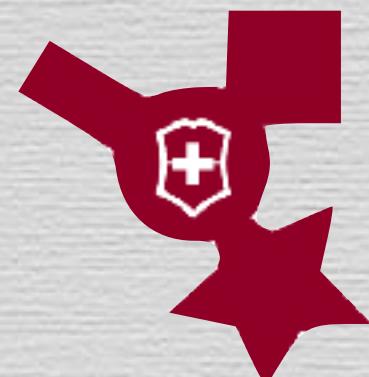


LOOSE COUPLING



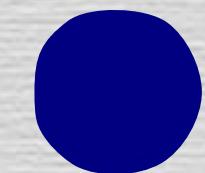
- easier to understand
- easier to compose

LOW COHESION



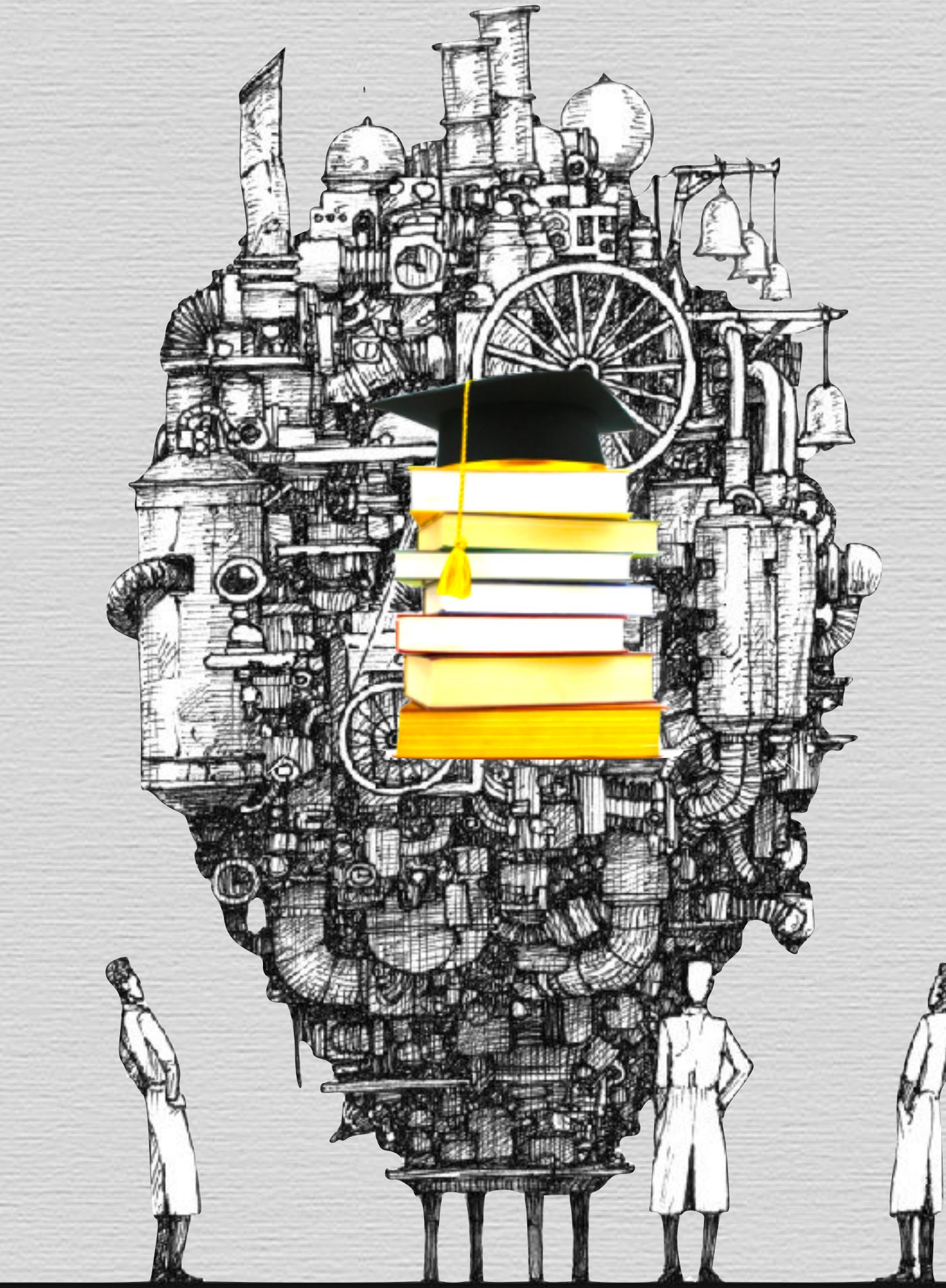
- difficult to maintain, test, read
- Swiss army knife modules

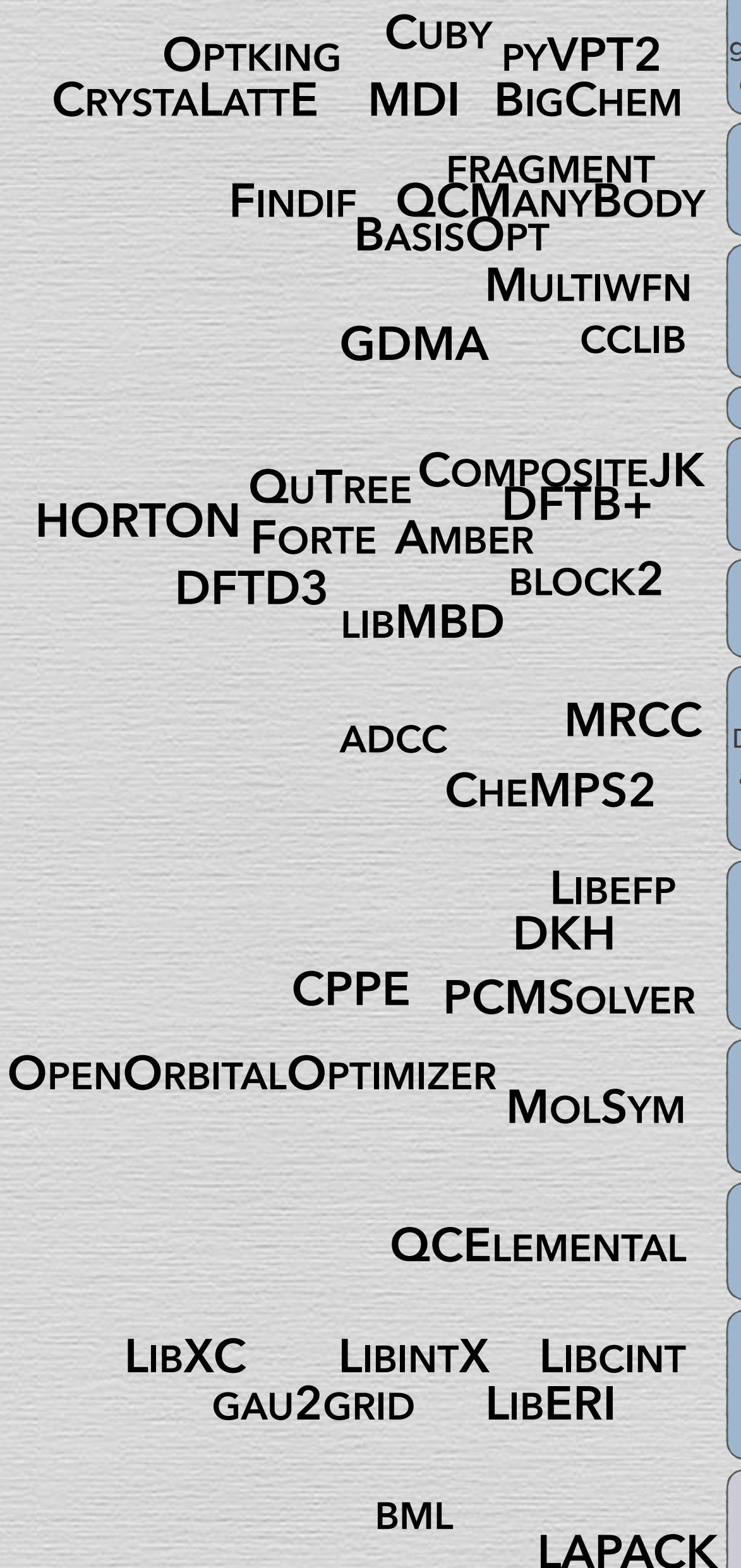
HIGH COHESION



- robust, reuseable, understandable
- stable APIs
- do one thing only and do it well
(e.g., Unix command line)

QC PROGRAMS





maximal observed monolithification

QC PROGRAMS

minimal desired monolithification

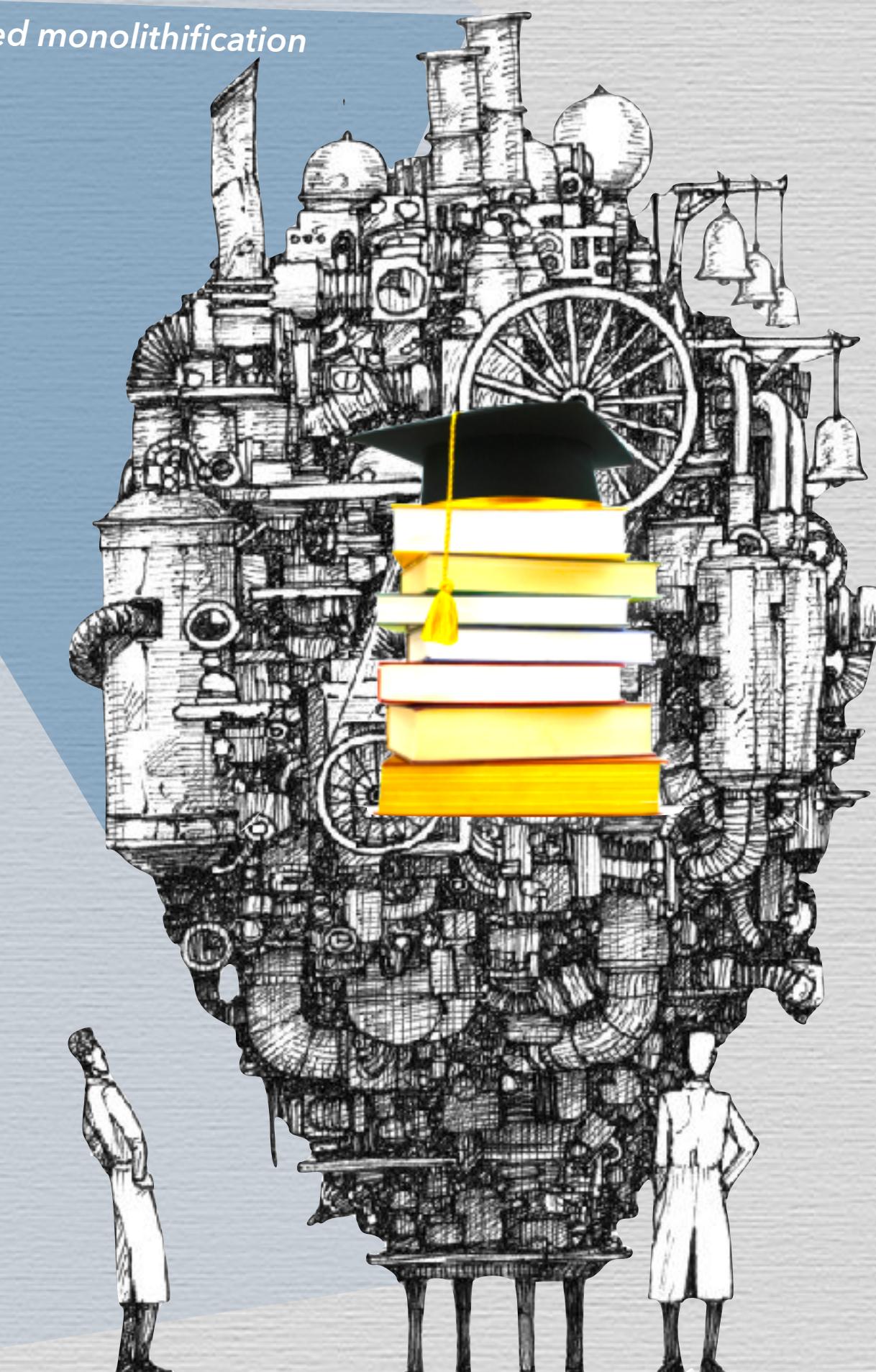
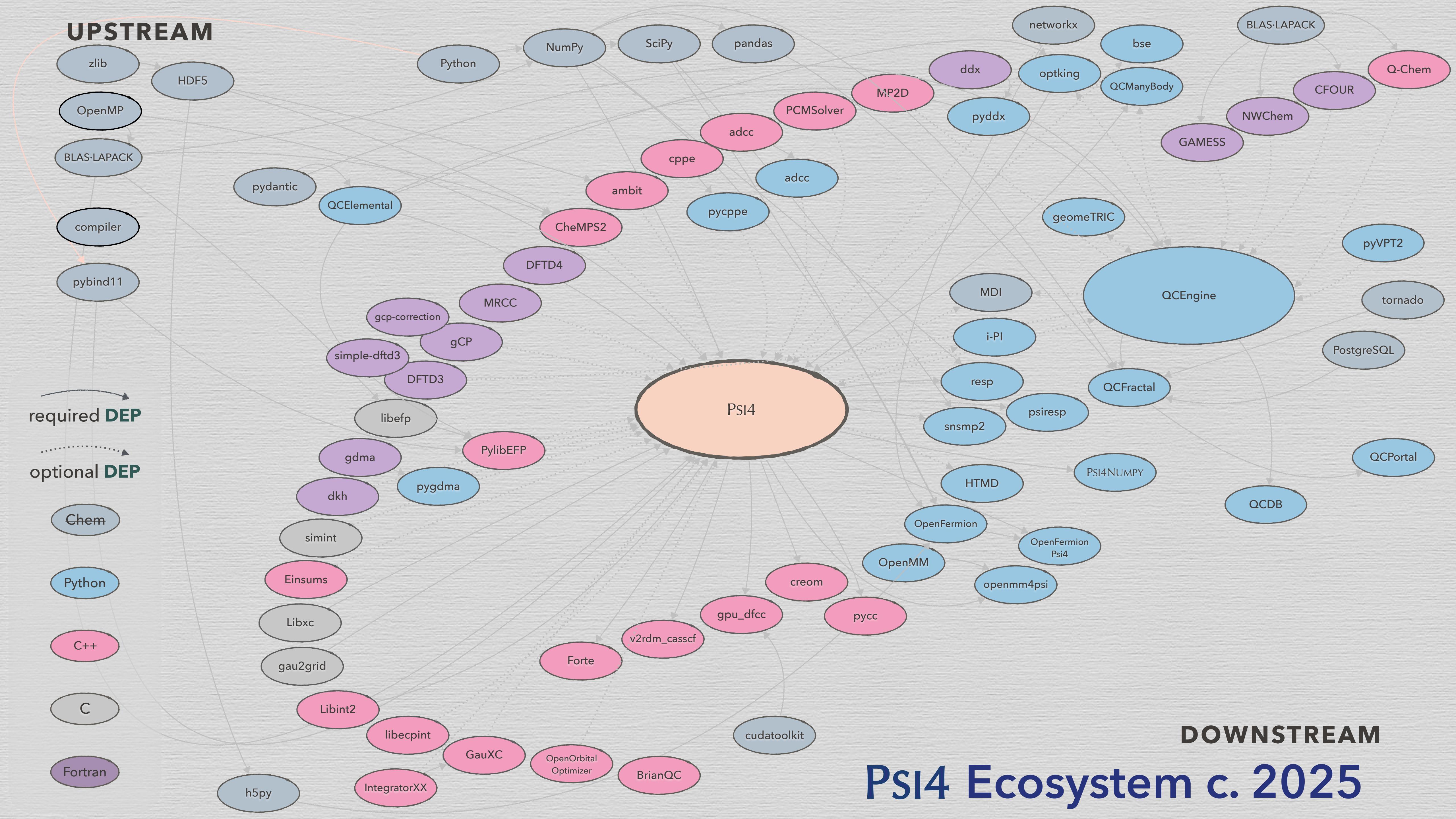
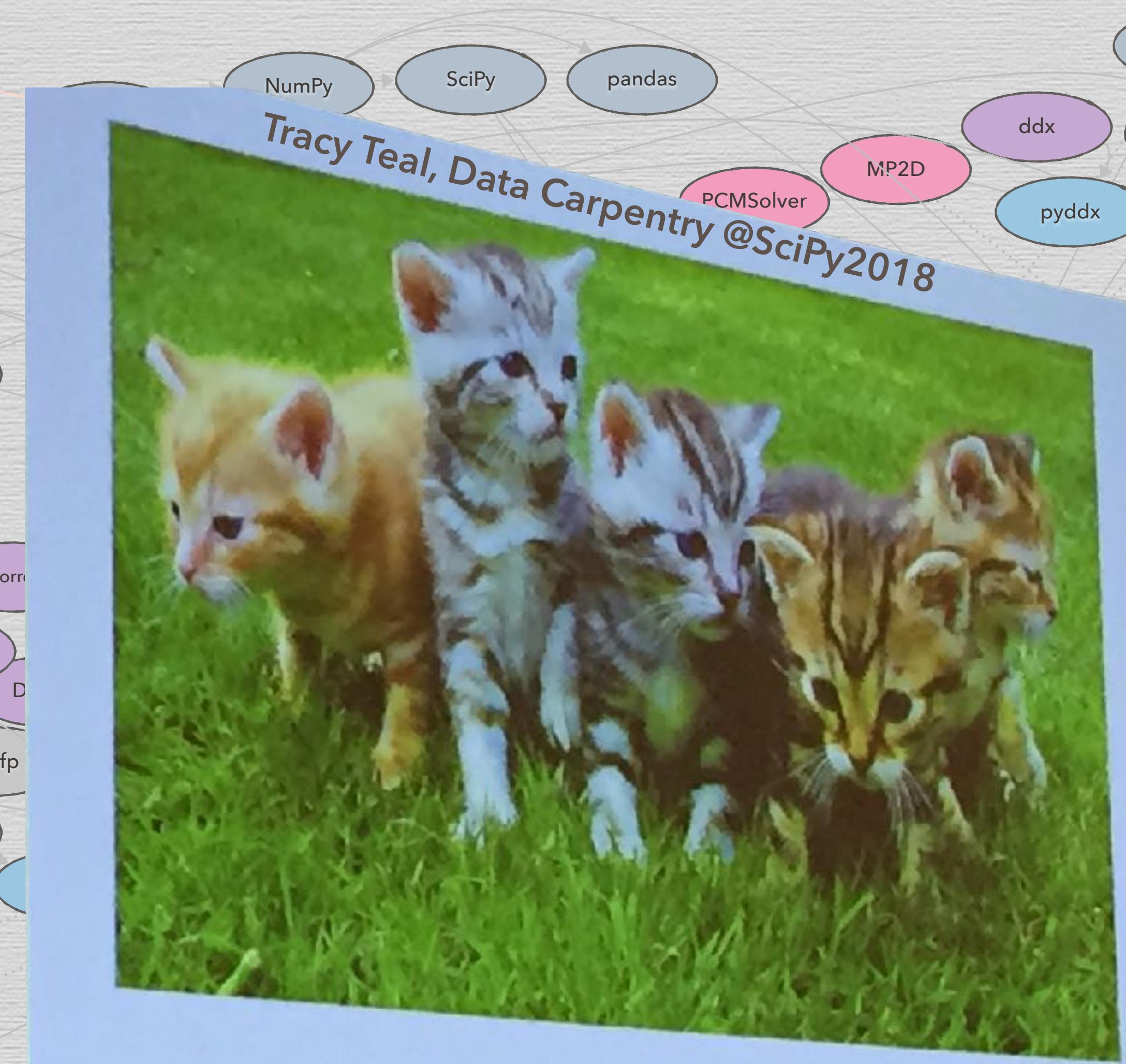
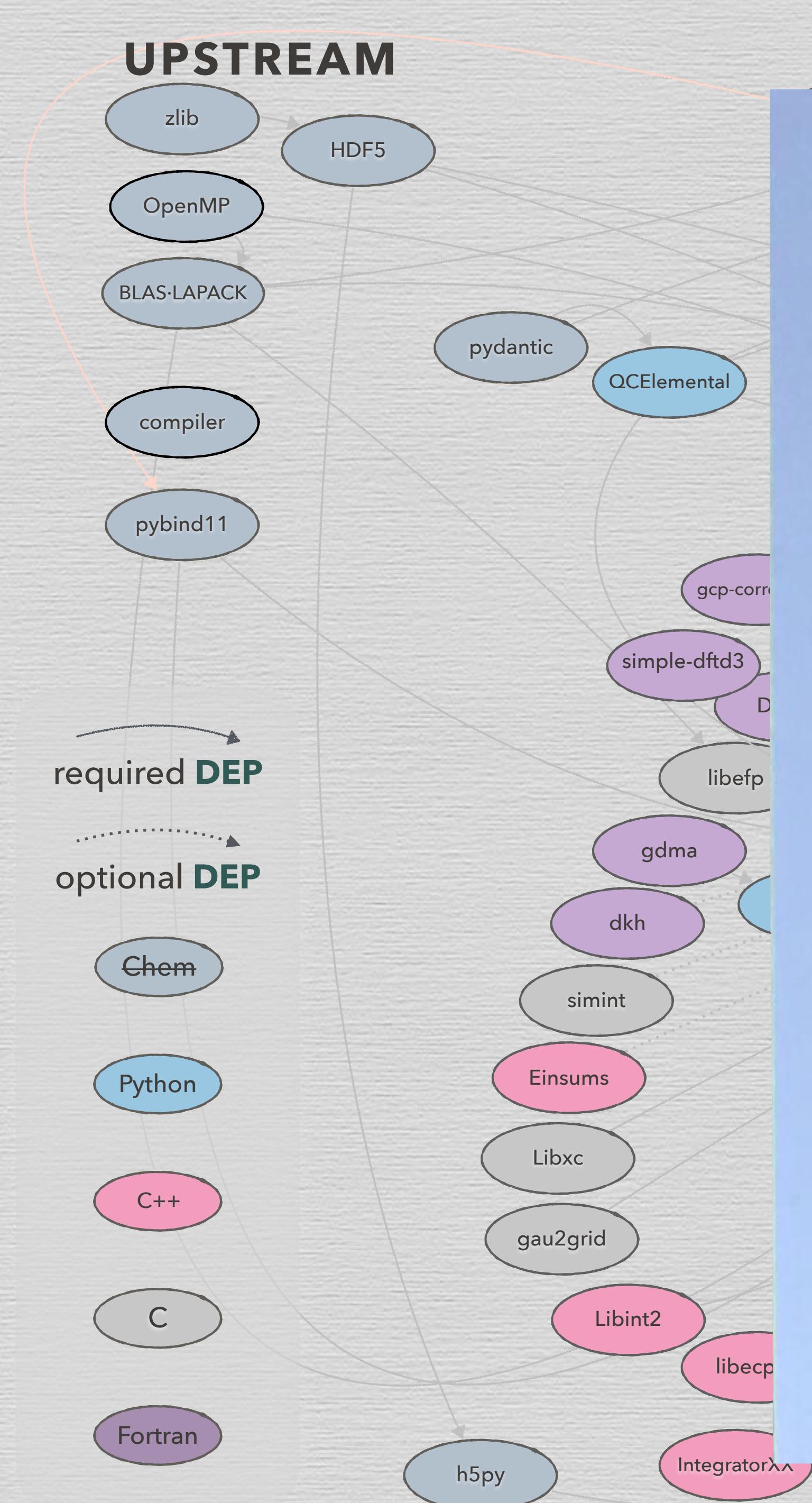


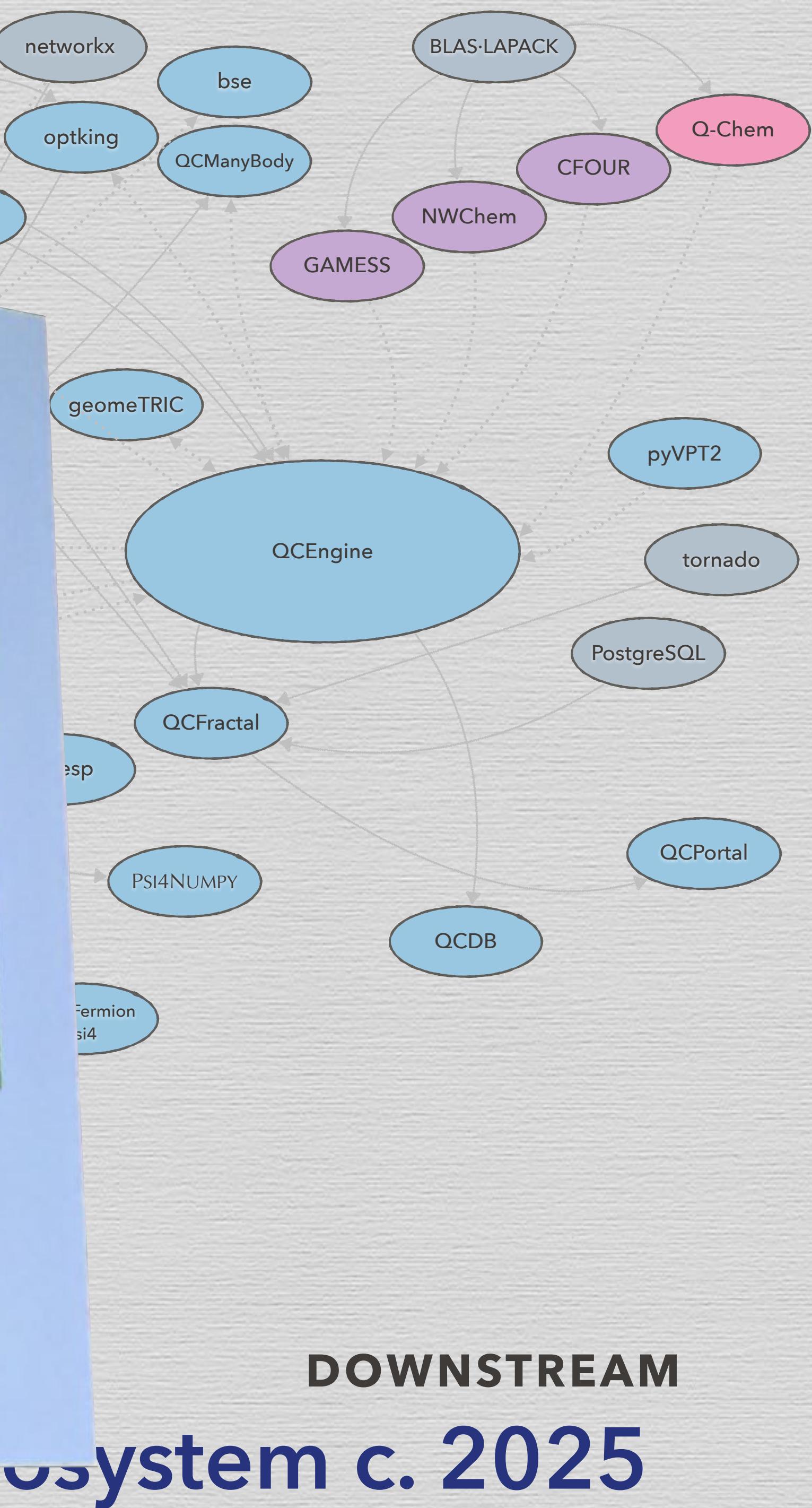
figure from *Electronic Structure (2024)* chapter

Blum, Windus, Burns, Oliveira, Pritchard, Slipchenko, et al.

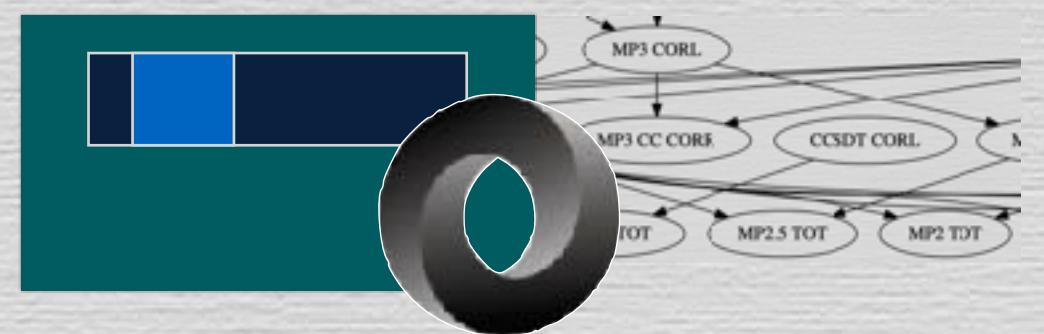




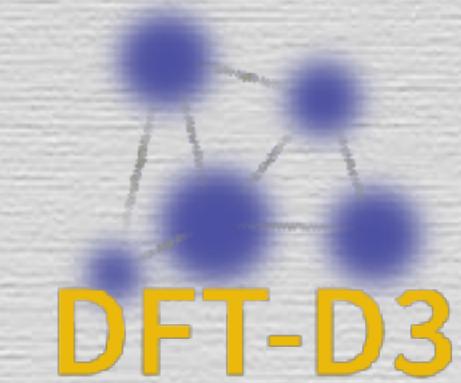
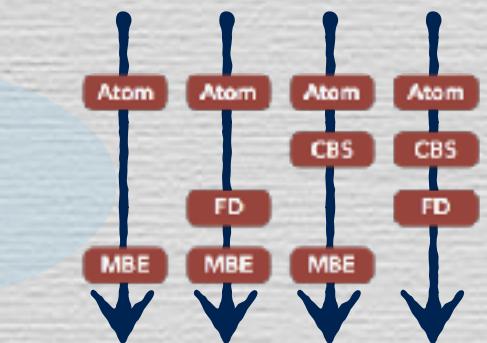
If you want to go fast, go alone.
If you want to go in a lot of different
uncoordinated directions at once, go together.



QC VARIABLES

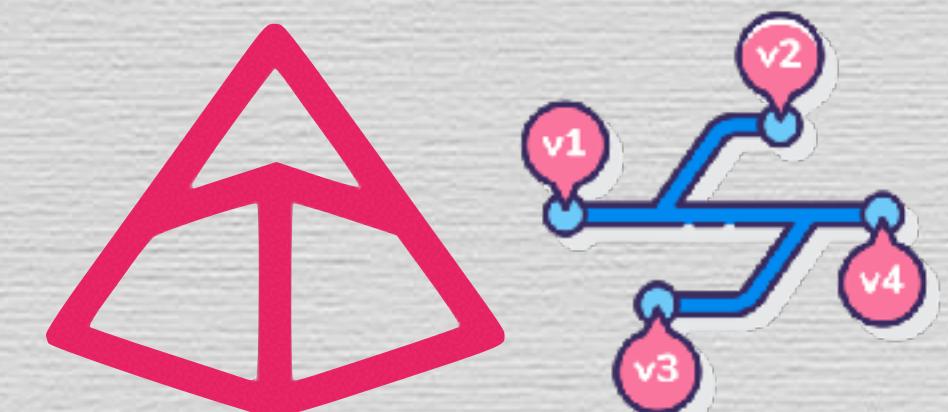
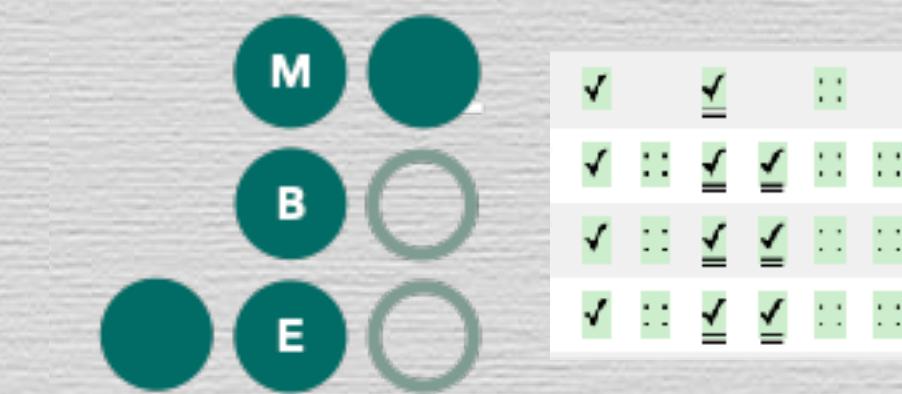
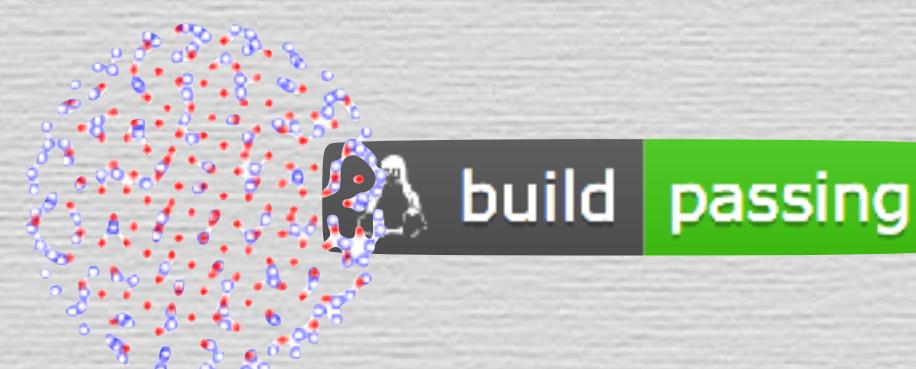


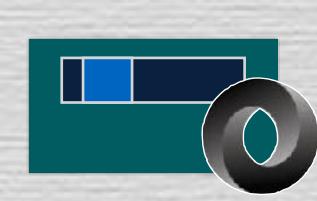
QC SCHEMA



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY

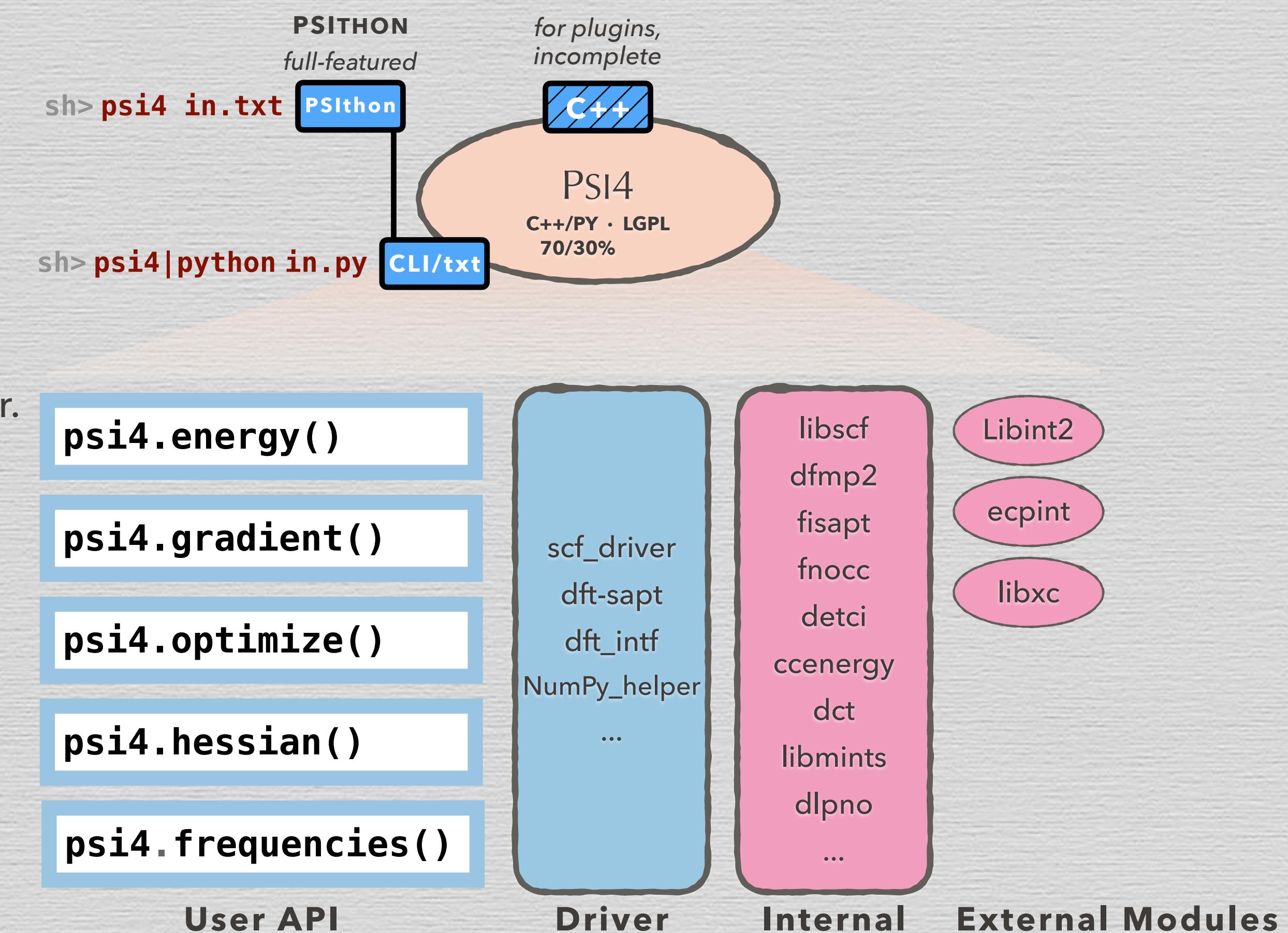


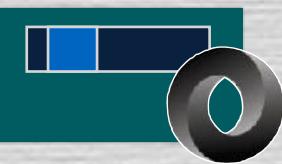


A QCSchema STORY

but first, an input mode overview

- **FULL-FEATURED** (lots of initialisms)
- **SMP** parallelism
- **HYBRID** C++/Python
- **INPUT** by DSL, API, or JSON
- **STRIVE** for easy production-quality user input with five driver “user-facing” functions through which 99+% of QC is run in Psi4, so easy to guess command. Minimal entry points.
- **PY/C++** interface layer below the user layer makes it simple to shift methods btwn languages without disturbing user.
 - **PY → C++** when a reference implementation is optimized in compiled language.
 - **C++ → PY** when a legacy code is refactored so that logic in Python and intensive parts in compiled.
- **SPECIALTIES** include density-fitting & intermolecular interactions





A QCSchema Story

PSI API

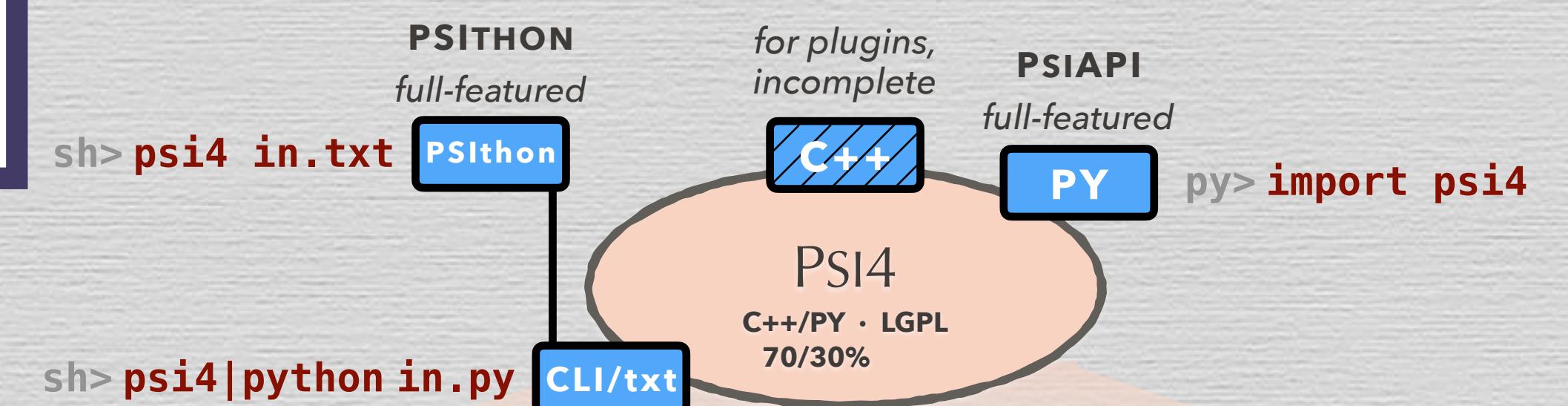
but first, an input mode overview

PSITHON

```
molecule {
Ne
Ne 1 3.0
}

set freeze_core True

energy('ccsd(t)/cc-pvtz')
```



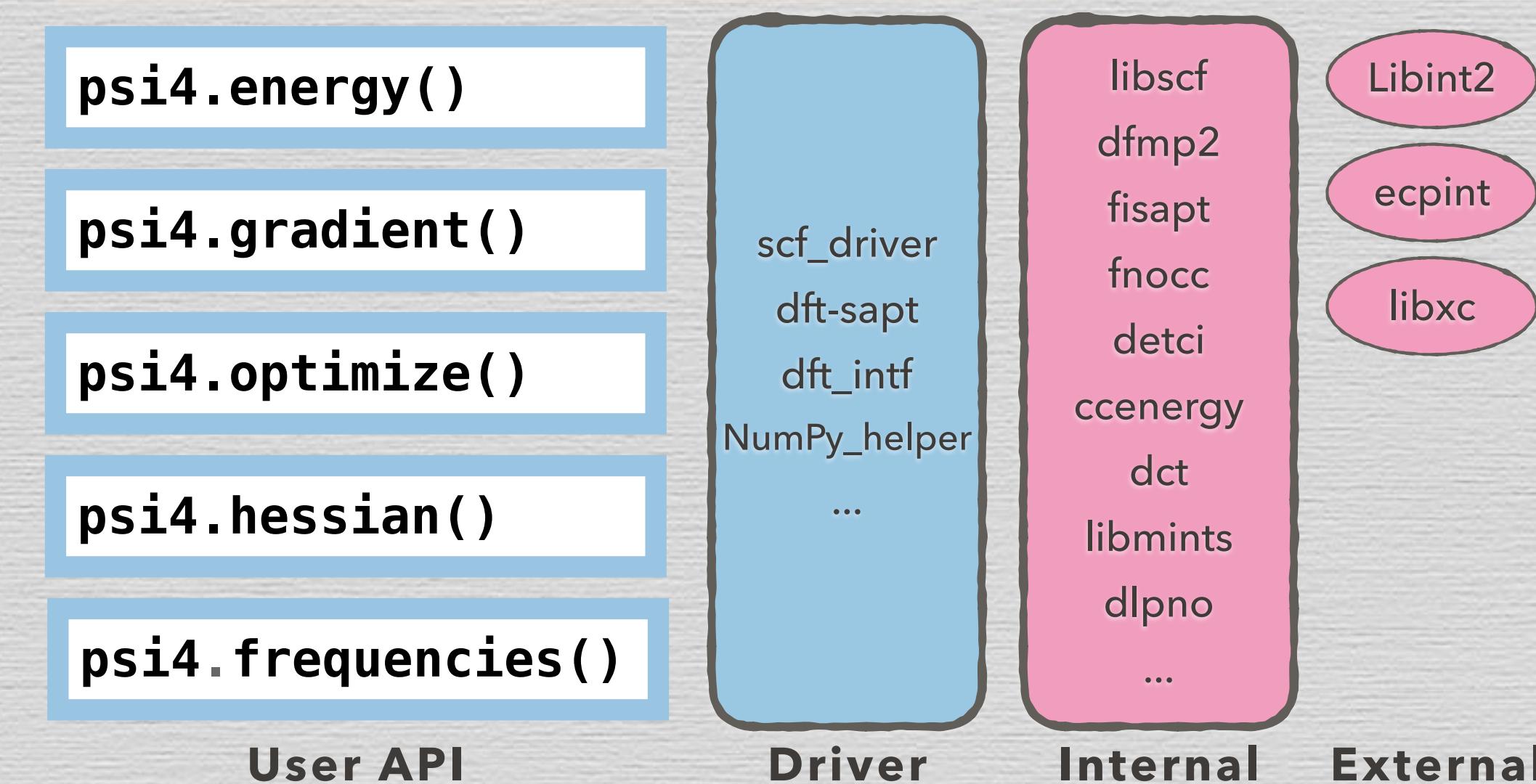
```
import psi4

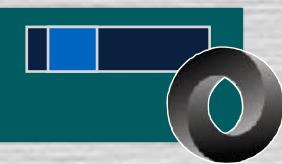
psi4.geometry'''
Ne
Ne 1 3.0
''')

psi4.set_options({
    'freeze_core': 'True'})

psi4.energy('ccsd(t)/cc-pvtz')
```

added in 2016





A QCSchema STORY

github.com/MolSSI/QCSchema

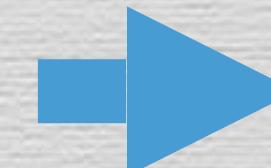
- **COMMUNICATION** channel between all pieces of the QCArchive ecosystem.
- **COMMUNITY** project useful for many aspects of quantum chem.
- **JSON** nominally, but any key/value/array language like MessagePack/BSON/XML/YAML ok.
- **TIMELINE** stable since 2019. Most changes for QCA. More community participation desired but little momentum. v2 soon.
- **SCHEMA** defined in Python with Pydantic in QCElemental module, then exported to JSON.

Molecule

```
{  
    "geometry": [0, 0, 0, 0, 0, 1],  
    "symbols": ["He", "He"],  
    ...  
}
```

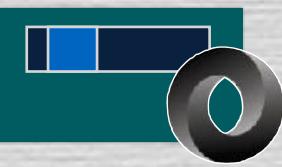
```
0 0 0 0  
H 2 0 0  
--  
@22Ne 5 0 0  
units bohr
```

Snippet 1:



```
{"atom_labels": ["", "", ""],  
 "atomic_numbers": [ 8,  1, 10],  
 "fix_com": False,  
 "fix_orientation": False,  
 "fragment_charges": [0.0, 0.0],  
 "fragment_multiplicities": [2, 1],  
 "fragments": [[0, 1], [2]],  
 "geometry": [[[0., 0., 0.],  
              [2., 0., 0.],  
              [5., 0., 0.]],  
             "mass_numbers": [16, 1, 22],  
             "masses": [15.99491462, 1.00782503, 21.99138511],  
             "molecular_charge": 0.0,  
             "molecular_multiplicity": 2,  
             "name": "HNeO",  
             "provenance": {"creator": "QCElemental",  
                           "routine": "qcelemental.molparse.from_schema",  
                           "version": "v0.8.0"},  
             "real": [ True,  True, False],  
             "schema_name": "qcschema_molecule",  
             "schema_version": 2,  
             "symbols": ["O", "H", "Ne"],  
             "validated": True}]
```

Snippet 2: QCSchema Molecule from Snippet 1.



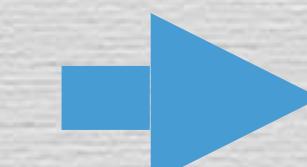
A QCSchema STORY

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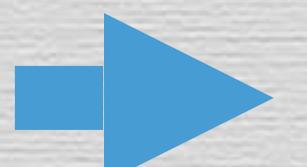
Molecule

```
{  
    "geometry": [0, 0, 0, 0, 0, 1],  
    "symbols": ["He", "He"],  
    ...  
}
```

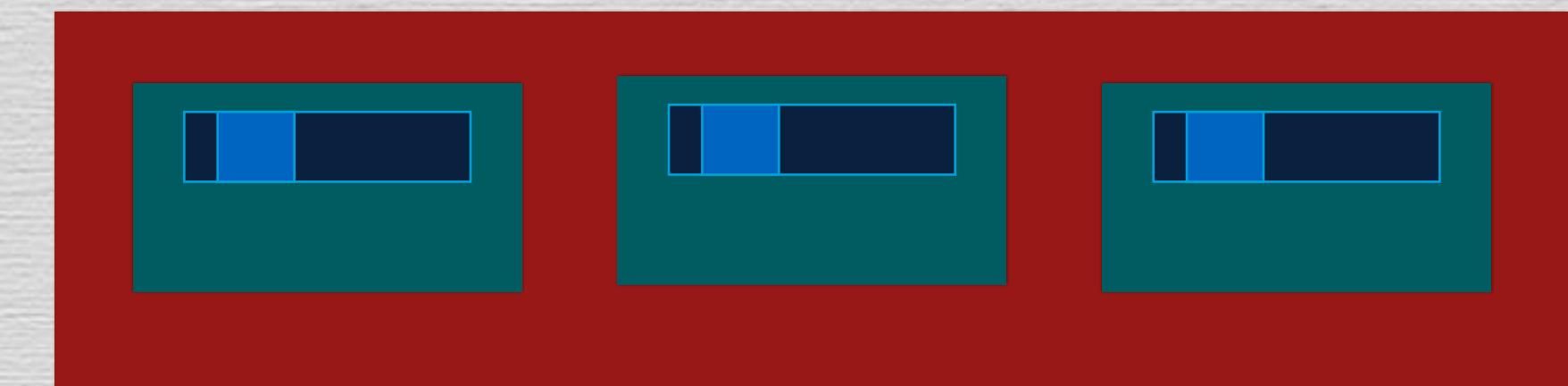


AtomicInput

```
{  
    "molecule": {  
        "geometry": [0, 0, 0, 0, 0, 1],  
        "symbols": ["He", "He"]  
    },  
    "driver": "energy",  
    "model": {  
        "method": "CCSD(T)",  
        "basis": "aug-cc-pVDZ",  
    },  
    "keywords": {}  
}
```

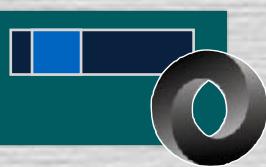


OptimizationResult



AtomicResult

```
{  
    ... AtomicInput ...  
    "provenance": {  
        "creator": "My QM Program",  
        "version": "1.1rc1",  
    },  
    "properties": {  
        "calcinfo_nalpha": 5,  
        "scf_total_energy": -5.433191881443323,  
        "nuclear_repulsion_energy": 2.11670883436,  
        "scf_iterations": 8.0,  
    },  
    "error": null,  
    "return_result": -6.5432123456,  
    "success": true,  
    "stdout": "...",  
    "wavefunction": "..."  
}
```



QCSchema ATOMIC

analytical single-point computations

- **AtomicInput** designed as a job directive to express any **SINGLE-GEOMETRY** analytic derivative or property calc. (as defined by driver) among CMS workhorses
- in practice, it can hide more complicated computations (e.g., finite difference frequencies), and advanced Result schema often seemingly inherit from **AtomicResult**
- note that QCSchema controls **LAYOUT** and "manages" values for **Molecule** and driver with model and enum respectively. All other values are free-form strings in CMS native syntax. See "Data Layouts" slide later.
- **AtomicResult APPENDS** to **AtomicInput** with output, simple scalar and array results, and wavefunction results.
- provenance tracks who wrote the data – program version, module, compute information
- note **FLAT** structure where Result inherits from Input
- overwhelmingly the most used of the calculation schema. These are stored by the **MILLIONS** at MolSSI
- Called "Atomic" for "fundamental"

AtomicResultProtocols

stdout: save text output
wavefunction: save orbital info
native_files: save raw prog. files
error_correction: auto-edit&rerun

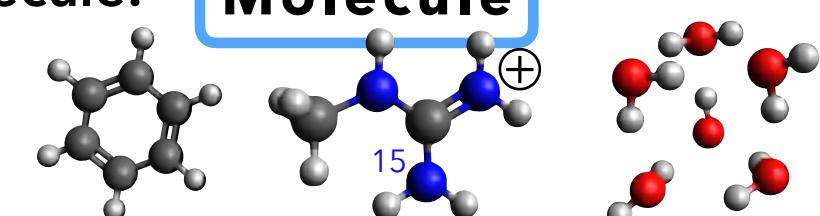
AtomicInput

schema_version: 1
provenance: creator, version, & RT info
QCElemental, 0.28.0

id: tracker for databases

extras: free-form storage

molecule: **Molecule**



protocols: customize return layout

AtomicResultProtocols

driver: single-point derivative

- energy
- gradient
- Hessian
- properties

E **G** **H**

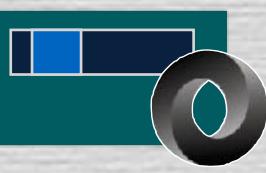
model: model chemistry or FF

method: B3LYP-D3, CCSD(T)

basis: 6-31G*, cc-pVQZ

keywords: knobs in SP program

{scftyp: uhf, cc_conv: 7, docc: [4,1]}



QCSchema ATOMIC

analytical single-point computations

```
'properties': {'calcinfo_nalpha': 10,
   'calcinfo_natom': 2,
   'calcinfo_nbasis': 28,
   'calcinfo_nbeta': 10,
   'calcinfo_nmo': 28,
   'nuclear_repulsion_energy': 16.666666666666668,
   'return_energy': -256.977621071098,
   'return_gradient': array([[0, 0, -6.54470201e-05],
                           [0, 0, 6.54470212e-05]]),
   'return_hessian': None,
   'scf_dipole_moment': array([0, 0, 4.97379915e-14]),
   'scf_iterations': 5,
   'scf_one_electron_energy': -398.5650589792168,
   'scf_total_energy': -256.977621071098,
   'scf_total_gradient': array([[0, 0, -6.54470201e-05],
                               [0, 0, 6.54470212e-05]]),
   'scf_total_hessian': None,
   'scf_two_electron_energy': 124.92077124145213},

'extras': {'qcvars': {'CURRENT DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
   'CURRENT ENERGY': -256.977621071098,
   'CURRENT GRADIENT': array([[0, 0, -6.54470201e-05],
                             [0, 0, 6.54470212e-05]]),
   'CURRENT REFERENCE ENERGY': -256.977621071098,
   'DD SOLVATION ENERGY': 0.0,
   'HF KINETIC ENERGY': 256.97713564856093,
   'HF POTENTIAL ENERGY': -513.9547567196589,
   'HF TOTAL ENERGY': -256.977621071098,
   'HF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
                             [0, 0, 6.54470212e-05]]),
   'HF VIRIAL RATIO': 2.000001888971701,
   'NUCLEAR REPULSION ENERGY': 16.666666666666668,
   'ONE-ELECTRON ENERGY': -398.5650589792168,
   'PCM POLARIZATION ENERGY': 0.0,
   'PE ENERGY': 0.0,
   'SCF DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
   'SCF ITERATION ENERGY': -256.977621071098,
   'SCF ITERATIONS': 5.0,
   'SCF TOTAL ENERGIES': array([-256.97759612, -256.97762005, -256.9776
                                 -256.97762107, -256.97762107, -256.9776
                                 -256.977621098,
   'SCF TOTAL ENERGY': -256.977621071098,
   'SCF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
                               [0, 0, 6.54470212e-05]]),
   'TWO-ELECTRON ENERGY': 124.92077124145213}}},
```

AtomicResultProperties (abr.)

calcinfo_natom

calcinfo_nmo

nuclear_repulsion_energy

return_energy:

return_gradient:

scf_total_gh: ref. energy, gradient, etc.

mtd_correlation_energy: post-scf energy

mtd_iterations: count to convergence

mtd_mpole_moment: dipole etc. array

AtomicResult

AtomicInput

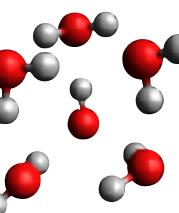
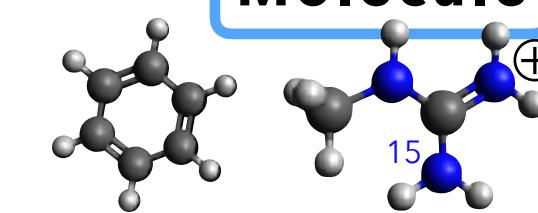
schema_version: 1

provenance: creator, version, & RT info
QCElemental, 0.28.0

id: tracker for databases

extras: free-form storage

molecule:



protocols: customize return layout

AtomicResultProtocols

driver: single-point derivative

- energy
- gradient
- Hessian
- properties



model: model chemistry or FF

method: B3LYP-D3, CCSD(T)

basis: 6-31G*, cc-pVQZ

keywords: knobs in SP program

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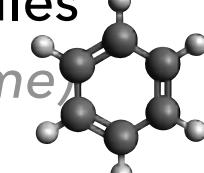
success: True

provenance: (overwrites input)

xtb, 1.0.0, psinet, CPU E5-2630

stdout/stderr: program's text output for ppl

native_files: requested raw program files



molecule: (overwrites inp w/result frame)

return_result: single-point derivative

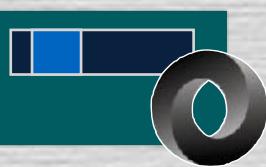


properties: key results for single-point

AtomicResultProperties

wavefunction: orbital-level results

WavefunctionProperties



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- **AtomicResult APPENDS** to **AtomicInput** with output, simple scalar and array results, and wavefunction results.
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- overwhelmingly the most used of the calculation schema. These are stored by the **MILLIONS** at MolSSI
- Called "Atomic" for "fundamental"

AtomicResult

AtomicInput

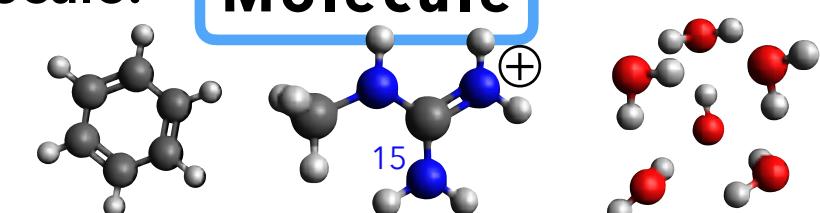
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provenance: creator, version, & RT info
QCElemental, 0.28.0

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AtomicResultProtocols

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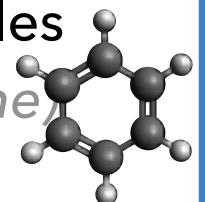
success: True

provenance: (overwrites input)

xtb, 1.0.0, psinet, CPU E5-2630

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molecule: (overwrites inp w/result frame)

return_result: single-point derivative



properties: key results for single-point

AtomicResultProperties

wavefunction: orbital-level results

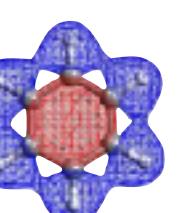
WavefunctionProperties

WavefunctionProperties (abr.)

basis: **BasisSet**

restricted: alpha == beta

orbitals_a/b

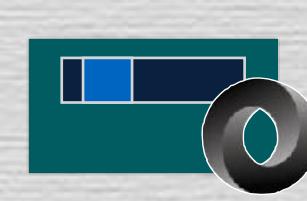


density_a/b

fock_a/b

eigenvalues_a/b

occupations_a/b



QCSCHEMA DEFINES DATA LAYOUTS

and text advice, nothing more; QCEngine will not standardize input

DOMAIN-SPECIFIC ASCII

```
N 0.000 0.000 -0.146 <CC|CC>
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
```

```
*CFOUR(REFERENCE=ROHF
BASIS=AUG-PVDZ,CALC_LEVEL=CCSD
COORDINATES=CARTESIAN,UNITS=BOHR
CHARGE=0,MULTIPLICITY=2)
```

all-electron restricted-open-shell CCSD/aug-cc-pVDZ energy of NH₂ molecule

```
$ccinp ncore=0 $end
$basis gbasis=accd $end
$contrl cctyp=ccsd coord=prinaxis
icharg=0 ispher=1 mult=2
runtyp=energy scftyp=rohf
units=bohr $end
$data

C1
N 7 0.000 0.000 -0.146
H 1 0.000 -1.511 1.014
H 1 0.000 1.511 1.014
$end
```



```
molecule {
0 2
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
units au
}

set reference rohf

energy('ccsd/aug-cc-pvdz')
```

```
geometry units bohr
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
end
charge 0
basis spherical
h library aug-cc-pvdz
n library aug-cc-pvdz
end
scf
rohf
nopen 1
end
tce
ccsd
end
task tce energy
```

DATA LAYOUT TRANSLATION



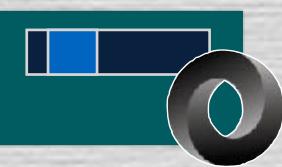
QCSchema: AtomicInput

```
{
'molecule': ,
'driver': ,
'model': {
'method': ,
'basis': ,
'keywords': {
}
},
```

```
{
'molecule': ,
'driver': ,
'model': {
'method': ,
'basis': ,
'keywords': {
}
},
```

```
{
'molecule': ,
'driver': ,
'model': {
'method': ,
'basis': ,
'keywords': {
}
},
```

```
{
'molecule': ,
'driver': ,
'model': {
'method': ,
'basis': ,
'keywords': {
}
},
```



QCSCHEMA DEFINES DATA LAYOUTS

and text advice, nothing more; QCEngine will not standardize input

DOMAIN-SPECIFIC ASCII

$\langle CC|CC \rangle$

```
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
```

```
*CFOUR(REFERENCE=ROHF
BASIS=AUG-PVDZ,CALC_LEVEL=CCSD
COORDINATES=CARTESIAN,UNITS=BOHR
CHARGE=0,MULTIPLICITY=2)
```

all-electron restricted-open-shell CCSD/aug-cc-pVDZ energy of NH₂ molecule

```
$ccinp ncore=0 $end
$basis gbasis=accd $end
$contrl cctyp=ccsd coord=prinaxis
icharg=0 ispher=1 mult=2
runtyp=energy scftyp=rohf
units=bohr $end
$data
C1
N 7 0.000 0.000 -0.146
H 1 0.000 -1.511 1.014
H 1 0.000 1.511 1.014
$end
```



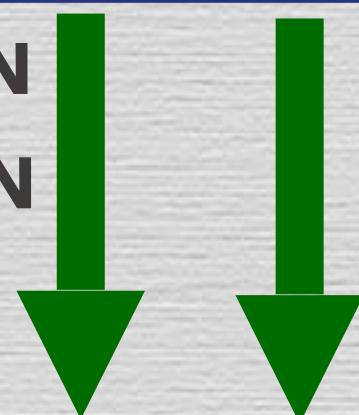
```
molecule {
0 2
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
units au
}
set reference rohf
energy('ccsd/aug-cc-pvdz')
```



```
geometry units bohr
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
end
charge 0
basis spherical
h library aug-cc-pvdz
n library aug-cc-pvdz
end
scf
rohf
nopen 1
end
tce
ccsd
end
task tce energy
```



DATA LAYOUT TRANSLATION
MOL & DRIVER STANDARDIZATION



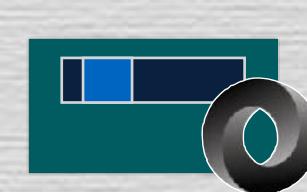
QCSchema: AtomicInput

```
{
'molecule': ,
'driver': 'energy',
'model': {
'method': ,
'basis': },
'keywords': {
},
'}
```

```
{
'molecule': ,
'driver': 'energy',
'model': {
'method': ,
'basis': },
'keywords': {
},
'}
```

```
{
'molecule': ,
'driver': 'energy',
'model': {
'method': ,
'basis': },
'keywords': {
},
'}
```

```
{
'molecule': ,
'driver': 'energy',
'model': {
'method': ,
'basis': },
'keywords': {
},
'}
```



QCSCHEMA DEFINES DATA LAYOUTS

and text advice, nothing more; QCEngine will not standardize input

DOMAIN-SPECIFIC ASCII

```
N 0.000 0.000 -0.146 <CC|CC>
H 0.000 -1.511 1.014
H 0.000 1.511 1.014

*CFOUR(REFERENCE=ROHF
BASIS=AUG-PVDZ,CALC_LEVEL=CCSD
COORDINATES=CARTESIAN,UNITS=BOHR
CHARGE=0,MULTIPLICITY=2)
```

all-electron restricted-open-shell CCSD/aug-cc-pVDZ energy of NH₂ molecule

```
$ccinp ncore=0 $end
$basis gbasis=accd $end
$contrl cctyp=ccsd coord=prinaxis
icharg=0 ispher=1 mult=2
runtyp=energy scftyp=rohf
units=bohr $end
$data

C1
N 7 0.000 0.000 -0.146
H 1 0.000 -1.511 1.014
H 1 0.000 1.511 1.014
$end
```



```
molecule {
0 2
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
units au
}
set reference rohf
energy('ccsd/aug-cc-pvdz')
```



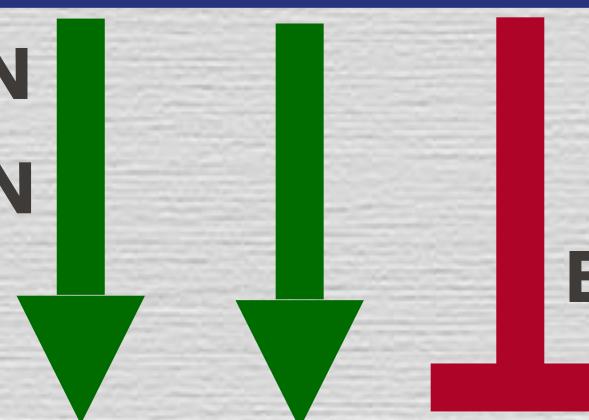
PSI4



```
geometry units bohr
N 0.000 0.000 -0.146
H 0.000 -1.511 1.014
H 0.000 1.511 1.014
end
charge 0
basis spherical
h library aug-cc-pvdz
n library aug-cc-pvdz
end
scf
rohf
nopen 1
end
tce
ccsd
end
task tce energy
```

DATA LAYOUT TRANSLATION

MOL & DRIVER STANDARDIZATION



BASIS & KEYWORDS STANDARDIZATION

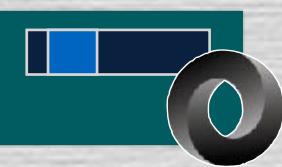
QCSchema: AtomicInput

```
{ 'molecule': NH2,
'driver': 'energy',
'model': {
  'method': 'ccsd',
  'basis': 'aug-pvdz' },
'keywords': {
  'reference': 'rohf'
}}
```

```
{ 'molecule': NH2,
'driver': 'energy',
'model': {
  'method': 'ccsd' ,
  'basis': 'accd' },
'keywords': {
  'contrl_ispher': 1,
  'contrl_scftyp': 'rohf',
  'ccinp_ncore': 0 }}
```

```
{ 'molecule': NH2,
'driver': 'energy',
'model': {
  'method': 'ccsd',
  'basis': 'aug-cc-pvdz' },
'keywords': {
  'reference': 'rohf' }}
```

```
{ 'molecule': NH2,
'driver': 'energy',
'model': {
  'method': 'ccsd',
  'basis': 'aug-cc-pvdz' },
'keywords': {
  'basis_spherical': True,
  'scf_rohf': True,
  'qc_module': 'tce' }}
```



A QCSchema Story

PSI API

PSITHON

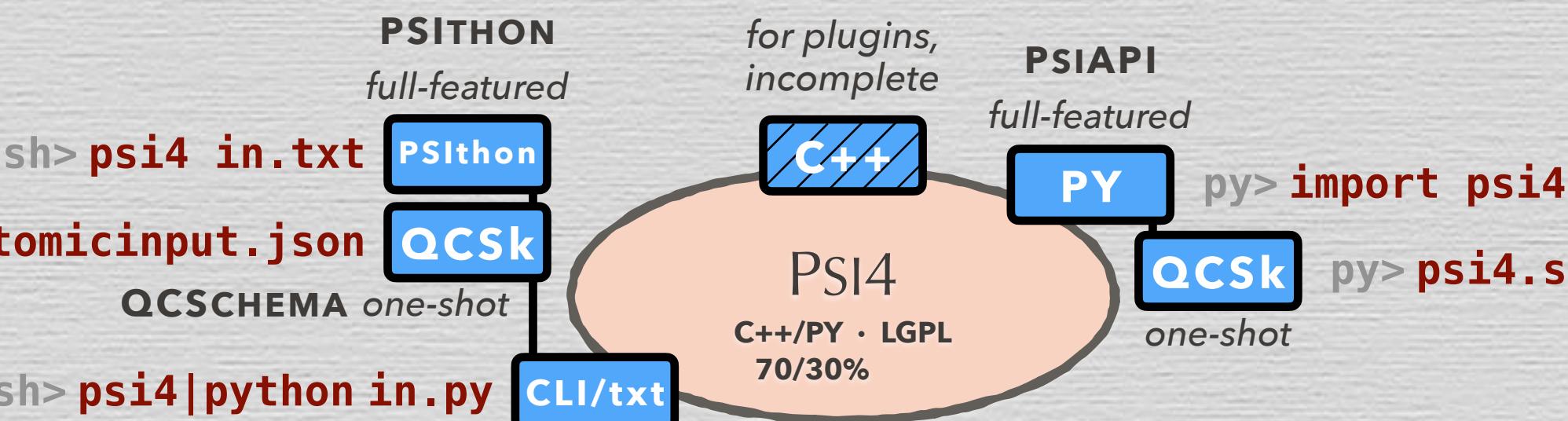
```
molecule {
Ne
Ne 1 3.0
}

set freeze_core True

energy('ccsd(t)/cc-pvtz')
```

sh> psi4 --qcschema atomicinput.json

```
{
'molecule': {
'symbols': ['Ne', 'Ne'],
'geometry':
[0, 0, 0, 5.67, 0, 0]
},
'driver': 'energy',
'model': {
'method': 'ccsd(t)',
'basis': 'cc-pvtz'
},
'keywords':
{'freeze_core': 'True'}
}
```



QCSchema

one-shot inputs now possible

import psi4

```
psi4.geometry(''  
Ne  
Ne 1 3.0  
...')
```

```
psi4.set_options({  
    'freeze_core': 'True'})
```

```
psi4.energy('ccsd(t)/cc-pvtz')
```

- **PAYOUT** PSI4 gets a structured, single-blob means of launching and recording the great majority of computations. And it's a community format, not our own.

- ★ Mapping calculation input into a structured format (harder for API-based than early QC programs) can open new ways of interacting with users & workflows.

QCSchema ATOMIC

analytical single-point computations

- As soon as move beyond Psi4 domain-specific output and start writing out QCSchema **AtomicResult** files, it's rather like entering into a contract that fields mean defined things.

```
'properties': {'calcinfo_nalpha': 10,
    'calcinfo_natom': 2,
    'calcinfo_nbasis': 28,
    'calcinfo_nbeta': 10,
    'calcinfo_nmo': 28,
    'nuclear_repulsion_energy': 16.66666666666668,
    'return_energy': -256.977621071098,
    'return_gradient': array([[0, 0, -6.54470201e-05],
        [0, 0, 6.54470212e-05]]),
    'return_hessian': None,
    'scf_dipole_moment': array([0, 0, 4.97379915e-14]),
    'scf_iterations': 5,
    'scf_one_electron_energy': -398.5650589792168,
    'scf_total_energy': -256.977621071098,
    'scf_total_gradient': array([[0, 0, -6.54470201e-05],
        [0, 0, 6.54470212e-05]]),
    'scf_total_hessian': None,
    'scf_two_electron_energy': 124.92077124145213},

'extras': {'qcvars': {'CURRENT DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
    'CURRENT ENERGY': -256.977621071098,
    'CURRENT GRADIENT': array([[0, 0, -6.54470201e-05],
        [0, 0, 6.54470212e-05]]),
    'CURRENT REFERENCE ENERGY': -256.977621071098,
    'DD SOLVATION ENERGY': 0.0,
    'HF KINETIC ENERGY': 256.97713564856093,
    'HF POTENTIAL ENERGY': -513.9547567196589,
    'HF TOTAL ENERGY': -256.977621071098,
    'HF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
        [0, 0, 6.54470212e-05]]),
    'HF VIRIAL RATIO': 2.000001888971701,
    'NUCLEAR REPULSION ENERGY': 16.66666666666668,
    'ONE-ELECTRON ENERGY': -398.5650589792168,
    'PCM POLARIZATION ENERGY': 0.0,
    'PE ENERGY': 0.0,
    'SCF DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
    'SCF ITERATION ENERGY': -256.977621071098,
    'SCF ITERATIONS': 5.0,
    'SCF TOTAL ENERGIES': array([-256.97759612, -256.97762005, -256.9776209,
        -256.97762107, -256.97762107, -256.97762107]),
    'SCF TOTAL ENERGY': -256.977621071098,
    'SCF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
        [0, 0, 6.54470212e-05]]),
    'TWO-ELECTRON ENERGY': 124.92077124145213}},
```

AtomicResultProperties (abr.)

calcinfo_natom

calcinfo_nmo

nuclear_repulsion_energy

return_energy:

return_gradient:

scf_total_gh: ref. energy, gradient, etc.

mtd_correlation_energy: post-scf energy

mtd_iterations: count to convergence

mtd_mpole_moment: dipole etc. array

AtomicResult

AtomicInput

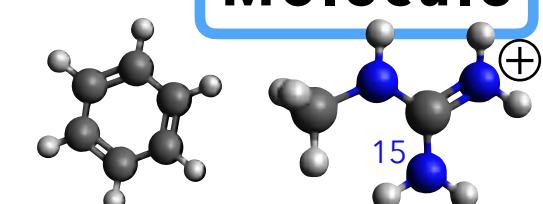
schema_version: 1

provenance: creator, version, & RT info
QCElemental, 0.28.0

id: tracker for databases

extras: free-form storage

molecule:



protocols: customize return layout

AtomicResultProtocols

driver: single-point derivative

- energy
- gradient
- Hessian
- properties



model: model chemistry or FF

method: B3LYP-D3, CCSD(T)

basis: 6-31G*, cc-pVQZ

keywords: knobs in SP program

{scftyp: uhf, cc_conv: 7, docc: [4,1]}

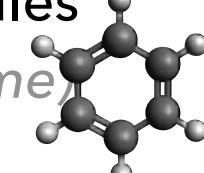
success: True

provenance: (overwrites input)

xtb, 1.0.0, psinet, CPU E5-2630

stdout/stderr: program's text output for ppl

native_files: requested raw program files



molecule: (overwrites inp w/result frame)

return_result: single-point derivative

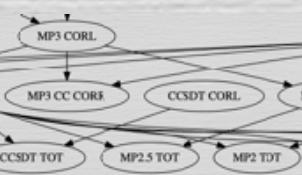


properties: key results for single-point

AtomicResultProperties

wavefunction: orbital-level results

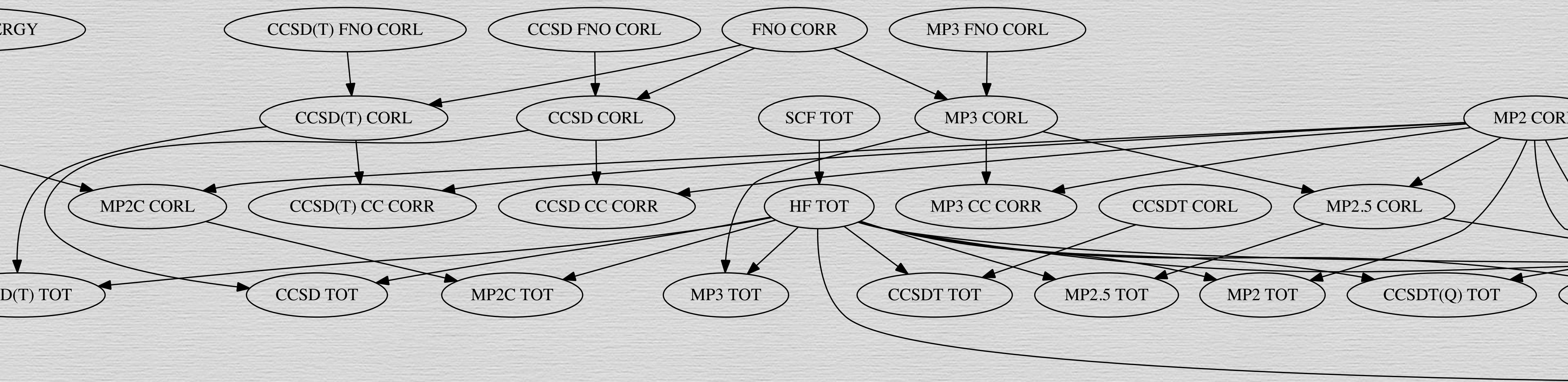
WavefunctionProperties

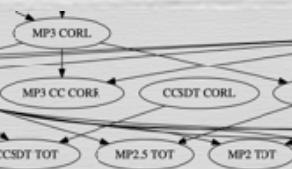


QCVARIABLES

common QC definitions & combining rules

- As soon as move beyond PSI4 domain-specific output and start writing out QCSchema **AtomicResult** files, it's rather like entering into a contract that fields mean defined things.

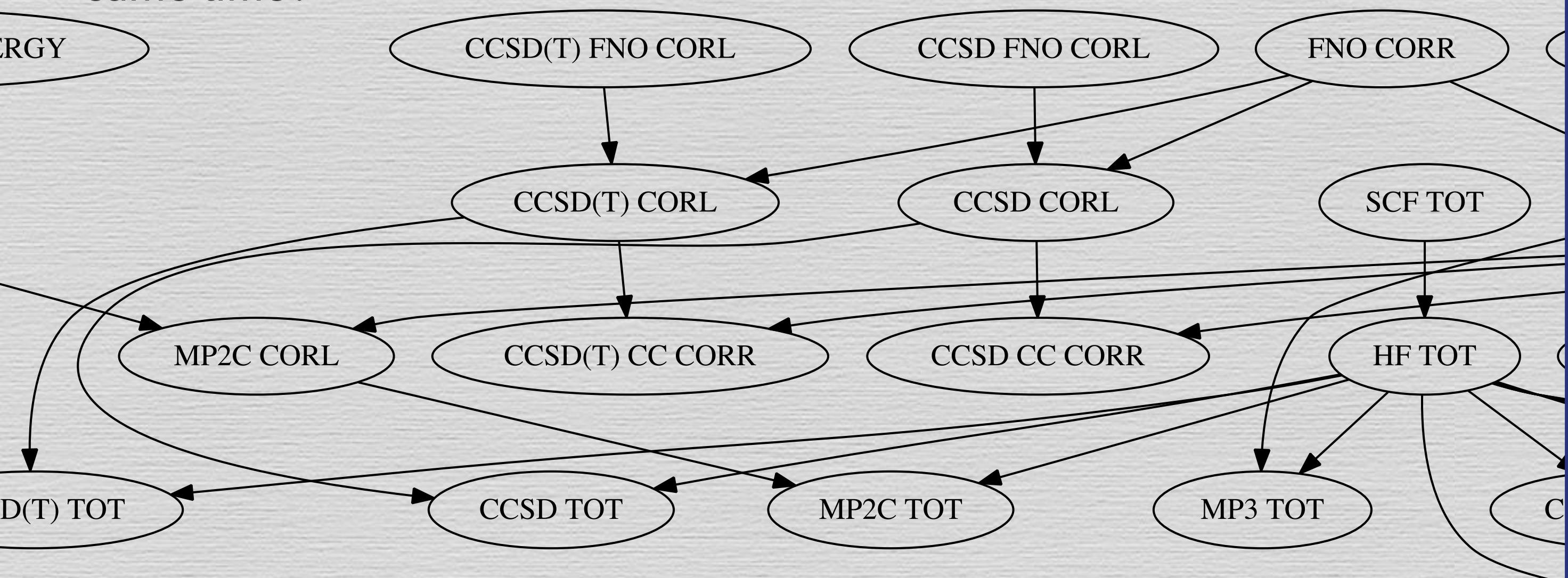




QC VARIABLES

common QC definitions & combining rules

- As soon as move beyond Psi4 domain-specific output and start writing out QCSchema **AtomicResult** files, it's rather like entering into a contract that fields mean defined things.
- When dealing with modular software, there's going to be certain modules with different capabilities and strengths, even for the same method. This is true for internal or external modules.
- Instituted "managed methods" so users could remain oblivious:
gradient('mp2') gets the most efficient module -or- intentional:
set qc_module dfocc; gradient('mp2') gets the directed module.
- How can we fulfill this contract and help manage complexity at the same time?



```

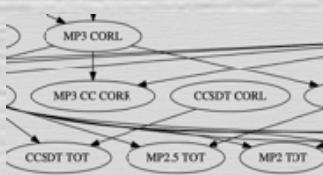
def select_mp2_gradient(name, **kwargs):
    """Function selecting the algorithm for a MP2 gradient call
    and directing to specified or best-performance default modules.
    """
    reference = core.get_option('SCF', 'REFERENCE')
    type_var, _, mtd_type = method_algorithm_type(name)
    module = core.get_global_option('QC_MODULE')
    all_electron = (core.get_global_option('FREEZE_CORE') == "FALSE")

    func = None
    if reference == 'RHF':
        if mtd_type == 'CONV':
            if all_electron:
                if module in ['', 'OCC']:
                    func = run_occ_gradient
        elif mtd_type == 'DF':
            if module == 'OCC':
                func = run_dfocc_gradient
        elif module in ['', 'DFMP2']:
            func = run_dfmp2_gradient
    elif reference == 'UHF':
        if mtd_type == 'CONV':
            if all_electron:
                if module in ['', 'OCC']:
                    func = run_occ_gradient
        elif mtd_type == 'DF':
            if module in ['', 'OCC']:
                func = run_dfocc_gradient

    if func is None:
        raise ManagedMethodError()

    return func(name, **kwargs)

```



QC VARIABLES

common QC definitions & combining rules

- As soon as move beyond Psi4 domain-specific output and start writing out QCSchema [AtomicResult](#) files, it's rather like entering into a contract that fields mean defined things.

★ Prefer and define programmatic output for data standards

```
'extras': {'qcvars': {'CURRENT DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
'CURRENT ENERGY': -256.977621071098,
'CURRENT GRADIENT': array([[0, 0, -6.54470201e-05],
[0, 0, 6.54470212e-05]]),
'CURRENT REFERENCE ENERGY': -256.977621071098,
'DD SOLVATION ENERGY': 0.0,
'HF KINETIC ENERGY': 256.97713564856093,
'HF POTENTIAL ENERGY': -513.9547567196589,
'HF TOTAL ENERGY': -256.977621071098,
'HF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
[0, 0, 6.54470212e-05]]),
'HF VIRIAL RATIO': 2.000001888971701,
'NUCLEAR REPULSION ENERGY': 16.666666666666668,
'ONE-ELECTRON ENERGY': -398.5650589792168,
'PCM POLARIZATION ENERGY': 0.0,
'PE ENERGY': 0.0,
'SCF DIPOLE': array([0.0000000e+00, 0.0000000e+00, 4.97379915e-14]),
'SCF ITERATION ENERGY': -256.977621071098,
'SCF ITERATIONS': 5.0,
'SCF TOTAL ENERGIES': array([-256.97759612, -256.97762005, -256.9776209,
-256.97762107, -256.97762107, -256.97762107]),
'SCF TOTAL ENERGY': -256.977621071098,
'SCF TOTAL GRADIENT': array([[0, 0, -6.54470201e-05],
[0, 0, 6.54470212e-05]]),
'TWO-ELECTRON ENERGY': 124.92077124145213}},
```

QCVariables GLOSSARY

DFT FUNCTIONAL TOTAL ENERGY

The total electronic energy (E_h) for the underlying functional of the requested DFT method, without any dispersion correction; the first four terms in Eq. (4) or (1). Quantity E_{FCFT} in Eqs. (4) and (1). Unless the method includes a dispersion correction, this quantity is equal to [SCF TOTAL ENERGY](#).

DFT TOTAL ENERGY

The total electronic energy (E_h) for the requested DFT method, E_{DFT} in Eq. (1).

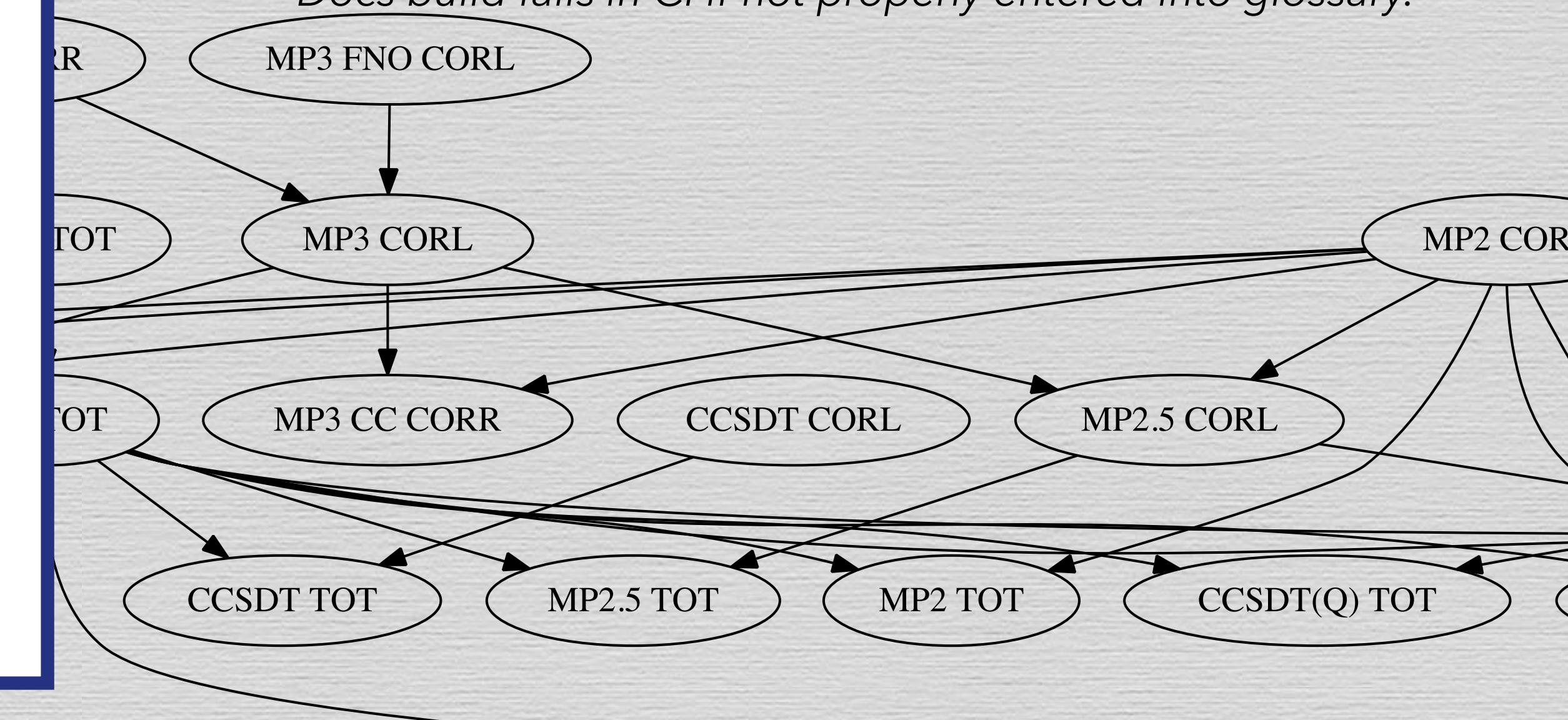
$$\begin{aligned} E_{\text{DFT}} &= E_{NN} + E_{1e^-} + E_{2e^-} + E_{xc} + E_{\text{D}} + E_{\text{DH}} \\ &= E_{\text{FCFT}} + E_{\text{D}} + E_{\text{DH}} \\ &= E_{\text{SCF}} + E_{\text{DH}} \end{aligned}$$

Unless the method is a DFT double-hybrid, this quantity is equal to [SCF TOTAL ENERGY](#). If the method is neither a double-hybrid, nor dispersion corrected, this quantity is equal to [DFT FUNCTIONAL TOTAL ENERGY](#).

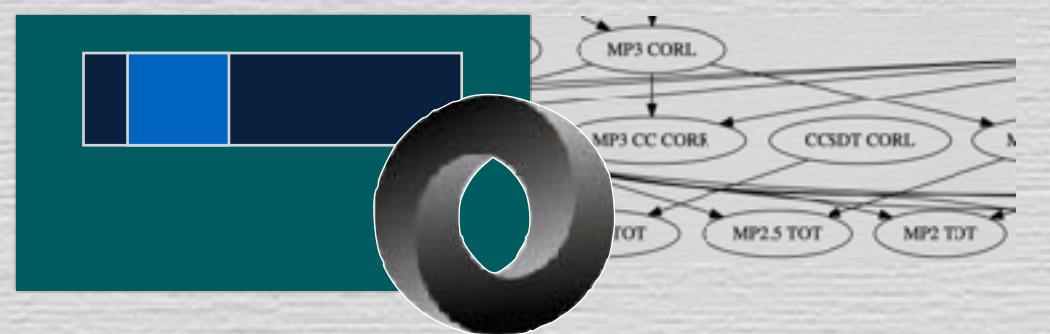
DFT TOTAL GRADIENT

The total electronic gradient [E_h/a_0] of the requested DFT method, $\langle \text{fnat}, 3 \rangle$.

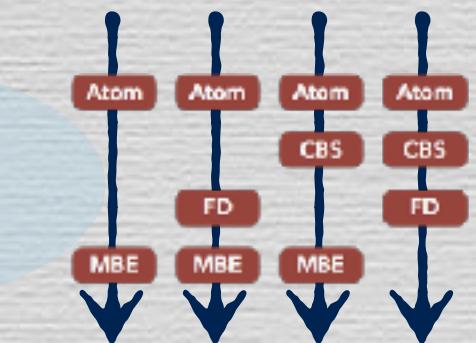
Modules register a var with # P::e MODULENAME . Docs build fails in CI if not properly entered into glossary.



METADATA

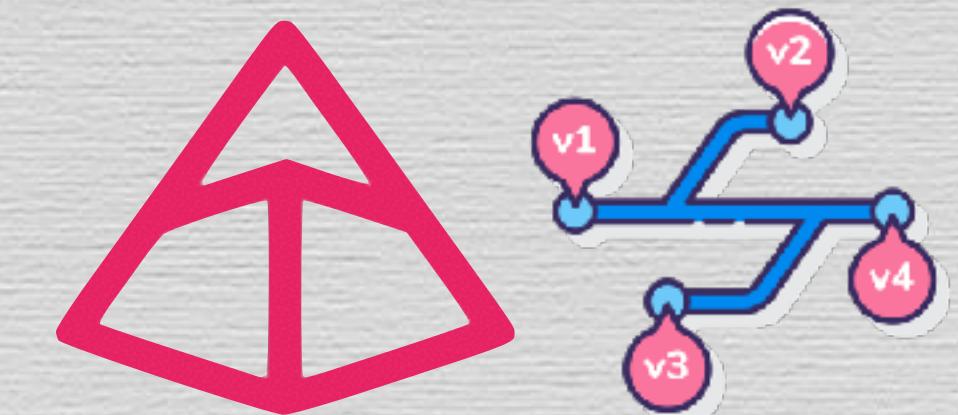
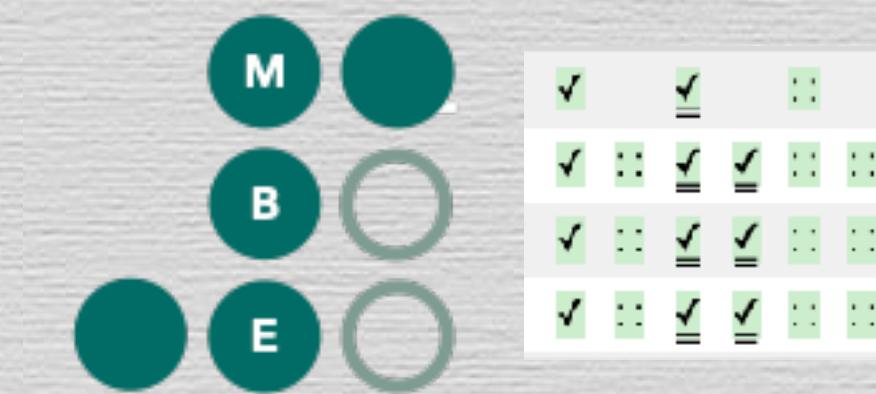
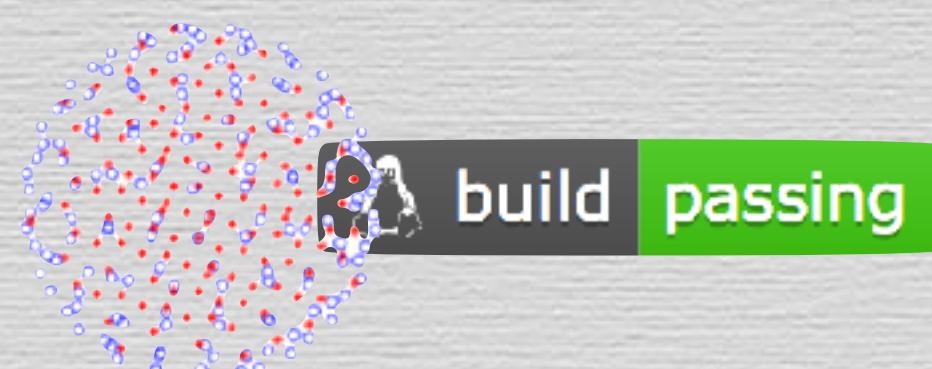


LIBINT



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY





Ed Valeev
VaTech



C

C++11

PY

Libint

C++ · LGPL

A LIBINT STORY

play nicely in the *runtime* software stack



Andy Simmonett
NIH → QC Ware



Maximilian Scheurer
Heidelberg → Covestro

In **2020** Psi4 started updating from 15yo Libint1 to maintained Libint2

step	status	libint ver & branch	Psi4	tarball ^[1]	order	component style
1 ^[3]	longstanding L1	L1 evaleev:5c89451	v1.3	—	gss	5
2 ^[4]	TEI L2	loriab:l2cmake evaleev/libint#148	20Nov20, after #1721, v1.4, 1.5	L: 7-7-4-7-7-5_1 , MW: 5-4-3-6-5-4_1	gss	95
3 ^[5]	OEI L2	ditto step 2	11Mar22, after #2388	L: 5-4-3-6-5-4_mm25f12ob2 , MW: 5-4-3-6-5-4_mm4ob2	gss	95
B ^[6]	upstream L2 cmake	loriab:new-cmake-harness-lab-rb1 evaleev/libint#233	23Mar22, after #2413, v1.6, v1.7	5-4-3-6-5-4_mm4f12ob2.tgz	gss	eri_c4_d1_l5
C ^[2]	McMurchie Davidson	any	31Mar22, after #2414, v1.6, v1.7			
A ^[7]	standardize ordering	ditto step B	#2537	ditto step B	sss	95
A' ^[8]	flex solid-harm ordering	loriab:new-cmake-2023-take2-b evaleev/libint#259	8May23, after #2861, v1.8, v1.9 (see also A'')	libint-2.7.2-post1-5-4-3-5-5-4_mm4f12ob2_1	sss	eri_c4_d1_l5
A'' ^[9]	flex solid-harm ordering, libtool-based	upstream master any point after evaleev/libint@ 10ca429	3Dec23, after #3047, v1.9 (see also A'')	libint-2.8.0-dev2-5-4-3-6-5-4_mm10f12ob2_0.tgz	ss	eri_hhhh_d1
A''' ^[10]	flex solid-harm ordering, libtool-based, released	any upstream release >=2.8.1	18Jan24, after #3107, v1.9.1		ss	eri_hhhh_d1

- **DEVELOPED** by E. Valeev for 20y
- **FULL** service integrals library
- **INTERFACES** in multiple languages/APIs
- **HIGHLY** configurable by integral classes, derivatives, orderings, & angular momentum levels
- **LARGE** size at production configurations, >GB generated source or built library

A LIBINT STORY

play nicely in the *runtime* software stack

both projects subject to own expectations:

Libint assumed each instance targeted toward a **SINGLE** QC consumer

Psi4 vulnerable to **SEGFAULT** if meagerly configured

Psi4 vulnerable to **WRONG RESULTS** if configured w/std orderings

Portable libint #190

Open susilehtola opened this issue on Sep 3, 2020 · 26 comments · May be fixed by #205



susilehtola commented on Sep 3, 2020

Contributor

Hi,

it looks like libint2 can be compiled in `Nsph x Ncart x Nshell_set = 2 x 5 x 2 = 20` different possible ways. Even though only 5 of these are currently used by major codes, this is a nightmare for distribution packaging since there's no way we can ship 20 mutually incompatible copies of a library. I'm horrified about shipping even two since it takes forever to compile libint on the Fedora build system.

Would it be possible to modify libint so that the orderings for the cartesians, solid harmonics, and shell sets are set by some function, and libint does all the necessary shuffles behind the scenes? This would be a minimal solution to the problem, since one could still set the default orderings at configure time, thus avoiding any extra shuffles if the compiled ordering matches the wanted one.



loria

Collaborator

psi is non-standard for solid harmonic and probably precipitated this thread. (below is harvested from libint wiki notes.) small mercy is that psi differs in the one dimension that can be chosen at library build time, not generator build time.

#	sh	cart	shell_set	used_by
#				
#	sss - search for standard	+ standard	+ standard	= mpqc4, cp2k
#	sso - search for			+ orca
#	sis - search for	+ intv3	+ standard	
#	sio - search for		+ orca	
#	sgs - search for	+ gamess	+ standard	= gamess
#	sgo - search for		+ orca	
#	sos - search for	+ orca	+ standard	
#	soo - search for		+ orca	= orca
#	sbs - search for	+ bagel	+ standard	= bagel
#	sbo - search for		+ orca	
#	gss - search for gaussian	+ standard	+ standard	= psi4
#	gso - search for		+ orca	
#	gis - search for	+ intv3	+ standard	
#	gio - search for		+ orca	
#	ggs - search for	+ gamess	+ standard	
#	ggo - search for		+ orca	
#	gos - search for	+ orca	+ standard	
#	goo - search for		+ orca	
#	gbs - search for	+ bagel	+ standard	
#	gbo - search for		+ orca	

A LIBINT STORY

play nicely in the *runtime* software stack

... so Psi4 has contributed back upstream

- expanded to Windows **ARCH** & CMake **BUILDSYS**
- added **LINK-TIME** & **RUN-TIME** configuration checking
- maintain a **HIGH-AM** (12), high-deriv build on conda
- enabled **RUN-TIME** ordering selection for OSS consumers

cart	shell_set	used_by
ss - library integrals use ordering	standard + standard	= mpqc4, cp2k, psi4
so - library integrals use ordering	+ orca	
is - library integrals use ordering	intv3 + standard	= mpqc3
io - library integrals use ordering	+ orca	
gs - library integrals use ordering	gamess + standard	= gamess
go - library integrals use ordering	+ orca	
os - library integrals use ordering	orca + standard	
oo - library integrals use ordering	+ orca	= orca
bs - library integrals use ordering	bagel + standard	= bagel
bo - library integrals use ordering	+ orca	

(psi4 requires runtime-setting of solid harmonic ordering to Gaussian)

- **PAYOUT** delete much home-grown code
- **PAYOUT** expand capabilities with new integrals types
- **PAYOUT** return to upstream as of Dec 2023 v2.8, so new Psi4 releases packageable w/other OSS QC
- **PAYOUT** conda-forge packaging

incompatibilities frustrate **PACKAGING**, so was stuck c. 2019

The screenshot shows the Anaconda.org website interface. At the top, there's a logo for ANACONDA.ORG and a search bar. Below that, a green header bar displays the URL "conda-forge / packages / psi4". To the right of the URL, the version "1.9.1" is circled in red. Below the header, a sub-header reads "Open-Source Quantum Chemistry - an electronic structure package in C++ driven by Python". The main content area lists several Fedora versions along with their corresponding Psi4 versions and build numbers:

Fedora Version	Psi4 Version
Fedora 42	2025
Fedora 41	1.9.1-3.fc41
Fedora 40	2024
Fedora 40	Psi4 c.2019 1.3.2-22.fc40

A LIBINT STORY

play nicely in the *runtime* software stack

Open Libint2 2022 Strategy #2442

step	status	libint ver & branch	Psi4	tarball ^[1]	order	component style
1 ^[3]	longstanding L1	L1 evaleev:5c89451	v1.3	—	gss	s
2 ^[4]	TEI L2	loriab:l2cmake evaleev/libint#148	20Nov20, after #1721, v1.4, 1.5	L: 7-7-4-7-7-5_1, MW: 5-4-3-6-5-4_1	gss	g5
3 ^[5]	OEI L2	ditto step 2	11Mar22, after #2388	L: 5-4-3-6-5-4_mm25f12ob2, MW: 5-4-3-6-5-4_mm4ob2	gss	g5
B ^[6]	upstream L2 cmake	loriab:new-cmake-harness-lab-rb1 evaleev/libint#233	23Mar22, after #2413, v1.6, v1.7	5-4-3-6-5-4_mm4f12ob2.tgz	gss	eri_c4_d1_l5
C ^[2]	McMurchie Davidson	any	31Mar22, after #2414, v1.6, v1.7			
A ^[7]	standardize ordering	ditto step B	#2537	ditto step B	sss	g5
A' ^[8]	flex solid-harm ordering	loriab:new-cmake-2023-take2-b evaleev/libint#259	8May23, after #2861, v1.8, v1.9 (see also A'')	libint-2.7.2-p0st1-5-4-3-6-5-4_mm4f12ob2_1	sss	eri_c4_d1_l5
A'' ^[9]	flex solid-harm ordering, libtool-based	upstream master any point after evaleev/libint@ 10ce429	3Dec23, after #3047, v1.9 (see also A'')	libint-2.8.0-dev2-5-4-3-6-5-4_mm10f12ob2_0.tgz	ss	eri_hhhh_d1
A''' ^[10]	flex solid-harm ordering, libtool-based, released	any upstream release >=2.8.1	18Jan24, after #3107, v1.9.1		ss	eri_hhhh_d1

- Libint1 to Libint2 conversion took many stages, not at all surprising for a change of this magnitude even to a cleaner API with negative thousands LOC
- lag and needing mutual changes and then consolidating those changes into tags/releases is a general trend, though.

TIME ELAPSED BEFORE PSI4 MASTER USING TAGGED DEPENDENCY

Libint2	3y	API switch; RT config
Libxc	2.5y	API switch; "tweaks" flexibility
CPPE	9m	ints efficiency upgrade; Py API switch
simple-dftd3	6m	handling defaults change; 3b details
QCManyBody	11m	qcvars switch; schema upgrade

A LIBINT STORY

play nicely in the *runtime* software stack

- Libint1 to Libint2 conversion took many stages, not at all surprising for a change of this magnitude even to a cleaner API with negative thousands LOC
- lag and needing mutual changes and then consolidating those changes into tags/releases is a general trend, though.

Friday, June 20th ▾

jturney 1:31 PM
Do you have any experience with using CMake FetchContent_* with the PATCH_COMMAND? I need to patch the CMakeLists.txt file from cpptrace.
I also wouldn't mind pinging the author of cpptrace to include Mac ARM on his conda feedstock.

Today ▾

jturney 2:33 PM
I got around the PATCH_COMMAND by gettin macOS ARM conda builds on cpptrace and its dependency libdwarf
😊

loriab 2:34 PM
great, cleaner too!

needing tweaks of upstream is ubiquitous in an ecosystem

TIME ELAPSED BEFORE PSI4 MASTER USING TAGGED DEPENDENCY

Libint2	3y	API switch; RT config
Libxc	2.5y	API switch; "tweaks" flexibility
CPPE	9m	ints efficiency upgrade; Py API switch
simple-dftd3	6m	handling defaults change; 3b details
QCManyBody	11m	qcvars switch; schema upgrade

★ interfacing two codes is rarely unilateral. Seize any opportunity to work with other devs in real time.



PROGRAMMATIC METADATA

cater to tools as well as people

REQUIREMENTS FOR PSI4 CONSUMING: CMakeLists.txt

- We've solved the uncertainty over ordering configuration of published Libint packages for OSS
- But there are several other things that consumers may need or want to know from their consumed software. Modular software needs to expose details to all of people, runtime artifact, and build system.
- Some may be needed at consumer **COMPILE-TIME**:
 - require an installation to supply certain derivatives or classes of integrals or compiled extensions
 - be sure to define your options via CMake components
 - packaging works now because packagers & project devs communicate (e.g., I tell Susi to enable 1-body Hessians). Give computers the tools to check without oversight.

```
find_package(gdma 2.3.3 CONFIG COMPONENTS Python)
find_package(Libxc 7.0.0 CONFIG COMPONENTS C)
find_package(Libint2 2.8.1 CONFIG
COMPONENTS
ss
CXX_ho
impure_sh
"eri_${amchar}_d0" eri_ggg_d0 eri_gg_d0 onebody_g_d0
eri_dddd_d1 eri_fff_d1 eri_ff_d1 onebody_f_d1
onebody_f_d2)
```

```
find_package(simint 0.7 CONFIG
COMPONENTS am${MAX_AM_ERI} der0)
```

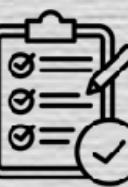
CONSUMED: simintConfig.cmake.in

```
...
set (simint_der0_FOUND 1) # always present
foreach(_d RANGE ${simint_MAXAM}) # simint_MAXAM
  set (simint_am${d}_FOUND 1)
endforeach()

check_required_components(simint)
...
```

PSI4 EXPORTED FEATURES

```
find_package(psi4)
message("psi4 components: ${psi4_FOUND_COMPONENTS}")
#> psi4 components: simint;einsums
```



PROGRAMMATIC METADATA

cater to tools as well as people

- We've solved the uncertainty over ordering configuration of published Libint packages for OSS
- But there are several other things that consumers may need or want to know from their consumed software. Modular software needs to expose details to all of people, runtime artifact, and build system.
- Some may be needed at consumer **COMPILE-TIME**:
 - require an installation to supply certain derivatives or classes of integrals or compiled extensions
 - be sure to define your options via CMake components
 - packaging works now because packagers & project devs communicate (e.g., I tell Susi to enable 1-body Hessians). Give computers the tools to check without oversight.
- Additionally, they may be needed at consumer **RUN-TIME**:
 - whether Hessian integrals are available or else redirect to finite difference or whether a rung of functionals is available or else disable those methods
 - be sure to provide accessors so consumers can apply logic.

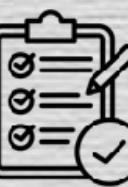
LIBINT2 CAPABILITIES

```
libint2::initialize();
printf("SHGShell: %d\n", libint2::solid_harmonics_ordering());
// if patched
printf("Configuration: %s\n",
       libint2::configuration_accessor().c_str());
printf("Supports: dddd=%d mmmm=%d\n",
       libint2::supports("eri_dddd_d0"),
       libint2::supports("eri_mmmm_d0"));
libint2::finalize();

#> SHGShell: 1
#> Configuration: eri_dddd_d0;eri_ffff_d0;ss;...
#> Supports: dddd=1 mmmm=0
```

PSI4 EXTRA CAPABILITIES

```
print(psi4.addons())
#> ['adcc', 'ambit', 'bse', 'eet3', 'efour', 'chemps2',
    'eppe', 'ddx', 'dftd3', 'dftd4', 'dkh', 'eepint',
    'einsums', 'forte', 'gauxc', 'gep', 'gdma', 'geometric',
    'gpu_dfee', 'integratorxx', 'ipi', 'tibefp', 'mdi',
    'mp2d', 'mfee', 'openfermionpsi4', 'pemsetver',
    'psi4fockei', 'psixas', 'resp', 'simint', 'snsmp2',
    'v2fdm_cassef']
```



PROGRAMMATIC METADATA

cater to tools as well as people

OPTKING

```
def welcome():
    return """
\t\t\t OPTKING 3.0: for geometry optimizations
\t\t\t By R.A. King, Bethel University
\t\t\t with contributions from
\t\t\t A.V. Copan, J. Cayton, A. Heide
\t\t"""

```

LIBINT

```
const char *libint_reference(void) {
    std::string ref =
        "Libint: @LIBINT_DESCRIPTION@, Version " +
        std::string(libint_version_string(true)) +
        " Edward F. Valeev, http://libint.valeev.net/";
    return ref;
}
```

LIBXC

```
=> LibXC <=
Version 6.2.2
S. Lehtola, C. Steigemann, M. J.T. Oliveira, and M. A.L.
Marques., SoftwareX 7, 1–5 (2018) (10.1016/j.softx.2017.11.002)
```

PSI4

```
print(psi4.citation_formatter())
#> Psi4 v1.10a1.dev86, J. Chem. Phys. 152(18) 184108 (2020)
#> (10.1063/5.0006002)

print(psi4.citation_formatter("DOI: {doi}"))
#> DOI: 10.1063/5.0006002
```

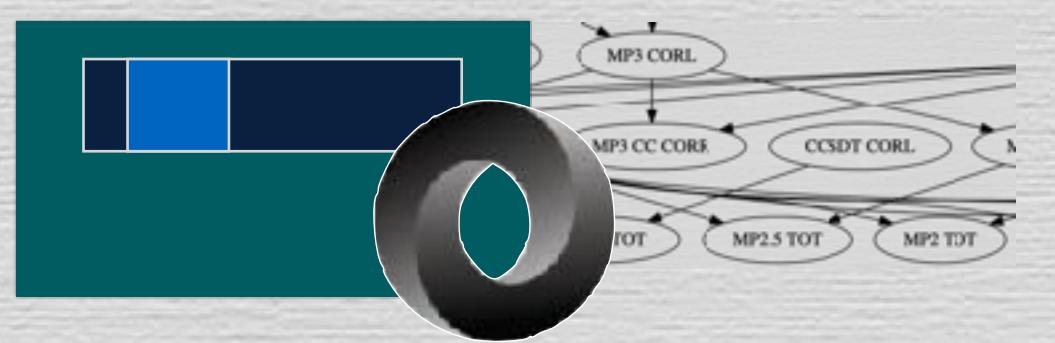


PROGRAMMATIC METADATA

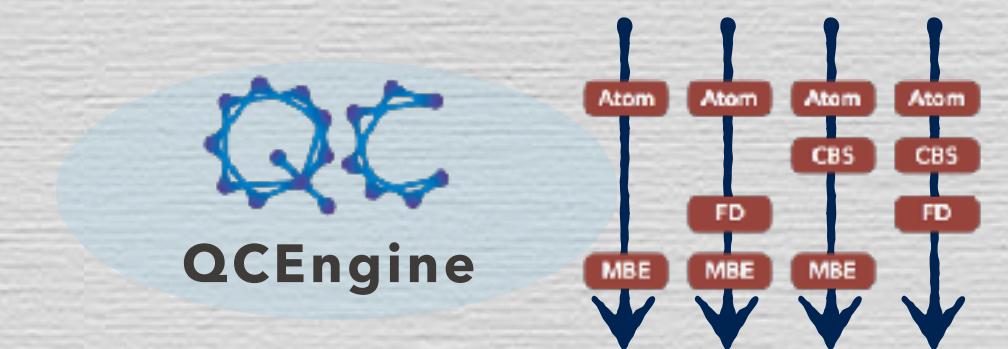
cater to tools as well as people **TORSIONDRIVE: CITATION.cff**

- We've solved the uncertainty over ordering configuration of published Libint packages for OSS
- But there are several other things that consumers may need or want to know from their consumed software. Modular software needs to expose details to all of people, runtime artifact, and build system.
- Some need to be brought to **PEOPLE**'s attention:
 - modular software can get buried behind the front-end – I'm sure Libxc has many users who have never heard of it. PSI4 uses some addons anonymously.
 - be sure to make your name, version, & citation accessible, so consuming programs can toast you.
- Projects increasingly good at recording their citation and authorship and license somewhere, and GitHub conventions have helped a lot. PSI4 uses codemeta.json, but I've increasingly used **CITATION.CFF** that facilitates both a journal and a repository citation.

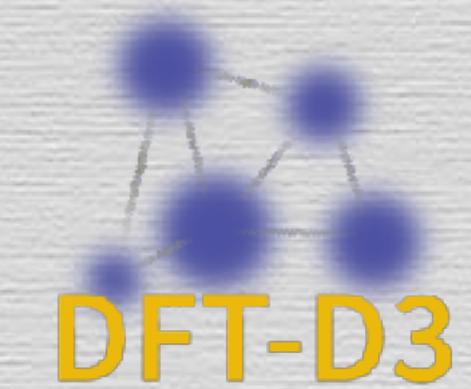
```
cff-version: 1.2.0
title: TorsionDrive
type: software
authors:
  - given-names: Lee-Ping
    family-names: Wang
    affiliation: University of California Davis
    orcid: 'https://orcid.org/0000-0003-3072-9946'
  - given-names: Yudong
    family-names: Qiu
    orcid: 'https://orcid.org/0000-0003-4345-8356'
repository-code: 'https://github.com/lpwgroup/torsiondrive'
abstract: Dihedral scanner with waveform propagation.
keywords:
  - molecular geometry optimization
  - molecular force-field fitting
license: MIT
version: 1.2.0
date-released: '2025-05-03'
preferred-citation:
  type: article
  authors:
    - family-names: "Yudong"
      given-names: "Qiu"
    - family-names: "Wang"
      given-names: "Lee-Ping"
doi: "10.1063/5.0009232"
journal: "The Journal of Chemical Physics"
month: 6
start: 244116
title: "Driving torsion scans with waveform propagation"
issue: 24
volume: 152
year: 2020
```



DISTRIBUTED DRIVER

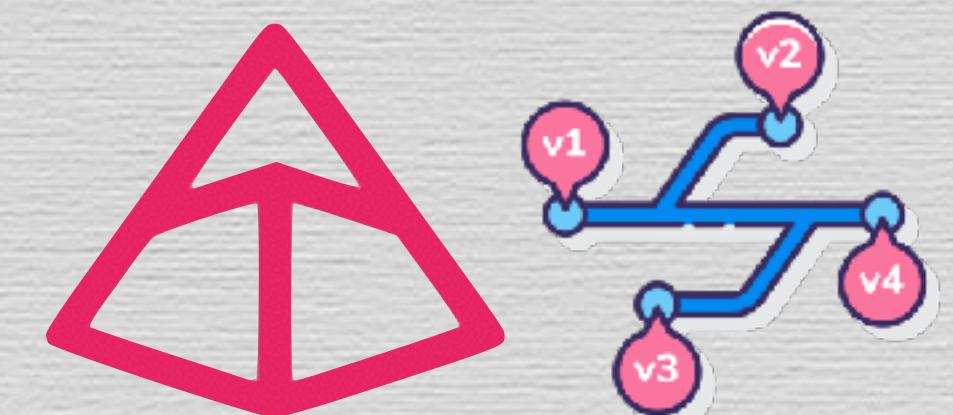
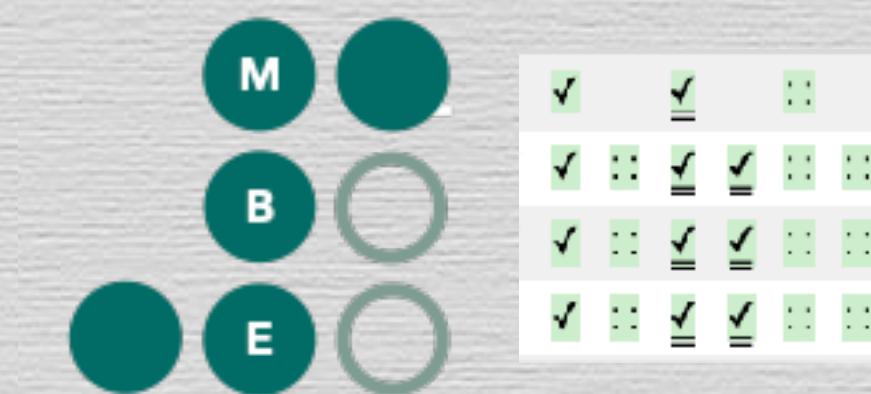
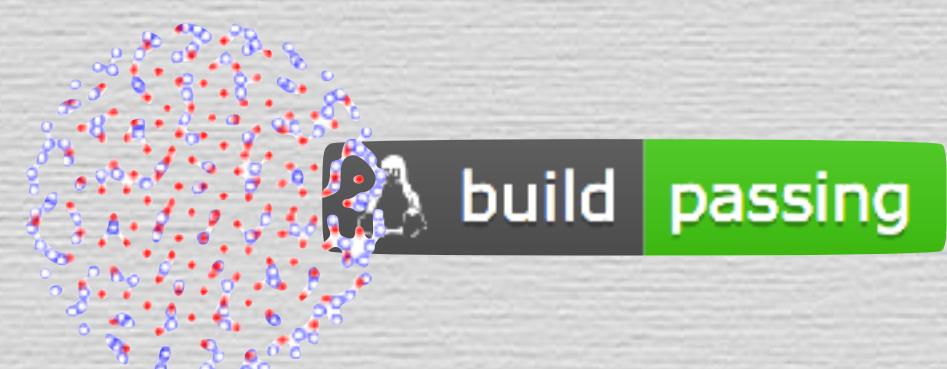


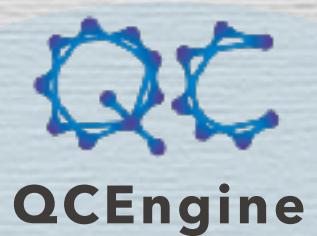
QCENGINE



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY

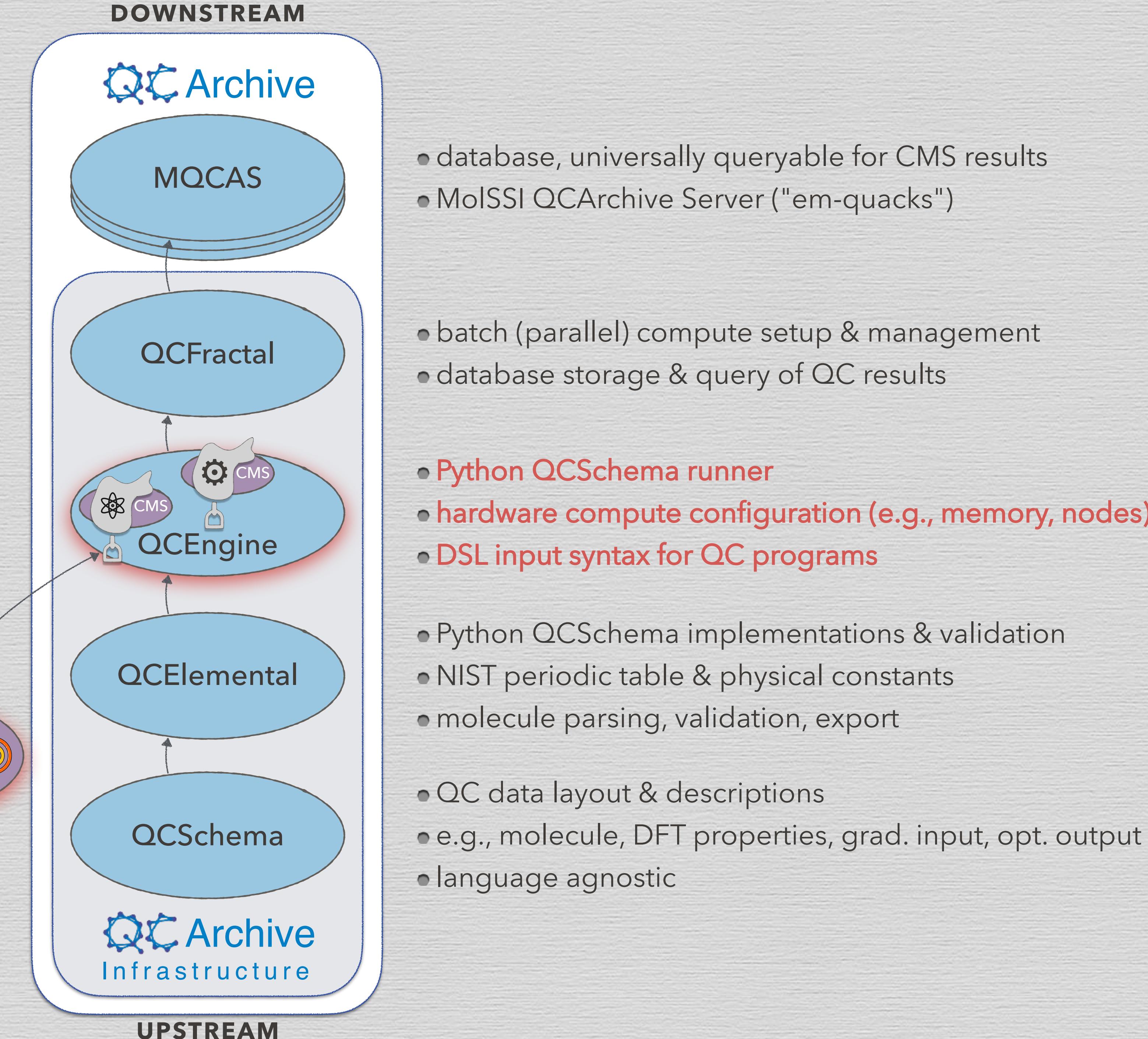
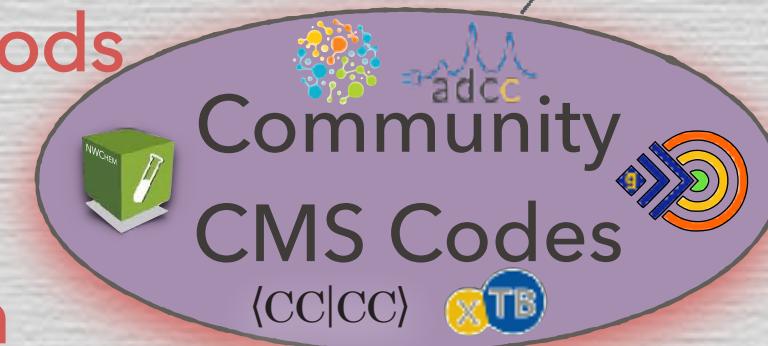




A QCENGINE STORY

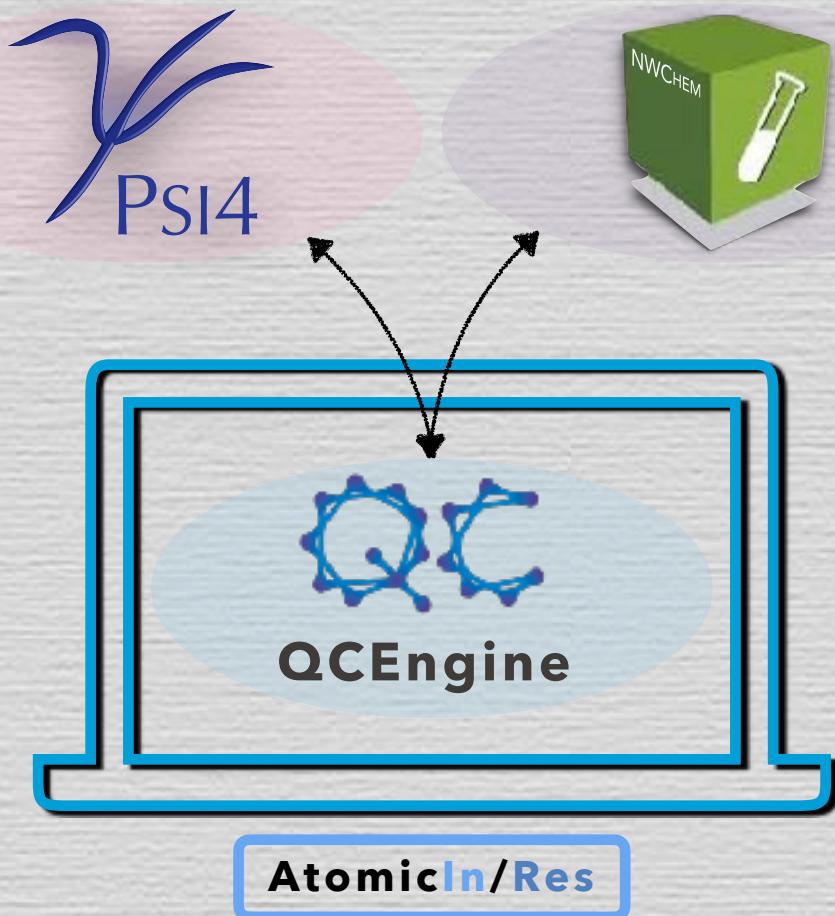
the QCArchive stack

- the difficult part – coded QC methods
- structured output uncommon
- DSL input; API/schema uncommon



A QCENGINE STORY: PYTHON QCSHEMA RUNNER

github.com/MoSSI/QCEngine



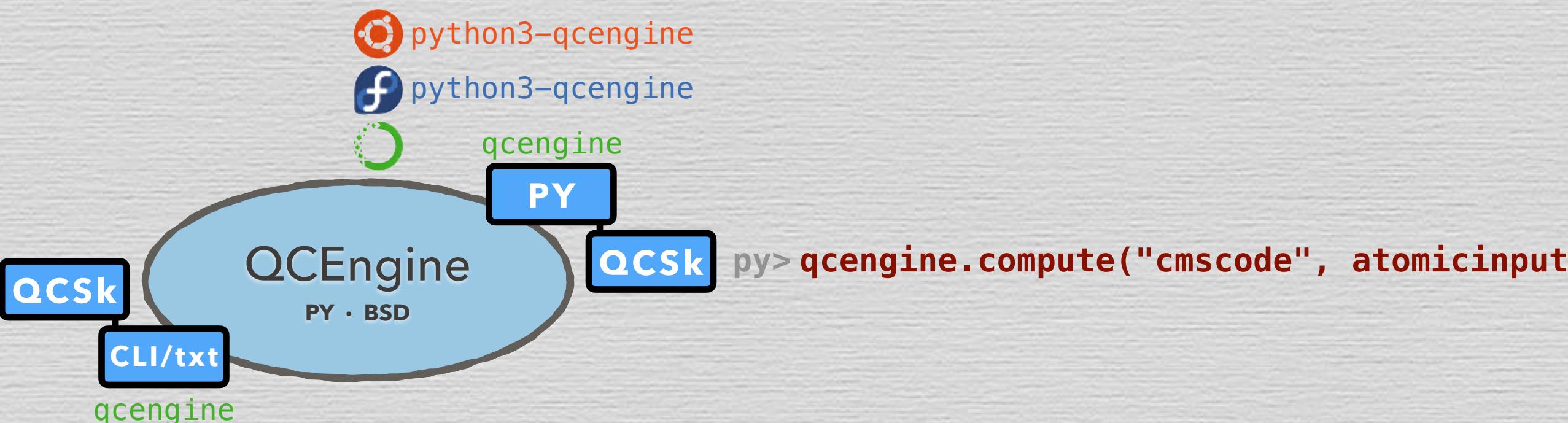
CMDLINE

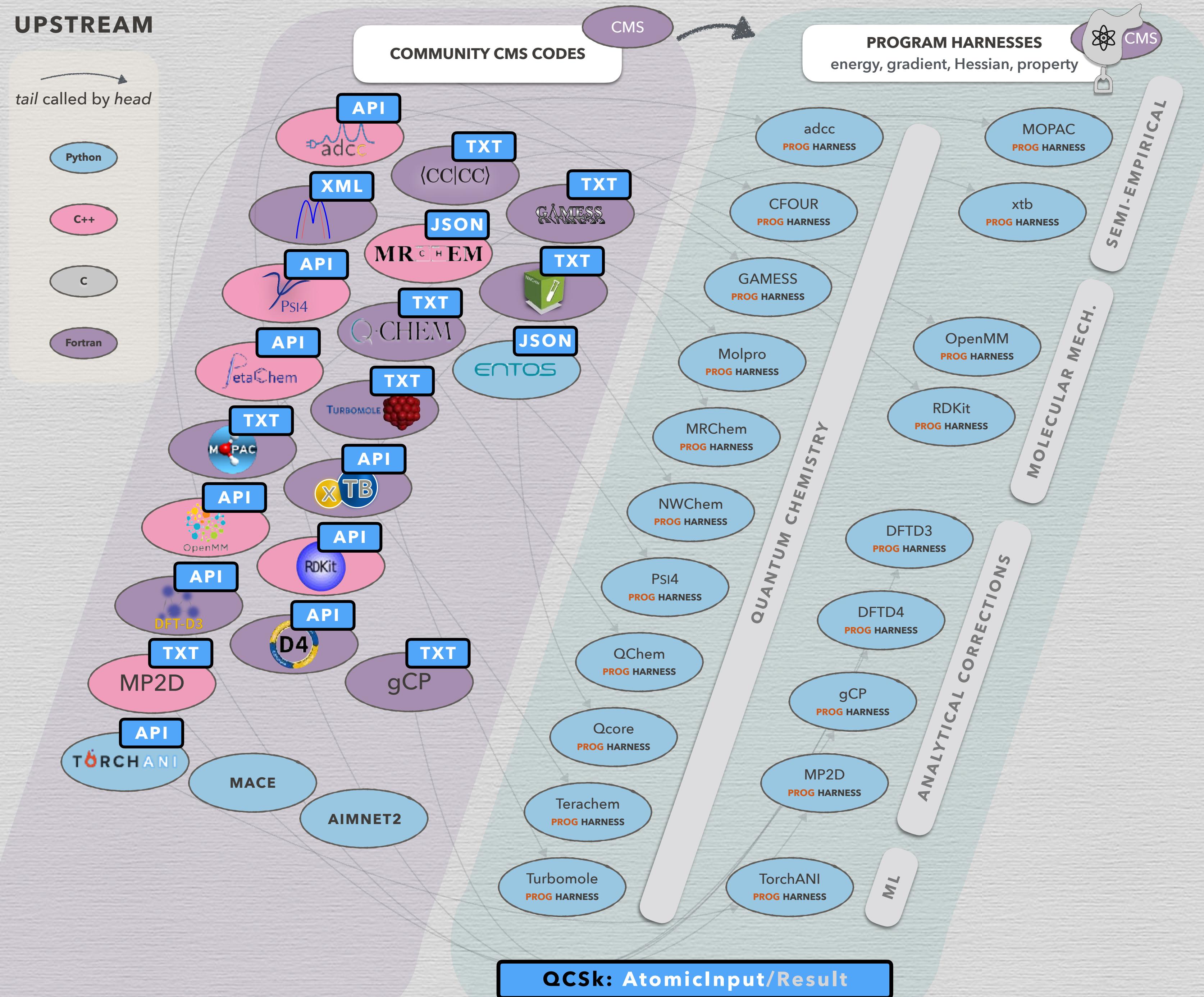
```
cfour  
> qcengine run molpro atomicinput  
psi4  
dftd4
```

PYAPI

```
qcengine.compute(atomicinput, "molpro", task_config({"nnodes": 4, "memory": 10})  
                  cfour  
                  psi4  
                  dftd4
```

sh> qcengine run cmscode atomicinput.json py> qcengine.compute("cmscode", atomicinput)



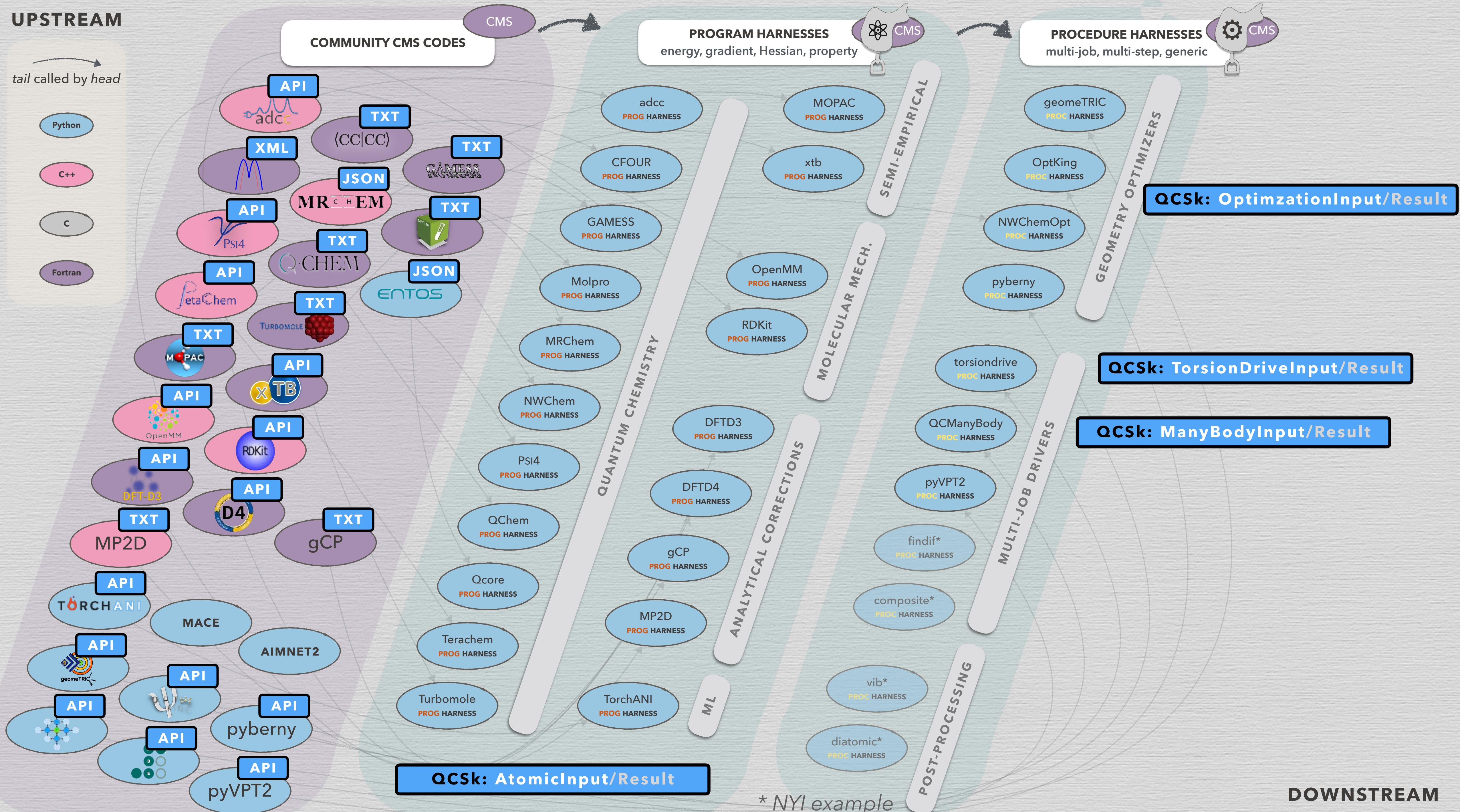


QCE ENGINE: INTERNAL MAP

github.com/MoISSI/QCEngine

- **SCHEMA** runner
 - **PROGRAMS** any analytic single point from CMS code
 - **PROCEDURES** anything except analytic single point. So far, mostly optimizers: geometric, optking, pyberny, NWChem (int.).
 - **PROGRAM CHECKS** indicate some methods accessible through QCSchema
 - **CMS** primarily QM but also SE, MM, partial, or ML
 - **INTERFACE** in variety of ways from API to structured data to regex. Former preferred for numerical precision.

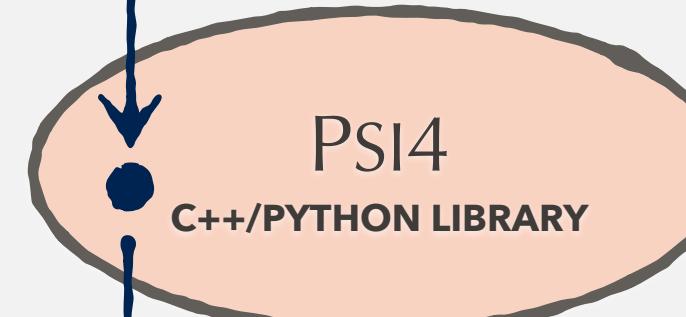
DOWNSTREAM



PSI4 DISTRIBUTED DRIVER

```
def energy(method): o def gradient(method): o def hessian(method): o
```

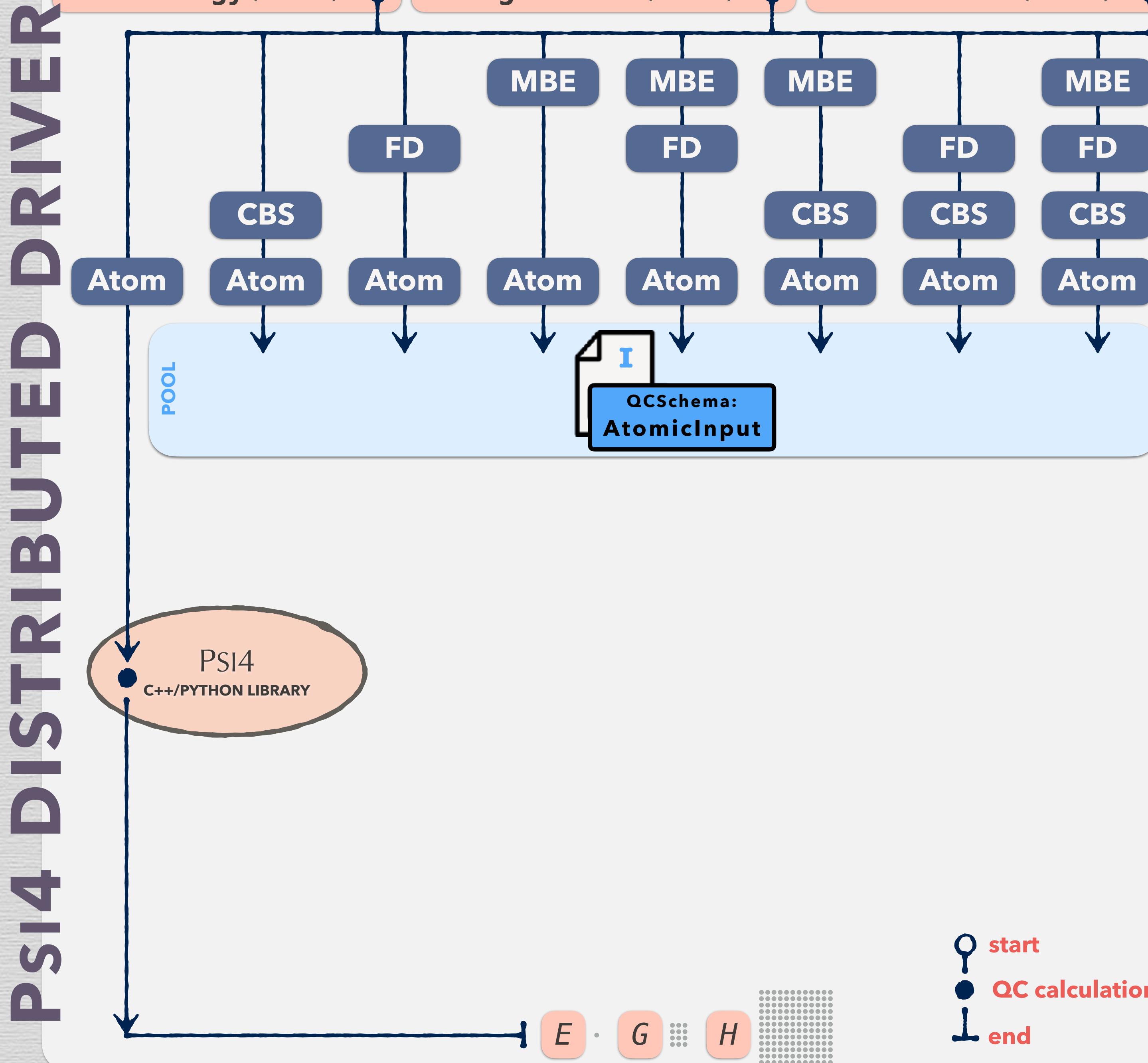
Atom = analytical single-point



E • G H

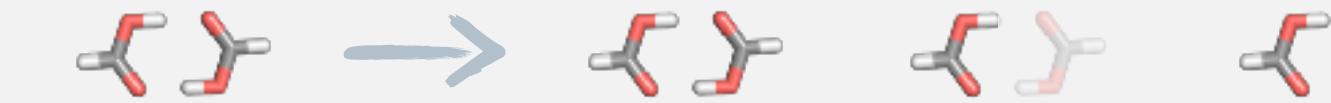
start
 QC calculation
 end

```
class AtomicComputer ():  
    PLAN molecule & method unchanged. return qcschema  
    ....  
    ASM Return analytic energy, gradient, or Hessian.
```



```
class ManyBodyComputer ():
```

AN Separate **molecule** into subsystems. CP, noCP, VMFC basis.
method unchanged.



```
for frag in fragments: return qcschema
```

SM Assemble n-body & interaction results from fragments.

```
class FiniteDifferenceComputer ():
```

AN Displace **molecule** according to stencil.
Reference **molecule & method** unchanged.



```
for disp in displacements: return qcschema
```

SM Assemble derivative results from displacements.

```
class CompositeComputer ():
```

AN Separate **method** into method, basis, & extrapolations.
molecule unchanged.



```
for mc in modelchems: return qcschema
```

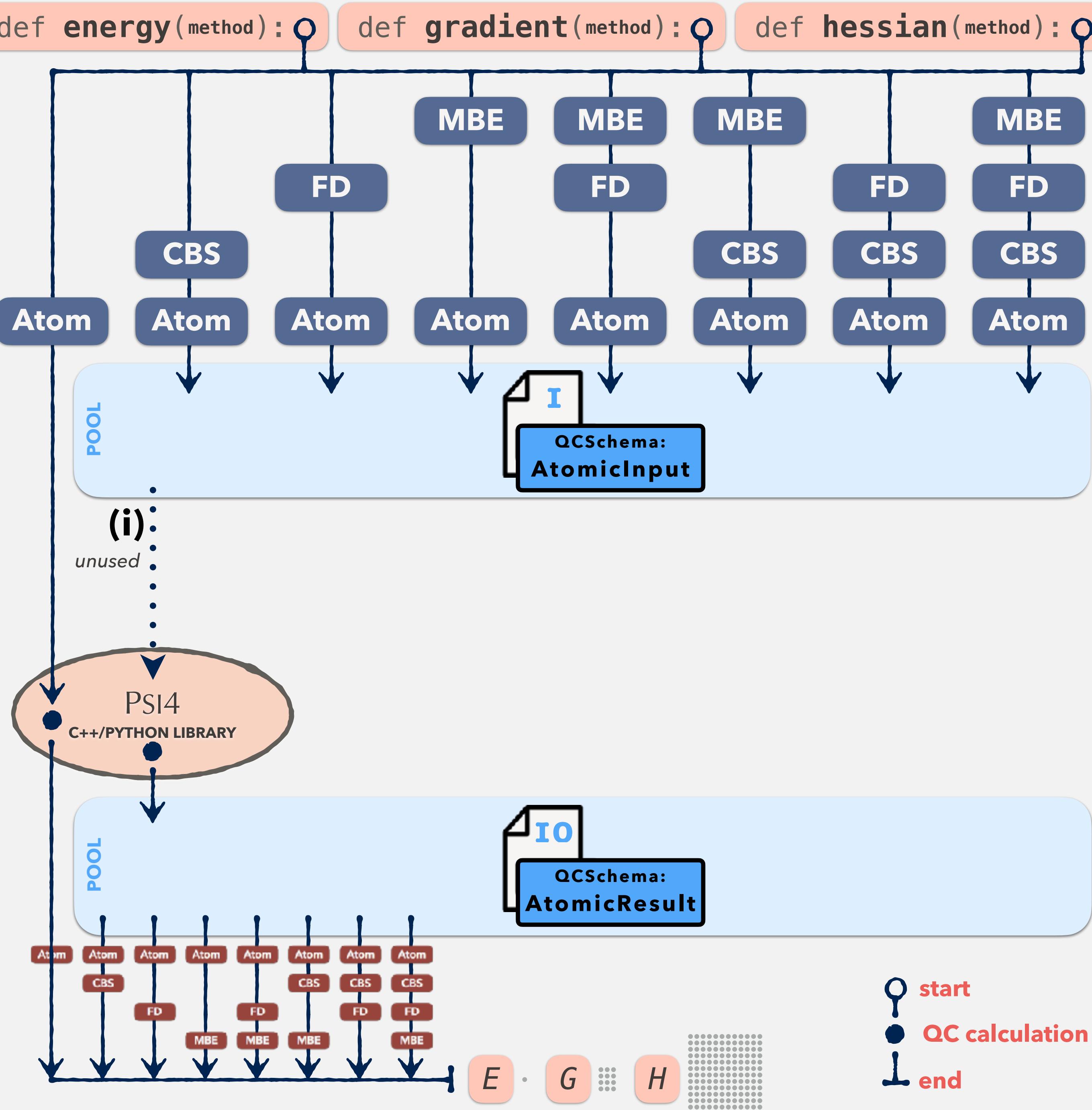
SM Assemble extrapolations & total results from modelchems

```
class AtomicComputer ():
```

AN molecule & method unchanged. return gcschema **gcschema**

SM Return analytic energy, gradient, or Hessian.

PSI4 DISTRIBUTED DRIVER



class ManyBodyComputer ():

Separate **molecule** into subsystems. CP, noCP, VMFC basis.
method unchanged.



for frag in fragments: return qcschema

.....
Assemble n-body & interaction results from fragments.

class FiniteDifferenceComputer ():

Displace **molecule** according to stencil.
Reference **molecule** & **method** unchanged.



for disp in displacements: return qcschema

.....
Assemble derivative results from displacements.

class CompositeComputer ():

Separate **method** into method, basis, & extrapolations.
molecule unchanged.



for mc in modelchems: return qcschema

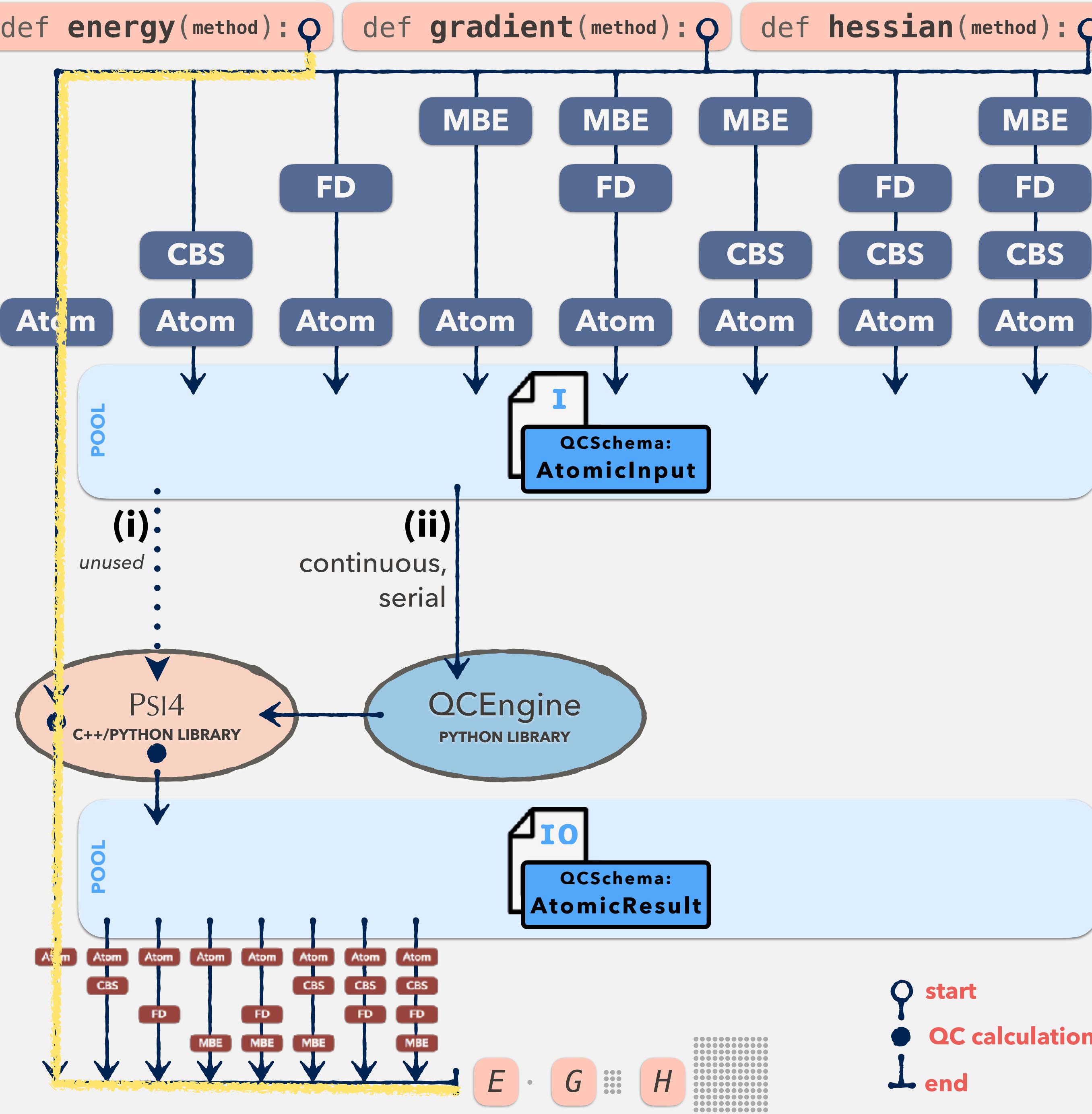
.....
Assemble extrapolations & total results from modelchems.

class AtomicComputer ():

molecule & **method** unchanged. return qcschema

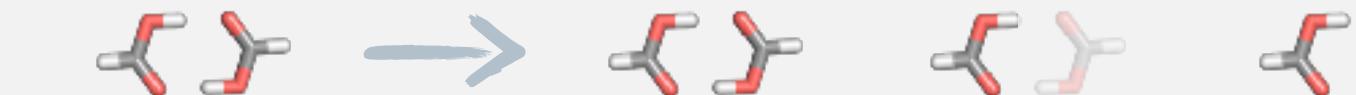
.....
Return analytic energy, gradient, or Hessian.

PSI4 DISTRIBUTED DRIVER



`class ManyBodyComputer ():`

Separate **molecule** into subsystems. CP, noCP, VMFC basis.
method unchanged.



for frag in fragments: return qcschema



Assemble n-body & interaction results from fragments.

`psi4.energy("mp2/cc-pvdz")`

`class CompositeComputer ():`

Separate **method** into method, basis, & extrapolations.
molecule unchanged.



for mc in modelchems: return qcschema



Assemble extrapolations & total results from modelchems.

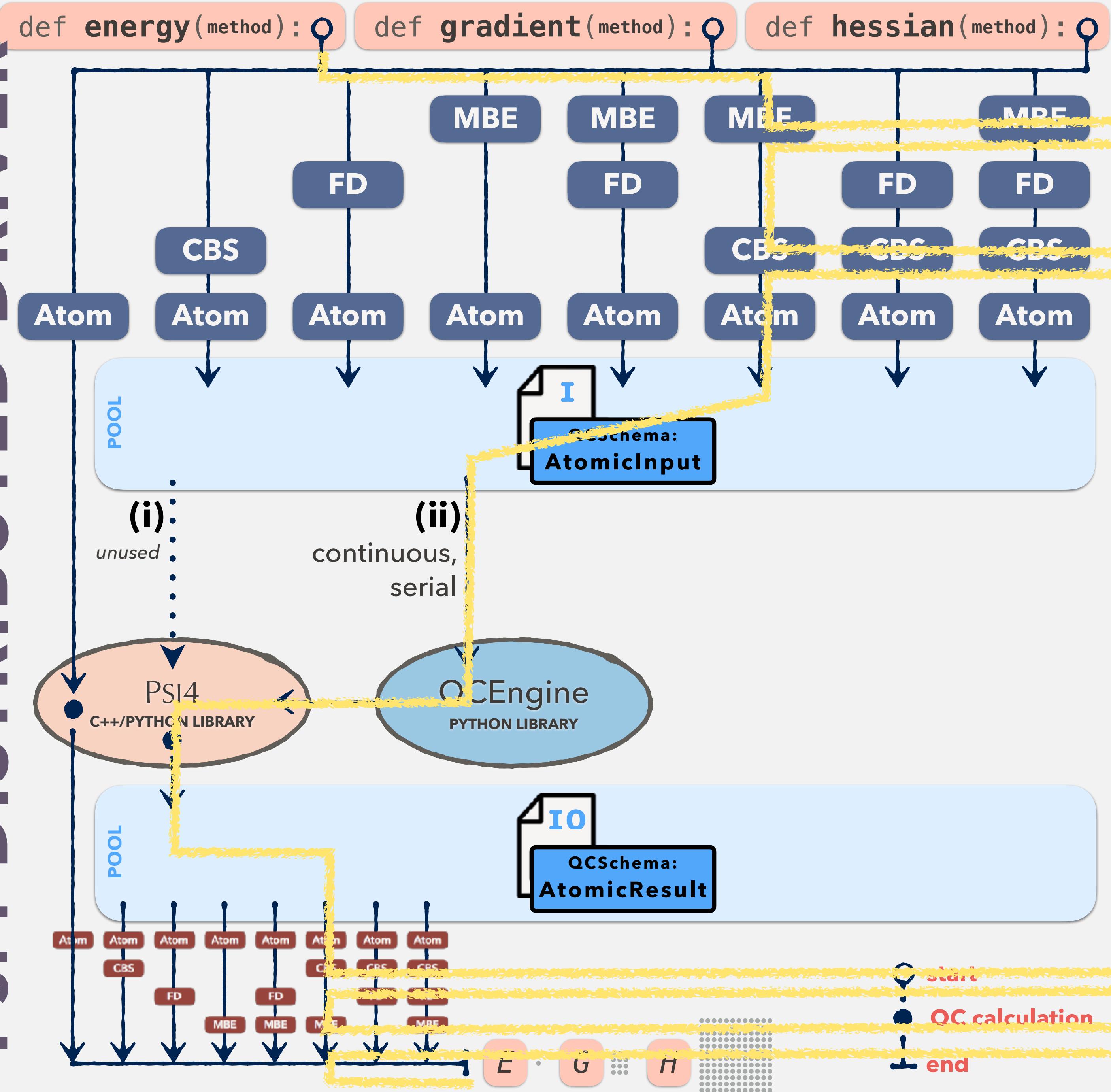
`class AtomicComputer ():`

molecule & **method** unchanged. return qcschema



Return analytic energy, gradient, or Hessian.

PSI4DISTIBUTEDDRIVER



```
class ManyBodyComputer ():
```

AN Separate molecule into subsystems. CP, noCP, VMFC basis.
method unchanged.

A chemical reaction diagram showing the conversion of two molecules with red and grey groups into one molecule with red and grey groups and one with pink and grey groups.

Erstpräfung im Praktikum / Praktikumsbericht

CH- Assemble in body & interaction testbeds and prototypes.

```
psi4.energy("mp2/cc-pvdz")
```

```
psi4.energy("mp2/cc-pv[dt]z", bsse_type="cp")
```

```
class CompositeComputer ():
```

AN Separate method into method, basis, & extrapolations.
molecule unchanged.

mp2/cc-pv[**tq**]z

MP2 TOTAL ENERGY/cc-pVTZ
MP2 TOTAL ENERGY/cc-pVQZ

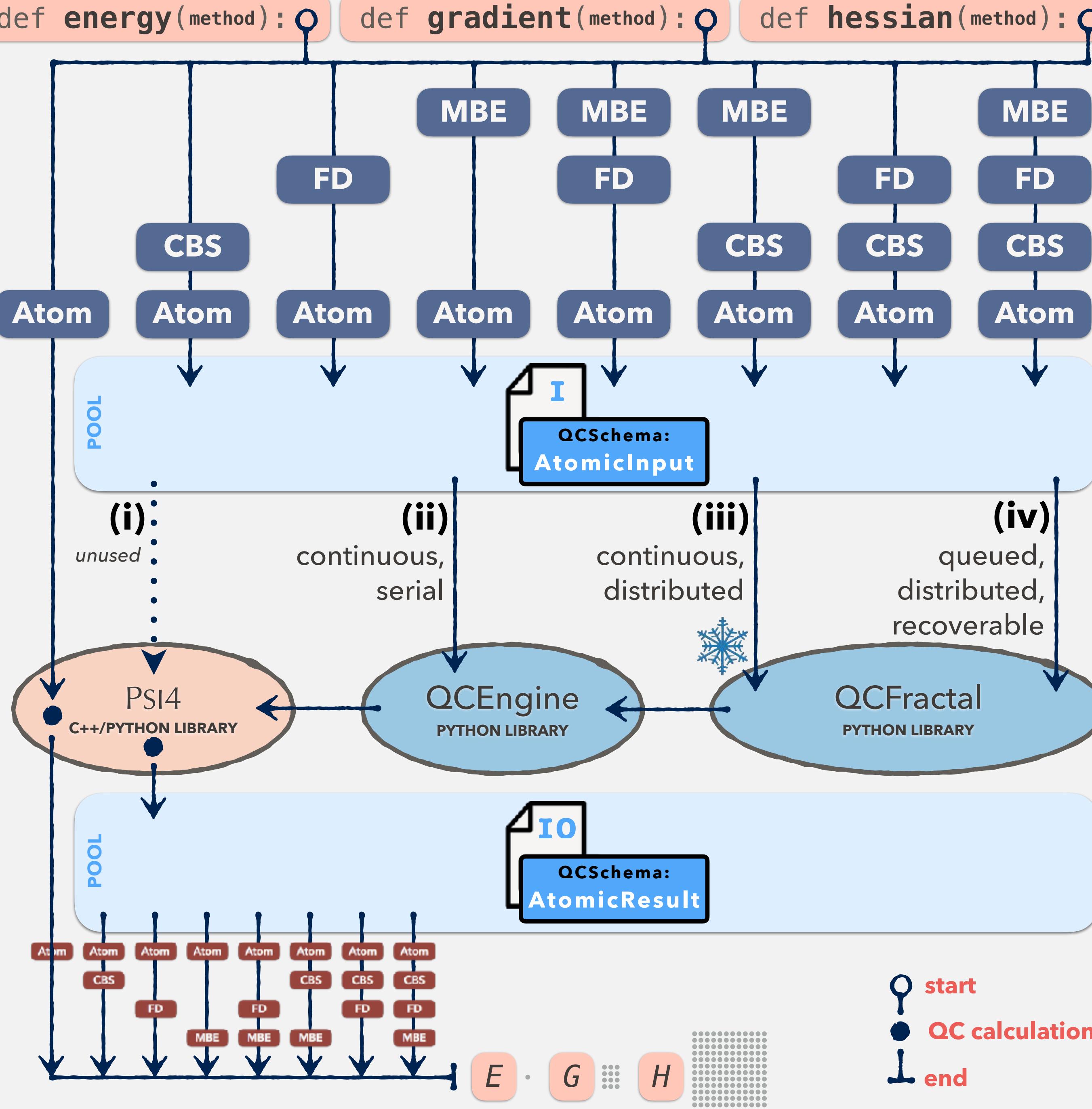
SM Assemble extrapolations & total results from modelchms

```
class AtomicComputer ():
```

LAN molecule & method unchanged. return qcschema 

SM Return analytic energy, gradient, or Hessian.

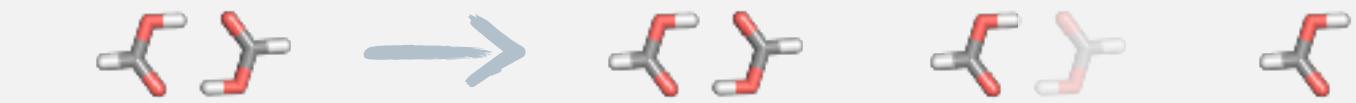
PSI4 DISTRIBUTED DRIVER



```
def energy(method): o
def gradient(method): o
def hessian(method): o
```

```
class ManyBodyComputer ():
```

Separate molecule into subsystems. CP, noCP, VMFC basis.
method unchanged.



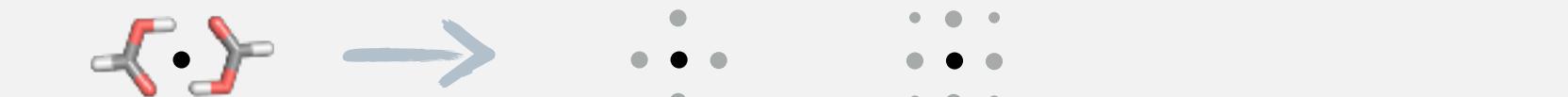
for frag in fragments: return qcschema



Assemble n-body & interaction results from fragments.

```
class FiniteDifferenceComputer ():
```

Displace molecule according to stencil.
Reference molecule & method unchanged.



for disp in displacements: return qcschema



Assemble derivative results from displacements.

```
class CompositeComputer ():
```

Separate method into method, basis, & extrapolations.
molecule unchanged.



for mc in modelchems: return qcschema



Assemble extrapolations & total results from modelchems.

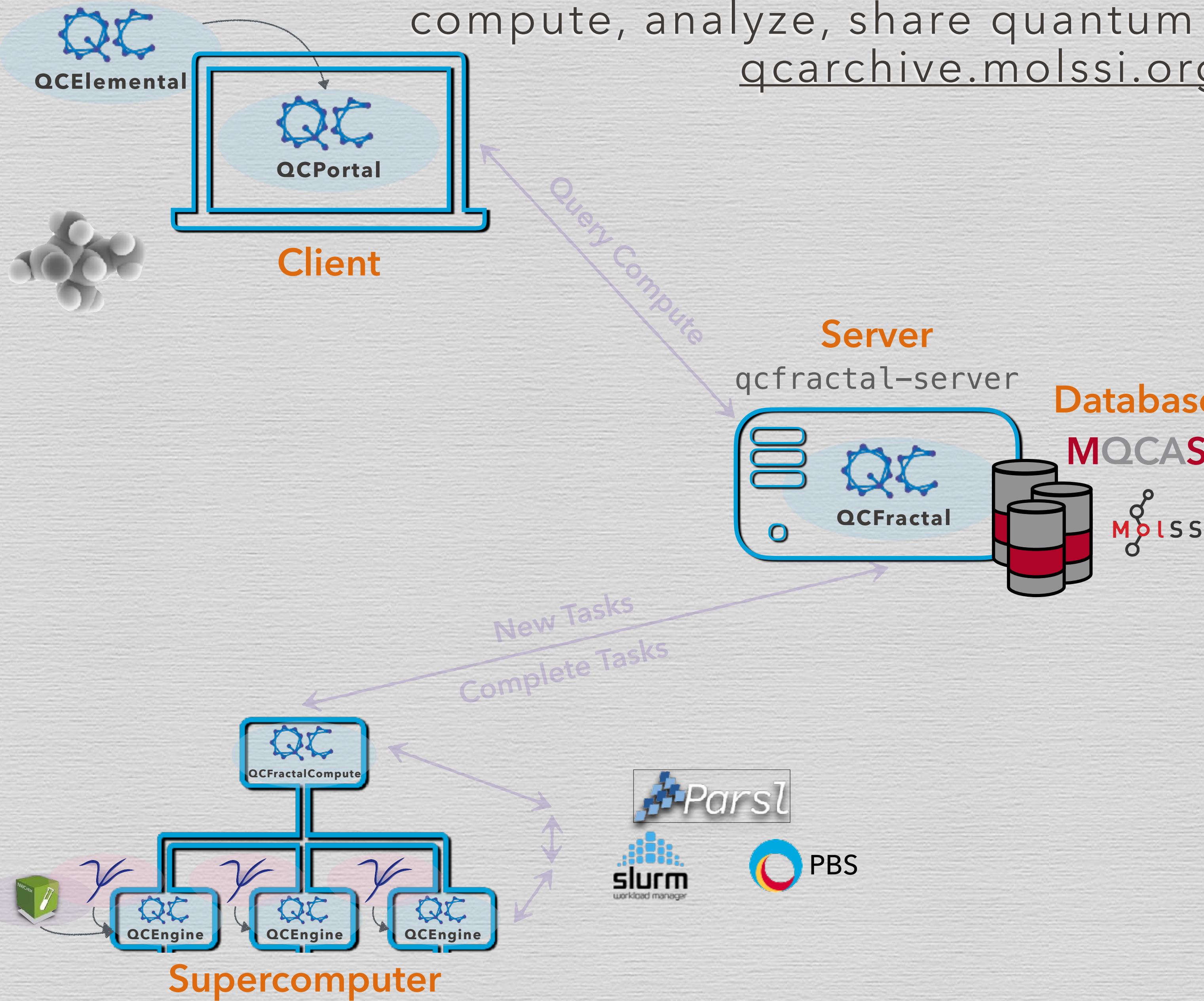
```
class AtomicComputer ():
```

molecule & method unchanged. return qcschema



Return analytic energy, gradient, or Hessian.

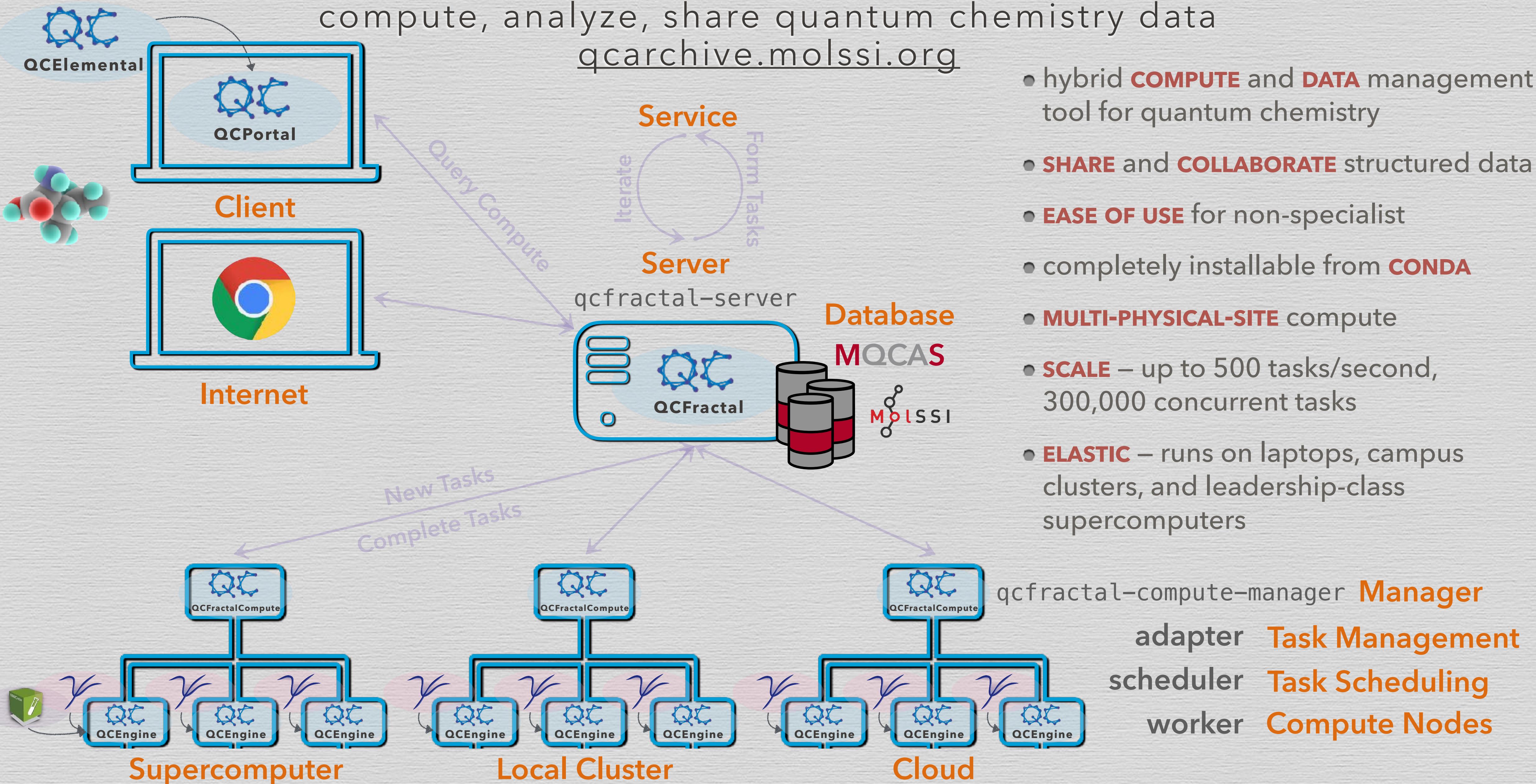
QCARCHIVE OVERVIEW



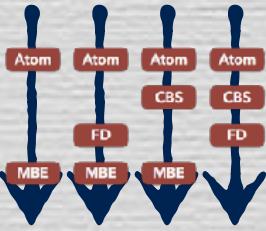
- hybrid **COMPUTE** and **DATA** management tool for quantum chemistry
- **SHARE** and **COLLABORATE** structured data
- **EASE OF USE** for non-specialist
- completely installable from **CONDA**
- **MULTI-PHYSICAL-SITE** compute
- **SCALE** – up to 500 tasks/second, 300,000 concurrent tasks
- **ELASTIC** – runs on laptops, campus clusters, and leadership-class supercomputers

qcfractal-compute-manager **Manager**
adapter **Task Management**
scheduler **Task Scheduling**
worker **Compute Nodes**

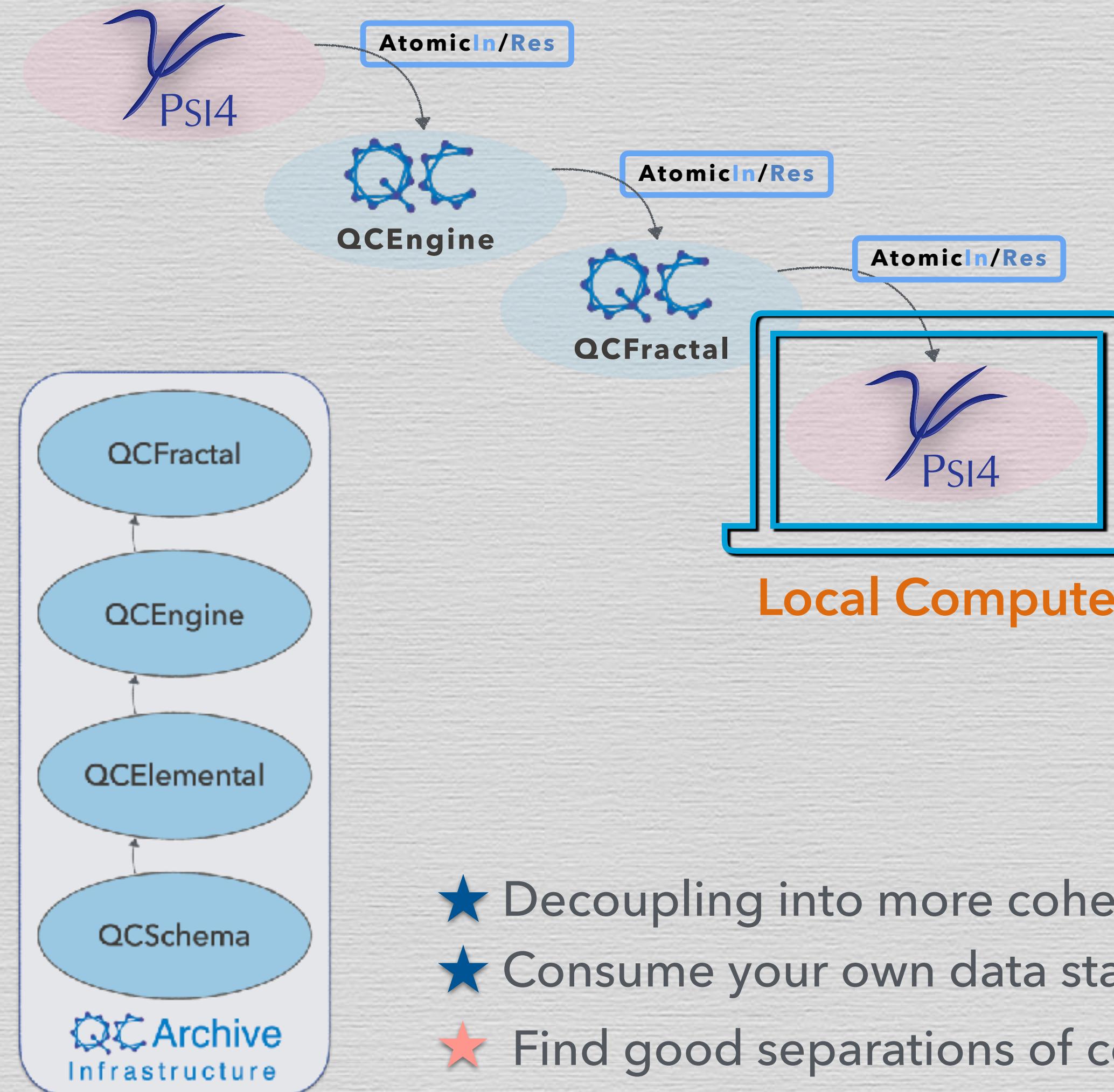
QCARCHIVE OVERVIEW



PSI4 DISTRIBUTED DRIVER + QCARCHIVE



distributed, searchable, and error-resistant

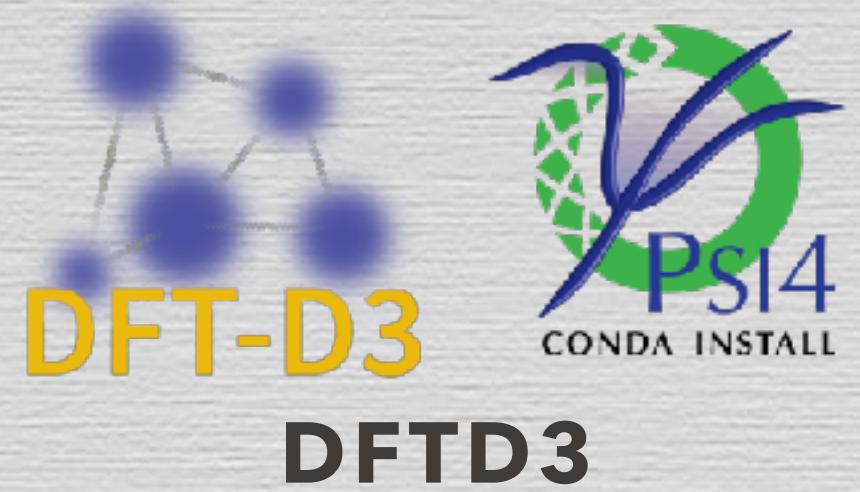
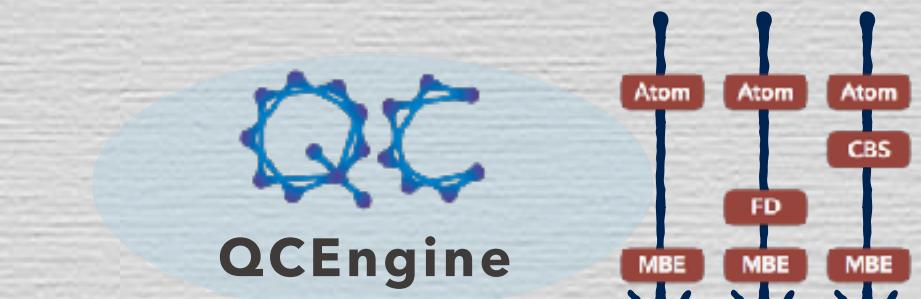
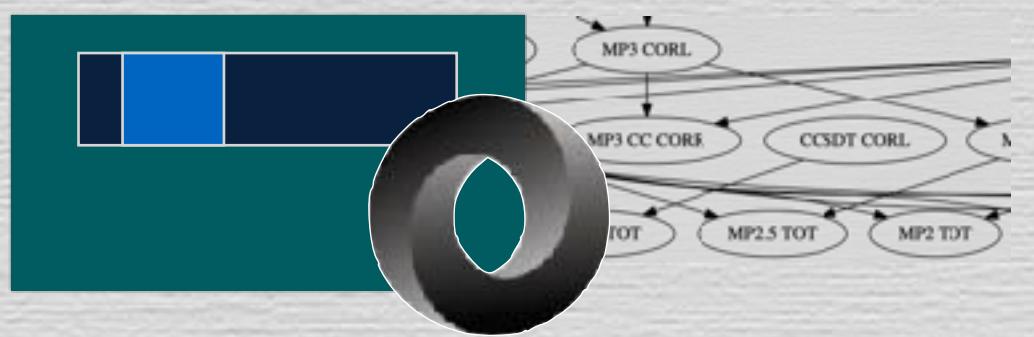


- (1) **PSI4** driver (post-proc. manager) plans single-point jobs from API input
- (2) **QCFractal** orchestrates job running among attached resources
- (3) **QCEngine** receives jobs as **AtomicInput QCSchema**
- (4) **PSI4** driver (core) runs single-point jobs
- (5) **QCEngine** receives jobs as **AtomicResult QCSchema**
- (6) **QCFractal** stores jobs in DB & retrieves on query
- (7) **PSI4** driver assembles results into usual API

- **PAYOUT** PSI4 gets parallelism we don't have to manage.
- **PAYOUT** we had to become more robust to handle PSI4 calling PSI4.

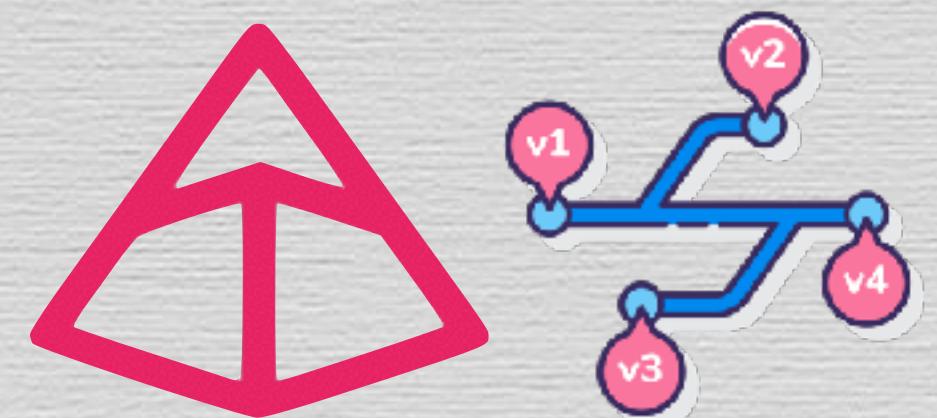
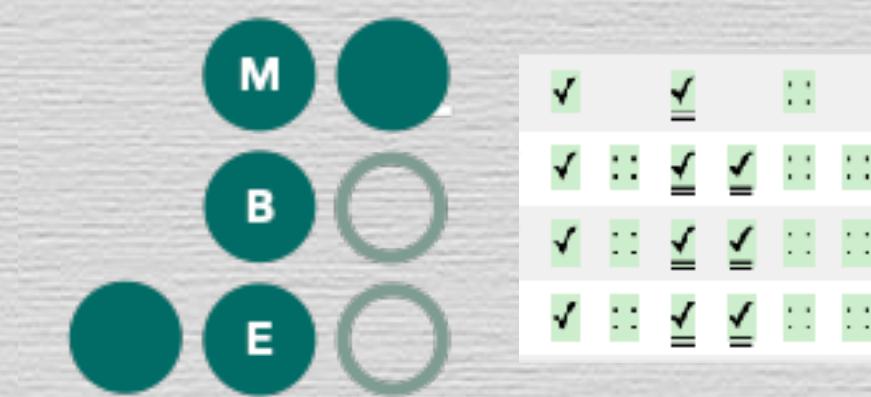
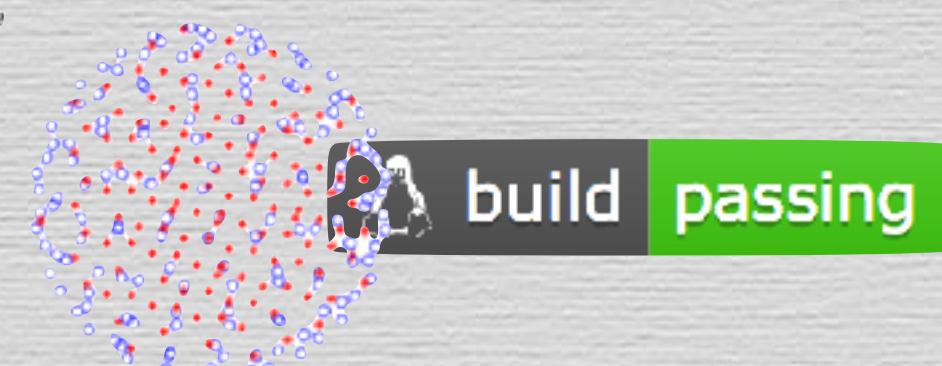
- ★ Decoupling into more cohesive internal modules is progress, too.
- ★ Consume your own data standards to exercise them.
- ★ Find good separations of concern for software stack & build proj. around those missions.

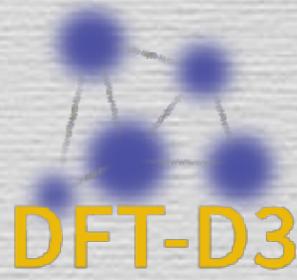
CONDA PACKAGING



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY





DFT-D3

A DFTD3 STORY

hold your interfaces and tests close

In the beginning was the **METHOD** publication



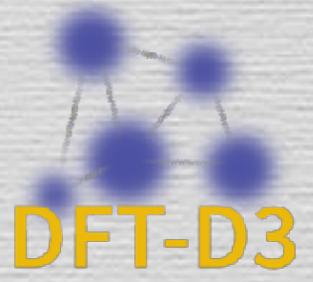
A consistent and accurate *ab initio* parametrization of density functional dispersion correction (DFT-D) for the 94 elements H-Pu

J. Chem. Phys. **132**, 154104 (2010); <https://doi.org/10.1063/1.3382344>

Stefan Grimme^{a)}, Jens Antony, Stephan Ehrlich, and Helge Krieg

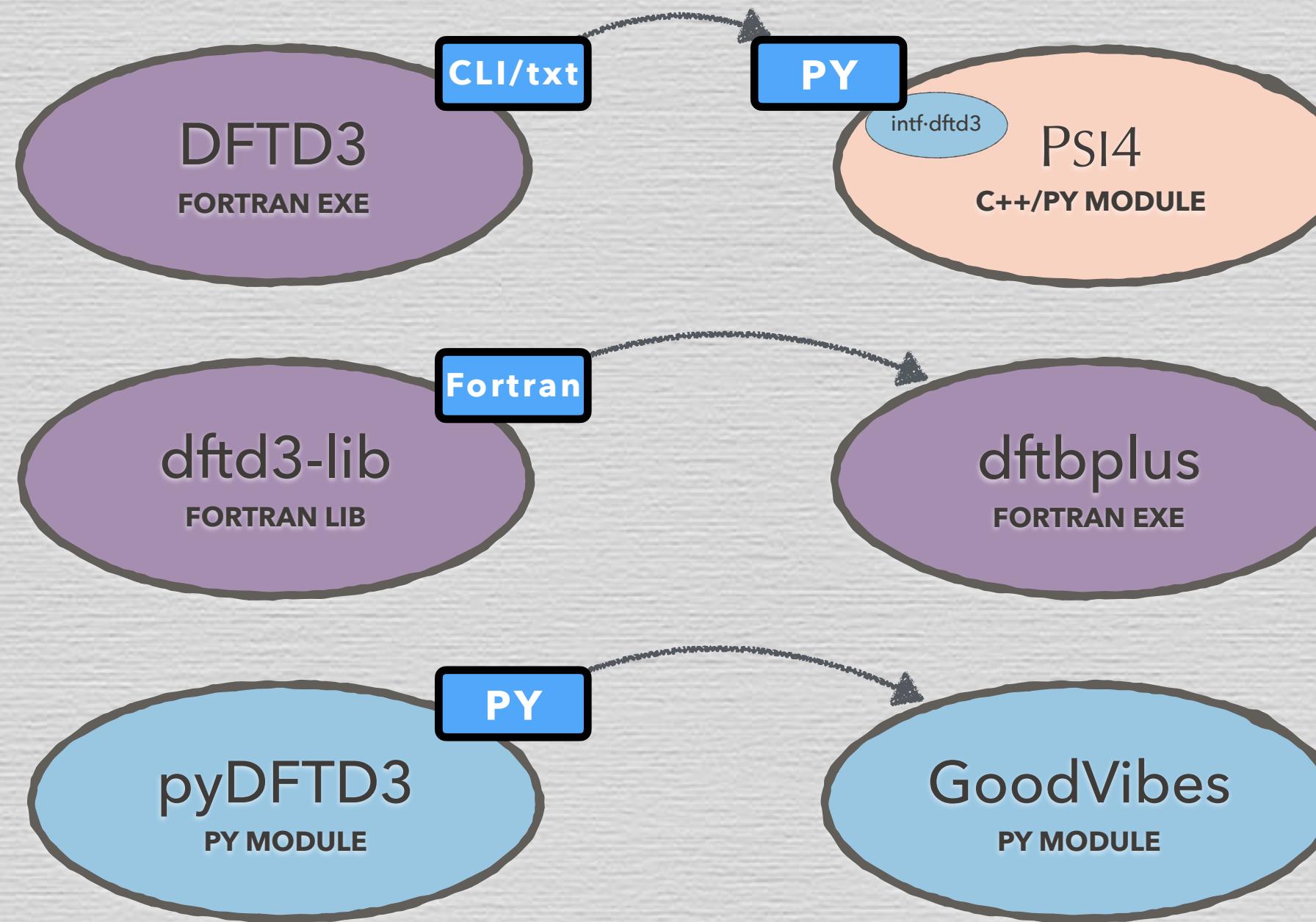
Still better, the **CODE** for a Fortran executable

The screenshot shows the homepage of the DFT-D3 software website. At the top, it features the University of Bonn logo and the Mulliken Center for Theoretical Chemistry logo. A navigation bar includes links for Workgroups, Research, Teaching, Seminar, Software, Contact, and History. Below the navigation bar, there is a sidebar titled "Software" under the "Grimme" section, listing various tools: DFT-D3, DFT-D4, FODplot, gCP, gCP-D3 Webservice, GMTKN, HF-3c, MOR41, QCEIM6, QMDFF, QMS-M, ROST61, sTDA, TMG145, AZIDE, and xtb. To the right of the sidebar, there is a main content area with a photograph of a street in Bonn. The content area includes a brief description of DFT-D3, a link to the article in JCP, and a note about the BJ-damping. It also mentions a related article by Sherrill et al. at J. Phys. Chem. Lett. and a note about the C6-coefficients. At the bottom of the page, there is a "Contents" section with a link to "Get DFT-D3 and the most recent coefficient file".



A DFTD3 STORY

hold your interfaces and tests close



OPEN-SOURCE projects proliferated until inevitably ...

Which dftd3 should one use? #4

Closed zerothi opened this issue on Nov 18, 2021 · 10 comments

zerothi commented on Nov 18, 2021 · edited

Thanks for the work on this!

We are bit in a dead-lock now, since there exists 3 different libraries which all say do the same thing? :)

So which one should we support :)

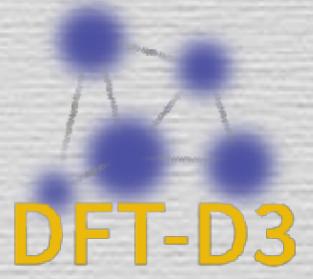
... at least ten **TEN IMPLEMENTATIONS** were identified

Which dftd3 should one use? #4
zerothi opened this issue on Nov 18, 2021 · 10 comments

awwgk commented on Nov 18, 2021 · edited

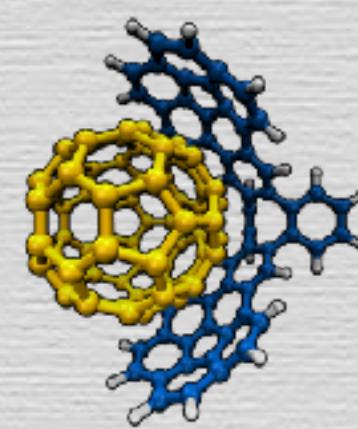
Here is a quick search for `dftd3` versions which are currently around:

repo	version	license	note
http://mctc.uni-bonn.de/software/dft-d3	v3.2.0	GPL-1.0-or-later	reference implementation
https://github.com/awwgk/simple-dftd3	v0.4.2	LGPL-3.0-or-later	this project
https://github.com/dftbplus/dftd3-lib	v0.10	GPL-1.0-or-later	CMake
https://github.com/ehermes/ased3	—	LGPL-3.0-or-later	ASE, f2py
https://github.com/pfnet-research/torch-dftd	v0.3.0	MIT	torch
https://github.com/cuanto/libdftd3	v3.1.1 (fork)	GPL-3.0-or-later	fork of dftd3-lib?, Python
https://github.com/cresset-group/dftd3	v3.2.0 (fork)	GPL-1.0-or-later	patched?
https://github.com/loriab/dftd3	v3.2.1 (fork)	GPL-1.0-or-later	CMake, patched
https://github.com/f3rmion/dftd3	v3.2.0 (fork)	GPL-1.0-or-later	patched
https://github.com/bobbypaton/pyDFTD3	v1.0.0	MIT	Python



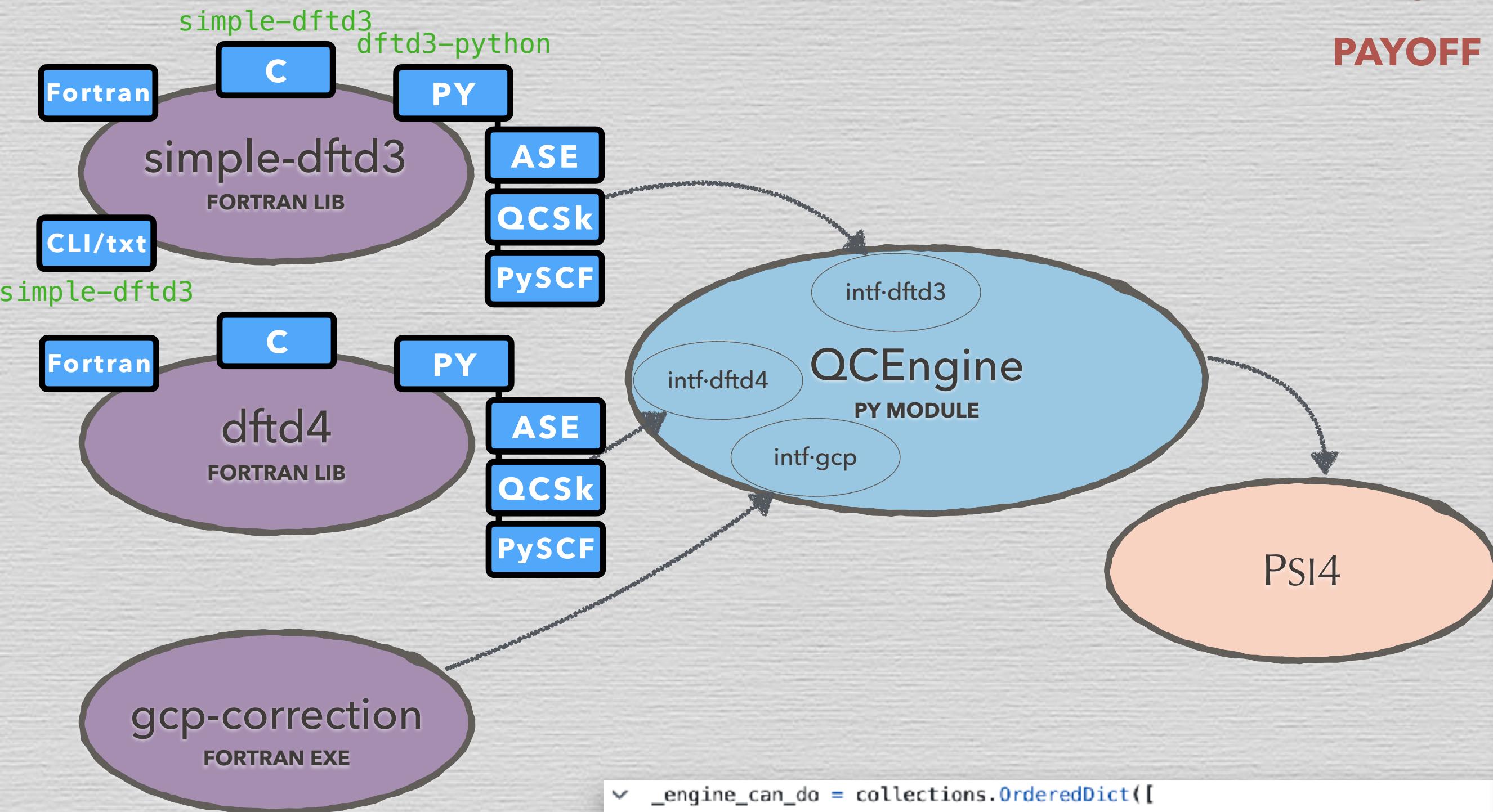
A DFTD3 STORY

★ hold your interfaces and tests close



Sebastian Ehlert Holger Kruse
Bonn → MS Czech Acad. Sci.
→ Pending AI

RE-IMPLEMENTED upstream with abundant interfaces



PAYOUT Psi4 no longer maintains and packages forks of D3, D4, & gCP
PAYOUT new DFT capabilities easy to add

```

249 + funcs.append({
250 +     "name": "R2SCAN3C",
251 +     "alias": ["R2SCAN-3C"],
252 +     "x_functionals": {
253 +         "MGGA_X_R2SCAN": {}
254 +     },
255 +     "c_functionals": {
256 +         "MGGA_C_R2SCAN": {}
257 +     },
258 +     "description": 'r2SCAN Meta-GGA based 3C composite method with a TZ basis set, gCP and D4\n',
259 +     "citation": 'S. Grimme, A. Hansen, S. Ehlert, J.-M. Mewes J. Chem. Phys. 154, 064103, 2021\n',
260 +     "doi": "10.1063/5.0040021",
261 +     "dispersion": {
262 +         "type": "d4bjeeqatm",
263 +         "params": {
264 +             'a1': 0.420,
265 +             'a2': 5.650,
266 +             's6': 1.000,
267 +             's8': 0.000,
268 +             's9': 2.000,
269 +             'ga': 2.000,
270 +             'gc': 1.000,
271 +         },
272 +     },
273 + })

```

Adds r2SCAN hybrids; r2SCAN-3c and B97-3c #2842
↳ Merged loriab merged 24 commits into psi4:master from hokru:r2scan on Dec 4, 2023

```

if self.engine in ["s-dftd3", 'dftd3', 'mp2d', "dftd4"]:
    resi = AtomicInput(
        **{
            'driver': 'energy',
            'model': {
                'method': self.fctldash,
                'basis': '(auto)',
            },
            'keywords': {
                'level_hint': self.dashlevel,
                'params_tweaks': self.dashparams,
                'dashcoeff_supplement': self.dashcoeff_supplement,
                'pair_resolved': self.save_pairwise_disp,
                'apply_qcengine_aliases': True, # for s-dftd3
                'verbose': 1,
            },
            'molecule': molecule.to_schema(dtype=2),
            'provenance': p4util.provenance_stamp(__name__),
        })
    jobrec = qcng.compute(
        resi,
        self.engine,
        raise_error=True,
        task_config={"scratch_directory": core.IOManager.shared_object().get_def(
            dashd_part = float(jobrec.extras['qcvars']['DISPERSION CORRECTION ENERGY'])

```

A DFTD3 STORY

smoothly replacing a much-used external feature

GUIDED PROGRESSION THROUGH ALTERNATE IMPLEMENTATIONS

pre-v1.0 **classic** dftd3 only

v1.7 **s-dftd3** preferred; **classic** through:

```
energy("b3lyp-d3", engine="dftd3")
```

v1.9 **s-dftd3** preferred; **classic** deprecated but runs:

(Note: apparently I forgot the warning, but this is the usual progression.)

```
FutureWarning("Using `engine='dftd3'` instead of `engine='s-dftd3'` is deprecated, and as soon as 1.10 it will stop working")
```

v1.10 **s-dftd3** only; **classic** errors:

```
UpgradeHelper("Using `engine='dftd3'` instead of `engine='s-dftd3'` is obsolete as of 1.10. Allow ... installing dftd3-python")
```

someday remove **classic UpgradeHelper** so errors unhelpfully

QCEngine soon remove **classic** harness

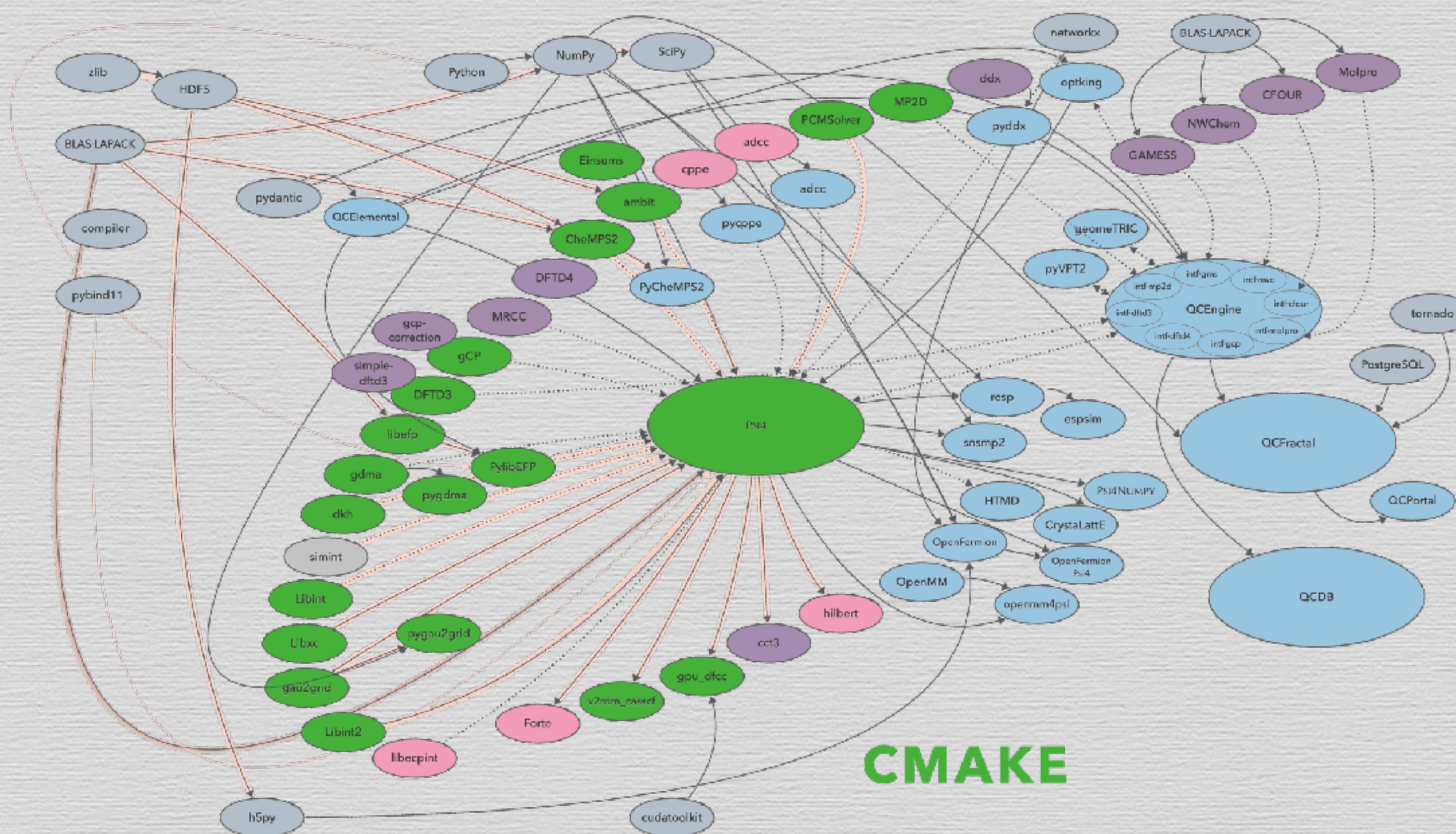
★ new syntax advice is more helpful at runtime than in docs. a grace period is a bonus

- Dispersion switches implementation w/i function call, and optimizers are similar: `optimize("b3lyp", engine="geometric|optking")`
- Alternate impl. of deeper functionality use **keywords**: `set integrals_package libint2|simint, set orbital_optimizer_package openorbitaloptimizer|internal`

```
#ifdef USING_OpenOrbitalOptimizer
    /*- Orbital optimizer package to use for SAD guess. -/
    options.add_str("ORBITAL_OPTIMIZER_PACKAGE", "OPENORBITALOPTIMIZER", "OPENORBITALOPTIMIZER INTERNAL");
#else
    /*- Orbital optimizer package to use for SAD guess. -/
    options.add_str("ORBITAL_OPTIMIZER_PACKAGE", "INTERNAL", "INTERNAL");
#endif
```

CONDA-FORGE PACKAGING

<https://anaconda.org/>



CMAKE

- **CONDA** is a cross-platform, cross-language binary package manager supporting community packaging & focused on the scientific stack
- **MKL LAPACK**, compilers, CUDA runtimes are among the production-quality tools available
- **CMAKE** has been contributed to a number of community projects to facilitate integration with Psi4 (and add Windows)
- **THIRTY** ecosystem projects are or have been packaged by Psi4
- **STARTING** 2015 Psi4 available from a custom `-c psi4` channel. Since 2023, have recc. **conda install psi4 --channel conda-forge**
- **CONDA-FORGE** is increasingly (**30k**) the home for downstream & dep. projects, so for robust environment solving, we've transitioned to c-f
- **PRESENTLY** Linux (Intel & ARM), Windows, and Mac (Intel & Silicon) Psi4 ecosystems available from conda



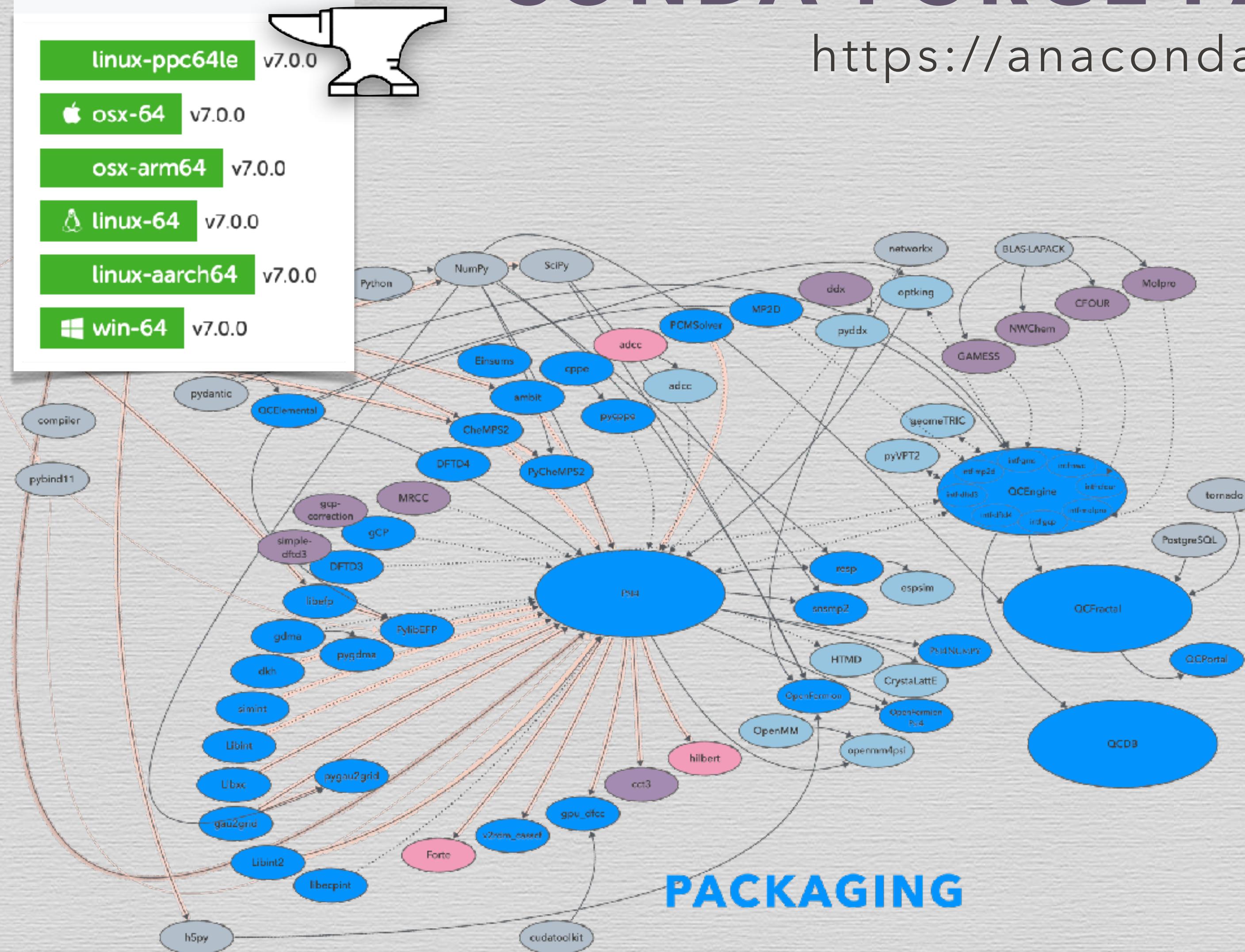
recipe libxc-c

CONDA-FORGE PACKAGING

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recipe/meta.yaml

CONDA-FORGE RECIPE: CPPE

a Py extension module

- A conda **RECIPE** contains project metadata and build directions. It's similar to a Homebrew formula (Linux/Mac) or a Fedora Spec file (Linux).
- To begin, make a PR with these files to **CONDA-FORGE/STAGED-RECIPES**. Packages will build in CI, so iterate until all passing, then request review.
- Once merged, you (and any co-**MAINTAINERS**) will have full permissions on your own project repository.
- **META** defines source, patches, expected API compat., build tools, build-time and run-time deps, anticipated files, testing, and licensing.
- **BUILD** scripts build & install for Unix & Windows with many defined variables.
- Multiple **OUTPUTS** (e.g., C, Fortran, Py) possible from one build.
- Everything is done in **GITHUB CI**, so arch are often cross-compiled.

```
package:
  name: cppe
  version: "0.3.2"

source:
  url: GH:maxscheurer/cppe/archive/v0.3.2.tar.gz
  patches:
    - 0001-update-math-call.patch

build:
  number: 3
  run_exports:
    - {{ pin_subpackage('cppe', max_pin='x.x') }}

requirements:
  build:
    - {{ stdlib("c") }}
    - {{ compiler('cxx') }}
    - python # [build_platform != target_platform]
    - cmake
    - eigen
  host:
    - pybind11
    - python
  run:
    - python

test:
  requires:
    - pytest
  imports:
    - cppe
  commands:
    - test -f $SP_DIR/cppe/pycppe*.so # [not win]
    - pytest --pyargs cppe

about:
  home: https://github.com/maxscheurer/cppe
  license: LGPL-3.0-or-later
  summary: "C++/Py library for polarizable embedding"

extra:
  recipe-maintainers:
    - robertodr
    - maxscheurer
```



build

passing

recipe/build.sh

```
cmake \
  -S "${SRC_DIR}" \
  -B build \
  -D CMAKE_INSTALL_PREFIX:PATH="${PREFIX}" \
  -D Python_EXECUTABLE:STRING="${PYTHON}" \
  -D PYMOD_INSTALL_FULLDIR:PATH="${SP_DIR}${PREFIX}/cppe"

cmake --build build --target install
```



build

passing

recipe/bld.bat

```
cmake ^
  -S "%SRC_DIR%" ^
  -B build ^
  -D CMAKE_INSTALL_PREFIX:PATH="%PREFIX%" ^
  -D CMAKE_CXX_COMPILER:STRING=clang-cl ^
  -D CMAKE_WINDOWS_EXPORT_ALL_SYMBOLS:BOOL=ON ^
  -D Python_EXECUTABLE:STRING="%PYTHON%" ^
  -D PYMOD_INSTALL_FULLDIR:PATH="Lib\site-packages\cppe"

if errorlevel 1 exit 1

cmake --build build --config Release --target install
if errorlevel 1 exit 1
```

CONDA-FORGE RECIPE: CPPE

CONDA-FORGE BUILD MATRIX & VERSION PINNINGS

[conda-forge-pinning-feedstock / recipe / conda_build_config.yaml](#)

Code Blame 983 lines (965 loc) · 16.3 KB · 

```

23   cxx_compiler:
24     - gxx           # [linux]
25     - clangxx      # [osx]
26     - vs2022       # [win]
27   cxx_compiler_version:
28     - 13           # [linux]
29     - 18           # [osx]
30     - 12           # [linux and ppc64le and ...
31     - 13           # [linux and not ppc64le]

...
189  blas_impl:
190    - openblas
191    - mkl          # [x86 or x86_64]
192    - blis          # [x86 or x86_64]

...
813  pybind11_abi:
814    - 4
815  python:
816    # part of a zip_keys: python, is_python_min
817    - 3.9.* *_cpython
818    - 3.10.* *_cpython
819    - 3.11.* *_cpython
820    - 3.12.* *_cpython
821  python_impl:
822    - cpython
...
980  zlib_ng:
981    - '2.2'
982  zstd:
983    - '1.5'

```

recipes can override as needed

a Py extension module

- A conda **RECIPE** contains project metadata and build directions. It's similar to a Homebrew formula (Linux/Mac) or a Fedora Spec file (Linux).
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- **BUILD** scripts build & install for Unix & Windows with many defined variables.
- Multiple **OUTPUTS** (e.g., C, Fortran, Py) possible from one build.
- Everything is done in **GITHUB CI**, so arch are often cross-compiled.

 build passing **recipe/build.sh**

```

cmake \
-S "${SRC_DIR}" \
-B build \
-D CMAKE_INSTALL_PREFIX:PATH="${PREFIX}" \
-D Python_EXECUTABLE:STRING="${PYTHON}" \
-D PYMOD_INSTALL_FULLDIR:PATH="${SP_DIR#${PREFIX}}/cppe"

cmake --build build --target install

```

 build passing **recipe/bld.bat**

```

cmake ^
-S "%SRC_DIR%" ^
-B build ^
-D CMAKE_INSTALL_PREFIX:PATH="%PREFIX%" ^
-D CMAKE_CXX_COMPILER:STRING=clang-cl ^
-D CMAKE_WINDOWS_EXPORT_ALL_SYMBOLS:BOOL=ON ^
-D Python_EXECUTABLE:STRING="%PYTHON%" ^
-D PYMOD_INSTALL_FULLDIR:PATH="Lib\site-packages\cppe"
if errorlevel 1 exit 1

cmake --build build --config Release --target install
if errorlevel 1 exit 1

```

CONDA-FORGE RECIPE: CPPE

anaconda.org/conda-forge/cppe

conda-forge / packages / cppe 0.3.2

C++ and Python library for Polarizable Embedding calculations

copied from [cf-staging / cppe](#)

Conda	Files	Labels	Badges
-------	-------	--------	--------

- 📄 License: [LGPL-3.0-or-later](#)
- 🏡 Home: <https://github.com/maxscheurer/cppe>
- </> Development: <https://github.com/maxscheurer/cppe>
- 📝 Documentation: <https://cppe.readthedocs.io>
- ⬇️ 242831 total downloads
- 📅 Last upload: 2 months and 18 days ago

Installers

🐧 linux-64	v0.3.2
🍎 osx-64	v0.3.2
osx-arm64	v0.3.2
linux-aarch64	v0.3.2
win-64	v0.3.2
linux-ppc64le	v0.3.2

conda install

To install this package run one of the following:

```
conda install conda-forge::cppe
```

a Py extension module

PACKAGES

Size	Name
2.9 MB	linux-aarch64/cppe-0.3.2-py39hebc598f_3.conda
3.0 MB	linux-aarch64/cppe-0.3.2-py312h279dd22_3.conda
3.0 MB	linux-aarch64/cppe-0.3.2-py311hef841b3_3.conda
3.0 MB	linux-aarch64/cppe-0.3.2-py310h37ef1f8_3.conda
2.9 MB	win-64/cppe-0.3.2-py310h9d296e7_3.conda
3.0 MB	win-64/cppe-0.3.2-py313hf4e7ab2_3.conda
3.0 MB	linux-ppc64le/cppe-0.3.2-py39h59d6d83_3.conda
2.9 MB	win-64/cppe-0.3.2-py312ha5776e6_3.conda
3.0 MB	linux-ppc64le/cppe-0.3.2-py311hcf87a8c_3.conda
3.0 MB	linux-ppc64le/cppe-0.3.2-py312hce74920_3.conda
3.0 MB	linux-ppc64le/cppe-0.3.2-py313h7ffc702_3.conda
3.0 MB	linux-ppc64le/cppe-0.3.2-py310hae48ae8_3.conda
3.0 MB	win-64/cppe-0.3.2-py39hbe34ff7_3.conda
3.0 MB	win-64/cppe-0.3.2-py311h038952a_3.conda
2.9 MB	osx-arm64/cppe-0.3.2-py310h1577cc6_3.conda
3.0 MB	osx-64/cppe-0.3.2-py39h1959004_3.conda
2.8 MB	osx-arm64/cppe-0.3.2-py312h382e28c_3.conda
3.0 MB	linux-aarch64/cppe-0.3.2-py313h2f34184_3.conda
3.0 MB	osx-64/cppe-0.3.2-py312h1701139_3.conda
2.9 MB	osx-arm64/cppe-0.3.2-py311h1b28dfe_3.conda
3.0 MB	osx-64/cppe-0.3.2-py310h7a5d817_3.conda
3.0 MB	osx-64/cppe-0.3.2-py313h1c70d2e_3.conda
2.9 MB	osx-arm64/cppe-0.3.2-py313h403398f_3.conda
2.9 MB	osx-arm64/cppe-0.3.2-py39hf608f95_3.conda
3.0 MB	osx-64/cppe-0.3.2-py311hcdf574e_3.conda

6 architectures x 5 pythons

CONDA-FORGE RECIPE: CPPE

anaconda.org/conda-forge/cppe

conda-forge / packages / cppe 0.3.2

C++ and Python library for Polarizable Embedding calculations

copied from cf-staging / cppe

Conda Files Labels Badges

License: LGPL-3.0-or-later
 Home: <https://github.com/maxscheurer/cppe>
 Development: <https://github.com/maxscheurer/cppe>
 Documentation: <https://cppe.readthedocs.io>
 242831 total downloads
 Last upload: 2 months and 18 days ago

Installers

linux-64 v0.3.2
 osx-64 v0.3.2
 osx-arm64 v0.3.2
 linux-aarch64 v0.3.2
 win-64 v0.3.2
 linux-ppc64le v0.3.2

conda install ?

To install this package run one of the following:

conda install conda-forge::cppe

a Py extension module

- From this setup, you get many built packages with indiv. & cumulative download **COUNTS**.
- With **30K** packages, your deps are probably already present and maintained.
- You can **SHARE** tasks with co-maintainers.
- You get prodded by **BOTS** to update for new Py, HDF5, etc. versions and for new tech: compilers, architectures, GPU.
- When you need a **BUGFIX FAST**, it can be fully published within 30m (+ build time)
- Enter into a **COMMUNITY** that will push along standards and best-practices and bring deficiencies to your attention.
- WHEN?** IMO, once you have the buildsys working on one arch, you start developing GHA CI for other arch. Once you have GHA CI for all arch and a small test, you start the packaging process.

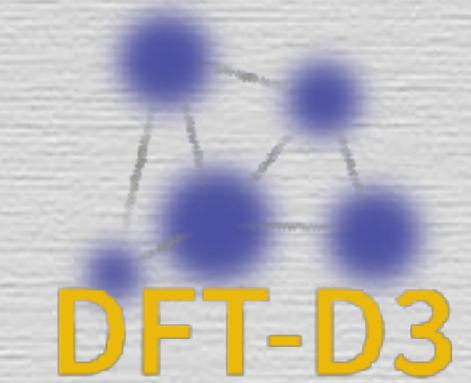
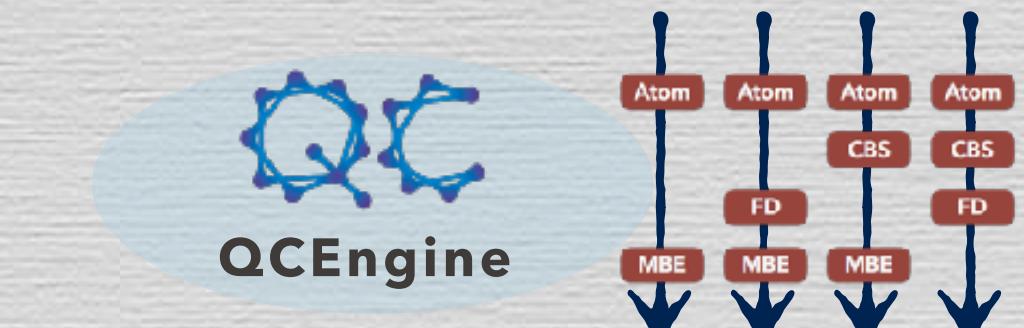
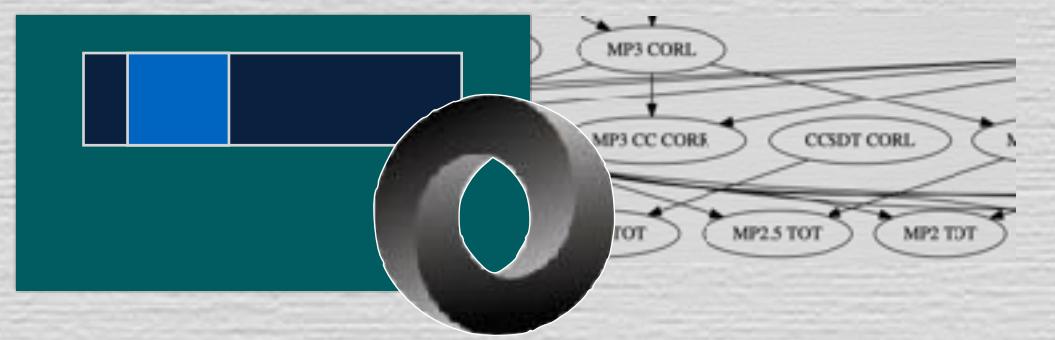
PR HISTORY

<input type="checkbox"/>	<input type="checkbox"/> ↗ Rebuild for python312
	#5 by regro-cf-autotick-bot was merged on Sep 28, 2023
<input type="checkbox"/>	<input type="checkbox"/> ↗ ARM OSX Migrator
	#4 by regro-cf-autotick-bot was merged on Jul 28, 2023
<input checked="" type="checkbox"/>	<input type="checkbox"/> ↗ Rebuild for PyPy3.8 and PyPy3.9
	#3 by regro-cf-autotick-bot was merged on Jul 29, 2023

bot prompts for version or c-f updates

★ Getting a production-quality binary to users for multiple architectures can be fairly easy with conda-forge.

★ Joining a packaging community can prod software engineering aspects.



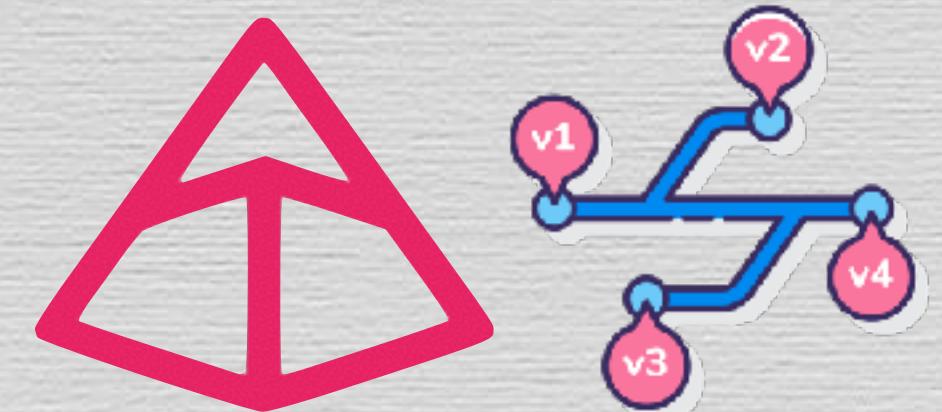
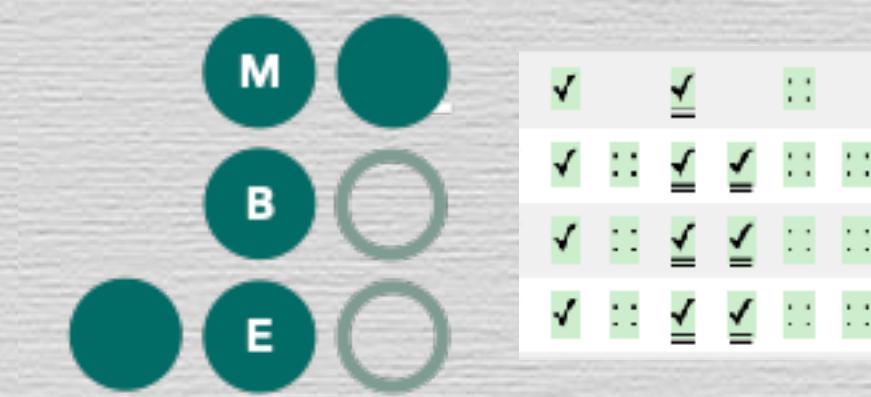
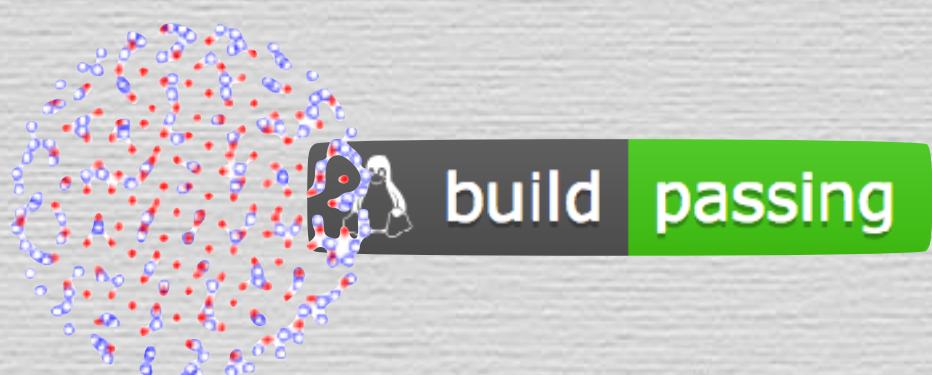
PSI4

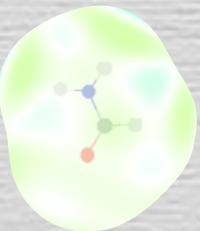
OPEN-SOURCE QUANTUM CHEMISTRY

DISTRIBUTION



GDMA





GDMA: DISTRIBUTED MULTIPOLE ANALYSIS

gitlab.com/anthonysjs/gdma

ANALYSIS technique & code publication ~2005

Journal of Chemical Theory and Computation > Vol 1/Issue 6 > Article
ARTICLE | September 13, 2005

Distributed Multipole Analysis: Stability for Large Basis Sets

Anthony J. Stone

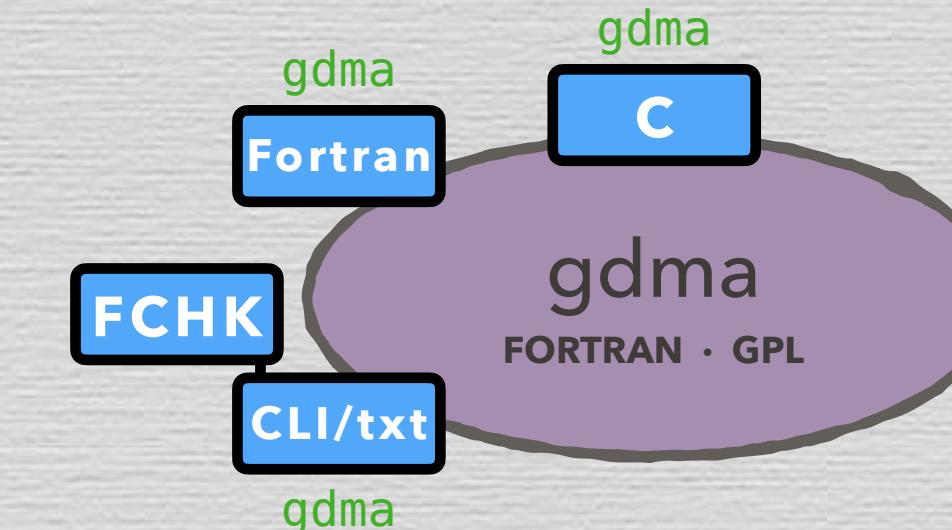
Anthony Stone: computer programs

- **GDMA 2.3.** GDMA is a Fortran program for performing Distributed Multipole Analysis of wavefunctions calculated using the [Gaussian program system](#). That is, it calculates electric multipole moments at the atomic positions, or at other specified sites, that can give a very accurate representation of the electrostatic field of the molecule. It uses the formatted checkpoint file produced by the Gaussian program. Version 2.3 handles checkpoint files from any version of Gaussian. It can also use checkpoint files produced by the Psi4 program system. It is also possible to use GDMA with Dalton or NWChem or Psi4 as part of the CamCASP package. The GDMA package also includes the MULFIT program of Ferenczy et al.



Anthony Stone
Cambridge

- **DEVELOPED** by A. J. Stone (Ret.)
- stand-alone Fortran **EXECUTABLE**
- **FCHK** interface
- GPL v3 **LICENSE**



CONVERSION to C library in 2016 for Psi4

andysim commented on Mar 24, 2016

Because the code just calls GDMA, which I turned into a function call instead of an executable, I think we should distribute the GDMA manual with the Psi4. I dumped the PDF into doc/ext/manual for now. Any objections?

andysim commented on Mar 25, 2016

I added full Sphinx documentation for both the FCHK writer and for GDMA, in commit [f2aca76](#), @CDSherrill. I heard back from Prof. Stone and he has kindly agreed to allow us to use GDMA in the way that we want to, so this PR is ready to go. In

C INTERFACE (+ I/O WORK)

gdma / src / dma.f90

Code Blame 3100 lines (2832 loc) · 118 KB

```

327
328 CONTAINS
329 !-
330 ! Some getter functions to access the DMA information from C
331 INTEGER(C_INT) FUNCTION get_nsites() BIND(c, name='get_nsites')
332     get_nsites = ns
333 END FUNCTION get_nsites
334
335 INTEGER(C_INT) FUNCTION get_order(site) BIND(c, name='get_order')
336     INTEGER(C_INT), VALUE, INTENT(IN) :: site
337     get_order = limit(site)
338 END FUNCTION get_order
339
340 REAL(C_DOUBLE) FUNCTION get_dma_value(site, addr) BIND(c, name='get_dma_value')
341     INTEGER(C_INT), VALUE, INTENT(IN) :: site, addr
342     get_dma_value = q(addr,site)
343 END FUNCTION get_dma_value
344
345 REAL(C_DOUBLE) FUNCTION get_tot_value(addr) BIND(c, name='get_tot_value')
346     INTEGER(C_INT), VALUE, INTENT(IN) :: addr
347     get_tot_value = qt(addr)
348 END FUNCTION get_tot_value
349 !-
350

```

DMA



Andy Simmonett
NIH → QC Ware



GDMA: DISTRIBUTED MULTIPOLE ANALYSIS

adapting established code to new use

noticed the **LICENSE** when updating to upstream

Update gdma to v2.3; convert to runtime #2968

Merged loriab merged 4 commits into psi4:master from loriab:pygdma on Aug 11, 2023

Conversation 4 Commits 4 Checks 0 Files changed 14

loriab commented on May 24, 2023 · edited

A perhaps obscure point is that psi hasn't been using exactly upstream GDMA, which is a Fortran executable. Instead, Andy Simmonett did lots of I/O conversion and turned it into a library and bound some functions to C so psi could extract results w/o parsing. That's great, but it's meant we've been fixed at upstream v2.2.06.

I was startled to realize gdma is GPL. For background, roughly speaking, GPL probably does adhere to psi4 if libgdma statically linked, might adhere if dynamically linked (FSF thinks yes; everyone else thinks no; result is that any GPL makes ppl nervous), doesn't adhere if dlopen'ed or runtime optional. Conda-package-wise, we've always dynamically linked to libgdma, and scouring my memory, there was a plan to have an alternate gdma- and chemps2-free conda build of psi4 to remove doubt; that never happened.

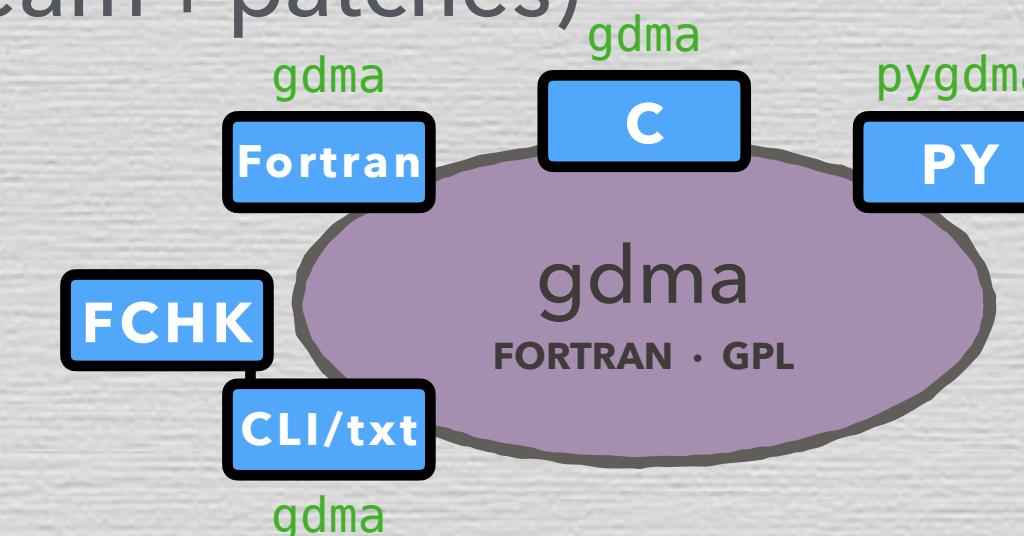
CONVERSION to Python module in 2023 for license

PYBIND11 INTERFACE

```
4     extern "C" int get_nsites();
5     extern "C" int get_order(int site);
6     extern "C" double get_dma_value(int site, int addr);
7     extern "C" double get_tot_value(int addr);
8
9     PYBIND11_MODULE(gdma, m) {
10
11     m.def("run_gdma", &run_gdma,
12           "Execute the GDMA analysis with input directives from the file named in the second argument (FCHK file");
13     m.def("get_nsites", &get_nsites, "Return count of sites at which multipoles are placed in DMA analysis.");
14     m.def("get_order", &get_order, "Return multipole order for site in first argument (1-indexed).");
15     m.def("get_dma_value", &get_dma_value, "Return the spherical harmonic DMA for the site in the first argument");
16     m.def("get_tot_value", &get_tot_value, "Return the total multipoles translated to the origin for component in");
17     m.attr("__version__") = "@pygdma_VERSION@";
18 }
```

- **PAYOFF** library accesses results w/o parsing & output merged with Psi4's
- **PAYOFF** module defers enabling to runtime, avoiding entanglement of linking Psi4 with GPL (c.f. CheMPS2)
- **PAYOFF** conda-forge packaging (upstream + patches)

- ★ adapt established modular code with new interfaces
- ★ beware choice of license can limit uptake
- ★ defer loading to runtime for licensing and modularity



DISTRIBUTION

<https://psicode.org/install/latest>

Get Started with Psi4

Select Preferences

LINUX LINUX ARM MAC MAC SILICON WINDOWS

INSTALLER CONDA DOCKER SOURCE

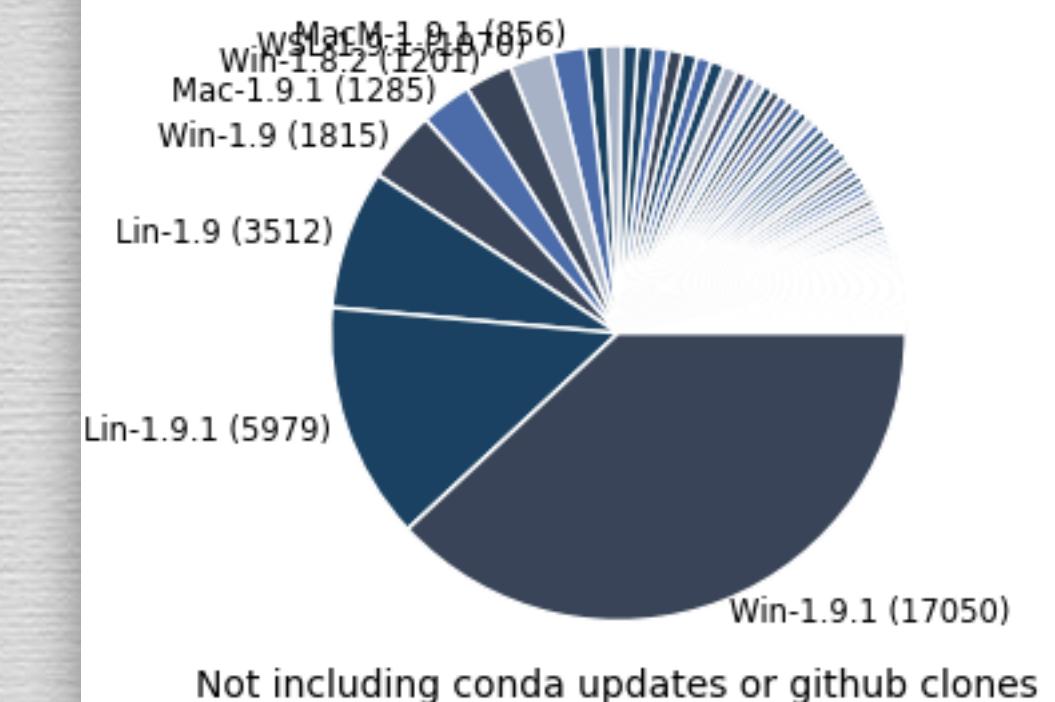
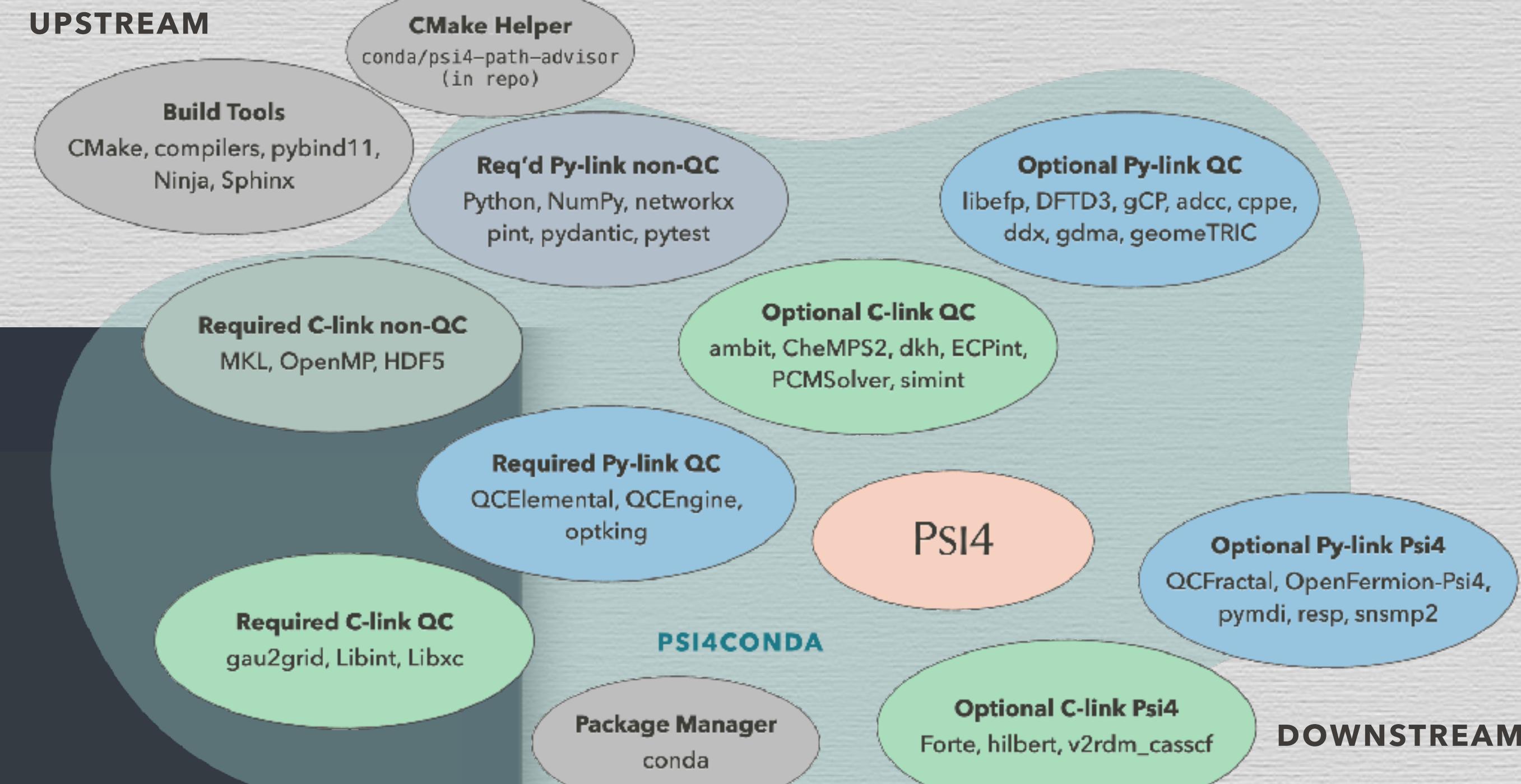
3.9 3.10 3.11 3.12 3.13

PREV RELEASE, v1.9.1 STABLE RELEASE, v1.10.0 NIGHTLY BUILD

Run this command

[DOWNLOAD PSI4CONDA INSTALLER \(ALT. CURL CMD BELOW\)](#)

```
# download via button above -OR- following line
> curl "https://vergil.chemistry.gatech.edu/psicode-download/Psi4conda-1.10-py312-Linux-x86_64.sh"
> bash Psi4conda-1.10-py312-Linux-x86_64.sh -b -p $HOME/psi4conda
bash > echo '$ . $HOME/psi4conda/etc/profile.d/conda.sh\nconda activate' >> ~/.bashrc
tcsh > echo "source $HOME/psi4conda/etc/profile.d/conda.csh\nconda activate" >> ~/.tcshrc
# log out, log back in so conda and psi4 in path
> psi4 --test
```



**PY-WARY USERS:
CONDA INSTALLER**
Psi4 · dependencies · add-ons

PSI4CONDA 633K
downloads

DISTRIBUTION

<https://psicode.org/install/latest>

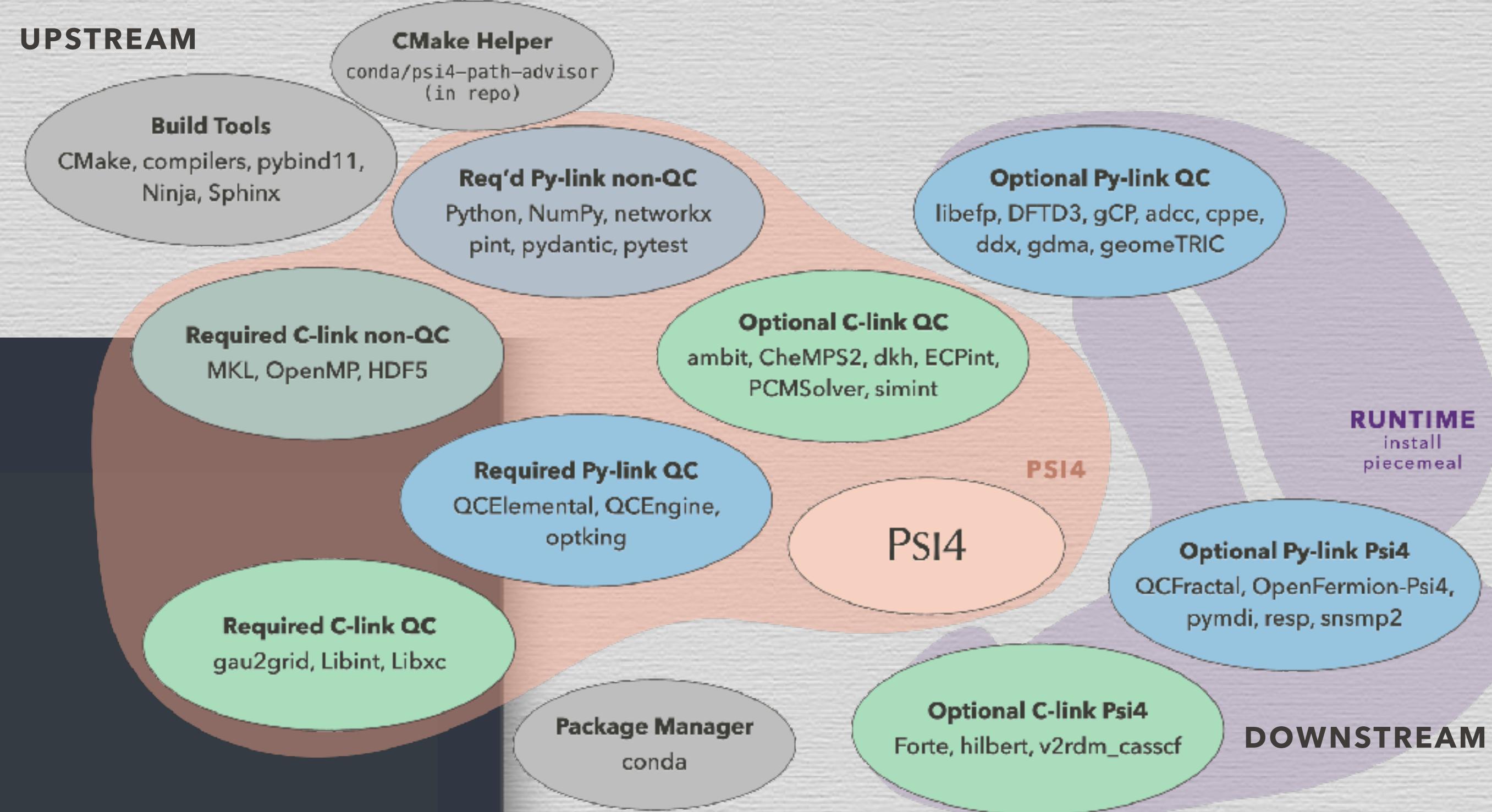
Get Started with PSI4

Select Preferences

LINUX LINUX ARM MAC MAC SILICON WINDOWS
INSTALLER CONDA DOCKER SOURCE
3.9 3.10 3.11 3.12 3.13
PREV RELEASE, v1.9.1 STABLE RELEASE, v1.10.0 NIGHTLY BUILD
GOTO MINICONDA INSTALLERS

Run this command

```
>_ conda install psi4 python=3.12 -c conda-forge
# -OR- create environment ("p4dev" name is arbitrary)
>_ conda create -n p4dev psi4 python=3.12 -c conda-forge && conda activate p4dev
```



**PY-FRIENDLY USERS:
CONDA PACKAGE**
PSI4 · dependencies · add-ons

**PY-WARY USERS:
CONDA INSTALLER**
PSI4 · dependencies · add-ons

628K
downloads

PSI4CONDA 633K
downloads

DISTRIBUTION

<https://psicode.org/install/latest>

Get Started with Psi4

Select Preferences

LINUX LINUX ARM MAC MAC SILICON WINDOWS

INSTALLER CONDA DOCKER SOURCE

3.9 3.10 3.11 3.12 3.13

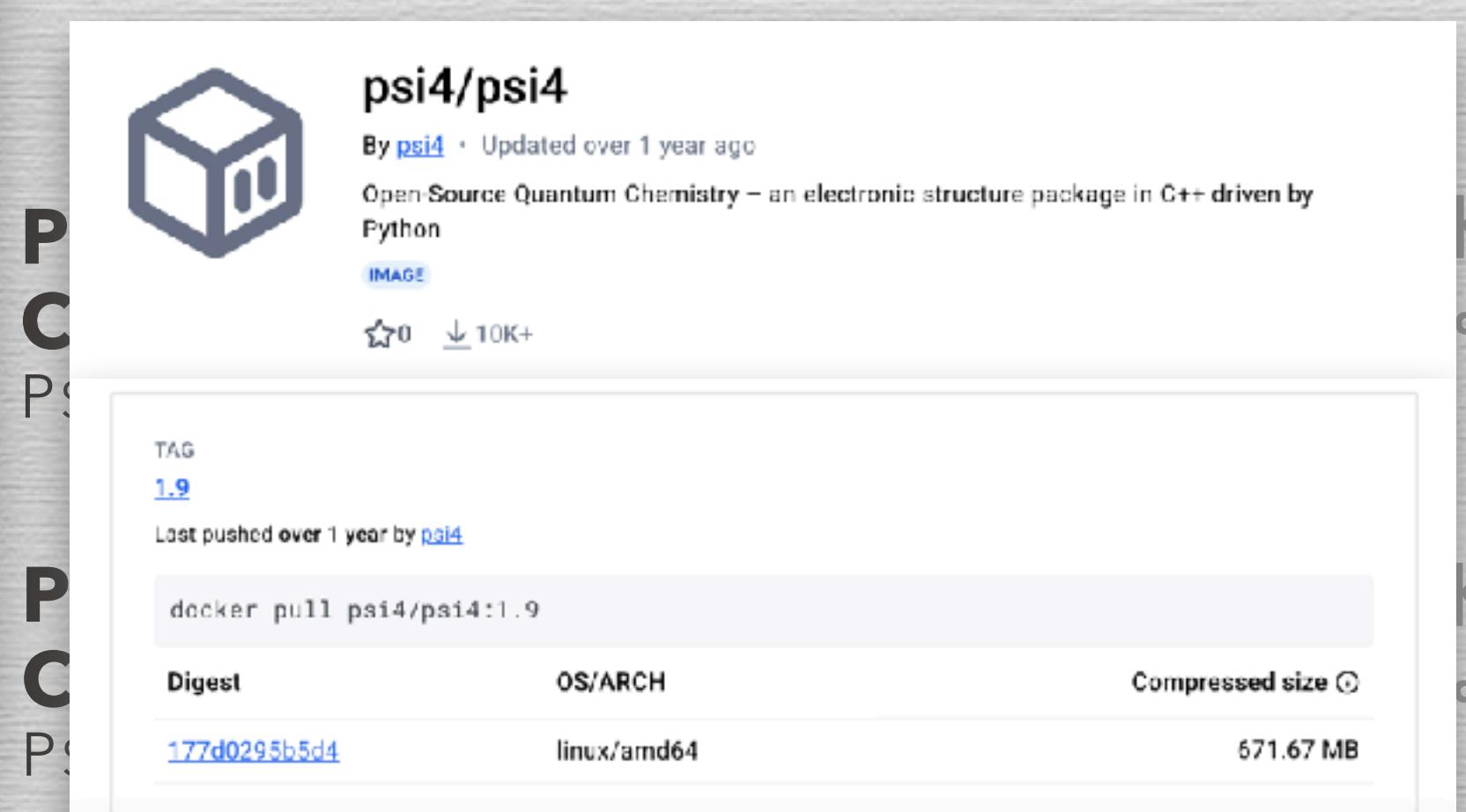
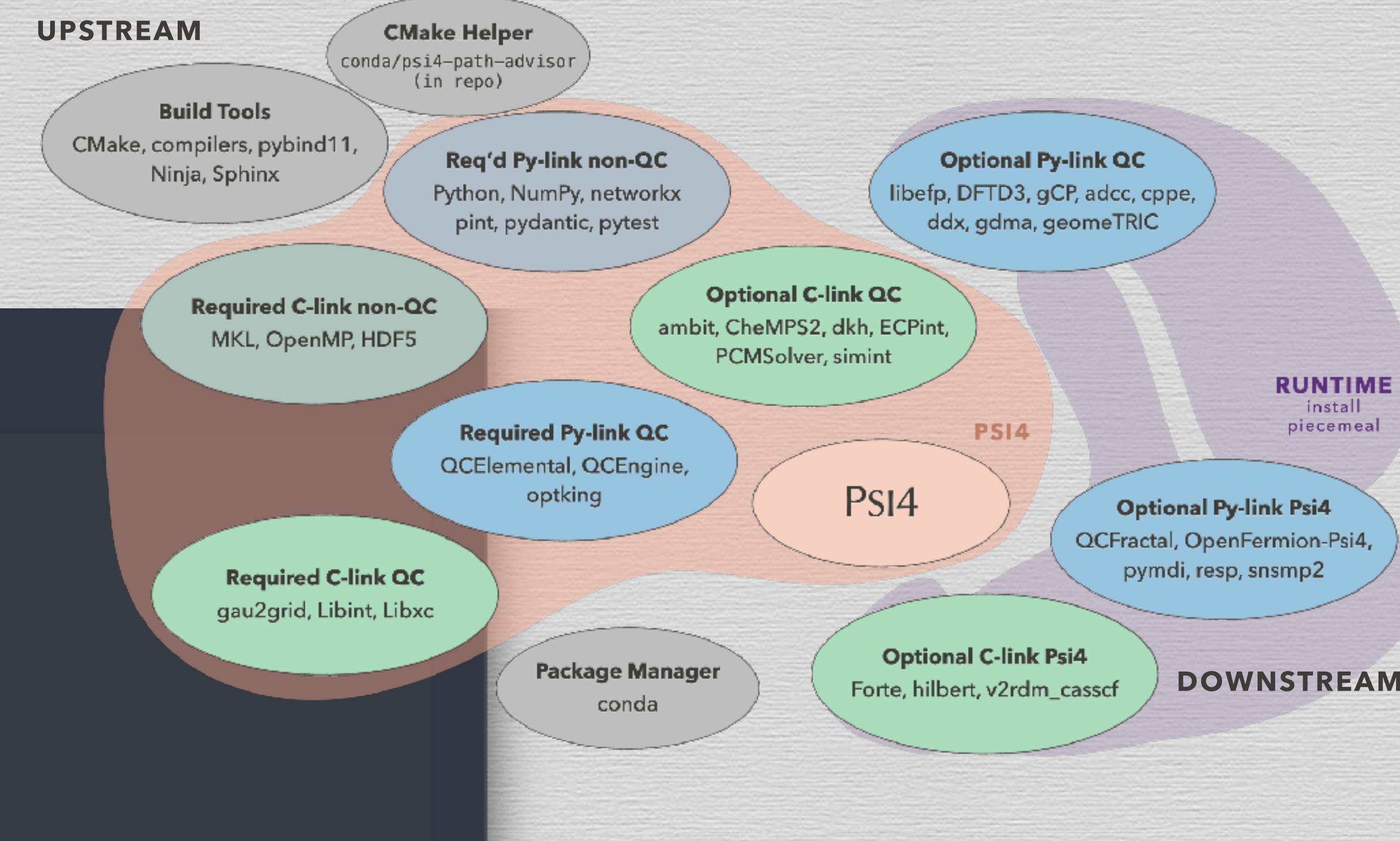
PREV RELEASE, v1.9.1 STABLE RELEASE, v1.10.0 NIGHTLY BUILD

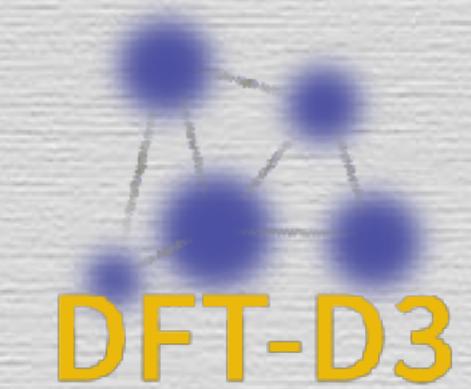
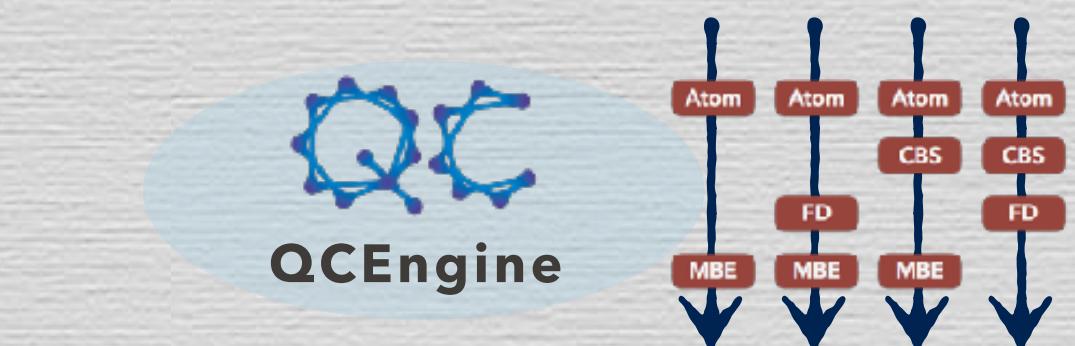
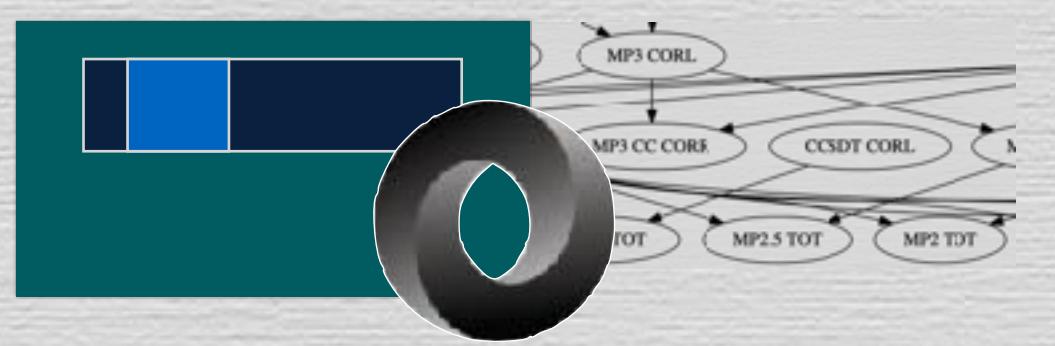
Run this command

GOTO MINICONDA INSTALLERS

```
>_ docker pull psi4/psi4:1.10.0
```

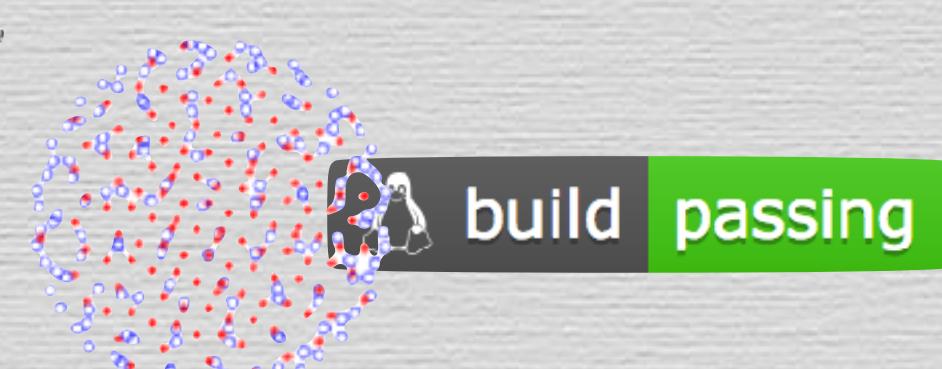
★ Provide a variety of installation routes & consolidate the choices.



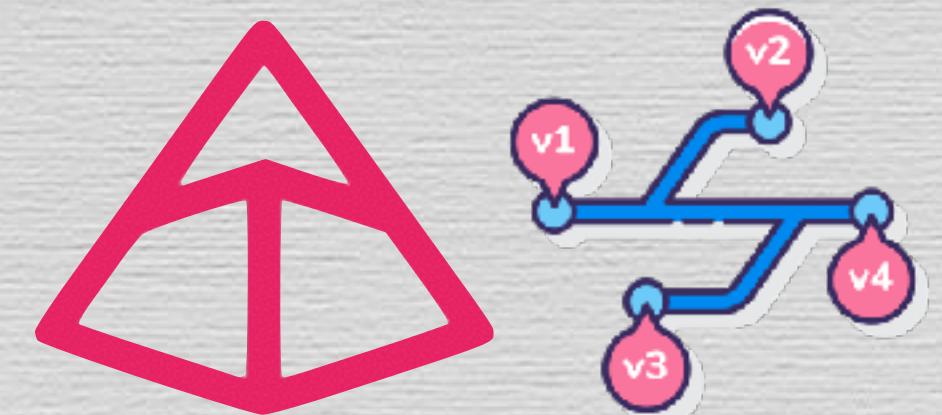
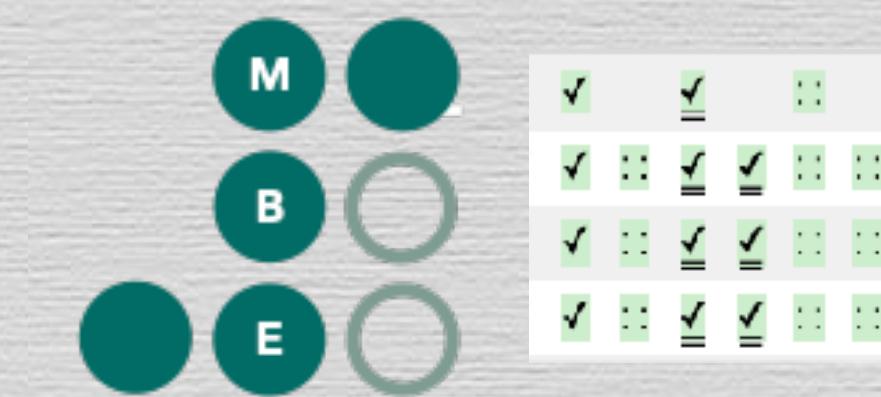


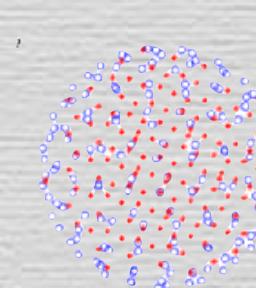
PSI4

OPEN-SOURCE QUANTUM CHEMISTRY



GAU2GRID





GAU2GRID: COLLOCATION SPECIALIST

github.com/dgasmith/gau2grid (now psi4/gau2grid)

Identified **BOTTLENECK** in internal code

- **COLLOCATION** matrix btwn basis functions & DFT grid inefficient in Psi4 as simple non-vectorizable loops so extracted ops.

$$\phi_{mp} = Y_\ell^m(\mathbf{r}_p - \widehat{\mathbf{r}_{\text{center}}}) \sum_i c_i e^{-\alpha_i (\mathbf{r}_{\text{center}} - \mathbf{r}_p)^2}$$

- Daniel Smith, Rob Parrish, Andy Simmonett for Psi4 **v1.2**
- Python generates a standalone **C LIBRARY** of unrolled loops. Algorithm adjusted for AM, deriv, hardware at gen-time
- **ARBITRARY** (specified) AM, up to 3rd derivatives
- C and Python (ref. impl.) **APIs**, no dependencies
- **RUNTIME** ordering switching w/o loss of efficiency (v2)
- Much easier to share **MAINTENANCE** on GitHub orgs
- **PAYOUT** enhanced efficiency for Psi4 DFT

dgasmith commented on Dec 5, 2017

Switched over Psi4 grid generation to `libgg`. Looks like about a 4x speedup in general for collocation generation. However, it spends ~65% of the time transposing the collocation matrices back to Psi4 order otherwise this would be killer (~12x faster). I can look at a few other transpose options in `libgg`, but it would be tough to spit them out correctly. Still helps dramatically for small molecules/bases and large grids.

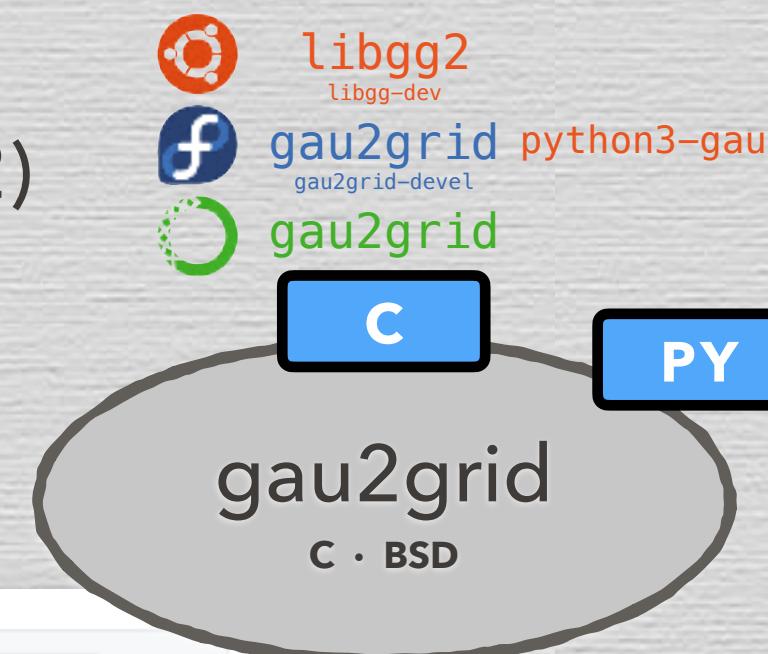
Wrote new **C LIBRARY** in 2017 for Psi4

```
#include "gau2grid/gau2grid.h"

#if psi4_SHGSHELL_ORDERING == LIBINT_SHGSHELL_ORDERING_STANDARD
const int order = (int)puream_ ? GG_SPHERICAL_CCA : GG_CARTESIAN_CCA;
#elif psi4_SHGSHELL_ORDERING == LIBINT_SHGSHELL_ORDERING_GAUSSIAN
const int order = (int)puream_ ? GG_SPHERICAL_GAUSSIAN : GG_CARTESIAN_CCA;
#endif

// Compute collocation
if (deriv_ == 0) {
    gg_collocation(L, npoints, xyz.data(), 1, nprim, norm, alpha,
                  center.data(), order, phi_start);
} else if (deriv_ == 1) {
    gg_collocation_deriv1(L, npoints, xyz.data(), 1, nprim, norm, alpha,
                          center.data(), order, phi_start,
                          phi_x_start, phi_y_start, phi_z_start);
}
```

Using Gau2grid in Psi4. Note the flexible ordering.



- ★ extract code into a module to concentrate performance optimizations
- ★ ship a Py reference implementation and/or a Py MWE to aid downstream adoption
- ★ defer ordering configuration to runtime & build broadly for consumers

CENTRALIZING ECOSYSTEM PROJECTS

SSOT

- Packaging – providing artifacts compiled under fixed conditions – is easier than providing a flexible buildsys plus enough **GUIDANCE** for users to cope with an ecosystem build.
- Build guidance shouldn't be overly **PREScriptive** and add another constraint to a developer's chosen workflow.
- After much floundering, I'm pleased with a single-source-of-truth file, **codedeps.yaml**, plus a mediating script **psi4-path-advisor**.
- **CODEDEPS** accumulates in YAML format all information on all the projects for building, augmenting, testing, documenting, or interfacing to Psi4. It runs to 1500 lines.

codedeps.yaml

1500 LOC

```
python -c "import yaml, pathlib; ydict = yaml.safe_load(pathlib.Path('..')
codedeps.yaml').read_text());print([itm['project'] for itm in ydict['data']])
#> ['c', 'cxx', 'fortran', 'python', 'cmake', 'generator', 'cfour', 'chemps2', 'dftd3',
'dkh', 'erd', 'gdma', 'lapack', 'libint', 'libefp', 'mrcc', 'openmp', 'pcmsolver',
'v2rdm_casscf', 'doxygen', 'python-graphviz', 'sphinx', 'sphinx-psi-theme', 'gcp',
'pybind11', 'pytest', 'nbsphinx', 'numpy', 'simint', 'ambit', 'gau2grid',
'memory_profiler', 'libxc', 'openfermionpsi4', 'qcelemental', 'adcc', 'brianqc',
'cppe', 'geometric', 'i-pi', 'mdi', 'mp2d', 'qcengine', 'dftd4', 'ecpint', 'qcfractal',
'scipy', 'ddx', 'optking', 'bse', 'einsums', 'integratorxx', 'gauxc', 'h5py']
```

CENTRALIZING ECOSYSTEM PROJECTS

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- Packaging – providing artifacts compiled under fixed conditions – is easier than providing a flexible buildsys plus enough **GUIDANCE** for users to cope with an ecosystem build.
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- Two years ago it was experimental because I don't like another invented standard (does have a basic **SCHEMA**), but I'm increasingly pleased with the scheme.

codedeps.yaml

1500 LOC

```
# - project: !!str          # Field required. Project <dummy> is replaced by the Psi4 internal label used for CMake (e.g., TODO FALSE ENABLE_<dummy>, external/upstream/<dummy>/CMakeLists.txt)
#   use: !!map              # Section required. Important Psi4 dependency usage information.
#     added: !!str          # Field required. Minor version at which <dummy> added to Psi4 ecosystem. v1.0 is earliest value.
#     deprecated: !!str      # Field required. Minor version at which <dummy> removed from Psi4 ecosystem. Further fields not present.
#     required: !!bool        # Field required. Whether <dummy> is a required Psi4 dependency.
#     buildable: !!bool       # Field required. If <dummy> is required or enabled, `true` if the Psi4 build system can supply it by building from source or `false` if it must be built upstream.
#     buildtime: !!bool        # Field required. Whether <dummy> is compile-linkage (`true`) or runtime-linkage (`false`); some runtime are marked `true` if required to detect at runtime.
#     cms: !!bool             # Field required. Whether <dummy> is a computational molecular sciences project vs. generic compsci.
#     test_required: !!bool    # Field required if used in Psi4 testing. Whether <dummy> is a required dependency for testing.
#     docs_required: !!bool   # Field required if used in building Psi4 documentation. Whether <dummy> is a required dependency for docs build.
#   repository: !!map         # Section required if use/buildable=true. Where to acquire source code to build.
#     host: !!str              # Field required. (github | gitlab | url)
#     url: !!str              # Field required if host=url. Full url to tarball for source.
#     account: !!str           # Field required if host!=url. Personal or organization account name on host.
#     name: !!str or !!set      # Field required if host!=url. Project name on host/account. Use list for optional dependencies.
#     commit: !!str             # Field required if host!=url. The tag, branch, or hash of the repository at which to pull source. Usually latest compatible release, not earliest commit.
#   cmake: !!map                # Section required if use/buildable=true. How/what to detect <dummy> with `find_package(<cmake/name> <cmake/constraint> COMPONENTS <cmake/components>)`.
#     name: !!str               # Field required. CMake project name in `find_package` call. Often has capital letters.
#     constraint: !!str          # Field required. Version of earliest compatible release. May also include ATLEAST or EXACT. Remember CMake has only crude spec for version constraints.
#     target: !!str              # Field required. Target <cmake/name>::<cmake/target> used by Psi4. Field may be `null` if no target, like for CMake-built python package.
#     components: !!set          # Field required. Required components for pre-built to be accepted. Field may be `null` if no components.
#   conda: !!map                # Section required if use/required=true.
#     channel: !!str             # Field required. Preferred channel on anaconda.org suitable for `-c`. Use the broadest if conda/name on multiple channels or subchannels.
#     name: !!str or !!map        # Field required. The conda package name. Use minimal required package (e.g., libxc-c (C intf) rather than libxc (C & Fortran intf)). Occasionally mismatched.
#     constraint: !!str or !!map  # Field required. Version or build number restrictions on package. Field should be `null` if most recent settings will do or needed to solve environment conflicts.
#     aux_run_names: !!set        # Field optional. Use to list add'l packages needed to run dummy and Psi4 together that aren't deps of dummy (e.g., postgresql for qcfractal).
#     aux_build_names: !!set      # Field optional. Use to list add'l packages needed to build Psi4 against dummy that aren't deps of dummy (e.g., mkl-devel for mkl).
#     cmake: !!map                # Field required. Use to specify CMake variables relevant to this package. Those starting with // aren't usually set. May involve substitutions.
#     skip_win: !!bool            # Field optional. Use to list packages not adapted for Windows. Perhaps should be in 'use' block?
```

IMPROVISED CODEDEPS SCHEMA

CENTRALIZING ECOSYSTEM PROJECTS

SSOT

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- Build guidance shouldn't be overly **PREScriptive** and add another constraint to a developer's chosen workflow.
- After much floundering, I'm pleased with a single-source-of-truth file, **codedeps.yaml**, plus a mediating script **psi4-path-advisor**.
- **CODEDEPS** accumulates in YAML format all information on all the projects for building, augmenting, testing, documenting, or interfacing to PSI4. It runs to 1500 lines.
- Two years ago it was experimental because I don't like another invented standard (does have a basic **SCHEMA**), but I'm increasingly pleased with the scheme.
- Main **GOALS**:
 - Users need to know if an addon is **INTERFACED** to PSI4
 - Packagers need to know what **DEPS** are new/retired/updated
 - Developers need to know how to **BUILD** with mixed binary and from-source dependencies
 - I need to not **SYNC** version/deps changes among multiple CMake, 2 Azure, 3 GHA, docs, and env spec files.
 - I need to know if the ecosystem that worked yesterday **BROKE** overnight.

codedeps.yaml

1500 LOC

CODEDEPS.YAML

TOOL

```

- project: cxx
  use:
    added: "1.0"

  required: true
  required_note: "Must support c++20 standard."
  buildable: false
  buildtime: true
    cms: false
  repository: null

cmake: null

conda:
  channel: conda-forge
  name: cxx-compiler
  constraint: null
  aux_build_names:
    - //dpcpp_linux-64
cmake:
  GNU_linux_64:
    CMAKE_CXX_COMPILER:
      "${CONDA_PREFIX}/bin/${HOST}-g++"
    //CMAKE_CXX_FLAGS: null
  MSVC_win-64:
    CMAKE_CXX_COMPILER: "clang-cl"
    //CMAKE_CXX_FLAGS: null
  IntellLVM_linux-64:
    CMAKE_CXX_COMPILER:
      "\\"${CONDA_PREFIX}/bin/icpx;
      --target=${HOST};
      --gcc-toolchain=${CONDA_PREFIX};
      --sysroot=${CONDA_PREFIX}/.${HOST}/sysroot\\""
    //CMAKE_CXX_FLAGS: null

```

OPT'L COMPILED

```

- project: gdma
  use:
    added: "1.0"
    added_note: "Lib to pygdma at 1.9. Ext at 1.1."
    required: false
    required_note: "Allow Stone multipole analysis"
    buildable: true
    buildtime: false
      cms: true
  repository:
    host: github
    account: psi4
    name: gdma
    commit: v2.3.3
    commit_note: "GL upstream c2e0b5 + lib, I/O, Py"
  cmake:
    name: gdma
    constraint: 2.3.3
    components:
      - Python
      target: gdma::pygdma
      enable: ENABLE_gdma
  conda:
    channel: conda-forge
    name: pygdma
    constraint: null
  cmake:
    ENABLE_gdma: true
    gdma_DIR:
      unix: ${CONDA_PREFIX}/share/cmake/gdma
      win-64: ${CONDA_PREFIX}/Library/share/cmake/gdma
    //CMAKE_INSIST_FIND_PACKAGE_gdma: true
    //CMAKE_DISABLE_FIND_PACKAGE_gdma: true
  cmake_note: "Primarily OTF runtime detected.
    CMake enabling only relevant for CTest."

```

REQ'D PY

```

- project: qcelemental
  use:
    added: 1.3

  required: true
  buildable: true
  buildtime: true
  buildtime_note: "BT for cmake checks; use is RT"
  cms: true
  repository:
    host: github
    account: MolSSI
    name: QCElemental
    commit: v0.29.0

  cmake:
    name: qcelemental
    constraint: "ATLEAST 0.28.0"

  target: null

  conda:
    channel: conda-forge
    name: qcelemental
    constraint: null
    aux_run_names:
      - msgpack-python
      - networkx
      - setuptools
      - pydantic>=1.10.17
    aux_run_names_note:
      "setuptools": "Needed for cmake detection"
      "pydantic>=1.10.17": "Uniform Pydantic v1 API"
  cmake:
    //CMAKE_INSIST_FIND_PACKAGE_qcelemental: true
    //CMAKE_DISABLE_FIND_PACKAGE_qcelemental: true
  cmake_note: "Primarily OTF runtime detected."

```

- **USE:** info like when added to PSI4, req'd/opt'l, and buildable, along with formal notes.

CODEDEPS.YAML

```

- project: cxx
  use:
    added: "1.0"

  required: true
  required_note: "Must support c++20 standard."
  buildable: false
  buildtime: true

  cms: false
  repository: null

cmake: null

conda:
  channel: conda-forge
  name: cxx-compiler
  constraint: null
  aux_build_names:
    - //dpcpp_linux-64
  cmake:
    GNU_linux_64:
      CMAKE_CXX_COMPILER:
        "${CONDA_PREFIX}/bin/${HOST}-g++"
      //CMAKE_CXX_FLAGS: null
    MSVC_win-64:
      CMAKE_CXX_COMPILER: "clang-cl"
      //CMAKE_CXX_FLAGS: null
    IntellLVM_linux-64:
      CMAKE_CXX_COMPILER:
        "\`${CONDA_PREFIX}/bin/icpx;
        --target=${HOST};
        --gcc-toolchain=${CONDA_PREFIX};
        --sysroot=${CONDA_PREFIX}/${HOST}/sysroot\`"
      //CMAKE_CXX_FLAGS: null

```

TOOL

```

- project: gdma
  use:
    added: "1.0"
    added_note: "Lib to pygdma at 1.9. Ext at 1.1."
    required: false
    required_note: "Allow Stone multipole analysis"
    buildable: true
    buildtime: false

    cms: true
    repository:
      host: github
      account: psi4
      name: gdma
      commit: v2.3.3
      commit_note: "GL upstream c2e0b5 + lib, I/O, Py"
    cmake:
      name: gdma
      constraint: 2.3.3
      components:
        - Python
        target: gdma::pygdma
        enable: ENABLE_gdma
    conda:
      channel: conda-forge
      name: pygdma
      constraint: null
    cmake:
      ENABLE_gdma: true
      gdma_DIR:
        unix: ${CONDA_PREFIX}/share/cmake/gdma
        win-64: ${CONDA_PREFIX}/Library/share/cmake/gdma
      //CMAKE_INSIST_FIND_PACKAGE_gdma: true
      //CMAKE_DISABLE_FIND_PACKAGE_gdma: true
    cmake_note: "Primarily OTF runtime detected.
      CMake enabling only relevant for CTest."

```

OPT'L COMPILED

```

- project: qcelemental
  use:
    added: 1.3

  required: true
  buildable: true
  buildtime: true
  buildtime_note: "BT for cmake checks; use is RT"
  cms: true
  repository:
    host: github
    account: MolSSI
    name: QCElemental
    commit: v0.29.0

  cmake:
    name: qcelemental
    constraint: "ATLEAST 0.28.0"

  target: null

  conda:
    channel: conda-forge
    name: qcelemental
    constraint: null
    aux_run_names:
      - msgpack-python
      - networkx
      - setuptools
      - pydantic>=1.10.17
    aux_run_names_note:
      "setuptools": "Needed for cmake detection"
      "pydantic>=1.10.17": "Uniform Pydantic v1 API"
    cmake:
      //CMAKE_INSIST_FIND_PACKAGE_qcelemental: true
      //CMAKE_DISABLE_FIND_PACKAGE_qcelemental: true
    cmake_note: "Primarily OTF runtime detected."

```

REQ'D PY

- **USE:** info like when added to PSI4, req'd/opt'l, and buildable, along with formal notes.
- **REPOSITORY:** info to build from source: repository and find_package call.

CODEDEPS.YAML

```

- project: cxx
  use:
    added: "1.0"

  required: true
  required_note: "Must support c++20 standard."
  buildable: false
  buildtime: true

  cms: false
  repository: null

cmake: null

conda:
  channel: conda-forge
  name: cxx-compiler
  constraint: null
  aux_build_names:
    - //dpcpp_linux-64
  cmake:
    GNU_linux_64:
      CMAKE_CXX_COMPILER:
        "${CONDA_PREFIX}/bin/${HOST}-g++"
      //CMAKE_CXX_FLAGS: null
    MSVC_win-64:
      CMAKE_CXX_COMPILER: "clang-cl"
      //CMAKE_CXX_FLAGS: null
    IntellLLVM_linux-64:
      CMAKE_CXX_COMPILER:
        "\`${CONDA_PREFIX}/bin/icpx;
        --target=${HOST};
        --gcc-toolchain=${CONDA_PREFIX};
        --sysroot=${CONDA_PREFIX}/${HOST}/sysroot\`"
      //CMAKE_CXX_FLAGS: null

```

TOOL

```

- project: gdma
  use:
    added: "1.0"
    added_note: "Lib to pygdma at 1.9. Ext at 1.1."
    required: false
    required_note: "Allow Stone multipole analysis"
    buildable: true
    buildtime: false

    cms: true
    repository:
      host: github
      account: psi4
      name: gdma
      commit: v2.3.3
      commit_note: "GL upstream c2e0b5 + lib, I/O, Py"
    cmake:
      name: gdma
      constraint: 2.3.3
      components:
        - Python
      target: gdma::pygdma
      enable: ENABLE_gdma
  conda:
    channel: conda-forge
    name: pygdma
    constraint: null
  cmake:
    ENABLE_gdma: true
    gdma_DIR:
      unix: ${CONDA_PREFIX}/share/cmake/gdma
      win-64: ${CONDA_PREFIX}/Library/share/cmake/gdma
    //CMAKE_INSIST_FIND_PACKAGE_gdma: true
    //CMAKE_DISABLE_FIND_PACKAGE_gdma: true
  cmake_note: "Primarily OTF runtime detected.
              CMake enabling only relevant for CTest."

```

OPT'L COMPILED

```

- project: qcelemental
  use:
    added: 1.3

  required: true
  buildable: true
  buildtime: true
  buildtime_note: "BT for cmake checks; use is RT"
  cms: true
  repository:
    host: github
    account: MolSSI
    name: QCElemental
    commit: v0.29.0

  cmake:
    name: qcelemental
    constraint: "ATLEAST 0.28.0"

  target: null

  conda:
    channel: conda-forge
    name: qcelemental
    constraint: null
    aux_run_names:
      - msgpack-python
      - networkx
      - setuptools
      - pydantic>=1.10.17
    aux_run_names_note:
      "setuptools": "Needed for cmake detection"
      "pydantic>=1.10.17": "Uniform Pydantic v1 API"
    cmake:
      //CMAKE_INSIST_FIND_PACKAGE_qcelemental: true
      //CMAKE_DISABLE_FIND_PACKAGE_qcelemental: true
    cmake_note: "Primarily OTF runtime detected."

```

REQ'D PY

- **USE:** info like when added to PSI4, req'd/opt'l, and buildable, along with formal notes.
- **REPOSITORY:** info to build from source: repository and find_package call.
- **CONDA:** info to use pre-built, including any constraints and additional BT dependencies.

CODEDEPS → CONDA

GENERATED env_p4dev.yaml

```

name: p4dev
channels:
- conda-forge
- nodefaults
dependencies:
# build
- c-compiler
- cmake
- cxx-compiler
#- dpcpp_linux-64          # opt'l with cxx-compiler
- ninja
# non-qc buildtime required
- blas-devel                # req'd with libblas
- eigen                      # req'd with libint
- libblas==*=mkl             # req'd with libint
- libboost-headers
- llvm-openmp
- numpy
- pip
- pybind11
- python
# qc buildtime required
- gau2grid
#- psi4::libint            # opt'l with libint; Optionally for linux-64, an AM=7 build here.
- libint
- libxc-c
- optking
- qcelemental
- qcengine
# buildtime optional
- dkh
- einsums>=1.0.3
- fortran-compiler
- hdf5
- integratorxx
- libecpint
- pcmsolver
- range-v3
- zlib
# runtime required
- msgpack-python
- networkx
- pydantic>=1.10.17
- scipy
- setuptools
# runtime optional
- adcc
- basis_set_exchange
- cppe
- dftd3-python>=1.0
- dftd4-python
- gcp-correction
- geometric
- i-pi==3.1.0
- postgresql
- pyddx
- pygmda
- pylibefp
- pymdi
- qcfractal>=0.60
- toml
# test
- pytest
- pytest-xdist
# req'd with qcfractal; Easy DB or supply your own.
# req'd with dftd4-python
# req'd with pytest; Parallel runner handy, !req'd.

```

psi4-path-advisor.py env

analyze codedeps and generate a Conda environment specification file w/ build reqs & optional full eco

```

conda/psi4-path-advisor.py env --help
#> usage: psi4-path-advisor env
#>   [-h] [-n NAME] [--python PYTHON]
#>   [--lapack {mkl,openblas,accelerate,blis,netlib}]
#>   [--disable {compilers,lapack,addons,test,docs}]
#>   [--platform {linux-64,osx-64,osx-arm64,win-64}]
#>   [--offline-conda]

```

REQ'D PY

```

- project: qcelemental
  use:
    added: 1.3
    required: true
    buildable: true
    buildtime: true
    buildtime_note: "BT for cmake checks; use is RT"
    cms: true
  repository:
    host: github
    account: MolSSI
    name: QCElemental
    commit: v0.29.0
  cmake:
    name: qcelemental
    constraint: "ATLEAST 0.28.0"
  target: null
  conda:
    channel: conda-forge
    name: qcelemental
    constraint: null
    aux_run_names:
      - msgpack-python
      - networkx
      - setuptools
      - pydantic>=1.10.17
    aux_run_names_note:
      "setuptools": "Needed for cmake detection"
      "pydantic>=1.10.17": "Uniform Pydantic v1 API"
    cmake:
      //CMAKE_INSIST_FIND_PACKAGE_qcelemental: true
      //CMAKE_DISABLE_FIND_PACKAGE_qcelemental: true
    cmake_note: "Primarily OTF runtime detected."

```

CODEDEPS → CMAKE

```
- project: cxx
use:
  added: "1.0"

required: true
required_note: "Must support c++20 standard."
buildable: false
buildtime: true

cms: false
repository: null
```

TOOL

psi4-path-advisor.py cache

analyze current conda env and generate likely & optional cache variables to seed CMake configuration

```
conda:
channel: conda-forge
name: cxx-compiler
constraint: null
aux_build_names:
- //dpcpp_linux-64
cmake:
  GNU_linux_64:
    CMAKE_CXX_COMPILER:
      "${CONDA_PREFIX}/bin/${HOST}-g++"
    //CMAKE_CXX_FLAGS: null
  MSVC_win-64:
    CMAKE_CXX_COMPILER: "clang-cl"
    //CMAKE_CXX_FLAGS: null
  IntellLVM_linux-64:
    CMAKE_CXX_COMPILER:
      "\\"${CONDA_PREFIX}/bin/icpx;
      --target=${HOST};
      --gcc-toolchain=${CONDA_PREFIX};
      --sysroot=${CONDA_PREFIX}/${HOST}/sysroot\""
    //CMAKE_CXX_FLAGS: null
```

```
# <<< cxx-compiler
set(CMAKE_CXX_COMPILER /home/miniconda/envs/p4dev/bin/x86_64-conda-linux-gnu-g++ CACHE STRING "")
# set(CMAKE_CXX_FLAGS <placeholder> CACHE STRING "")

# <<< pygdma
# Primarily OTF runtime detected. With package present, CMake enabling only relevant for CTest.
set(ENABLE_gdma ON CACHE BOOL "")
set(gdma_DIR /home/miniconda/envs/p4dev/share/cmake/gdma CACHE PATH "")
# set(CMAKE_INSIST_FIND_PACKAGE_gdma ON CACHE BOOL "")
# set(CMAKE_DISABLE_FIND_PACKAGE_gdma ON CACHE BOOL "")

# <<< libblas
set(LAPACK_LIBRARIES blas-devel
set(LAPACK_INCLUDE_DIRS /home/miniconda/envs/p4dev/lib/libmkl_rt.so CACHE STRING "")
# /home/miniconda/envs/p4dev/include CACHE STRING "")

# <<< libint
set(BOOST_ROOT /home/miniconda/envs/p4dev CACHE STRING "")
set(Eigen3_ROOT /home/miniconda/envs/p4dev CACHE STRING "")
set(Libint2_DIR /home/miniconda/envs/p4dev/lib/cmake/libint2 CACHE PATH "")
# set(MAX_AM_ERI 5 CACHE STRING "")
# set(CMAKE_INSIST_FIND_PACKAGE_Libint2 ON CACHE BOOL "")
# set(CMAKE_DISABLE_FIND_PACKAGE_Libint2 ON CACHE BOOL "")

# <<< qcelemental           msgpack-python, networkx, setuptools
# Primarily OTF runtime detected.
# set(CMAKE_INSIST_FIND_PACKAGE_qcelemental ON CACHE BOOL "")
# set(CMAKE_DISABLE_FIND_PACKAGE_qcelemental ON CACHE BOOL "")

...
# Sections skipped because packages detected on-the-fly, not through CMake
# <<< pytest                  Primarily OTF runtime detected.
# <<< geometric               Primarily OTF runtime detected.

# Sections skipped because packages absent from current environment
# <<< ~pylibefp~
# <<< ~integratorxx~
```

**GENERATED
cache_p4dev.cmake**

```
conda/psi4-path-advisor.py cache --help
#> usage: psi4-path-advisor cache
#>   [-h] [--objdir OBJDIR]
#>   [--compiler {byo, conda, GNU, IntellLVM, Intel}]
#>   [--lapack {byo, conda, mkl}] [--insist]
```

CODEDEPS → REPO

psi4-path-advisor.py deploy

copy repository source info into many CMakeLists.txt. Also dry-run solve for all arch and write static conda env spec files.

external/gdma/CMakeLists.txt

```
if(${ENABLE_gdma})
  find_package(gdma 2.3.3 COMPONENTS Python
    CONFIG QUIET) # edit in codedeps

  if(${gdma_FOUND})
    get_property(_loc TARGET gdma::pygdma PROPERTY LOCATION)
    message(STATUS "${Cyan}Found gdma${ColourReset}: ${_loc}")
    add_library(gdma_external INTERFACE) # dummy

  else()
    if(${CMAKE_INSIST_FIND_PACKAGE_gdma})
      message(FATAL_ERROR "Suitable gdma could not be found")
    endif()
    message(STATUS "Suitable gdma could not be located")

    ExternalProject_Add(gdma_external
      DEPENDS pybind11_external
      URL https://github.com/psi4/gdma/archive/v2.3.3.tar.gz # Stone/upstream c2e0b5 + lib, I/O, Py # edit in codedeps
      CMAKE_ARGS -DCMAKE_INSTALL_PREFIX=${STAGED_INSTALL_PREFIX}
                 -DCMAKE_CXX_COMPILER=${CMAKE_CXX_COMPILER}
                 -Dgdma_ENABLE_PYTHON=ON
                 -DPython_EXECUTABLE=${Python_EXECUTABLE}
                 -Dpybind11_DIR=${pybind11_DIR}
      ...
    )
  endif()
else()
  message(FATAL_ERROR "gdma must be found or enabled")
endif()

# edit in codedeps
```

OPT'L COMPILED

```
- project: gdma
  use:
    added: "1.0"
    added_note: "Lib to pygdma at 1.9. Ext at 1.1."
    required: false
    required_note: "Allow Stone multipole analysis"
    buildable: true
    buildtime: false

    cms: true
    repository:
      host: github
      account: psi4
      name: gdma
      commit: v2.3.3
      commit_note: "GL upstream c2e0b5 + lib, I/O, Py"
    cmake:
      name: gdma
      constraint: 2.3.3
      components:
        - Python
        target: gdma::pygdma
        enable: ENABLE_gdma
    conda:
      channel: conda-forge
      name: pygdma
      constraint: null
      cmake:
        ENABLE_gdma: true
        gdma_DIR:
          unix: ${CONDA_PREFIX}/share/cmake/gdma
          win-64: ${CONDA_PREFIX}/Library/share/cmake/gdma
        //CMAKE_INSIST_FIND_PACKAGE_gdma: true
        //CMAKE_DISABLE_FIND_PACKAGE_gdma: true
      cmake_note: "Primarily OTF runtime detected.
                  CMake enabling only relevant for CTest."
```

psi4 / devtools / conda-envs /

- docs-cf.yaml
- linux-64-buildrun.yaml
- osx-64-buildrun.yaml
- osx-arm64-buildrun.yaml
- win-64-buildrun.yaml

PUTTING CODEDEPS TO USE FOR DEVS

conda-enabled superbuild



GitLab

research homepage

PyPI

GitHub

GitHub:psi4



- Sounds overly prescriptive, but the idea is to meet known tools at known configuration points.
- Above is the recc. build procedure for devs. Note steps (2.2) & (3.2) where you customize.
- Nowadays, use `fetch_content` instead of superbuild, as latter poor for non-top-level or shared deps.

```
=====
(B) flexible usage
=====

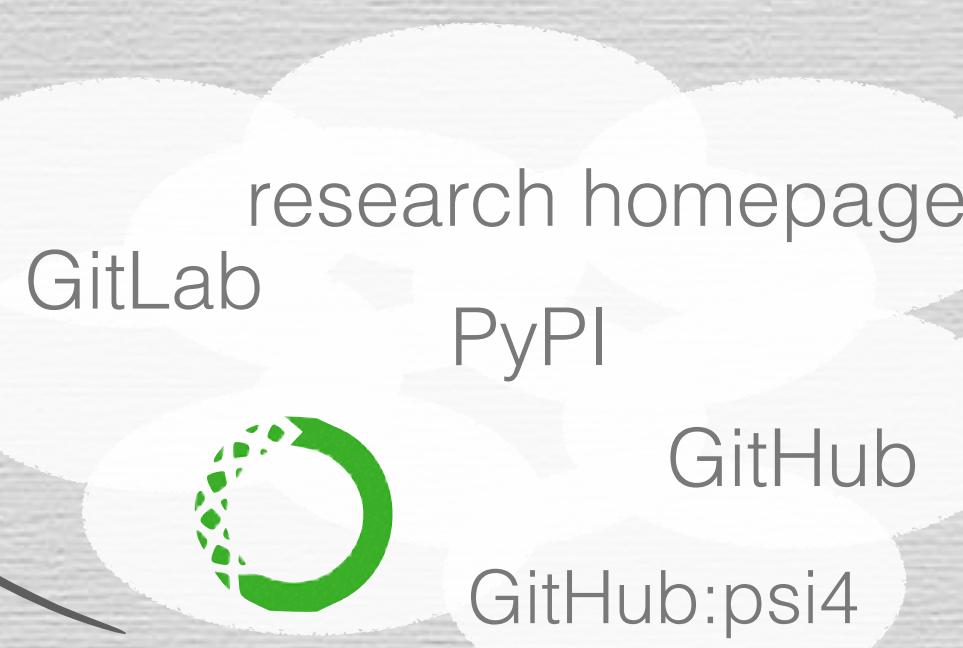
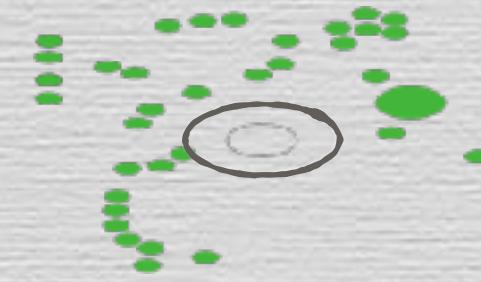
# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4

# (2.1) generate env spec file from codedeps.yaml.
conda/psi4-path-advisor.py env -n p4dev --python 3.12 --disable addons
#> conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev
# (2.2) edit env_p4dev.yaml to customize software packages.
# (2.3) issue suggested or customized command to create and activate conda env.
conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev

# (3.1) generate cmake cache file from conda env.
conda/psi4-path-advisor.py cmake
#> cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev && \
    cmake --build objdir_p4dev
# (3.2) edit cache_p4dev.cmake to customize build configuration.
# (3.3) issue suggested or customized command to configure and build with cmake.
cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev \
    -DCMAKE_INSTALL_PREFIX=/path/to/install-psi4 && cmake --build objdir_p4dev
```

PUTTING CODEDEPS TO USE FOR DEVS

conda-enabled superbuild



```
=====
(B) flexible usage
=====

# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4

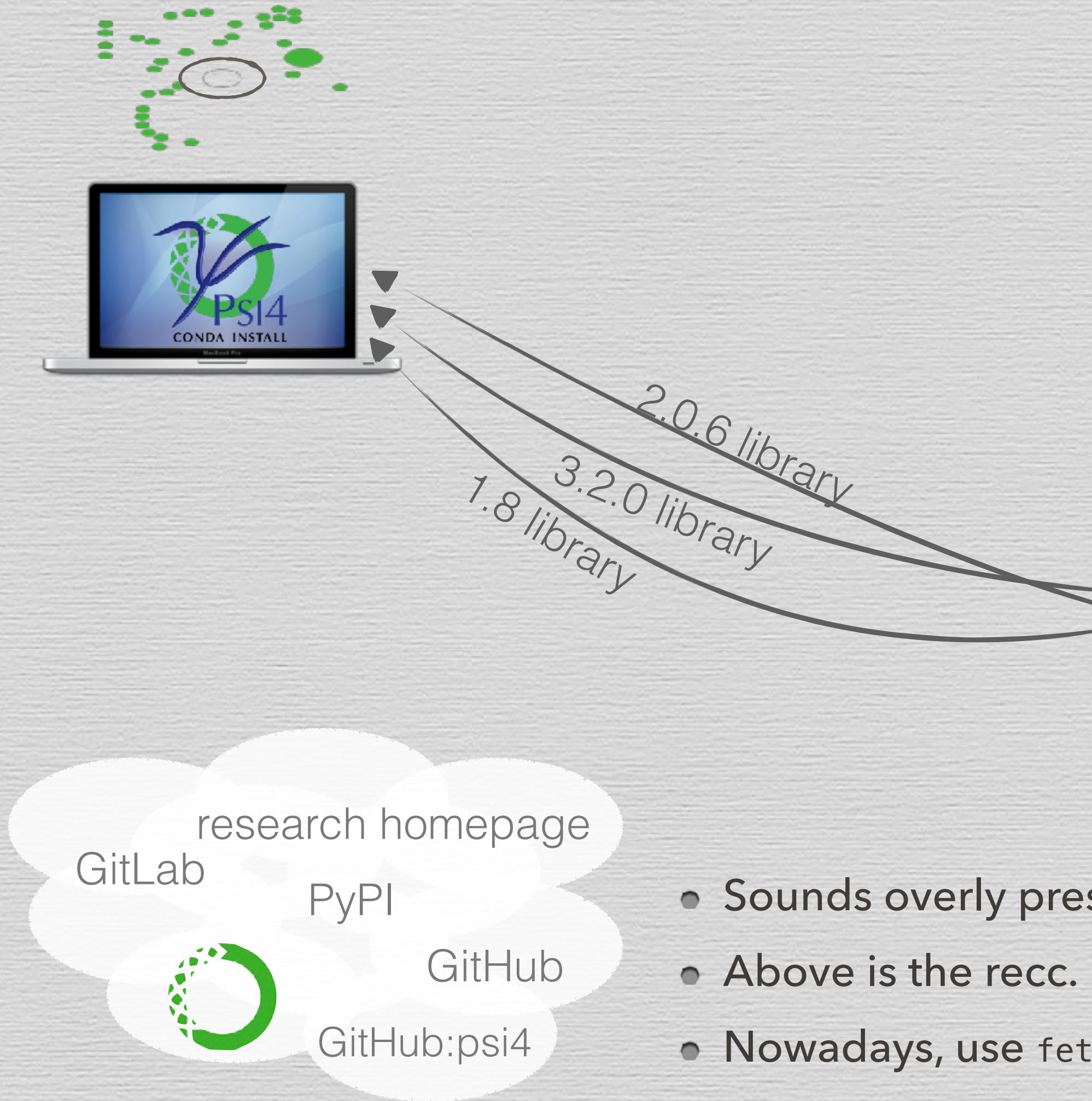
# (2.1) generate env spec file from codedeps.yaml.
conda/psi4-path-advisor.py env -n p4dev --python 3.12 --disable addons
#> conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev
# (2.2) edit env_p4dev.yaml to customize software packages.
# (2.3) issue suggested or customized command to create and activate conda env.
conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev

# (3.1) generate cmake cache file from conda env.
conda/psi4-path-advisor.py cmake
#> cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev && \
    cmake --build objdir_p4dev
# (3.2) edit cache_p4dev.cmake to customize build configuration.
# (3.3) issue suggested or customized command to configure and build with cmake.
cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev \
    -DCMAKE_INSTALL_PREFIX=/path/to/install-psi4 && cmake --build objdir_p4dev
```

- Sounds overly prescriptive, but the idea is to meet known tools at known configuration points.
- Above is the recc. build procedure for devs. Note steps (2.2) & (3.2) where you customize.
- Nowadays, use `fetch_content` instead of superbuild, as latter poor for non-top-level or shared deps.

PUTTING CODEDEPS TO USE FOR DEVS

conda-enabled superbuild



(B) flexible usage

```
=====
# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4

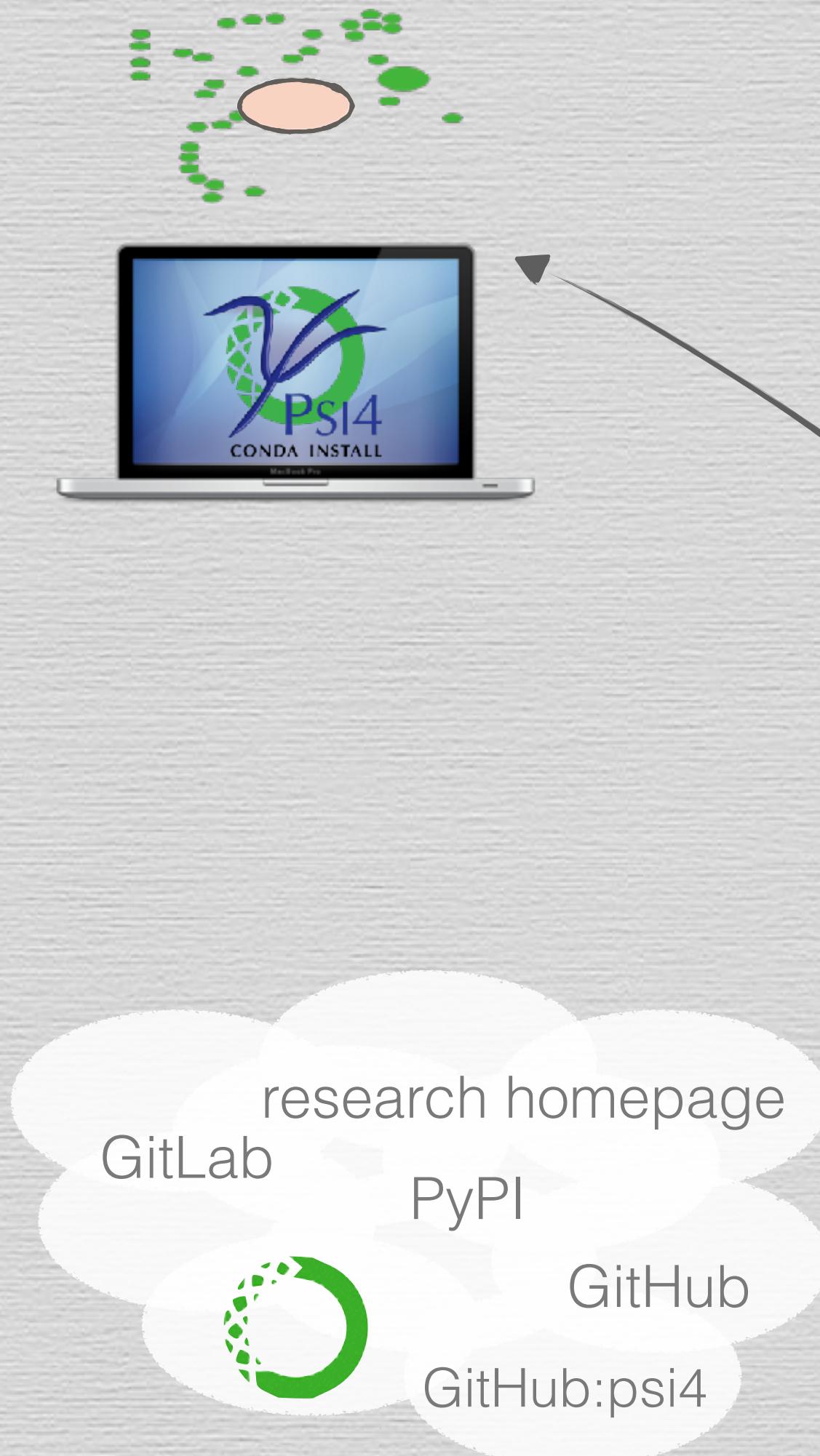
# (2.1) generate env spec file from codedeps.yaml.
conda/psi4-path-advisor.py env -n p4dev --python 3.12 --disable addons
#> conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev
# (2.2) edit env_p4dev.yaml to customize software packages.
# (2.3) issue suggested or customized command to create and activate conda env.
conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev

# (3.1) generate cmake cache file from conda env.
conda/psi4-path-advisor.py cmake
#> cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev && \
    cmake --build objdir_p4dev
# (3.2) edit cache_p4dev.cmake to customize build configuration.
# (3.3) issue suggested or customized command to configure and build with cmake.
cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev \
    -DCMAKE_INSTALL_PREFIX=/path/to/install-psi4 && cmake --build objdir_p4dev
```

- Sounds overly prescriptive, but the idea is to meet known tools at known configuration points.
- Above is the recc. build procedure for devs. Note steps (2.2) & (3.2) where you customize.
- Nowadays, use `fetch_content` instead of superbuild, as latter poor for non-top-level or shared deps.

PUTTING CODEDEPS TO USE FOR DEVS

conda-enabled superbuild



(B) flexible usage

```
=====
# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4

# (2.1) generate env spec file from codedeps.yaml.
conda/psi4-path-advisor.py env -n p4dev --python 3.12 --disable addons
#> conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev
# (2.2) edit env_p4dev.yaml to customize software packages.
# (2.3) issue suggested or customized command to create and activate conda env.
conda env create -n p4dev -f /home/psi4/env_p4dev.yaml && conda activate p4dev

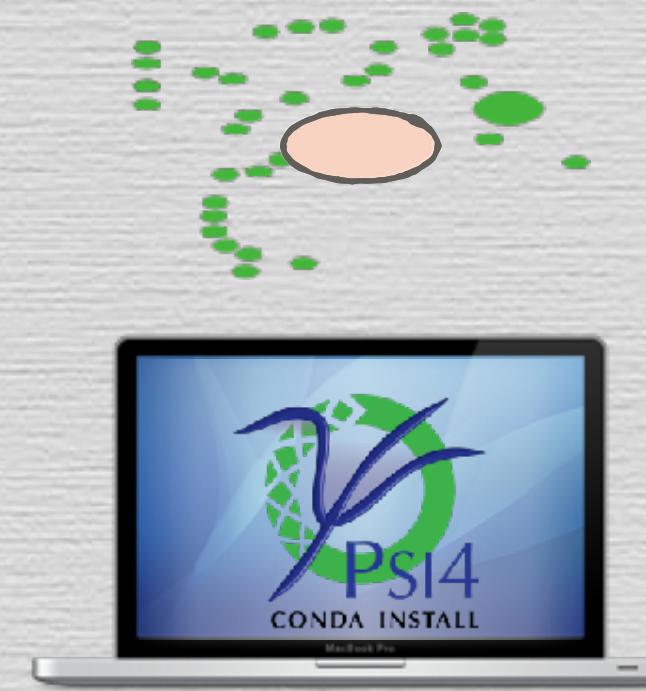
# (3.1) generate cmake cache file from conda env.
conda/psi4-path-advisor.py cmake
#> cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev && \
    cmake --build objdir_p4dev
# (3.2) edit cache_p4dev.cmake to customize build configuration.
# (3.3) issue suggested or customized command to configure and build with cmake.
cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev \
    -DCMAKE_INSTALL_PREFIX=/path/to/install-psi4 && cmake --build objdir_p4dev
```

- Sounds overly prescriptive, but that's the point.
- Above is the rec. build procedure.
- Nowadays, use `fetch_content` instead.

```
-- Found pybind11: /usr/share/miniconda/envs/p4build/include (found version 2.1.3)
-- Found ambit: /usr/share/miniconda/envs/p4build/lib/libambit.so (found version 0.6.0)
-- Found bse: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/basis_set_exchange (found version 0.11)
-- Found dkh: /usr/share/miniconda/envs/p4build/lib/libdkh.so (found version 1.2)
-- Found ecpint: /usr/share/miniconda/envs/p4build/lib/libecpint.so.1 (found version 1.6.7)
-- Found libefp: /usr/share/miniconda/envs/p4build/lib/libefp.so.1 (found version 1.5.0)
-- Found Einsums: /usr/share/miniconda/envs/p4build/lib/libEisums.so.1.0.4 (found version 1.0.4)
-- Found gau2grd: /usr/share/miniconda/envs/p4build/lib/libgau2grd.so.2 (found version 2.0.7)
-- Found gdma: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/gdma_cython-313-x86_64-linux-gnu.so (found version 2.3.3)
-- Found Boost: /usr/share/miniconda/envs/p4build/include (found suitable version "1.88.0", minimum required is "1.29")
-- Found Libint2: /usr/share/miniconda/envs/p4build/lib/libint2.so.2.9.0 (found version 2.9.0)
-- Found PCMSolver: /usr/share/miniconda/envs/p4build/lib/libpcmsolver.so.1 (found version 1.2.3.999)
-- Found pylibefp: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/pylibefp (found version undefined)
-- Found qclementel: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/qclementel (found version 0.29.0)
-- Found qcengine: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/qcengine (found version v0.32.0)
-- Found optking: /usr/share/miniconda/envs/p4build/lib/python3.13/site-packages/optking (found version 0.3.0)
-- Found Libxc: /usr/share/miniconda/envs/p4build/lib/libxc.so.15 (found version 7.0.0)
-- Suitable BrianQC could not be located, please contact info@brianqc.com to obtain BrianQC
-- Found IntegratorXX: /usr/share/miniconda/envs/p4build/lib/libintegratorxx.so (found version 1.2.0)
-- Suitable v2rdm_casscf could not be located, Building v2rdm_casscf instead.
-- No Doxygen, no docs.
-- No Sphinx, no docs. Pre-built documentation at http://psiccd.org/psi4manual/master/index.html
```

PUTTING CODEDEPS TO USE FOR DEVS

conda-enabled superbuild



```
=====
(B) flexible usage
=====

# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4

=====
(A) black-box usage (copy/paste-able)
=====

# (1) get code from GitHub
git clone https://github.com/psi4/psi4.git && cd psi4
# (2) generate env spec file from codedeps.yaml. "eval $(...)" creates and activates conda env.
eval $(conda/psi4-path-advisor.py env)
# (3) generate cmake cache file from conda env. "eval $(...)" configures and builds with cmake.
eval $(conda/psi4-path-advisor.py cmake)

shows up in p4dev
```

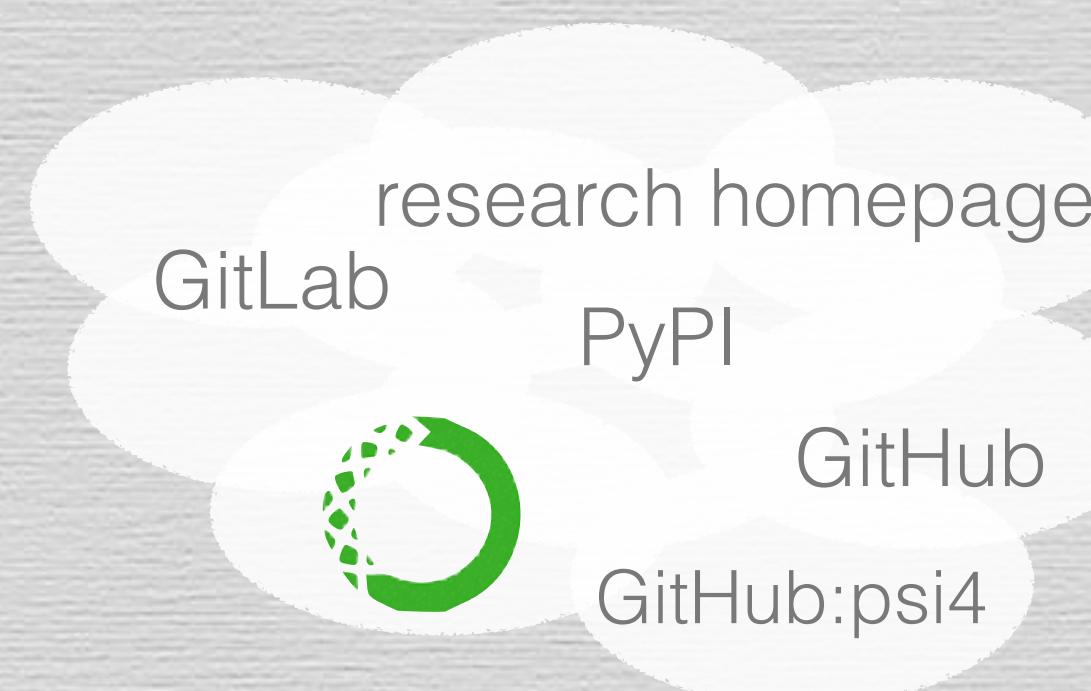
(1) activate p4dev
(2) conda env.
(3) create p4dev

```
# (3.2) edit cache_p4dev.cmake to customize build configuration.
# (3.3) issue suggested or customized command to configure and build with cmake.
cmake -S. -C/home/psi4/cache_p4dev.cmake -Bobjdir_p4dev \
-DCMAKE_INSTALL_PREFIX=/path/to/install-psi4 && cmake --build objdir_p4dev
```

- Sounds overly prescriptive, but the idea is to meet known tools at known configuration points.
- Above is the recc. build procedure for devs. Note steps (2.2) & (3.2) where you customize.
- Nowadays, use `fetch_content` instead of superbuild, as latter poor for non-top-level or shared deps.
- In practice, though, the procedure is so steady, one can do the short route:

★ At the ecosys level, even simple dependencies logic (pytest is for testing) can get overwhelming, so consolidate & encode.

★ A SSOT deps file can be a path to recovery when the build breaks by broadcast fixes



DISTRIBUTION

<https://psicode.org/install/latest>

★ Aim for copy/paste install even for compiled software

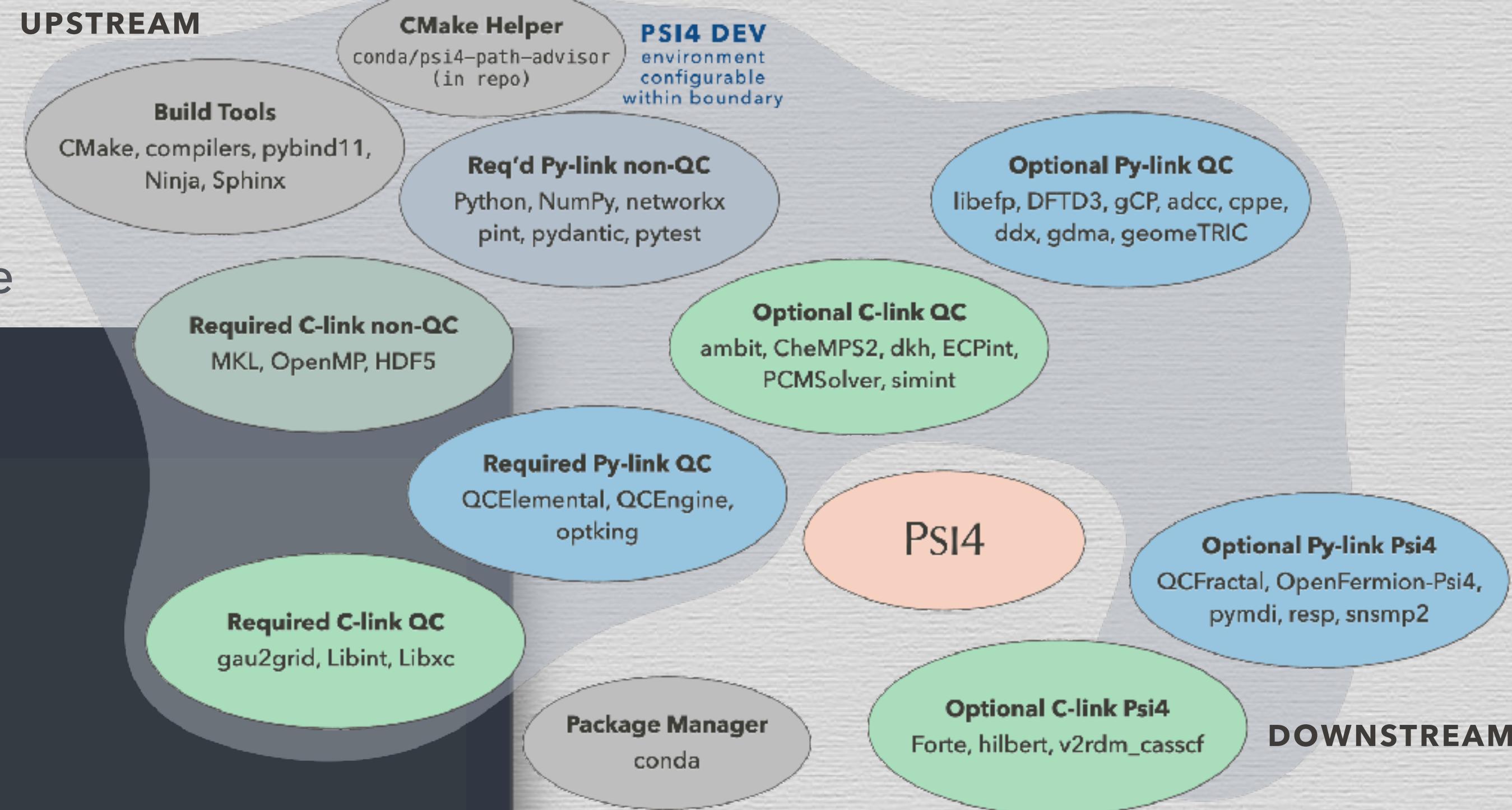
Get Started with PSI4

Select Preferences



Run this command

```
>_ git clone --branch 1.10.x --single-branch https://github.com/psi4/psi4.git && cd psi4
>_ eval $(conda/psi4-path-advisor.py env --python 3.12 --name p4dev --disable addons docs)
>_ eval $(conda/psi4-path-advisor.py cmake)
>_ eval $(objdir_p4dev/stage/bin/psi4 --psiapi)
>_ psi4 --test
>_ # run `conda/psi4-path-advisor.py -h` for guide and options
>_ # -or- see top-level CMakeLists.txt for generic (non-conda) guide
```



CORE DEVELOPERS:
CONDA-ENABLED SUPERBUILD
deps · dev tools · add-ons · cmake cmd

PY-FRIENDLY USERS:
CONDA PACKAGE
PSI4 · dependencies · add-ons

PY-WARY USERS:
CONDA INSTALLER
PSI4 · dependencies · add-ons

628K

PSI4CONDA

633K

CONTINUOUS INTEGRATION TESTING

a repository's lifeline and anchor

Strongly recommended characteristics of a **TEST RUNNER**

- **RUN-TIME DETECTION:** optional dependencies bringing new capabilities may be runtime-loaded, so a runner should detect current environment dependencies without reconfiguration (like CTest).
- QCElemental & QCEngine provide detection utilities
- Pytest marks can run, skip, & select: **pytest -m dftd3**

MARK TESTS TO RUN OR SKIP

```
@pytest.mark.scf
@pytest.mark.dft
@using("libxc")
@pytest.mark.parametrize("func,expected", [
    pytest.param('wM06-D3', 0.4563, marks=using("dftd3")), # Libxc 180
    pytest.param('X3LYP', 0.4549), # Q-Chem
    pytest.param('KMLYP', 0.4665), # ERKALE
], ids=name_dft_test)
def test_dft_bench_ionization(func, expected, dft_bench_systems):
    """functionals ionization energies vs. Q-Chem"""
    ...
```

DETECT ADDONS

```
{
    "mrcc": which("dmrcc", return_bool=True),
    "qcfractal": is_qcfractal_new_enough("0.60"),
    "bse": which_import("basis_set_exchange", return_bool=True),
    "gauxc": psi4.addons("gauxc"),
}
```

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- easy testing is another use for Py intf to a compiled lib

INSTALLED TESTING

```
>>> psi4 --test
test_ddx.py::test_ddx_fock_build[h2lpb] SKIPPED (ddx !found)
test_mdi.py::test_mdi_water SKIPPED (mdi !found)
test_option.py::test_spacious_option PASSED
test_psi4.py::test_psi4_basic PASSED
test_psi4.py::test_psi4_cc PASSED
test_psi4.py::test_psi4_cas PASSED
test_psi4.py::test_psi4_dfmp2 PASSED
test_psi4.py::test_psi4_sapt PASSED
test_psi4.py::test_psi4_scfproperty PASSED
test_psi4_qcschema.py::test_psi4_basic PASSED
test_psi4_qcschema.py::test_psi4_cc XFAIL (no AtomicInput opt)
test_psi4_qcschema.py::test_psi4_cas PASSED
test_psi4_qcschema.py::test_psi4_dfmp2 PASSED
test_psi4_qcschema.py::test_psi4_sapt PASSED
test_qcfractal.py::test_qcf_cbs_mbe[internal] PASSED
test_qcfractal.py::test_qcf_cbs_mbe[snowflake] PASSED
test_thc_eri.py::test_ls_thc_exact PASSED
tu1-h2o-energy/test_input.py::test_tu1_h2o_energy PASSED
===== 49 passed, 60 skipped, 7934 deselected, 1 xfailed in 4m
=====
```

CONTINUOUS INTEGRATION TESTING

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 - easy testing is another use for Py intf to a compiled lib
- **ERROR TESTING**: error & warning pathways & deprecated features should be as easy to test as numerical results. QC isn't run only by specialists anymore.
 - fend off nonsense jobs before resources run out
 - nearly as easy to enforce advice in code as via docs
 - check code paths that helpful error is being raised

CHECK ERROR PATHWAYS FUNCTIONING

```
def test_jumbledzmat_error():
    subject = """He
    He 1 2. 2 100. 3 35.
    He 1 2.
    ....
    """

    with pytest.raises(qcelemental.ValidationError) as e:
        qcelemental.molparse.from_string(subject)

    assert 'aim for lower triangular' in str(e)
```

CONTINUOUS INTEGRATION TESTING

a repository's lifeline and anchor

Strongly recommended characteristics of a **TEST RUNNER**

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GOBBLEDYGOOK OF EXPECTED ERRORS

```
_p42 = (psi4.MissingMethodError, r"Method=(mp4\$(dq\)|mp4|qcisd|qcisd\$(t\)) is not available for any derivative level under conditions (CI|MP)_TYPE=CONV, REFERENCE=(U|R0)HF, FREEZE_CORE=(TRUE|FALSE), QC_MODULE=\(auto\)", "no open-shell mp4(dq)/mp4/qcisd/qcisd(t) by fnocc")
_p43 = (psi4.ValidationError, r"(CD|DF) is not a valid choice", "no df/cd ci by psi4")
_p44 = (psi4.ValidationError, r"Invalid type (DF|CD) for FN0CC energy through `run_(fnocc|cepa)`.", "no df/cd except ccisd/ccsd(t) by fnocc")
_p45 = (psi4.ValidationError, r"Reference UHF for DETCI is not available", "no uhf by detci")
_p46 = (psi4.ValidationError, r"Invalid type (DF|CD) for DETCI energy through `run_detci`", "no df/cd by detci")
_p47 = (psi4.UpgradeHelper, r"Replace method ZAPT with method MP for RHF reference", "retire rhf zapt by detci")
_p48 = (psi4.MissingMethodError, r"Method=cisd is not available for any derivative level under conditions CI_TYPE=CONV, REFERENCE=UHF, FREEZE_CORE=(TRUE|FALSE), QC_MODULE=\(auto\)", "no uhf cisd by psi4")
_p98 = (psi4.MissingMethodError, r"Method=ccsd\$(t\)) is not available for requested derivative level \$(reqd=1 > avail=0) under conditions CC_TYPE=CONV, REFERENCE=(R|U)HF, FREEZE_CORE=(TRUE|FALSE), QC_MODULE=\(auto\)", "temporary: cc(t) disabled w/o qc_module=ccenergy for conv rhf/uhf gradients for ccisd(t) by ccenergy until scaling reworked")
_p99 = (psi4.MissingMethodError, r"Method=(ccsd\$(t\)|a-ccsd\$(t\)) is not available for any derivative level under conditions CC_TYPE=(DF|CD), REFERENCE=UHF, FREEZE_CORE=(TRUE|FALSE), QC_MODULE=\(auto\)", "temporary: cc(t) disabled w/o qc_module=occ in dfocc until further optimization ")
_w1 = ("CCSD TOTAL GRADIENT", "wrong semicanonical ae rohf ccisd gradients by ccenergy")
```

CONTINUOUS INTEGRATION TESTING

INTERNAL

```
conda/psi4-path-advisor.py \
env \
--name p4env \
--python ${PYTHON_VER} \
--disable compilers addons docs

# create conda env from yaml spec

conda/psi4-path-advisor.py \
cmake \
--objdir build \
--insist

# configure cmake from cache
```

DOCS

```
code/conda/psi4-path-advisor.py \
env \
--name p4docs \
--python ${matrix.python-ver} \
--disable addons \
--offline-conda

# env, cache, configure as above
```

ECO

```
conda/psi4-path-advisor.py \
env \
--name p4build \
--python ${matrix.python-ver} \
--disable docs \
--offline-conda

# env, cache, configure as above
```

a repository's lifeline and anchor

PSI4 INTERNAL CI, AZURE



psi4 / psi4 / Pipelines / psi4.psi4 / 20250618.3

Name	Status	Duration
Windows VS2022	Success	⌚ 1h 43m 18s
Linux Builds gcc_11	Success	⌚ 1h 5m 50s
Linux Builds gcc_latest	Success	⌚ 1h 57m 12s
Linux Builds clang_latest	Success	⌚ 1h 19m 26s

PSI4 ECOSYSTEM CI, GHA

- ✓ Docs / Propose Docs (ubuntu-latest, 3.10) (pull_request) Successful in 28m
- ✓ Eco / Eco · 3.10 · Conda Clang (M-Silicon) (pull_request) Successful in 37m
- ✓ Eco / Eco · 3.11 · Conda Clang (M-Intel) (pull_request) Successful in 103m
- ✓ Eco / Eco · 3.12 · LLVM clang-cl (W) (pull_request) Successful in 60m
- ✓ Eco / Eco · 3.13 · Conda GNU (L) (pull_request) Successful in 45m
- ✓ psi4.psi4 – #20250618.3 succeeded Required

2296 tests of internal Psi4 capabilities, Azure is cmdline-like for users, non-conda compilers

- quick
- quick + 1/3 full, earliest GCC, Py
- quick + 1/3 full, latest GCC, Py
- quick + 1/3 full

<386 tests integrating with external software, architecture variety, all-conda

- addon + quick
- addon + quick
- addon
- addon

- ★ Bring the changing env of diff arch, compilers, dep ver into freq. contact w/code.
- ★ Use CI to assure developers always getting a full & proved working environment.

CONTINUOUS INTEGRATION TESTING

★ a repository's lifeline and anchor

LIFELINE

- **CONSOLIDATE** and define gains
- **OUTSOURCE** fixing problems or discussing changes to PRs that introduce conflicts
- **FACILITATE** refactoring and improving codebase
- **ATTUNED** to version & packaging changes in your own deps



WHAT'S MISSING?

- testing against **DEV** (not release) versions of eco projects
- **PERFORMANCE** regression testing
- new arch or **HARDWARE** test suite

This is not a apology,
this is a warning:

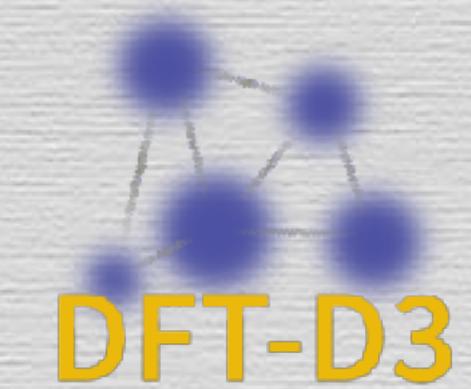
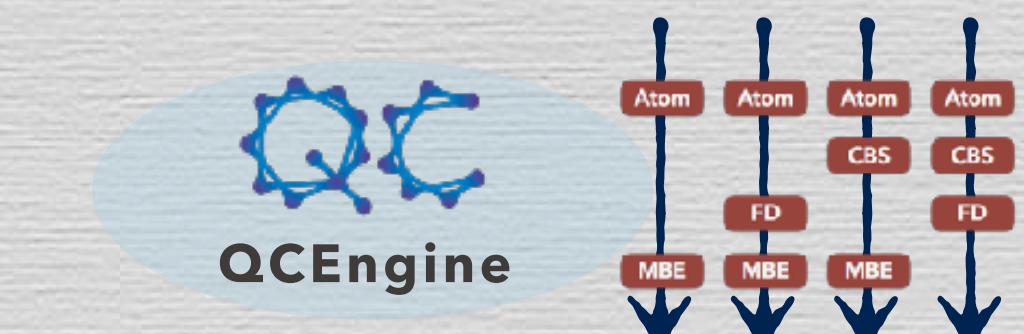
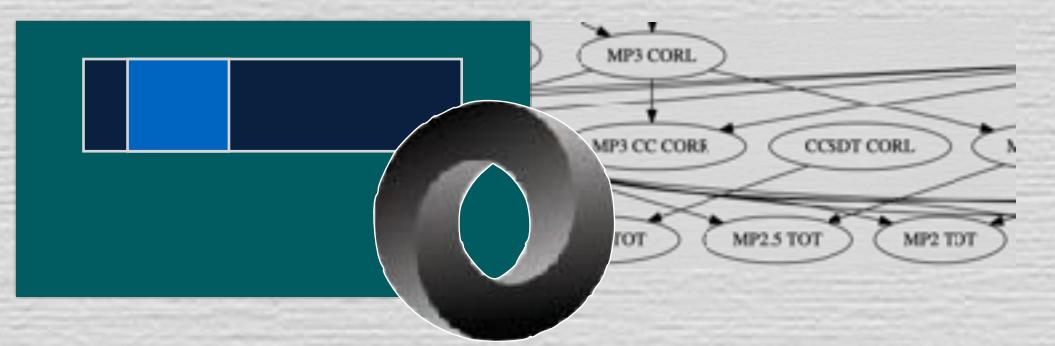
If it's not tested, it's broken

– Bruce Eckel

- GHA using a Windows image with different compiler & STL version; add define
- NumPy v2 needs to be used with latest Pybind11; seen as "atoms too close" error
- new compilers insert a breakpoint for oob memory access: sudden "Illegal Instruction"

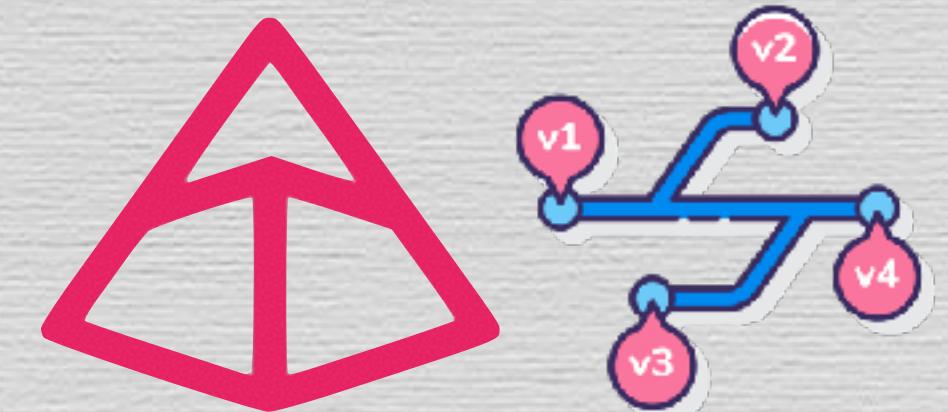
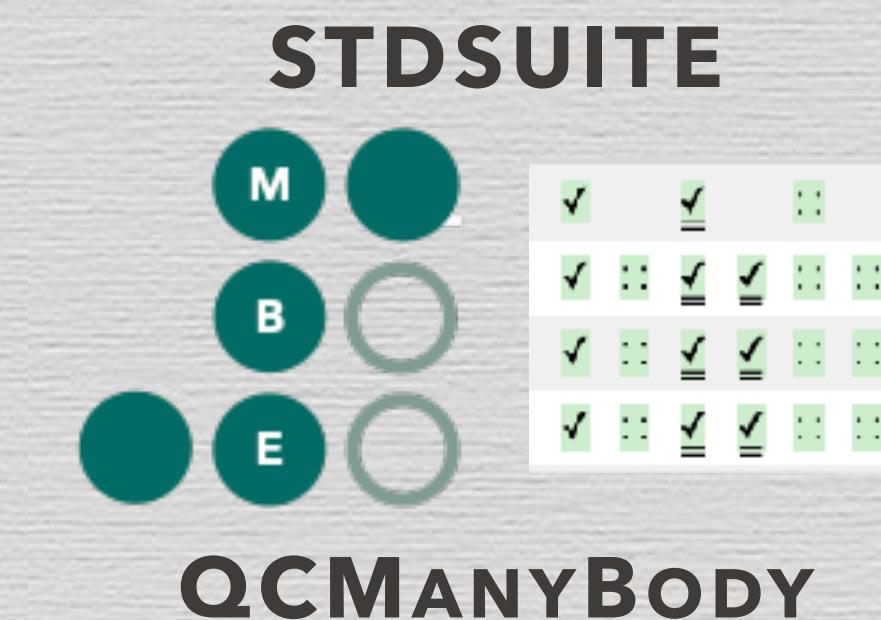
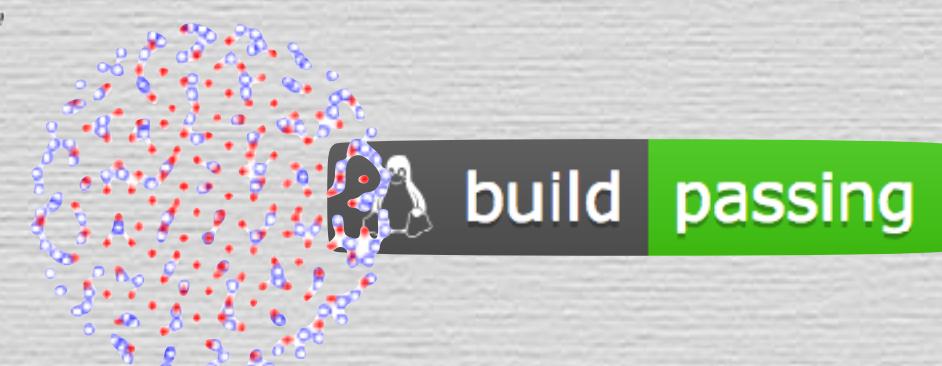
ANCHOR

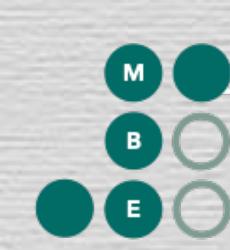
- **CODE** doesn't rot but envs sure do
- **ECO BUFFETTED** by every problem of upstream, most of them not QC
- As other packages are better maintained, their SE virtues (upgrades & deprecations) hit you.
- **FEATURE** Using CI, you're arranging trouble to hit the manager, not users & arranging for tweaks to reach users (pathadvisor). Try to consider it a feature, not a bug.
- **EXAMPLES** from the past year
 - stray pkg_resources usage in an addon triggered a printed setup-tools warning when importing Psi4, which printing interfered with qcng QC detection, breaking 1/3 of Psi4
 - syntax warning in Clang v18 collided with MSVC header Libint includes, repositioned in unreleased v19, so fallback to v17



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY





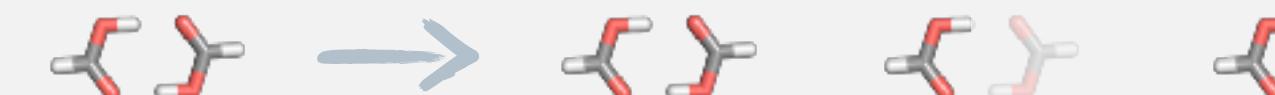
EXTRACTING THE QC MANYBODY PROCEDURE

multiple entry points for flexibility



class ManyBodyComputer ():

PLAN Separate molecule into subsystems. CP, noCP, VMFC basis.
method unchanged.



for frag in fragments: return qcschema



ASM Assemble n-body & interaction results from fragments.



- noCP, SSFC, VMFC
- indep. or schema interf.
- energy, gradient, Hess.
- intermolecular only
- IE or MBE
- multiple model chemistry MBE



Ben Pritchard
MolSSI



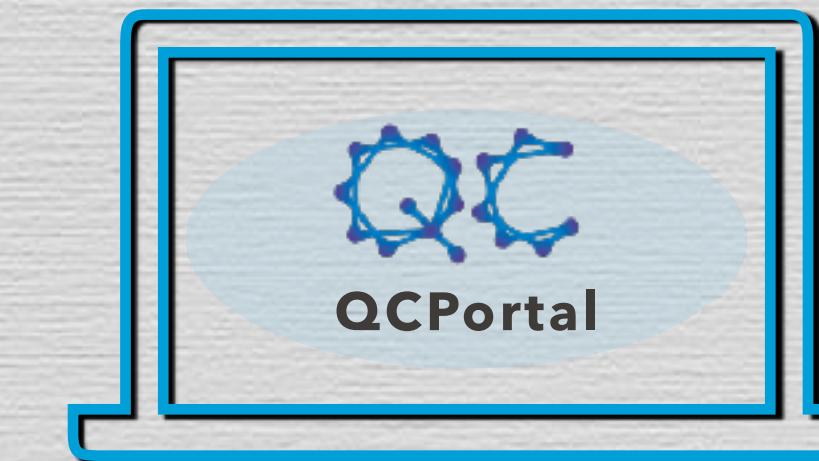
Asem Alenaizan
GaTech → KFUPM



Daniel Smith
MolSSI → Abiologics



QCManyBody direct



QCFractal service



MBE backend



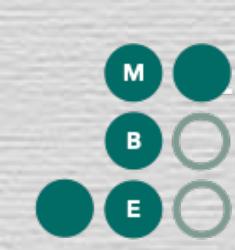
QCEngine procedure



PSI4 distributed driver



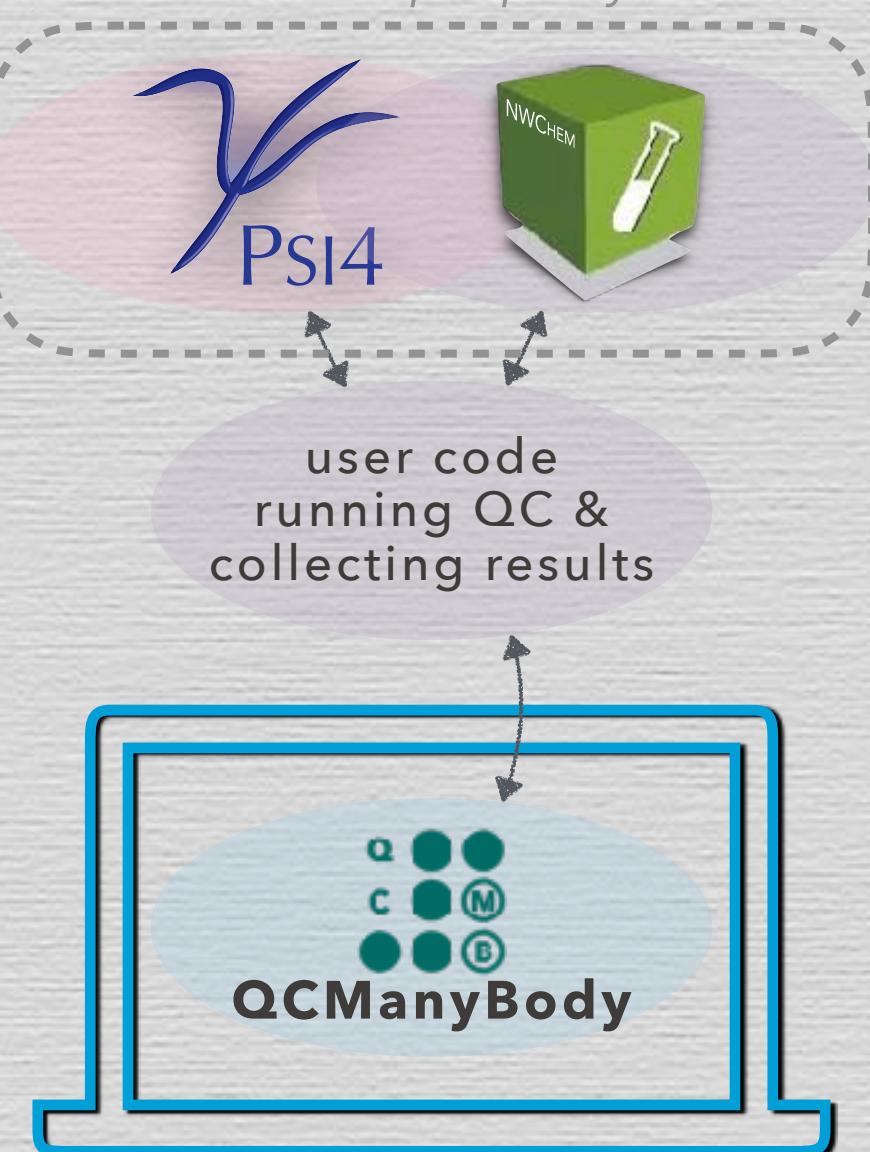
Optimizer backend



TWO INTERFACES FOR QC MANYBODY

independent or integrated with QC Archive

CORE INTERFACE

```
any CMS program producing  
E/G/H/property  
  
from qcmanybody import ManyBodyCore  
  
mbc = ManyBodyCore(  
    molecule= Ne - Ne - Ne,  
    bsse_type=["cp"],  
    levels={1: "ccsd/cc-pvtz",  
            2: "mp2/cc-pvdz",  
            3: "mp2/cc-pvdz"},  
    return_total_data=True,  
    supersystem_ie_only=False,  
    embedding_charges=None)  
  
calculation_results = {}  
for chem, lbl, imol in mbc.iterate_molecules():  
    # use chem, lbl, imol to construct inputs  
  
    # run them & fill in results  
    calculation_results[lbl] = {"energy": None}  
#> calculation_results = {  
#>   '["mp2/cc-pvdz", [1], [1, 2]]': {'energy': -128.6814},}  
  
results = mbc.analyze(calculation_results)  
  
print(results["ret_energy"])  
#> -384.1286
```

QCManyBody core interface

- user-prov. serial/parallel
- user runs CMS backends

HIGH-LEVEL INTERFACE

```
from qcmanybody import ManyBodyComputer  
from qcmanybody.models import ManyBodyInput  
  
mbin = ManyBodyInput(**{  
    "molecule": Ne - Ne - Ne,  
    "specification": {  
        "driver": "energy",  
        "keywords": {  
            "bsse_type": ["cp"],  
            "levels": {  
                1: "nwc-ccsd/tz",  
                3: "p4-mp2/dz"}},  
        "specification": {  
            "nwc-ccsd/tz": {  
                "program": "nwchem",  
                "model": {  
                    "method": "ccsd",  
                    "basis": "cc-pvtz"},  
            "p4-mp2/dz": {  
                "program": "psi4",  
                "model": {  
                    "method": "mp2",  
                    "basis": "cc-pvdz"},  
            }}  
    }  
    # runs with QC Engine  
    ret = ManyBodyComputer.from_manybodyinput(mbin)  
  
    print(ret.return_result)  
    #> -384.1286
```

QCManyBody high-level interface

- serial
- multi CMS backends



QCSchema MANYBODY

driving the many-body expansion

$$E^M = E^1 + E^{(2)} + E^{(3)} + \dots + E^{(M)},$$
$$= \sum_{I \in \mathcal{P}_1} E^I + \sum_{IJ \in \mathcal{P}_2} E^{(IJ)} + \sum_{IJK \in \mathcal{P}_3} E^{(IJK)} + \dots,$$

- **RECENT** added in 2024
- Counterpoise, VMFC, and no-CP arbitrary-order manybody expansion (**MBE**) extracted from Psi4 program
- **SUITABLE** for intermolecular fragments, no bond-breaking

ManyBodyKeywords

bsse_type: any combination of

- nocp/mbe
- cp/ssfc
- vmfc

embedding_charges
return_total_data: IE or total levels:

- {2: hi, 4: lo}
- {1: hi, 2: md, supersystem: lo}

max_nbody: truncate MBE early
supersystem_ie_only: no MBE analysis

ManyBodySpecification

schema_version: 1
program: any MBE speaking QCSchema



protocols: customize MBE return layout

ManyBodyProtocols

component_results: save cluster data

extras: free-form storage

driver: MBE target derivative

- energy
- gradient
- Hessian

E

G

H

keywords: knobs in the MBE program

ManyBodyKeywords

specification: how to run MBE subsystems

- single

AtomicSpecification

• mapping

{hi: **AtomicSpecification**,

lo: **AtomicSpecification**}

ManyBodyInput

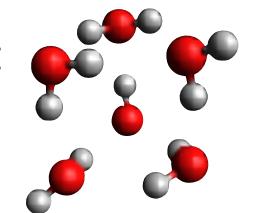
schema_version: 1
provenance: creator name and version info

QCManyBody, 0.3.0

id: tracker for databases

extras: free-form storage

molecule:



Molecule

specification: how to run MBE single-point

ManyBodySpecification



QCSHEMA MANYBODY

driving the many-body expansion

$$E^M = E^1 + E^{(2)} + E^{(3)} + \dots + E^{(M)},$$
$$= \sum_{I \in \mathcal{P}_1} E^I + \sum_{IJ \in \mathcal{P}_2} E^{(IJ)} + \sum_{IJK \in \mathcal{P}_3} E^{(IJK)} + \dots,$$

- **RECENT** added in 2024
- Counterpoise, VMFC, and no-CP arbitrary-order manybody expansion (**MBE**) extracted from Psi4 program
- **SUITABLE** for intermolecular fragments, no bond-breaking

AtomicSpecification

schema_version: 1

program: any SP in QCEngine



protocols: customize return layout

AtomicProtocols

stdout: save text output

wavefunction: save orbital info

native_files: save raw prog. files

error_correction: auto-edit&rerun

extras: free-form storage

driver: single-point derivative

- energy

- gradient

- Hessian

- properties

model: model chemistry or FF

method: B3LYP-D3, CCSD(T)

basis: 6-31G*, cc-pVQZ

keywords: knobs in SP program

{scftyp: uhf, cc_conv: 7, docc: [4,1]}

ManyBodySpecification

schema_version: 1

program: any MBE speaking QCSchema



protocols: customize MBE return layout

ManyBodyProtocols

component_results: save cluster data

extras: free-form storage

driver: MBE target derivative

- energy
- gradient
- Hessian

keywords: knobs in the MBE program

ManyBodyKeywords

specification: how to run MBE subsystems

- single

AtomicSpecification

- mapping

{hi: **AtomicSpecification**,

lo: **AtomicSpecification**}

ManyBodyInput

schema_version: 1

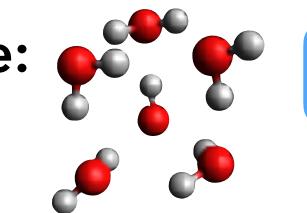
provenance: creator name and version info

QCManyBody, 0.3.0

id: tracker for databases

extras: free-form storage

molecule:

**Molecule**

specification: how to run MBE single-point

ManyBodySpecification

QCSchema MANYBODY

driving the many-body expansion

$$E^M = E^1 + E^{(2)} + E^{(3)} + \dots + E^{(M)},$$

$$= \sum_{I \in \mathcal{P}_1} E^I + \sum_{IJ \in \mathcal{P}_2} E^{(IJ)} + \sum_{IJK \in \mathcal{P}_3} E^{(IJK)} + \dots,$$

- **RECENT** added in 2024
- Counterpoise, VMFC, and no-CP arbitrary-order manybody expansion (**MBE**) extracted from Psi4 program
- **SUITABLE** for intermolecular fragments, no bond-breaking

ManyBodyProperties (abr.)

calcinfo_nfr: M

return_energy: E •

return_gradient: G 

bsse_corrected_interaction_egh_through_m_body:

- defined as E_{IE}^m

- defined for all bsse as bsse_type allows

- defined for all egh as driver allows

- def. for all m as max_nbody/levels allows

bsse_corrected_total_egh_through_m_body:

- defined as E^m

bsse_corrected_m_body_contribution_to_egh:

- defined as $E^{(m)}$

ManyBodyResult

schema_version: 1

input_data: record of starting conditions

ManyBodyInput

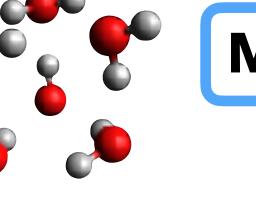
schema_version: 1

provenance: creator name and version info

QCManyBody, 0.3.0

id: tracker for databases

extras: free-form storage

molecule:  **Molecule**

specification: how to run MBE single-point

ManyBodySpecification

success: True

provenance: creator, version, & RT info

QCManyBody, 0.4.0, psinet, CPU E5-2630

id: tracker for databases

stdout: program's text output for humans

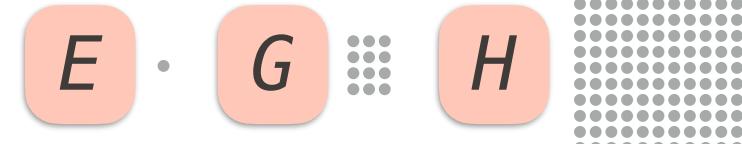
stderr: anything in error channel

native_files: requested raw program files

extras: free-form storage

molecule: orientation & frame for results

return_result: MBE derivative



properties: key results for MBE

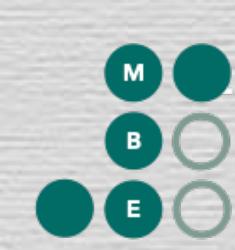
ManyBodyProperties

cluster_properties: key results for subsystems

{cluster id: **AtomicProperties**}

cluster_results: full results for subsystems

{cluster id: **AtomicResult**}

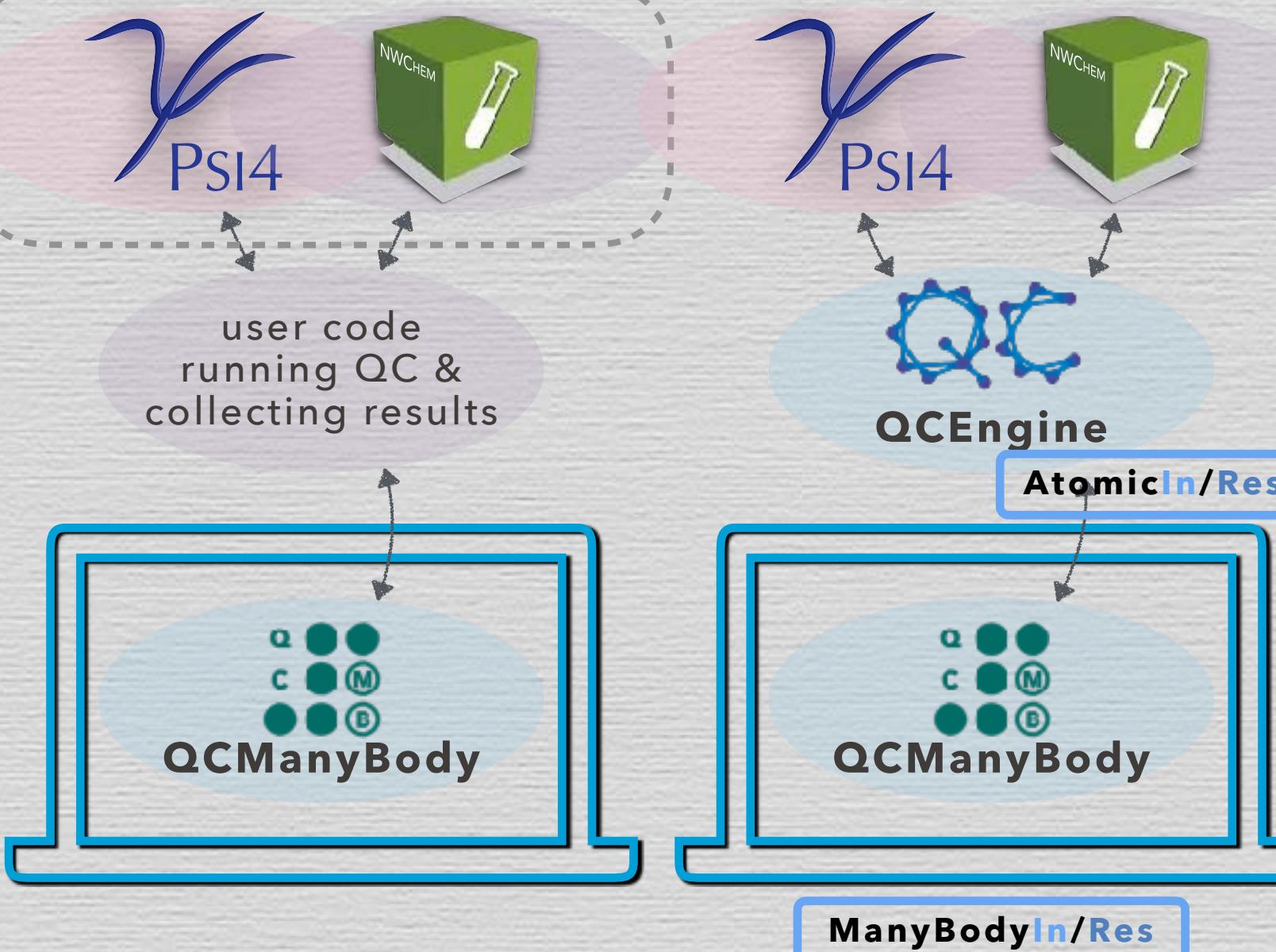


INTEGRATING QC MANYBODY

multiple entry points for flexibility

- ★ prepare multiple entry points – API, driver, driven
- ★ find your good, targeted code & set it free for broader use
- ★ avoid QCA or Aiida workflow silos

any CMS program producing
E/G/H/property

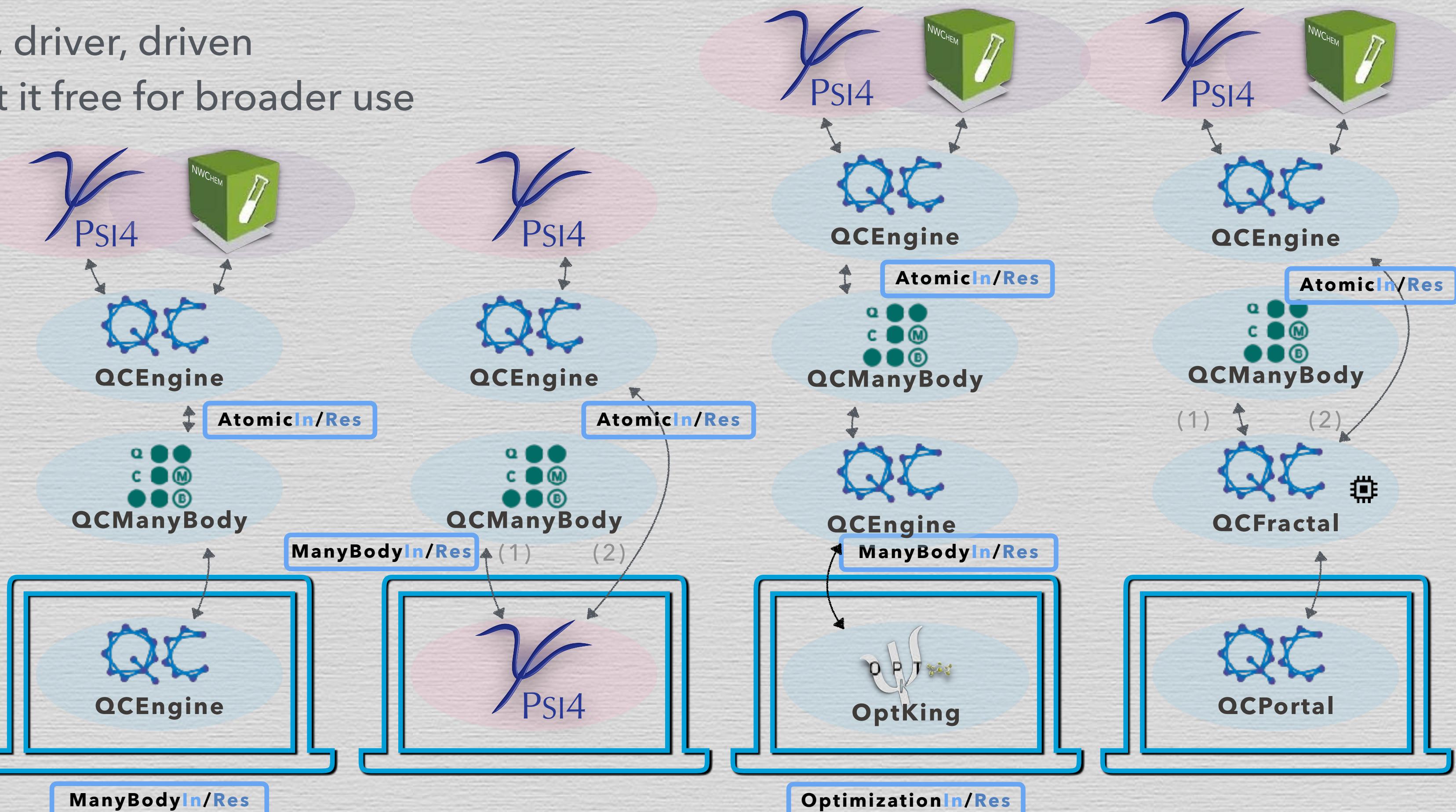


QCManyBody core interface

- user-prov. serial/parallel
- user runs CMS backends

QCManyBody high-level interface

- serial
- multi CMS backends



QC Engine procedure

- serial
- multi CMS backends
- only Psi4 backend

Psi4 distributed driver

- serial or parallel
- multi CMS backends

Optimizer backend

- serial
- multi CMS backends

QC Fractal service

- parallel
- multi CMS backends



STANDARD SUITE

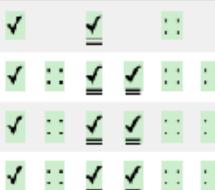
a framework for systematic testing

SOMEONE CONTRIBUTES A MODULE ADDING OR AFFECTING A METHOD

- Does it adhere to **QCSchema** inputs & returns?
- Is it returning as many **PROPERTIES** as possible?
- Does it **MATCH** other modules or programs?
- Does it work with all **REFERENCES** or give a sensible error message?
- Does it work for all **DERIVATIVES** and match finite difference or give a sensible error message?
- Is the intended module getting called by **DEFAULT**?

MOST QC TESTS ARE FOR CORRECTNESS OF SENSIBLE INPUTS

- Many QC calcs aren't run by specialists anymore. Could be naive users, workflows, or AI. Best to detect **UNFEASIBLE** calcs & exit ASAP.
- Role for tests with **SYSTEMATIC** coverage of many QC methods
- Help new contributions check correctness & **INTEGRATION** with Psi4 plumbing. Currently 3 mols; could add more to handle edge cases like ghost atoms or no beta e- or new modules like ECP.
- **REFERENCE** values & QCSchema properties contracts shared by Psi4, QCEngine, QCDB



STANDARD SUITE

PROGRAM METHOD & DERIV



5132 + 671

(CC|CC)

1158



642



496



66 tests

```
# <<< HF Energy
# <<< HF Gradient
# <<< HF Hessian
# <<< MP2 Energy
# <<< MP2 Gradient
# <<< MP2 Hessian
# <<< MP2.5 Energy
# <<< MP2.5 Gradient
# <<< MP3 Energy
# <<< MP3 Gradient
# <<< MP4(SDQ) Energy
# <<< MP4 Energy
# <<< ZAPT2 Energy
# <<< CISD Energy
# <<< QCISD Energy
# <<< QCISD(T) Energy
# <<< FCI Energy
# <<< REMP2 Energy
# <<< LCCD Energy
# <<< LCCD Gradient
# <<< LCCSD Energy
# <<< CEPA(1) Energy
# <<< CEPA(3) Energy
# <<< ACPF Energy
# <<< AQCC Energy
# <<< CCD Energy
# <<< CCD Gradient
# <<< BCCD Energy
# <<< CC2 Energy
# <<< CC2 Gradient
# <<< CCSD Energy
# <<< CCSD Gradient
# <<< CCSD(T) Energy
# <<< CCSD(T) Gradient
# <<< a-CCSD(T) Energy
# <<< BCCD(T) Energy
# <<< CC3 Energy
# <<< OMP3 Energy
# <<< OMP3 Gradient
# <<< OREMP2 Energy
# <<< OREMP2 Gradient
# <<< OLCCD Energy
# <<< OLCCD Gradient
# <<< MP2-F12 Energy
# <<< SVWN Energy
# <<< SVWN Gradient
# <<< SVWN Hessian
# <<< PBE Energy
# <<< PBE Gradient
# <<< B3LYP Energy
# <<< B3LYP Gradient
# <<< wB97X Energy
# <<< wB97X Gradient
# <<< B2PLYP Energy
# <<< B2PLYP Gradient
```

58 mtd+deriv

REFERENCE, INTEGRALS, MODULE

[ccsd rhf conv fc : * ccenergy-adz-ene0]	PASSED	[2%]
[ccsd uhf conv fc : * ccenergy-adz-ene0]	PASSED	[4%]
[ccsd rohf conv fc sd: * ccenergy-adz-ene0]	PASSED	[6%]
[ccsd rohf conv fc sc: ccenergy-adz-ene0]	PASSED	[8%]
[ccsd rhf conv ae : * ccenergy-adz-ene0]	PASSED	[10%]
[ccsd uhf conv ae : * ccenergy-adz-ene0]	PASSED	[13%]
[ccsd rohf conv ae sd: * ccenergy-adz-ene0]	PASSED	[15%]
[ccsd rohf conv ae sc: ccenergy-adz-ene0]	PASSED	[17%]
[ccsd rhf conv fc : mrcc -adz-ene0]	SKIPPED (No mrcc)	[19%]
[ccsd uhf conv fc : mrcc -adz-ene0]	SKIPPED (No mrcc)	[21%]
[ccsd rohf conv fc sc: mrcc -adz-ene0]	SKIPPED (No mrcc)	[23%]
[ccsd rhf conv ae : mrcc -adz-ene0]	SKIPPED (No mrcc)	[26%]
[ccsd uhf conv ae : mrcc -adz-ene0]	SKIPPED (No mrcc)	[28%]
[ccsd rohf conv ae sc: mrcc -adz-ene0]	SKIPPED (No mrcc)	[30%]
[ccsd rhf conv fc : fnocc -adz-ene0]	PASSED	[32%]
[ccsd rhf conv ae : fnocc -adz-ene0]	PASSED	[34%]
[ccsd rhf df fc : * fnocc -adz-ene0]	PASSED	[36%]
[ccsd rhf df ae : * fnocc -adz-ene0]	PASSED	[39%]
[ccsd rhf pk/df fc : fnocc -adz-ene0]	PASSED	[41%]
[ccsd rhf pk/df ae : fnocc -adz-ene0]	PASSED	[43%]
[ccsd rhf cd/df fc : fnocc -adz-ene0]	PASSED	[45%]
[ccsd rhf cd/df ae : fnocc -adz-ene0]	PASSED	[47%]
[ccsd rhf cd fc : * fnocc -adz-ene0]	PASSED	[50%]
[ccsd rhf cd ae : * fnocc -adz-ene0]	PASSED	[52%]
[ccsd rhf pk/df fc : fnocc -adz-ene0]	PASSED	[54%]
[ccsd rhf pk/df ae : fnocc -adz-ene0]	PASSED	[56%]
[ccsd rhf df fc : dfocc -adz-ene0]	PASSED	[58%]
[ccsd uhf df fc : dfocc -adz-ene0]	PASSED	[60%]
[ccsd rhf df ae : dfocc -adz-ene0]	PASSED	[63%]
[ccsd uhf df ae : dfocc -adz-ene0]	PASSED	[65%]
[ccsd rhf pk/df fc : dfocc -adz-ene0]	PASSED	[67%]
[ccsd uhf pk/df fc : dfocc -adz-ene0]	PASSED	[69%]
[ccsd rhf pk/df ae : dfocc -adz-ene0]	PASSED	[71%]
[ccsd uhf pk/df ae : dfocc -adz-ene0]	PASSED	[73%]
[ccsd rhf cd/df fc : dfocc -adz-ene0]	PASSED	[76%]
[ccsd uhf cd/df fc : dfocc -adz-ene0]	PASSED	[78%]
[ccsd rhf cd/df ae : dfocc -adz-ene0]	PASSED	[80%]
[ccsd uhf cd/df ae : dfocc -adz-ene0]	PASSED	[82%]
[ccsd rhf cd fc : dfocc -adz-ene0]	PASSED	[84%]
[ccsd uhf cd fc : dfocc -adz-ene0]	PASSED	[86%]
[ccsd rhf cd ae : dfocc -adz-ene0]	PASSED	[89%]
[ccsd uhf cd ae : dfocc -adz-ene0]	PASSED	[91%]
[ccsd rhf pk/df fc : dfocc -adz-ene0]	PASSED	[93%]
[ccsd uhf pk/df fc : dfocc -adz-ene0]	PASSED	[95%]
[ccsd rhf pk/df ae : dfocc -adz-ene0]	PASSED	[97%]
[ccsd uhf pk/df ae : dfocc -adz-ene0]	PASSED	[100%]

5132 pytests

RETURN, CALCINFO, PROPERTIES

[ccsd rhf pk/cd ae] wfn.....	PASSED
[ccsd rhf pk/cd ae] return.....	PASSED
[ccsd rhf pk/cd ae] nbasis wfn.....	PASSED
[ccsd rhf pk/cd ae] nmo wfn.....	PASSED
[ccsd rhf pk/cd ae] nalpha wfn.....	PASSED
[ccsd rhf pk/cd ae] nbeta wfn.....	PASSED
[ccsd rhf pk/cd ae] wfn HF TOTAL ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn HF TOTAL ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 CORRELATION ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 CORRELATION ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 TOTAL ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 TOTAL ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 SAME-SPIN CORRELATION ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 SAME-SPIN CORRELATION ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 SINGLES ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 SINGLES ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 DOUBLES ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 DOUBLES ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn MP2 OPPOSITE-SPIN CORRELATION ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn MP2 OPPOSITE-SPIN CORRELATION ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn HF TOTAL ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn HF TOTAL ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CCSD CORRELATION ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CCSD CORRELATION ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CCSD TOTAL ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CCSD TOTAL ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CCSD SAME-SPIN CORRELATION ENERGY	SKIP
[ccsd rhf pk/cd ae] wfn CCSD SINGLES ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CCSD SINGLES ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CCSD DOUBLES ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CCSD DOUBLES ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CCSD OPPOSITE-SPIN CORRELATION ENERGY	SKIP
[ccsd rhf pk/cd ae] wfn CURRENT CORRELATION ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CURRENT CORRELATION ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CURRENT ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CURRENT ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn SCF TOTAL ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn SCF TOTAL ENERGY v. CONV	PASSED
[ccsd rhf pk/cd ae] wfn CURRENT REFERENCE ENERGY	PASSED
[ccsd rhf pk/cd ae] wfn CURRENT REFERENCE ENERGY v. CONV	PASSED

7700 reference values



COMMUNICATING QC CAPABILITIES

a modular framework complicates the question and the answer

USER: IS A METHOD AVAILABLE?

MP2?
yes

open-shell?
sometimes

density-fitted?
most always

frozen-core?
usually

gradients?
often

by fast-coder?
depends

ECPs ok?
usually

w/ solvation?
possibly

any special
keyword to
access?

LORI: ANSWERS BASED ON PSI4 AS A WHOLE



COMMUNICATING QC CAPABILITIES

a modular framework complicates the question and the answer

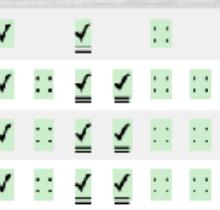
USER: IS A METHOD AVAILABLE?



LORI: ANSWERS BASED ON PSI4 AS A WHOLE

USER: NO, THE INTERSECTION OF MY REQUIREMENTS

- Answers to this usually belong in documentation, but local code PRs rarely revise broad documentation.
- Since the Standard Suite loops over many of these QC characteristics, as well as analytic vs. findif derivatives and default vs. directed modules, PSI4 leverages the large quantity of runs to robustly report capabilities.



COMMUNICATING QC CAPABILITIES

auto-generate documentation tables from Standard Suite tests

Detailed capabilities of the DFMP2 module. "✓" runs analytically. Single underline "✓" is default module when [QC_MODULE](#) unspecified. Double underline "✓" is default algorithm type when type selector (e.g., [CC_TYPE](#)) unspecified.

		QC_MODULE=DFMP2 Capabilities																	
		Restricted (RHF)						Unrestricted (UHF)						Restricted Open (ROHF)					
REFERENCE →		energy()			gradient() ^[3]			energy()			gradient() ^[3]			energy()			gradient() ^[3]		
name ↓ →		CV	DF	CD	CV	DF	CD	CV	DF	CD	CV	DF	CD	CV	DF	CD			
type ^[1] ↓ →		A	F	A	F	A	F	A	F	A	F	A	F	A	F	A			
FREEZE_CORE ^[2] →		A	F	A	F	A	F	A	F	A	F	A	F	A	F	A			
mp2 ^[4]	MP2_TYPE	✓	✓		✓	✓		✓	✓		✓	✓		✓	✓				

A focused module. Double-underlines indicate DF ints and this module are always default.

- A Standard Suite run dumps info from each test into a file, then a script processes these into various human-readable & reStructuredText tables. Hook into docs build.
- Functions as single-source-of-truth for non-exotic mtd cap. in Psi4.



COMMUNICATING QC CAPABILITIES

auto-generate documentation tables from Standard Suite tests

Detailed capabilities of the SCF module. "✓" runs analytically. Single underline "≤" is default.

QC_MODULE=SCF Capabilities										
REFERENCE →		Restricted (RHF)								
name ↓ →				energy()		gradient() ^[3]		hessian() ^[4]		
type ^[1] ↓ →		CV	DF	CD	CV	DF	CD	CV	DF	CD
FREEZE_CORE ^[2] →		A	F	A	F	A	F	A	F	A
hf	SCF_TYPE	✓	≤	✓	✓	≤	✓	✓	≤	
svwn, LSDA DFT	SCF_TYPE	✓	≤	✓	✓	≤	✓	✓	≤	
pbe, GGA DFT	SCF_TYPE	✓	≤	✓	✓	≤	✓			
b3lyp, Hybrid DFT	SCF_TYPE	✓	≤	✓	✓	≤	✓			
wb97x, LRC DFT	SCF_TYPE	✓	≤			✓	✓			
b2plyp, DH DFT ^[5]	SCF_TYPE	✓	≤	✓	≤	✓	✓			

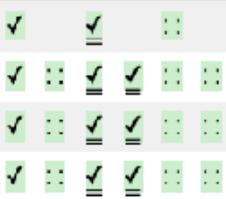
Energies available for all DFT rungs. DF default. No CD for LRC.

- A Standard Suite run dumps info from each test into a file, then a script processes these into various human-readable & reStructuredText tables. Hook into docs build.
- Functions as single-source-of-truth for non-exotic mtd cap. in Psi4.

COMMUNICATING QC CAPABILITIES

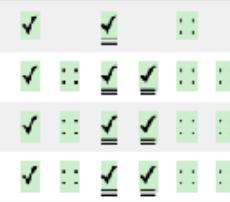
auto-generate documentation tables from Standard Suite tests

Per-module and cumulative capabilities reveal default and users' options.



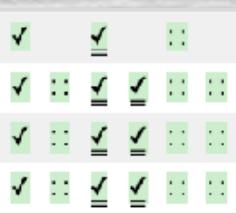
COMMUNICATING QC CAPABILITIES

auto-generate documentation tables from Standard Suite tests



I've more confidence in this whole-Psi4 capabilities table than a hand-updated one.

- A Standard Suite run dumps info from each test into a file, then a script processes these into various human-readable & reStructuredText tables. Hook into docs build.
 - Functions as single-source-of-truth for non-exotic mtd cap. in PSI4.



COMMUNICATING QC CAPABILITIES

auto-generate documentation tables from Standard Suite tests

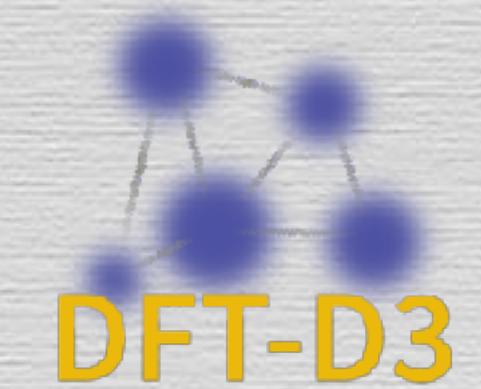
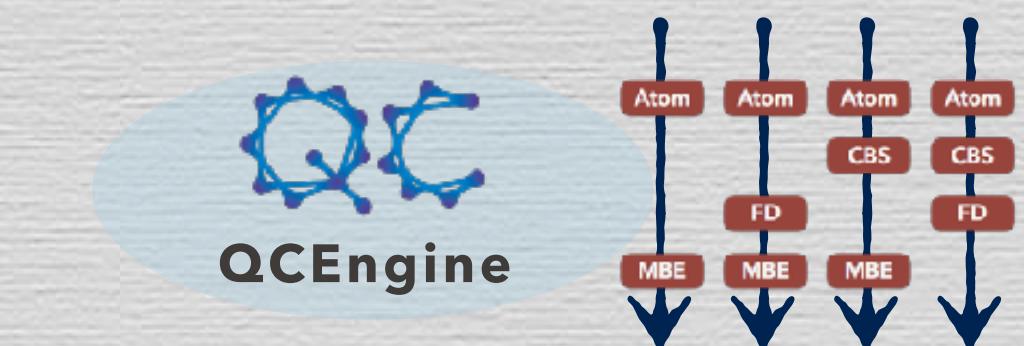
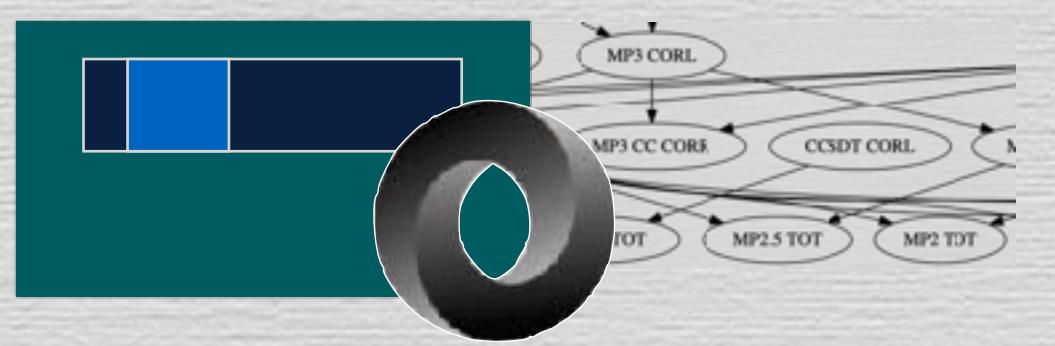
Capabilities of QCDB, including details of overlapping modules. "✓" runs analytically. "!" produces nonstandard results.									
	qc_module ↓	scf_reference →	QCDB Capabilities						
			Restricted (RHF)			Unrestricted (UHF)			Restricted Open (ROHF)
			E	G	H	E	G	H	E
		name ↓ →	A	F	A	F	A	F	A
		freeze_core ^[1] →	A	F	A	F	A	F	A
mp2	CFOUR		✓	✓	✓	✓	✓	✓	✓
	GAMESS-DDI		✓	✓	✓	✓	✓	✓	✓
	GAMESS-IMS		✓	✓	✓	✓			
	GAMESS-SERIAL		✓	✓	✓	✓	✓	✓	✓
	NWChem-DIRECTMP2		✓	✓					
	NWChem-MP2GRAD		✓	✓	✓	✓	✓	✓	✓
	NWChem-TCE		✓	✓		✓	✓		!
	Psi4-FNOCC		✓	✓					
	Psi4-OCC		✓	✓	✓	✓	✓	✓	✓
ccsd(t)	CFOUR-ECC		✓	✓	✓	✓	✓	✓	✓
	CFOUR-NCC		✓	✓	✓	!			
	CFOUR-VCC		✓	✓	✓	✓	✓	✓	✓
	GAMESS		✓	✓					
	NWChem-CC		✓	✓					
	NWChem-TCE		✓	✓		✓	✓		!
	Psi4-CCENERGY		✓	✓		✓	✓		✓
	Psi4-FNOCC		✓	✓					
	Psi4-MRCC		✓	✓		✓	✓		✓

[1] Active orbital values to the right: all-electron A and frozen-core F.

Also useable across QC programs to check answers and show interfaced capabilities.

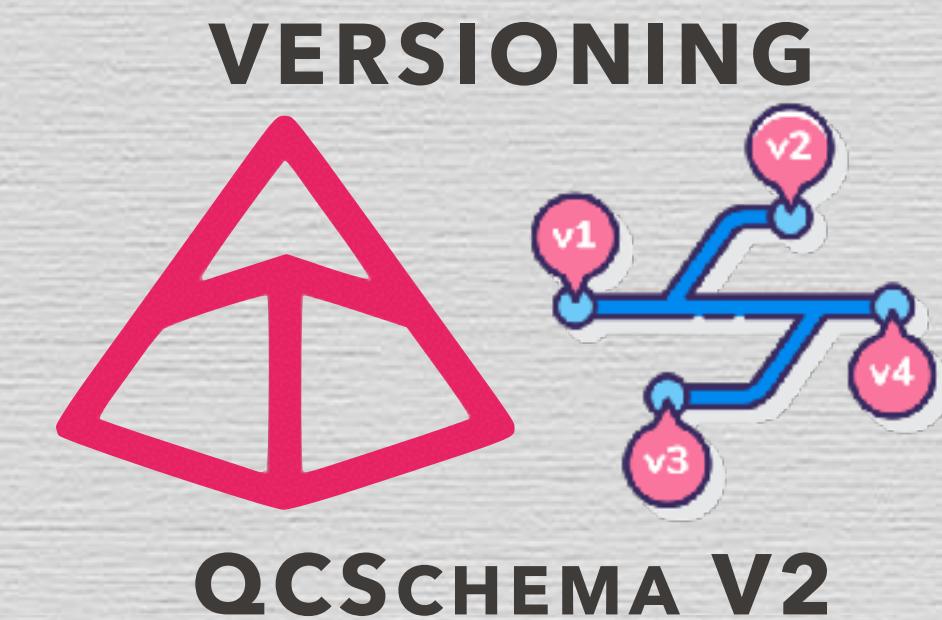
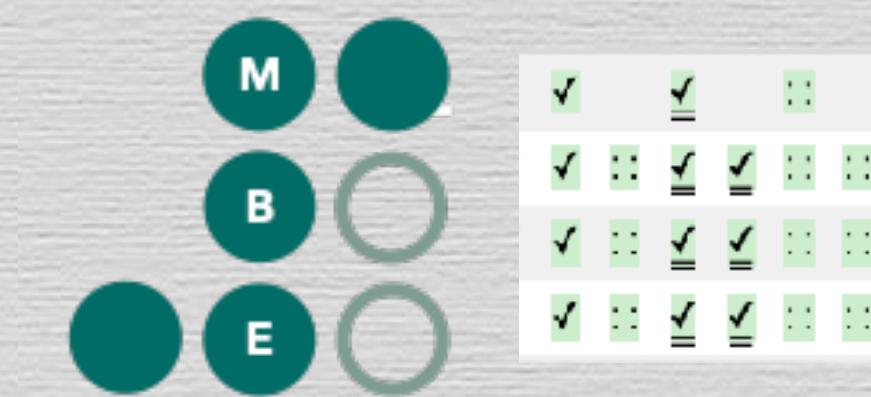
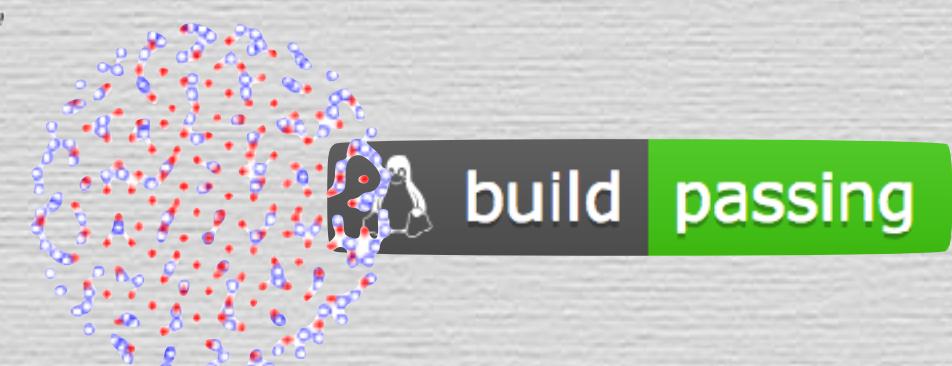
★ Automate the assessment & conveyance of detailed capabilities to users, lest contrib. from alternate modules become overwhelming

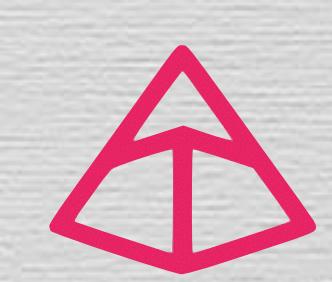
- A Standard Suite run dumps info from each test into a file, then a script processes these into various human-readable & reStructuredText tables. Hook into docs build.
- Functions as single-source-of-truth for non-exotic mtd cap. in Psi4.



PSI4

OPEN-SOURCE QUANTUM CHEMISTRY

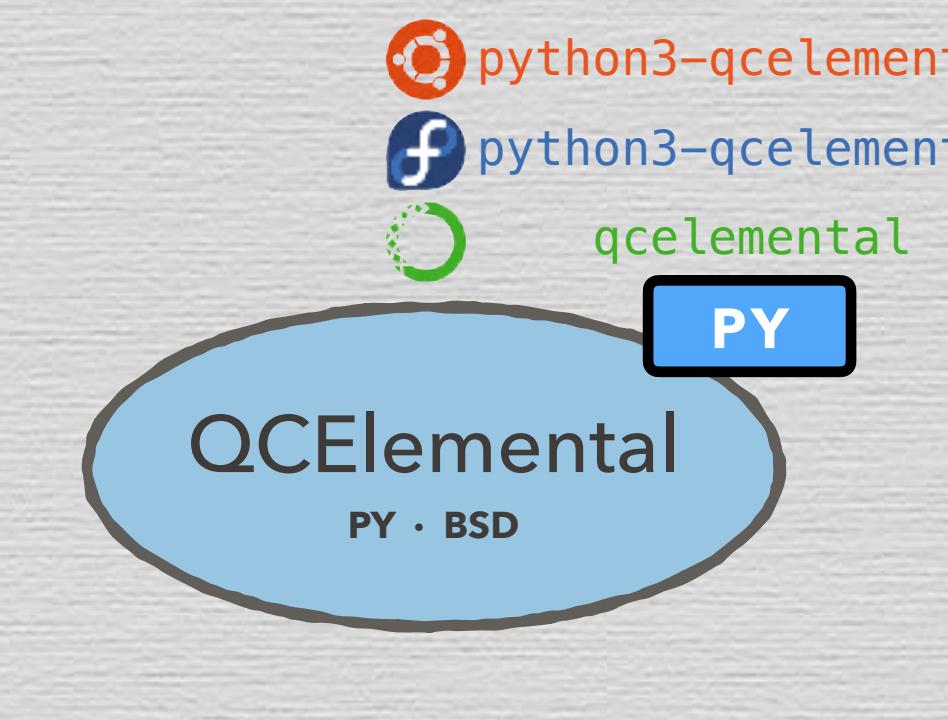




A QCSchema UPGRADE STORY

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AtomicResult

```

AtomicInput
schema_version: 1
provenance: creator, version, & RT info
@QCElemental, 0.28.0
id: tracker for databases
extras: free-form storage
molecule: CN(C)C CC(=O)OC CC1=CC=C1 CC1=CC=C1

protocols: customize opt return layout
  AtomicResultProtocols
    stdout: save text output
    wavefunction: save orbital info
    native_files: save raw prog. files
    error_correction: auto-edit&rerun

driver: single-point derivative
  - energy
  - gradient: E G H
  - Hessian
  - properties

model: model chemistry or FF
  method: B3LYP-D3, CCSD(T)
  basis: 6-31G*, cc-pV5Z BasisSet
  keywords: knobs in SP program
    (scf_type: ulf, cc_conv: 7, docc: [4,1])

success: True
provenance: (overwrites input)
  xtb, 1.0.0, psinet, CPU E5-2630

stdout: program's text output for humans
stderr: anything in error channel
native_files: requested raw program files
extras: (merges with input)
molecule: (overwrites input w/result from previous)
return_results: single-point derivative
  E G H

```

OptimizationInput

```

schema_version: 1
provenance: creator name and version info
@QCElemental, 0.28.0
id: tracker for databases
extras: free-form storage
initial_molecule: Molecule
  CN(C)C CC(=O)OC CC1=CC=C1 CC1=CC=C1

TDKeywords
dihedrals: dimensions to scan
dihedral_ranges: limits of scan
grid_spacing: 1E dihedral steps of scan
energy_decrease_thresh: point trigger
energy_upper_limit: new point trigger
optimization_specs: how to run geometry opts

OptimizationSpecification
schema_version: 1
procedure: any optimizer in QCEngine
  qchem, psi4, pyberry
protocols: customize opt return layout
  OptimizationProtocols
    trajectory: save fewer opt steps

keywords: knobs in the geo opt program
  (coordsys: tric, program: qchem, pyberry)
input_specification: how to run gradients for opt

QCInputSpecification
schema_version: 1

extra: free-form storage
driver: single-point derivative
  - energy
  - gradient: E G H
  - Hessian
  - properties

model: model chemistry or FF
  method: B3LYP-D3, CCSD(T)
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    (scf_type: ulf, cc_conv: 7, docc: [4,1])

extra: free-form storage
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  - energy
  - gradient: E G H
  - Hessian
  - properties

model: model chemistry or FF
  method: B3LYP-D3, CCSD(T)
  basis: 6-31G*, cc-pVQZ BasisSet
  keywords: knobs in SP program
    (scf_type: ulf, cc_conv: 7, docc: [4,1])

```

TorsionDriveInput

```

schema_version: 1
provenance: creator name and version info
@QCElemental, 0.28.0
ComputeError
extras: free-form storage
initial_molecule: Molecule
  CN(C)C CC(=O)OC CC1=CC=C1 CC1=CC=C1

TorsionDriveProperties (nyi)

calinfo_ngrid
nuclear_repulsion_energy
return_energy: E

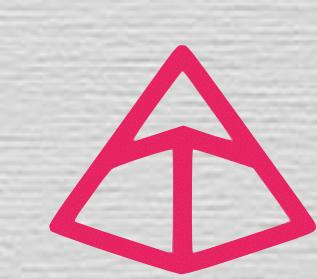
```

TDKeywords

OptimizationSpecification

QCInputSpecification

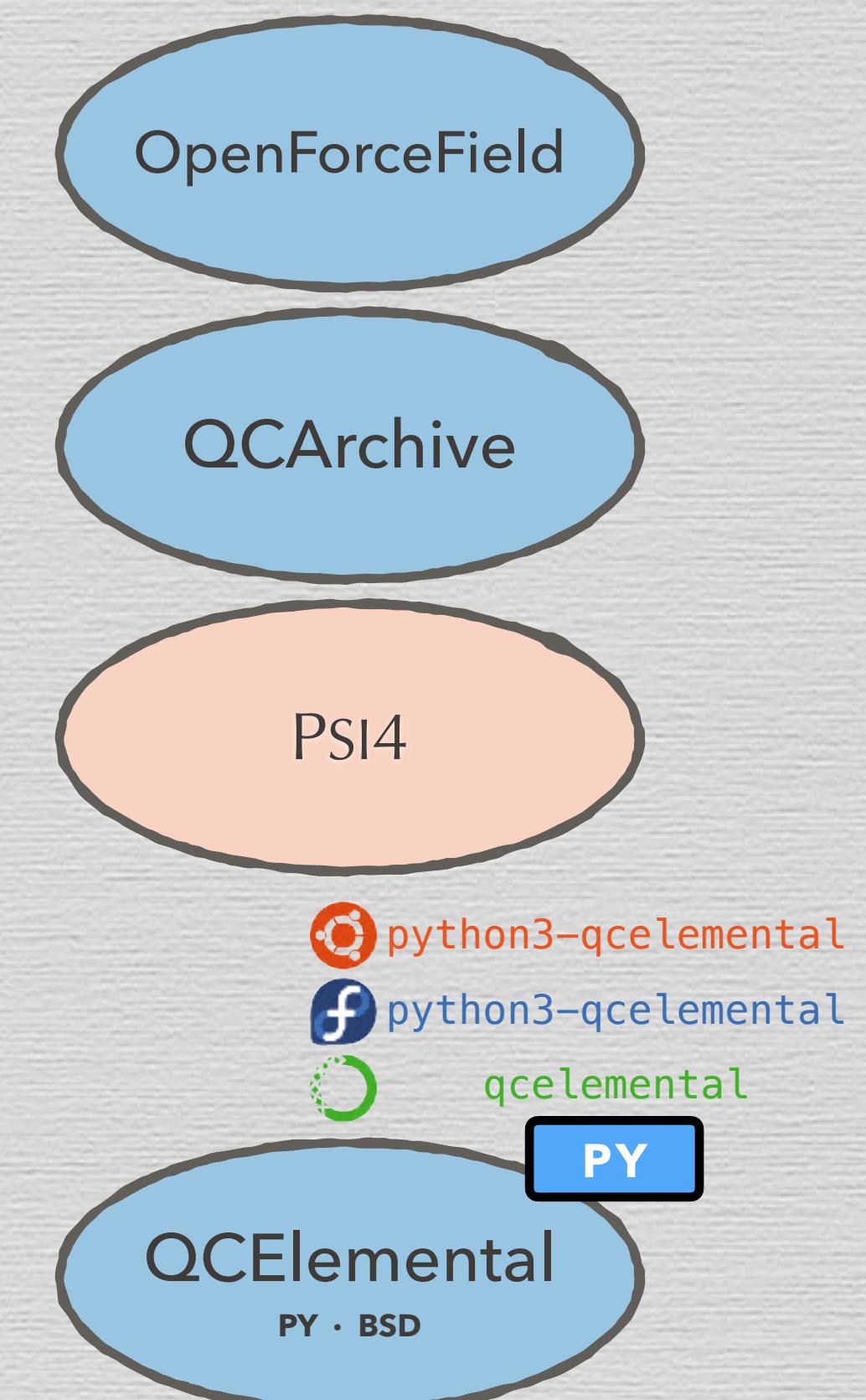
Overlapping or overwritten schema shown by color variations

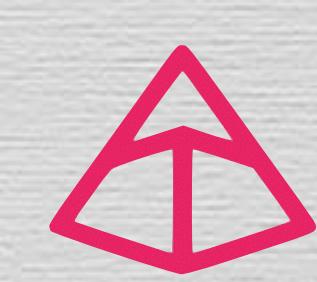


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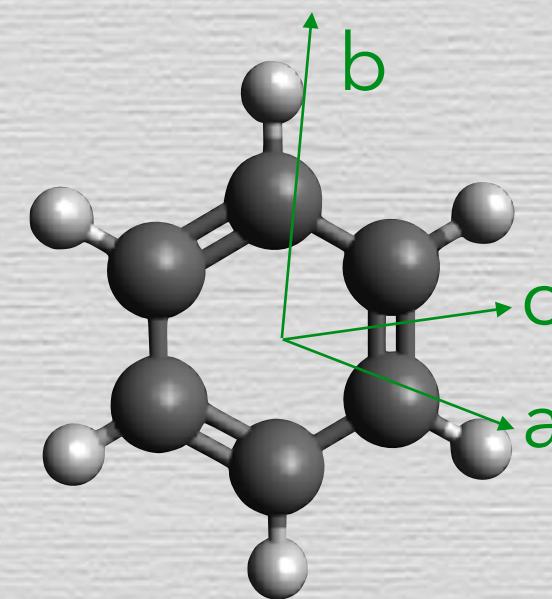
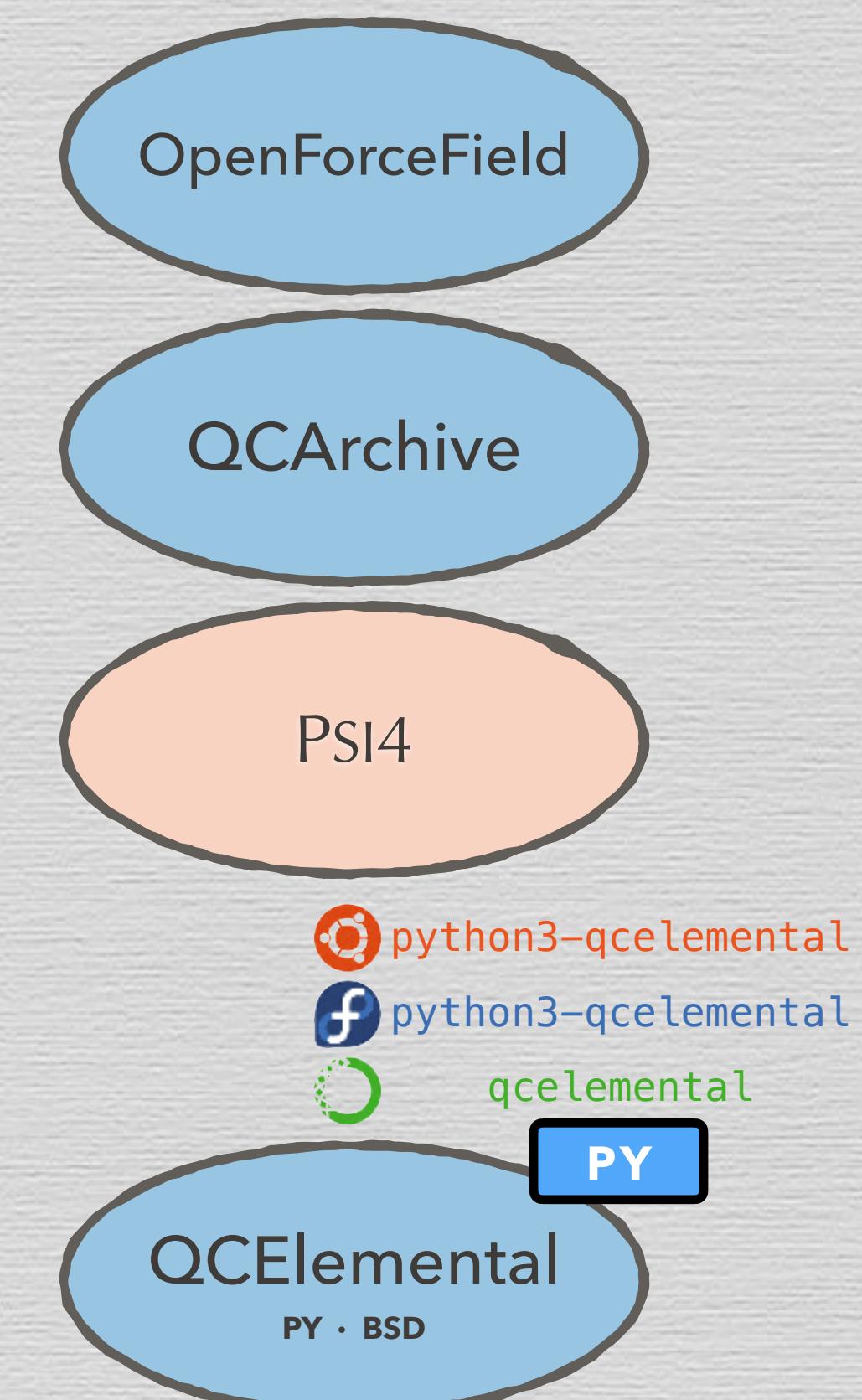


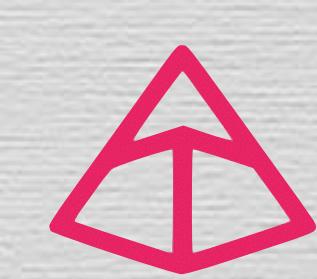


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- **PYDANTIC V2** API released June 2023, which is a mild upgrade except it forbids mixing v1/v2 models, requiring concurrent full-stack migration.
- At a CECAM meeting last year, I learned that QCSchema would be useful to a new community if only **Molecule** included **UNIT CELL** specification.



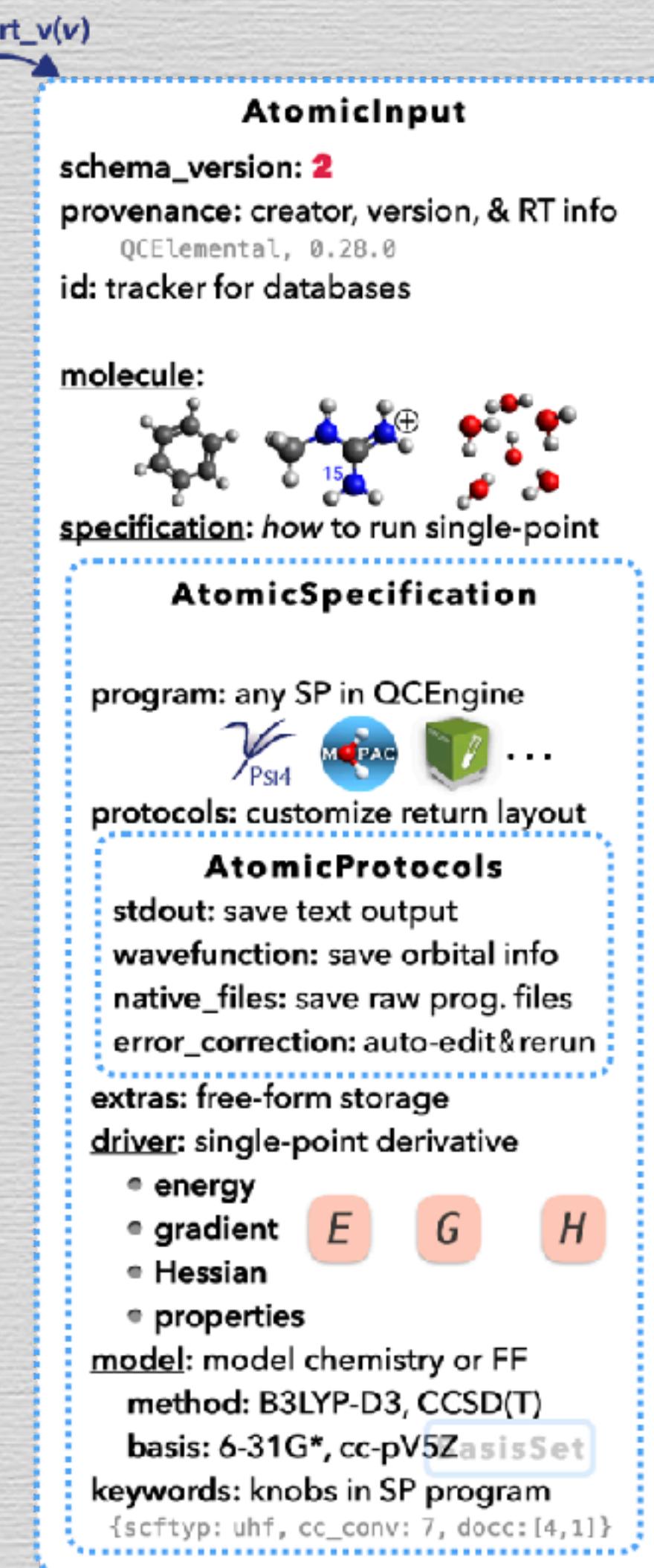
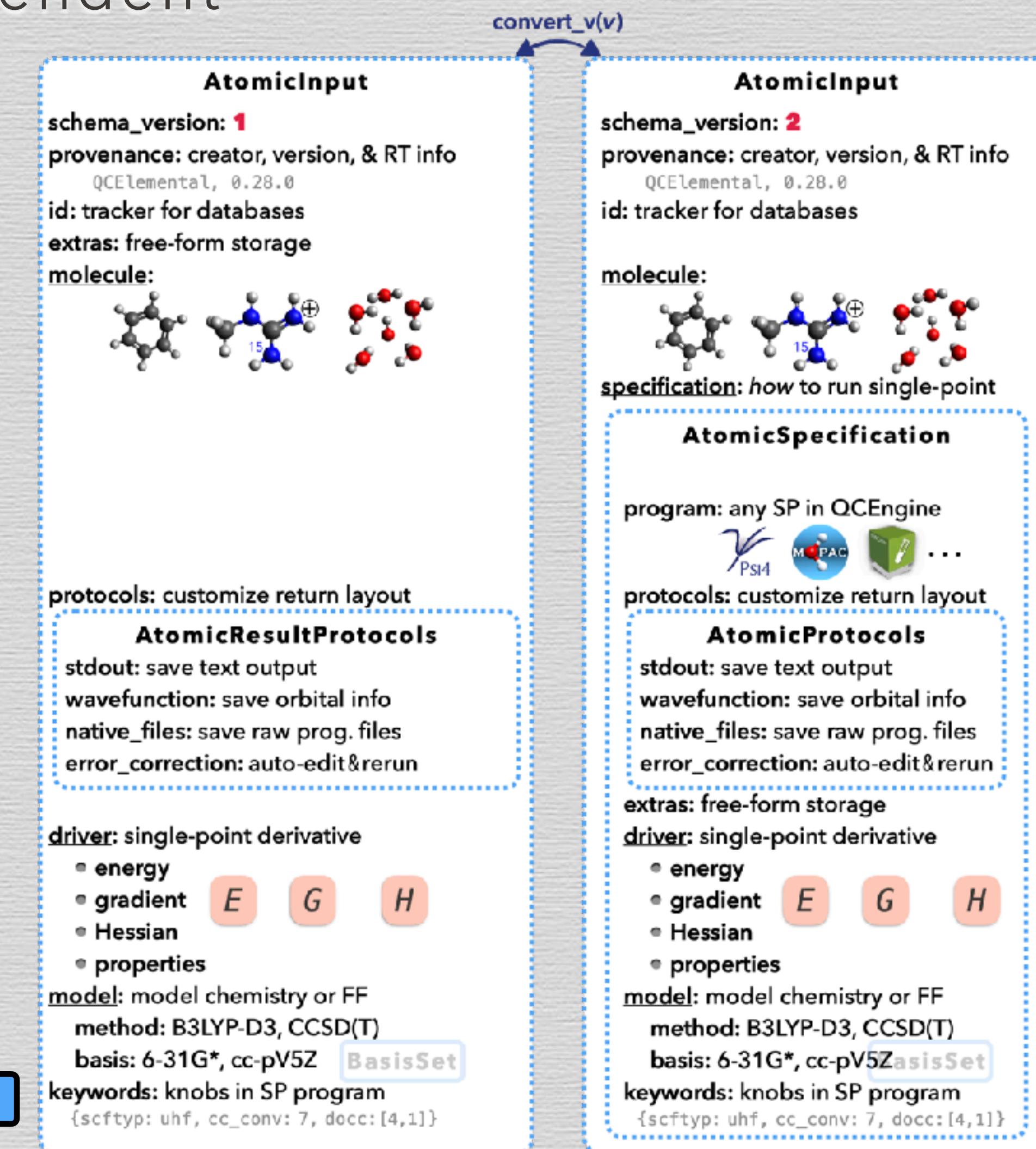


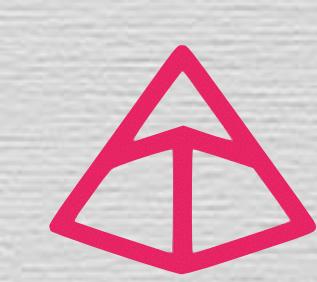
A QCSchema UPGRADE STORY

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Technical not content, upgrade to **v2** in Fall 2024

- Refactored **AtomicInput/Result** & procedure schema layouts to be more composable, reuseable, & intuitive.
- Updated schema v2 models to Pydantic v2 API and provided converter functions.





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ACCESS QCSchema v1

(PYDANTIC v1 API; c. 2019 field layout)

```
# longstanding until v0.70
> from qcelemental.models import Molecule

# longstanding until v0.70 but remove ASAP!
> from qcelemental.models.molecule import Molecule

# v0.50a1 until v1.0
> from qcelemental.models.v1 import Molecule
```

ACCESS QCSchema v2

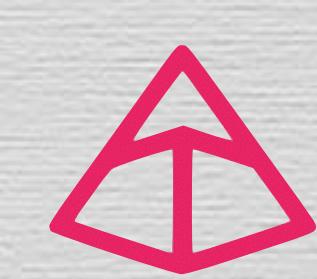
(PYDANTIC v2 API; c. 2024 field layout)

```
# v0.50a1 onwards
> from qcelemental.models.v2 import Molecule

# v0.70a1 onwards
> from qcelemental.models import Molecule
```

*Either v1 or v2 immediately accessible,
but no oblivious v1 code will break for 1y*



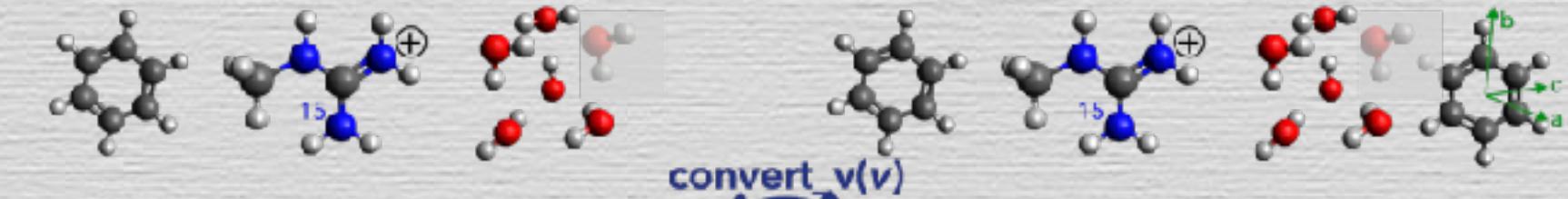


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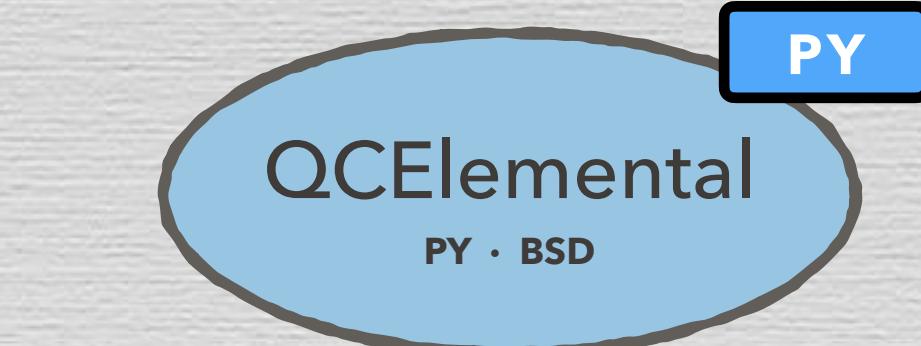
Technical note content, upgrade to **v2** in Fall 2024

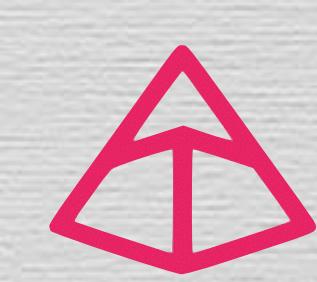
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- Foundational **Molecule** & **BasisSet** will receive small but useful additions from expressed need.



Molecule	Molecule
<code>schema_version: 2 (used w/QCSk v1)</code>	<code>schema_version: 3 (used w/QCSk v2)</code>
<code>provenance: creator info</code>	<code>provenance: creator info</code>
<code>id: tracker for databases</code>	<code>id: tracker for databases</code>
<code>extras: free-form storage</code>	<code>extras: free-form storage</code>
<code>name: user label</code>	<code>name: user label</code>
<code>comment: further user text</code>	<code>comment: further user text</code>
<code>symbols: elemental symb. for atoms</code>	<code>symbols: elemental symb. for atoms</code>
<code>geometry: Cartesian coord. in Bohr</code>	<code>geometry: Cartesian coord. in Bohr</code>
<code>molecular_charge: net elst. charge</code>	<code>molecular_charge: net elst. charge</code>
<code>molecular_multiplicity: total mult.</code>	<code>molecular_multiplicity: total mult.</code>
<code>masses: per atom atomic masses</code>	<code>masses: per atom atomic masses</code>
<code>real: per atom real vs. ghost/virtual</code>	<code>real: per atom real vs. ghost/virtual</code>
<code>atom_labels: further per atom labels</code>	<code>atom_labels: further per atom labels</code>
<code>atomic_numbers: per atom numbers</code>	<code>atomic_numbers: per atom numbers</code>
<code>mass_numbers: per atom isotope #</code>	<code>mass_numbers: per atom isotope #</code>
<code>fragments: group atoms into fragmts</code>	<code>fragments: group atoms into fragmts</code>
<code>fragment_charges: partitioned chg</code>	<code>fragment_charges: partitioned chg</code>
<code>fragment_multiplicities: ptn'd mult</code>	<code>fragment_multiplicities: ptn'd mult</code>
<code>fix_com: conserve input COM</code>	<code>fix_com: conserve input COM</code>
<code>fix_orientation: conserve input frame</code>	<code>fix_orientation: conserve input frame</code>
<code>fix_symmetry: use lesser point group</code>	<code>fix_symmetry: use lesser point group</code>
<code>connectivity: bond info</code>	<code>connectivity: bond info</code>
<code>validated: passed physics checks</code>	<code>validated: passed physics checks</code>
<code>identifiers: registration/classification</code>	<code>identifiers: registration/classification</code>
<code>Identifiers</code>	

For technical reasons, Molecule v2 needs a difference, and periodic box vectors will be ideal.





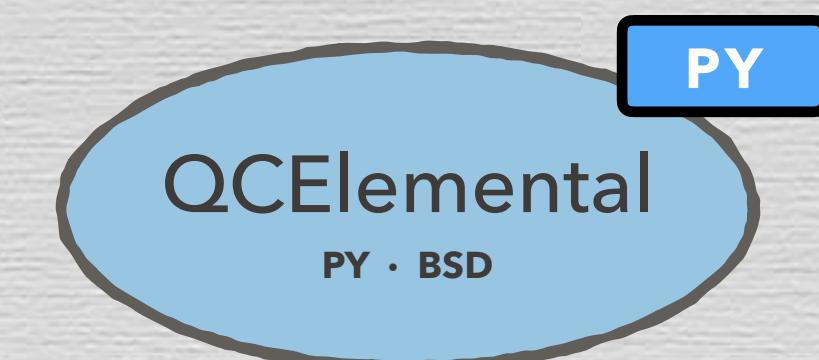
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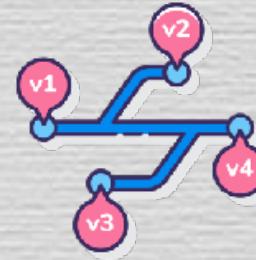
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- Foundational **Molecule** & **BasisSet** will receive small but useful additions from expressed need.
- **PAYOFF** every project gets unstuck from Pydantic stasis.
- **PAYOFF** after v2 roll-out, QCSchema will be ready to add community-requested features. Hopefully news soon.
- **PAYOFF** established pathway to upgrade in future.

- ★ modularity can mean dependence in deep software stacks that needs careful guiding
- ★ generous acceptance of feature requests can lead to high-cohesion module but beyond your comfort
- ★ research demands implementing some things quickly for downstream, lest they be blocked. Research also demands your own dependencies be stable and not interrupted by new nonsense fields. Need some governance to balance these interests.



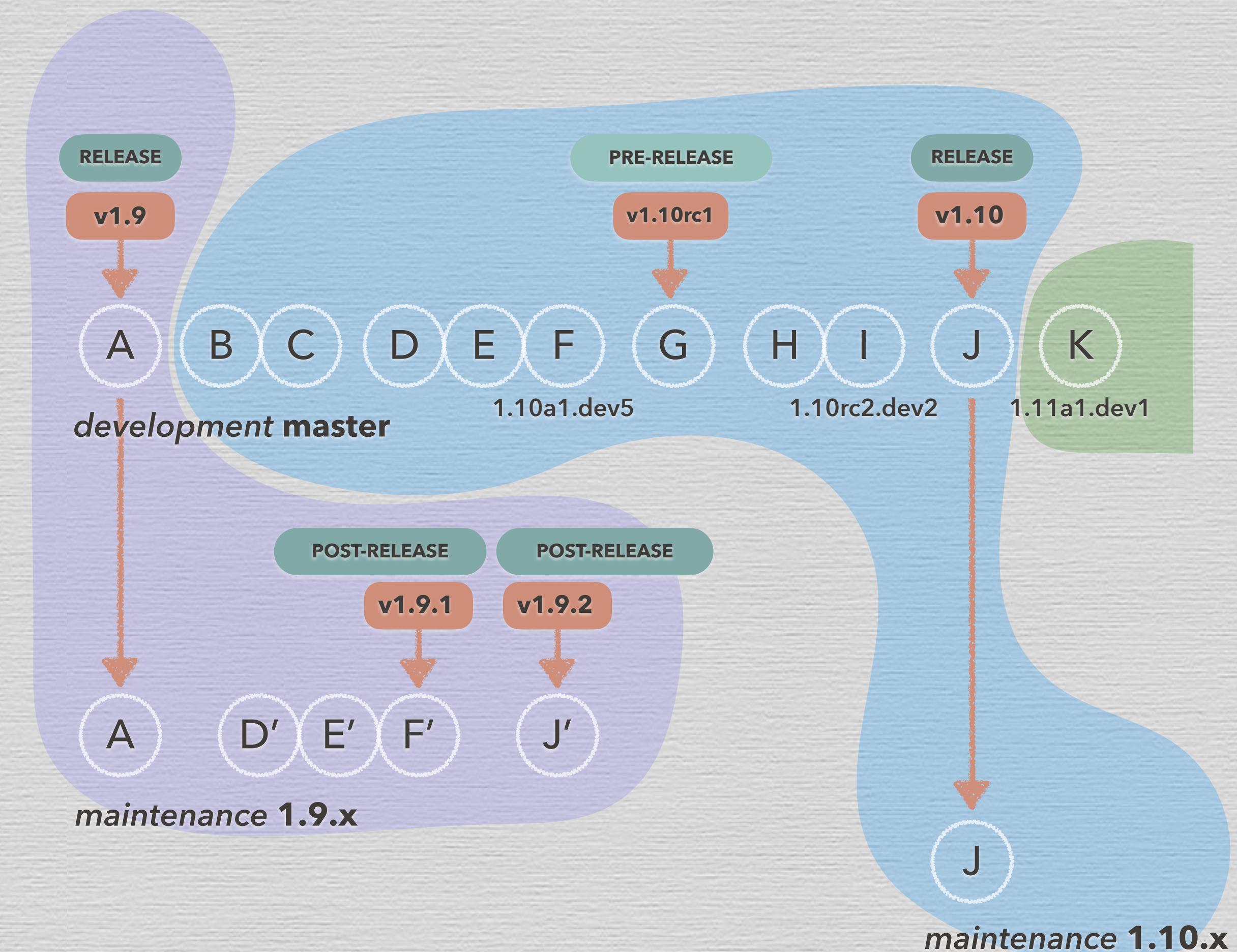
VERSIONING

ecosystems run on version compatibility



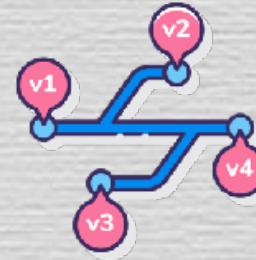
- **AUTOMATED** version reckoning after `git tag` signals a release. Upon `make`, info from `git describe` computes a unique, sortable version at every commit. Stamped into output.
- **CONDA** allows a package to advise any downstream consumers how many versions are binary compatible (`x` for semver or `x.x.x` more realistically).
- In an ecosystem arrangement, adequately tested and pushed to GitHub or even merged to master isn't sufficient for "**AVAILABLE**". A feature (or more importantly a bugfix or buildfix) isn't avail. until it's in a released version (or patched binary dist.) because far downstream it takes too much intervention to use a specialty changeset.
- Incline toward mercy on thine downstream and don't just fix things quickly but **PUBLISH** them.

VERSION COMPUTED EVERY `cmake --build` FROM GIT INFO



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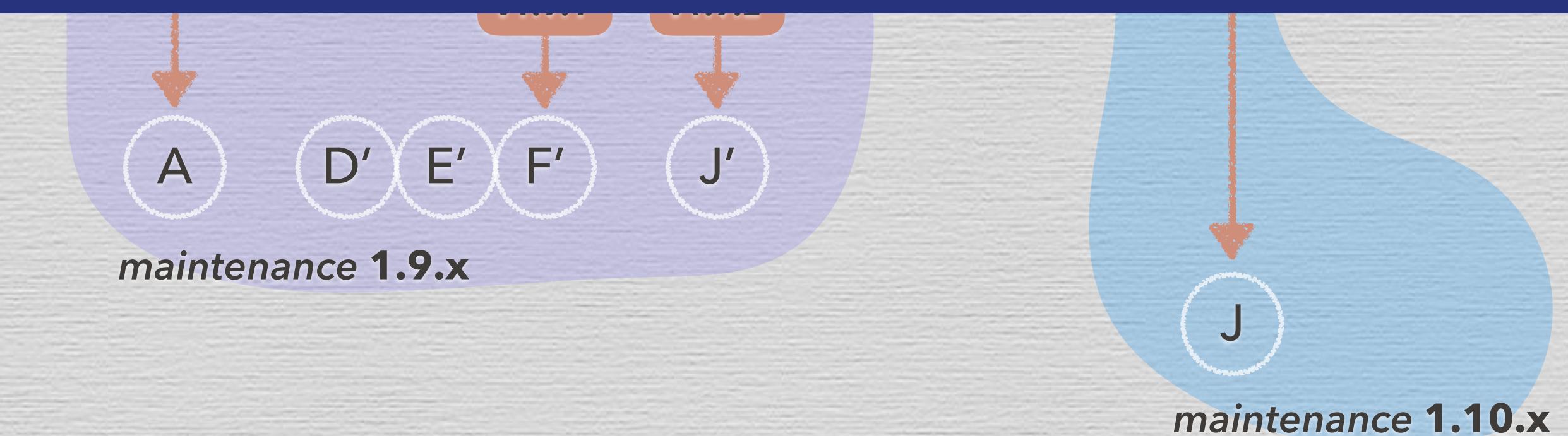
VERSION COMPUTED EVERY `cmake --build` FROM GIT INFO

```
import psi4
print(psi4.__version__)
#> 1.10a1.dev119

print(psi4.version_formatter())
#> 1.10a1.dev119

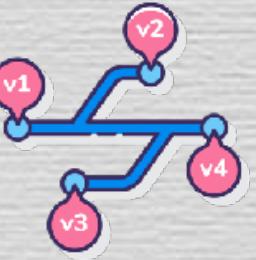
print(psi4.version_formatter('all'))
#> 1.10a1.dev119 {f12_draft_rb4} e2cc1fd 1.9.0.999 dirty 1.9 <-- 1.10a1.dev119+e2cc1fd

print(psi4.version_formatter('branch: {branch}\nsortable: {versionlong}'))
#> branch: f12_draft_rb4
#> sortable: 1.10a1.dev119+e2cc1fd
```

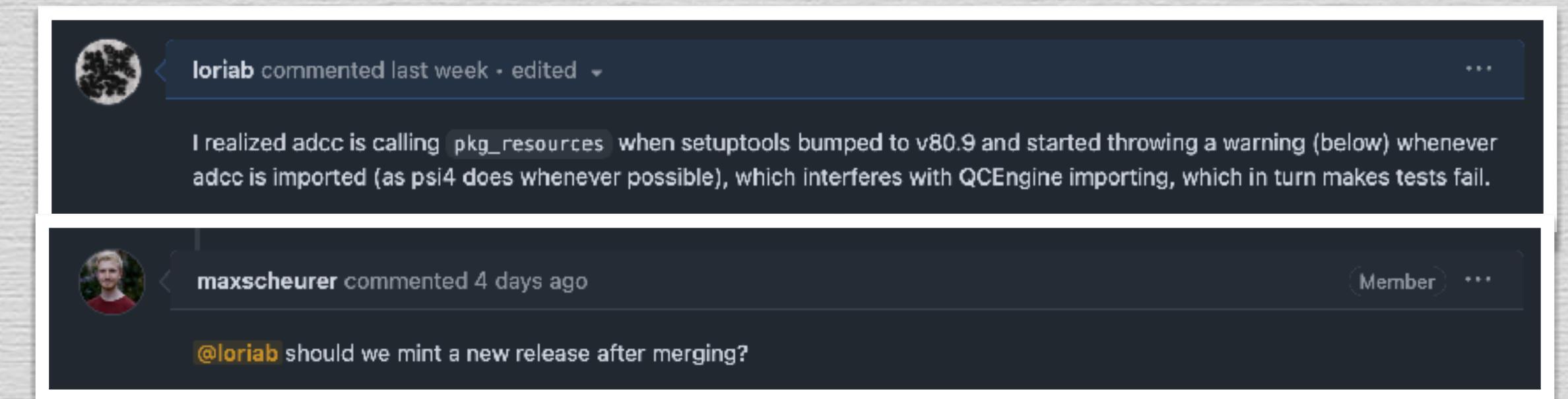


VERSIONING

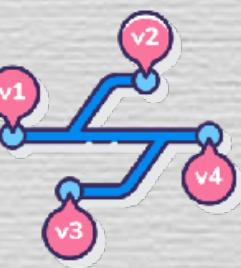
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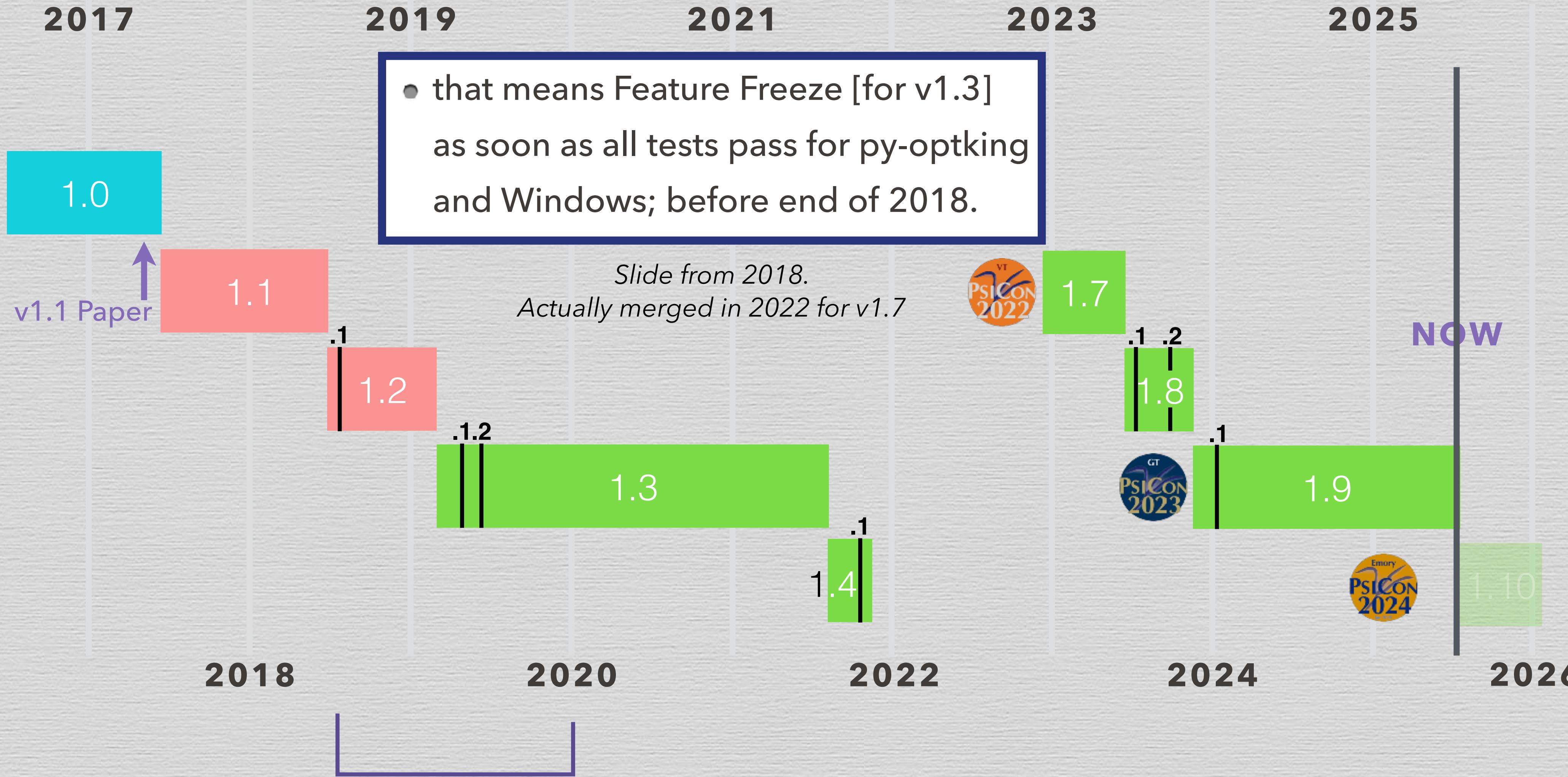


Bless the maintainer

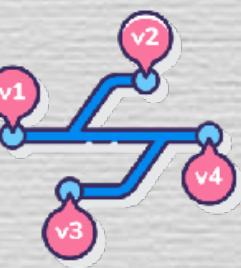


PSI4 RELEASE HISTORY

what not to do

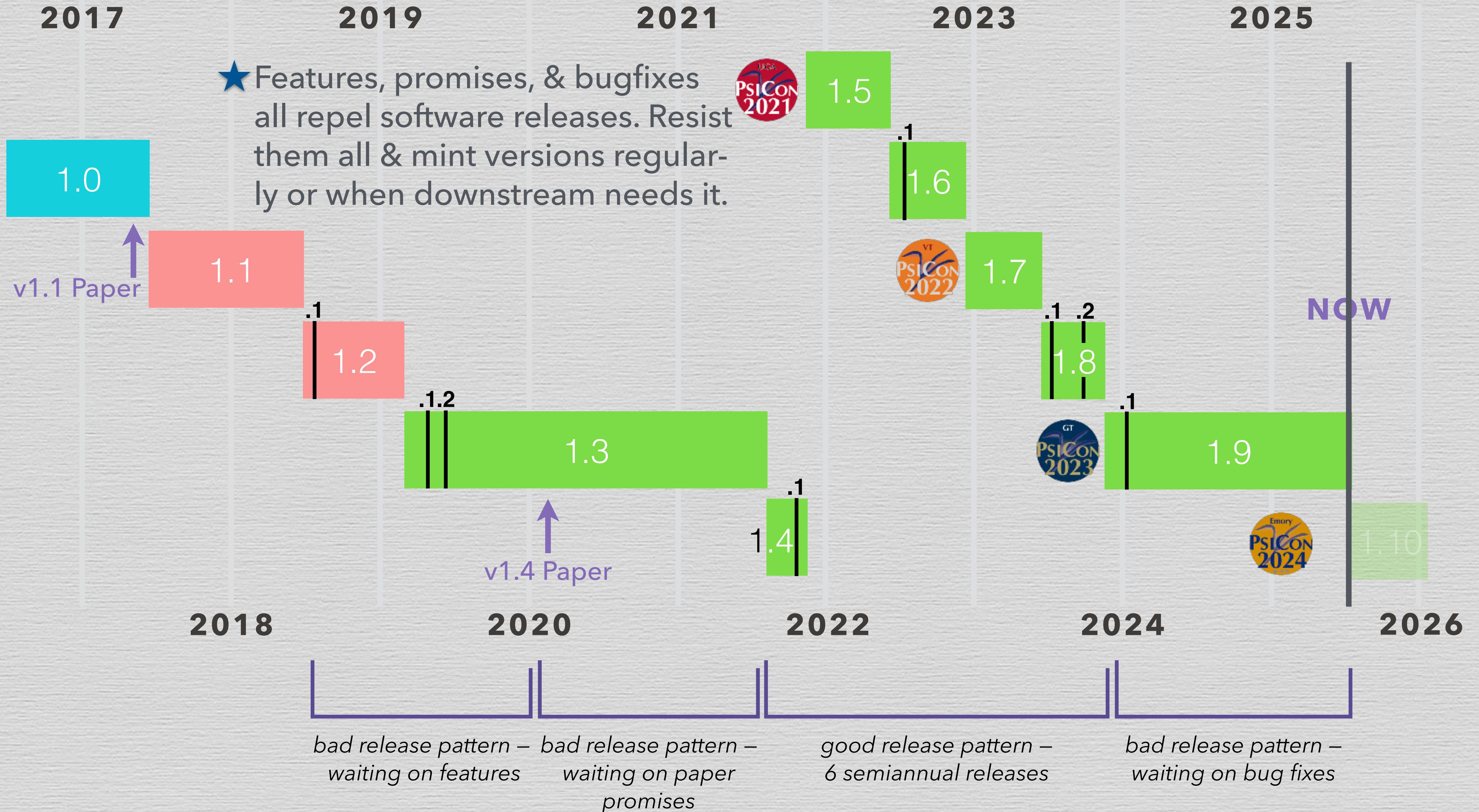


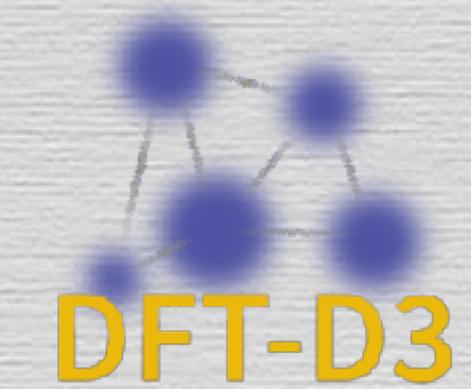
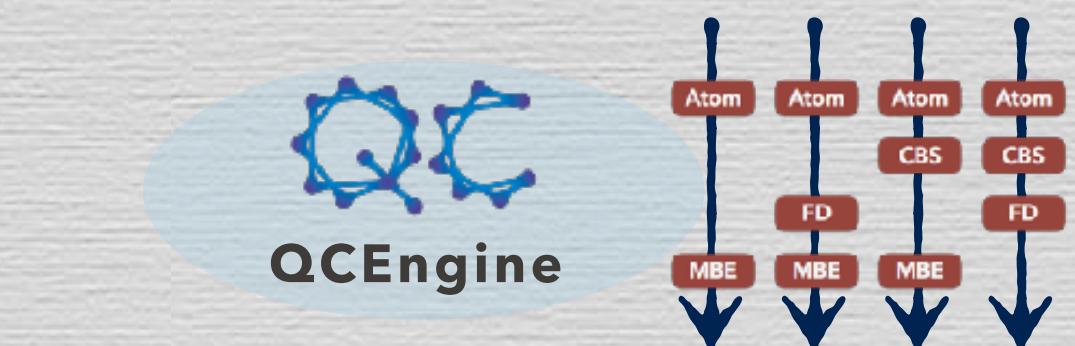
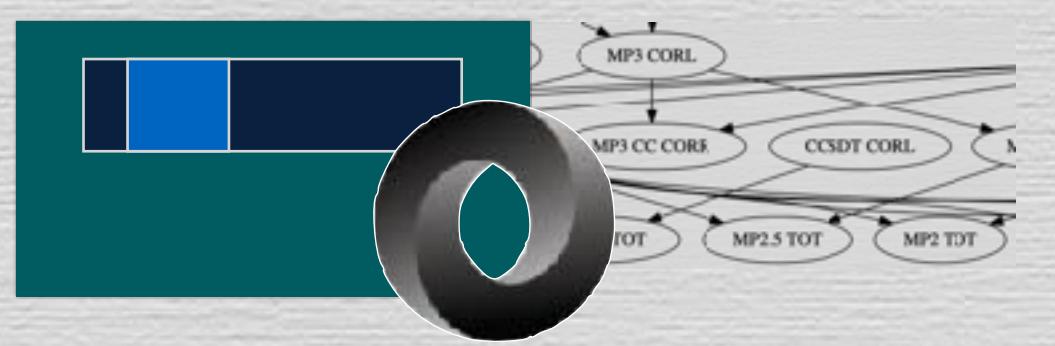
*bad release pattern –
waiting on features*



PSI4 RELEASE HISTORY

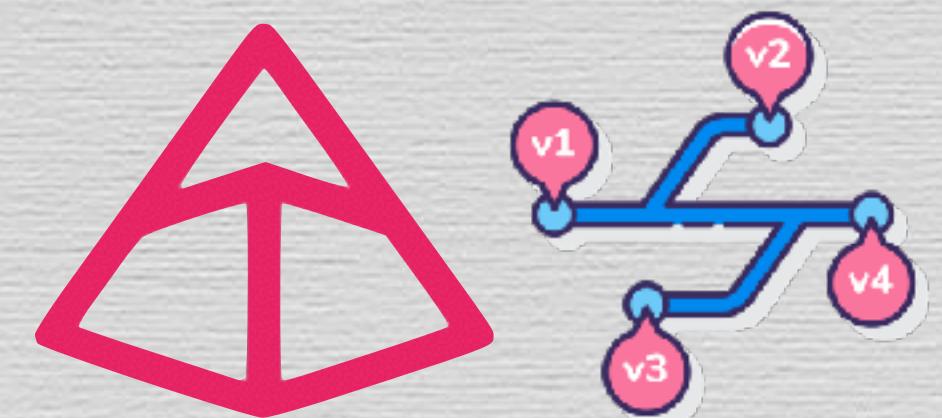
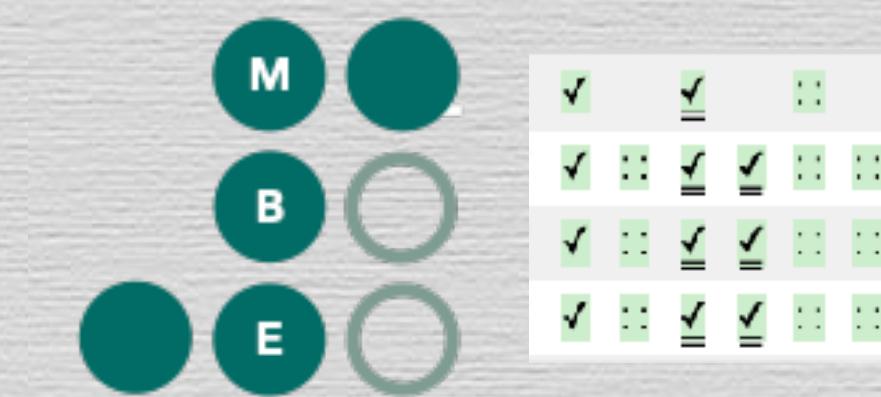
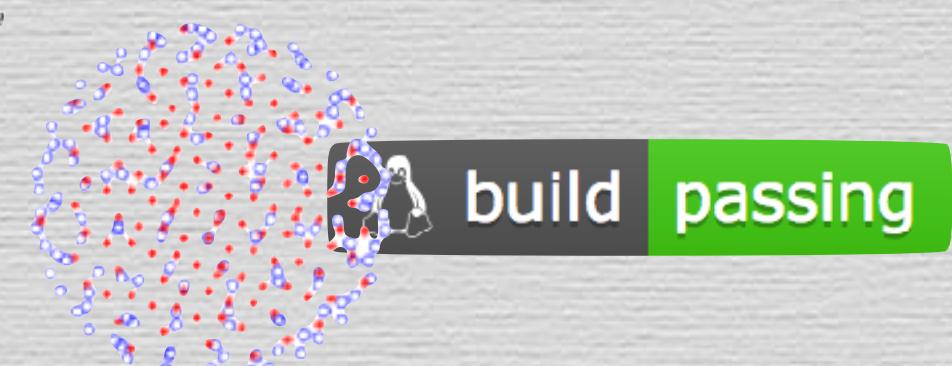
what not to do





PSI4

OPEN-SOURCE QUANTUM CHEMISTRY



★ Features, promises, & bugfixes all repel software releases. Resist them all & mint versions regularly or when downstream needs it.

★ Generous acceptance of feature requests can lead to a high-cohesion module but beyond your comfort.

★ Prepare multiple entry points – API, driver, driven. ★ Mapping calculation input into a structured format can open new ways of interacting with users & workflows.

★ Automate the assessment & conveyance of detailed cap-abilities to users, lest contrib. from alternate modules become overwhelming.

★ Ship a Py reference implementation or MWE to aid downstream adoption

★ Joining a packaging community can prod software engineering aspects.

★ CI testing is both a repository's lifeline and anchor.

★ Find good separations of concern for software stack & build proj. around those missions.

★ Adapt established modular code with new interfaces.

★ Give your module the tools to convey its requirements, capabilities, & attributions. ★ Hold your interfaces and tests close, lest others write them for you.

★ Consume your own data standards to exercise them.

★ At the ecosys level, even simple dependencies logic can get overwhelming, so consolidate & encode.

★ Extract code into a module to concentrate performance optimizations.

★ Automate systematic coverage & test-driven dev for correctness, schema compliance, completeness, & resilience for faster onboarding.

★ Beware that choice of license can limit uptake. ★ Avoid collecting into workflow silos like QCA or Aiida, as we have QC program silos.

★ Decoupling into more cohesive internal modules is progress, too.

★ Modularity can mean dependence in deep software stacks that needs careful guiding.

★ A SSOT deps file can broadcast fixes & be a path to recovery when the build breaks.

★ Getting a production-quality binary to users for multiple architectures can be fairly easy with conda-forge.

★ Find your good, targeted code & set it free for broader use. ★ Prefer and define programmatic output.

★ Provide a variety of installation routes & consolidate the choices.

★ Defer ordering configuration to runtime & build broadly for consumers.

★ Defer loading to runtime for licensing and modularity.

★ New syntax advice is more helpful at runtime than in docs. A grace period is a bonus.

★ Research demands implementing some things quickly for downstream, lest it be blocked. Research also demands one's dependencies be stable and not interrupted by new nonsense fields. Need some governance to balance these interests.

★ Bring the changing env of diff arch, compilers, dep ver into freq. contact w/code.

★ Aim for copy/paste install even for compiled software.

★ Use CI to assure developers always getting a full & proved working environment.

★ Interfacing two codes is rarely unilateral. Seize any opportunity to work with other devs in real time.

★ Features, promises, & bugfixes all repel software releases. Resist them all & mint versions regularly or when downstream needs it.

★ Generous acceptance of feature requests can lead to a high-cohesion module but beyond your comfort.

★ Prepare multiple entry points – API, driver, driven. ★ Mapping calculation input into a structured format can open new ways of interacting with users & workflows.

★ Automate the assessment & conveyance of detailed cap-abilities to users, lest contrib. from alternate modules become overwhelming.

★ Ship a Py reference implementation or MWE to aid downstream adoption.

★ Find good separations of concern for software.

★ Give your module the tools to convey its requirements.

★ At the ecosys level, even simple dependencies

★ Automate systematic dependency management.

★ Beware that choice of license.

★ Modularity can mean dependence in deep software stacks.

★ Getting a production-quality binary to users.

★ Find your good.

★ Defer ordering configuration to runtime & build broadly for consistency.

★ New syntax advice is more helpful at runtime than in docs. A grace period is a bonus.

★ Research demands implementing some things quickly for downstream, lest it be blocked. Research also demands one's dependencies be stable and not interrupted by new nonsense fields. Need some governance to balance these interests.

★ Use CI to assure developers always getting a full & proved working environment.

★ Joining a packaging community can prod software engineering aspects.

★ CI testing is both a repository's lifeline and anchor.

★ Adapt established modular code with new interfaces.

★ Add your interfaces and tests close, lest others write them for you. Exercise them.

★ Concentrate performance optimizations.

★ Aim for completeness, & resilience for faster onboarding.

★ More cohesive internal modules is progress, too. Like QCA or Aiida, as we have QC program silos.

★ Cast fixes & be a path to recovery when the build breaks.

★ Define and define programmatic output.

★ Consolidation routes & consolidate the choices.

★ Defer loading to runtime for licensing and modularity.

★ Bring the changing env of diff arch, compilers, dep ver into freq. contact w/code.

★ Aim for copy/paste install even for compiled software.

★ Interfacing two codes is rarely unilateral. Seize any opportunity to work with other devs in real time.



★ Features, promises, & bugfixes all repel software releases. Resist them all & mint versions regularly or when downstream needs it.

★ Generous acceptance of feature requests can lead to a high-cohesion module but beyond your comfort.

★ Prepare multiple entry points – API, driver, driven. ★ Mapping calculation input into a structured format can open new ways of interacting with users & workflows.

★ Automate the assessment of dependencies. ★ Linting can catch many bugs before they become engineering aspects.
users, lest contrib. from alternate sources. ★ Linting can catch many bugs before they become engineering aspects.

★ Ship a Py reference implementation.

★ Find good code examples.

★ Give your project a clear anchor.

★ At the ecosys level, even the most modular projects have dependencies.



★ Modularity can mean dependency hell.

★ Getting a product ready for release is a challenge.

★ Defer ordering constraints until you know what's needed.



★ Research demands implementing some things quickly for downstream, lest it be blocked. Research also demands one's dependencies be stable and not interrupted by new nonsense fields. Need some governance to balance these interests.

★ Use CI to assure developers always getting a full & proved working environment.

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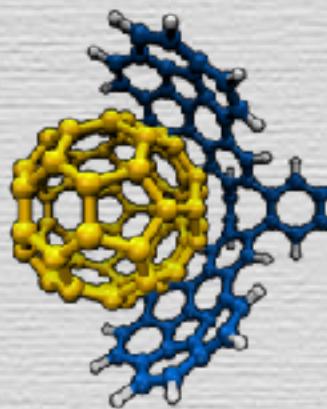


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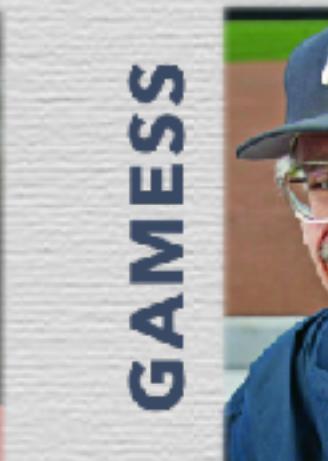
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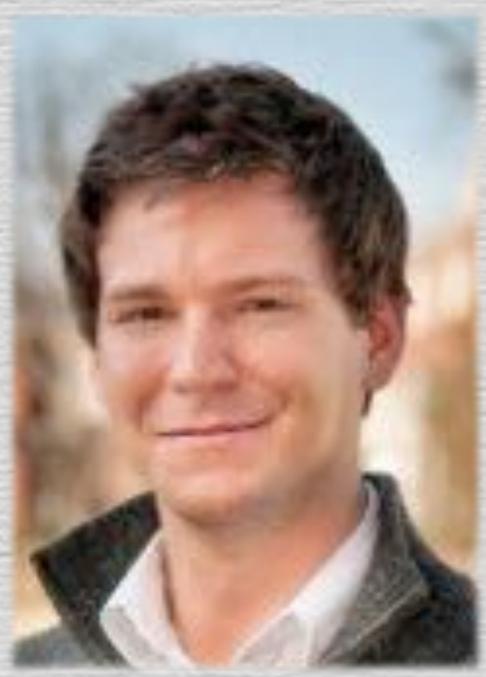
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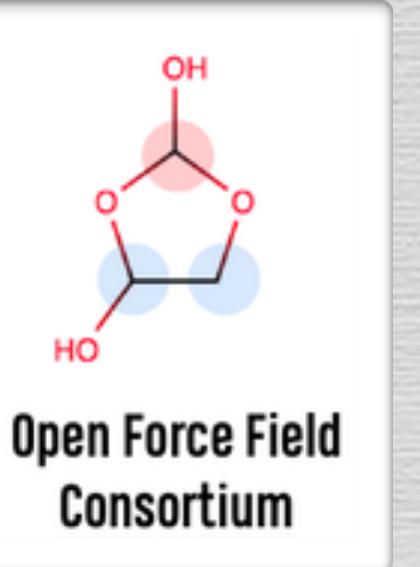
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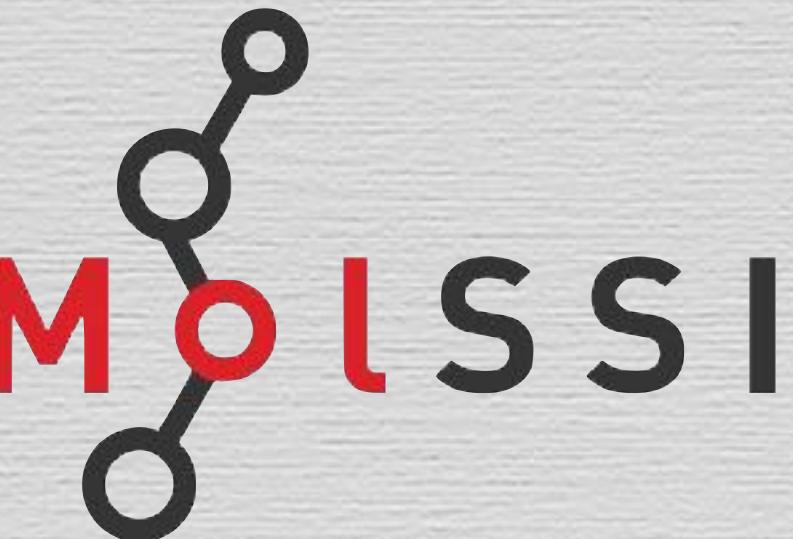


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molssi.github.io/QCFractal

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