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# Molecular Calculations driven by Nexus

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## Goal of this Tutorial

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- How to drive the calculations using Nexus
- How to define a trial wave function
- How to optimize Jastrow factors
- Considerations for DMC calculations

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## Beryllium dimer

- Using Nexus to drive pyscf and QMCPack
  - define the trial wave function
  - optimize the trial
  - explore parameters for production DMC

## Oxygen dimer

- a fully contained example

## Additional Considerations

- How to get started on your own calculations
- choice of trial wave functions

# Beryllium Dimer

# Our Test Case: Beryllium dimer

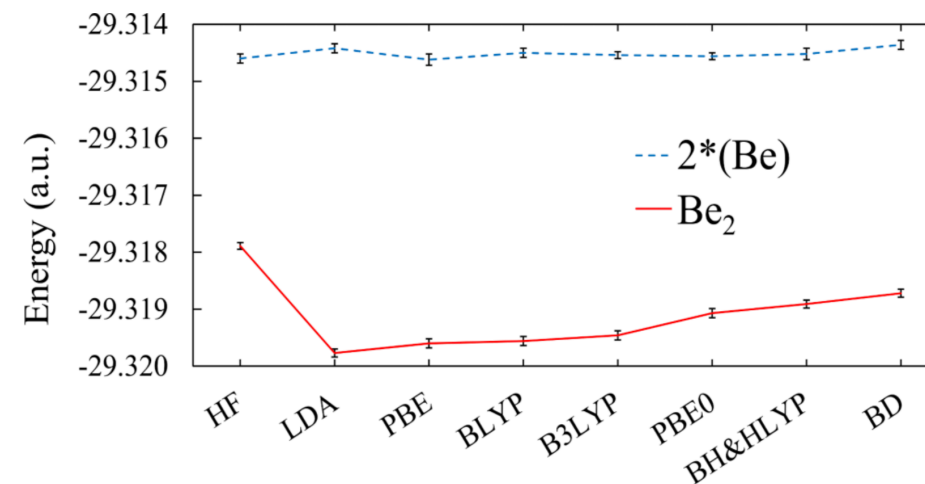
Quantum Monte Carlo calculation of the binding energy of the beryllium dimer

- Michael J. Deible, Melody Kessler, Kevin E. Gasperich, Kenneth D. Jordan, J. Chem. Phys. 143, 084116 (2015)

TABLE I. Total energies of Be and Be<sub>2</sub> and the Be<sub>2</sub> dissociation energy computed with DMC using various trial functions.

Trial function <sup>a</sup>	Total energy (a.u.)		D <sub>e</sub> (cm <sup>-1</sup> )
	Be <sup>b</sup>	Be <sub>2</sub>	
HF/QZ- <i>g</i>	-14.657 30(4)	-29.317 89(6)	724(21)
LDA/QZ- <i>g</i>	-14.657 21(4)	-29.319 77(7)	1174(25)
PBE/QZ- <i>g</i>	-14.657 31(5)	-29.319 60(8)	1094(26)
BLYP/QZ- <i>g</i>	-14.657 25(4)	-29.319 56(8)	1113(26)
B3LYP/QZ- <i>g</i>	-14.657 27(3)	-29.319 46(8)	1079(23)
PBE0/QZ- <i>g</i>	-14.657 28(3)	-29.319 07(8)	992(21)
BH&HLYP/QZ- <i>g</i>	-14.657 26(5)	-29.318 91(7)	966(26)
BD/QZ- <i>g</i>	-14.657 18(4)	-29.318 72(7)	955(24)
CAS(4,8)/QZ- <i>fg</i> <sup>c</sup>	-14.667 23(1)	-29.337 07(3)	573(8)
CAS(4,16)/QZ- <i>fg</i> <sup>c</sup>	-14.667 30(1)	-29.338 32(3)	819(8)
Ext. CAS(4,16)/QZ- <i>fg</i>	-14.667 30(1)	-29.338 41(2)	838(7)
CAS(4,16)/QZ- <i>g</i> <sup>c</sup>	-14.667 27(2)	-29.338 38(3)	845(8)
Ext. CAS(4,16)/QZ- <i>g</i>	-14.667 27(2)	-29.338 45(2)	857(9)
CI/QZ- <i>g</i> <sup>c</sup>	-14.667 25(1)	-29.338 48(2)	873(6)
Ext. CI/QZ- <i>g</i>	-14.667 25(1)	-29.338 64(2)	908(6)
Experimental <sup>d</sup>	-14.667 356	-29.338 97	934.9(4)

DMC energies for various trial wave functions



```
1 import numpy as np
2 from pyscf import df, scf, dft
3
4 from pyscf import gto as gto_loc
5 mol = gto_loc.Mole()
6 mol.verbose = 1
7 mol.atom = '''
8           Be    0.00000000    0.00000000    0.00000000
9           Be    2.45360300    0.00000000    0.00000000
10          '''
11 mol.basis = 'cc-pvtz'
12 mol.unit = 'A'
13 mol.charge = 0
14 mol.spin = 0
15 mol.symmetry = True
16 mol.build()
17
18 mf = scf.ROHF(mol).density_fit()
19 mf.max_cycle=200
20 mf.level_shift=0.0
21 mf.tol = '1e-10'
22 e_scf = mf.kernel()
```

① e\_scf = -29.133800914375186

①



## Nexus to drive Pyscf

be2\_nexus.py

```
1 from nexus import settings, job, run_project, ob
2 from nexus import ppset
3 from nexus import generate_physical_system
4 from nexus import generate_pyscf
5
6 XC=["LDA", "PBE", "PBE0", "SCAN"]
7 MyBasis=["cc-pvdz", "cc-pvtz", "cc-pvqz"]
8 for y in MyBasis:
9     # perform Hartree Fock!
10    scf = generate_pyscf(
11        identifier = 'scf',
12        path       = 'Be2/'+y+'/hf/scf',
13        job        = job(serial=True, app='pyt
14        system     = system,
15        mole       = obj(
16            basis   = y,
17            symmetry = True,
18            verbose  = 5,
19        ),
20        calculation = obj(
21            method   = 'ROHF',
22            df_fitting = True,
```

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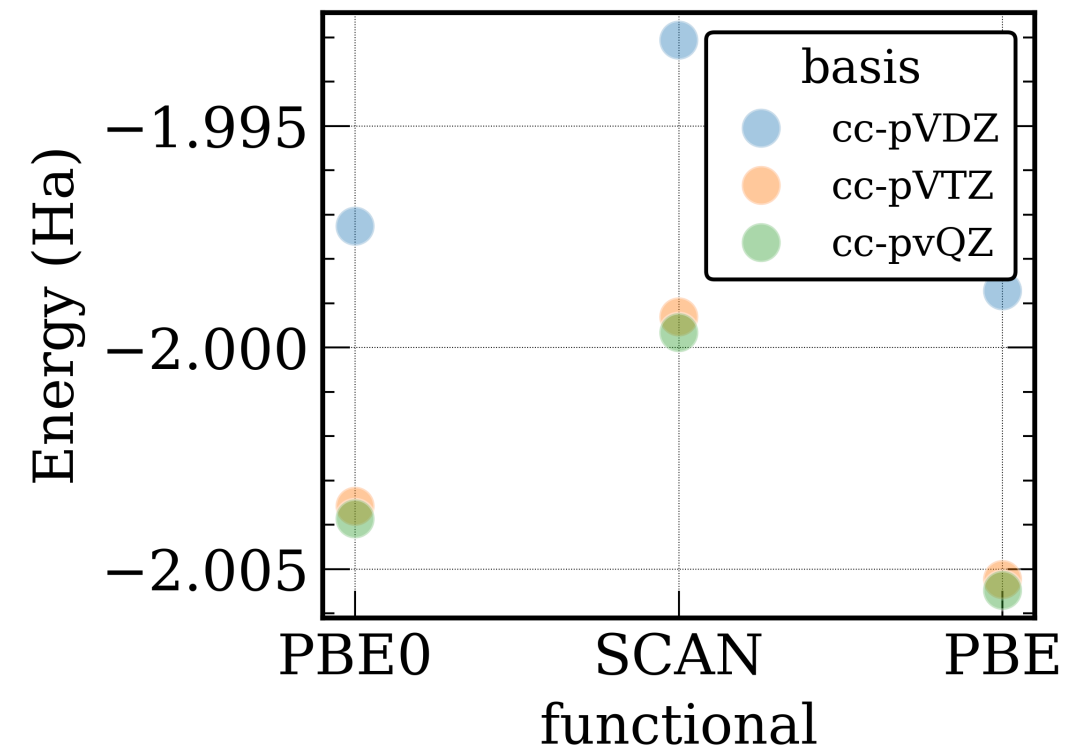
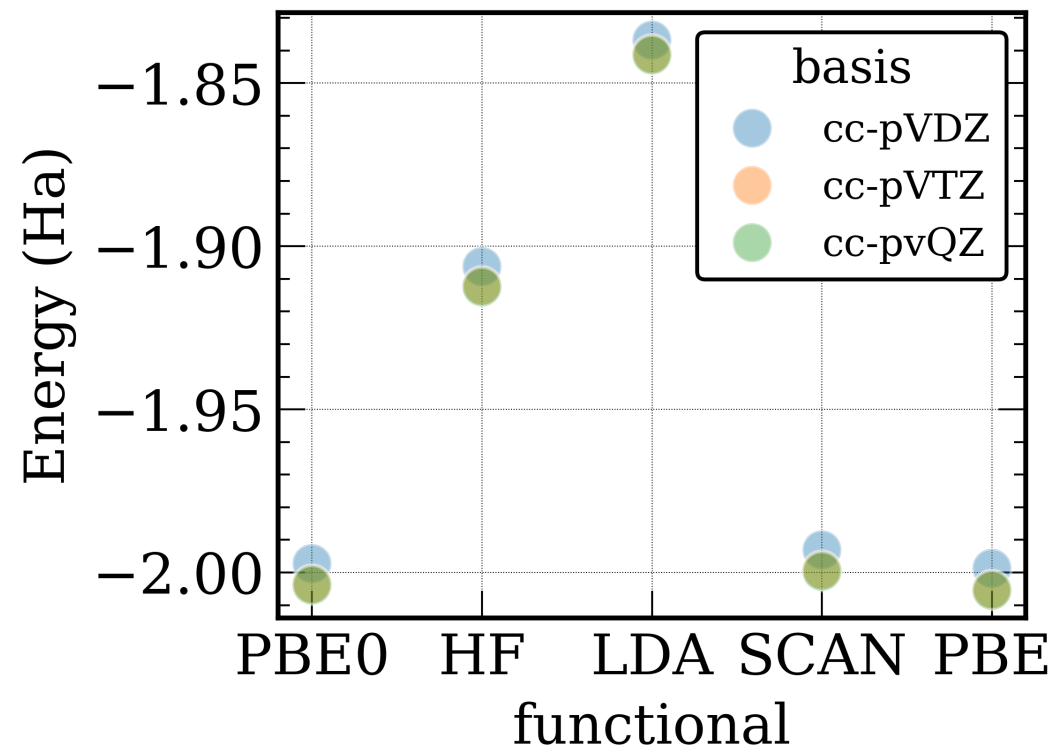
be2\_nexus.py

```
8 for y in MyBASIS:
9     # perform Hartree Fock!
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14        system     = system,
15        mole       = obj(
16            basis   = y,
17            symmetry = True,
18            verbose  = 5,
19        ),
20        calculation = obj(
21            method   = 'ROHF',
22            df_fitting = True,
23            max_cycle = 200,
24            level_shift = 0.0,
25            tol       = '1e-10',
26        ),
27    )
28 for x in XC:
29     # perform DFT
30     scf = generate_pyscf(
```



## Nexus to drive Pyscf

```
be2_nexus.py
29 # perform DFT
30     scf = generate_pyscf(
31         identifier = 'scf',
32         path       = 'Be2/'+y+'/' + x + '/scf
33         job        = job(serial=True, app=
34         system     = system,
35         mole       = obj(
36             basis   = y,
37             symmetry = True,
38             verbose  = 5,
39         ),
40         calculation = obj(
41             method   = 'ROKS',
42             df_fitting = True,
43             max_cycle = 200,
44             level_shift = 0.0,
45             tol       = '1e-10',
46             xc        = x,
47         ),
48     )
49
50
51 run_project()
```



## Our wave function

---

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

1. Antisymmetric portion: Slater determinant
2. Symmetric portion: the Jastrow factor





$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

1. Antisymmetric portion: Slater determinant
2. Symmetric portion: the Jastrow factor

$$\Psi_{AS} = \sum_k^M C_k D_k^\uparrow(\phi) D_k^\downarrow(\phi)$$

$$D(\mathbf{r}_1, \mathbf{r}_2, \dots, \mathbf{r}_N) = \frac{1}{\sqrt{N!}} \begin{vmatrix} \chi_1(\mathbf{r}_1) & \chi_1(\mathbf{r}_2) & \cdots & \chi_1(\mathbf{r}_N) \\ \chi_2(\mathbf{r}_1) & \chi_2(\mathbf{r}_2) & \cdots & \chi_2(\mathbf{r}_N) \\ \vdots & \vdots & \ddots & \vdots \\ \chi_N(\mathbf{r}_1) & \chi_N(\mathbf{r}_2) & \cdots & \chi_N(\mathbf{r}_N) \end{vmatrix}$$

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

1. Antisymmetric portion: Slater determinant
2. Symmetric portion: the Jastrow factor

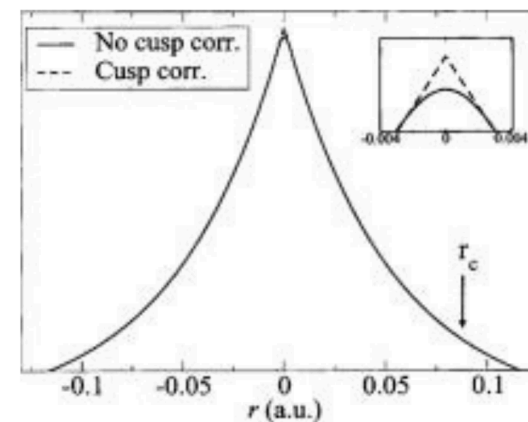
$$\Psi_{AS} = \sum_k^M C_k D_k^\uparrow(\phi) D_k^\downarrow(\phi)$$

$$D(\mathbf{r}_1, \mathbf{r}_2, \dots, \mathbf{r}_N) = \frac{1}{\sqrt{N!}} \begin{vmatrix} \chi_1(\mathbf{r}_1) & \chi_1(\mathbf{r}_2) & \cdots & \chi_1(\mathbf{r}_N) \\ \chi_2(\mathbf{r}_1) & \chi_2(\mathbf{r}_2) & \cdots & \chi_2(\mathbf{r}_N) \\ \vdots & \vdots & \ddots & \vdots \\ \chi_N(\mathbf{r}_1) & \chi_N(\mathbf{r}_2) & \cdots & \chi_N(\mathbf{r}_N) \end{vmatrix}$$

Linear Combination of Atomic Orbitals

$$\chi(\mathbf{r}) = \sum_{i=1}^N c_i \phi_i(\mathbf{r})$$

For all electron calculations, employ a correction for electron nuclear cusp



# QMC Workflow Overview



## 1. Wave function definition for QMCPack

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

## 2. Wave function optimization

## 3. Production parameter choices

Considerations	Files to explore
cusp correction	session4_molecules/01_Be2_dimer/run0_vmc_noj.py
optimize $\Psi_T$	session4_molecules/01_Be2_dimer/run1_qmc_wfopt.py
timestep error	session4_molecules/01_Be2_dimer/run2_dmc_timestep.py
error bar control	session4_molecules/01_Be2_dimer/run3_dmc_errorbars.py
population bias	session4_molecules/01_Be2_dimer/run4_dmc_population.py
production run	session4_molecules/01_Be2_dimer/run5_dmc_production.py

## Calculating the antisymmetric portion of trial

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

run0\_vmc\_noJ.py

```
1 scf = generate_pyscf(  
2     identifier = 'scf',           # log out  
3     ...  
4     calculation = obj(  
5         method      = 'R0KS',     # Restrict  
6         df_fitting   = True,       # Density  
7         ...  
8         xc           = 'pbe',     # Exchange  
9     ),  
10    save_qmc        = True ,       # Save the  
11    )
```

Run a restricted DFT calculation using PBE functional.  
**save\_qmc** option outputs orbitals in a usable way



## Calculating the antisymmetric portion of trial

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

run0\_vmc\_noJ.py

```
1 scf = generate_pyscf(
2     identifier = 'scf',          # log out
3     ...
4     calculation = obj(
5         method      = 'ROKS',    # Restrict
6         df_fitting   = True,      # Density
7         ...
8         xc           = 'pbe',    # Exchange
9     ),
10    save_qmc        = True,       # Save the
11)
```

```
1 # convert orbitals to QMCPACK format
2 c4q = generate_convert4qmc(
3     identifier      = 'c4q',
4     path            = 'Be2/' + y + '/' + x + '/SCF',
5     job             = job(cores=1),
6     dependencies    = (scf, 'orbitals'),
7 )
```

Run a restricted DFT calculation using PBE functional.  
**save\_qmc** option outputs orbitals in a usable way

use **convert4qmc** to generate input files for QMCPack

- supports a number of codes

## Results from convert4qmc

output files: c4q.orbs.h5, c4q.qmc.in-wfj.xml, c4q.structure.xml, c4q.wfj.xml

c4q.qmc.in-wfj.xml

```

1  <!--Example QMCPACK input file produced by convert4qmc
2  -->
3  <!--Name and Series number of the project.-->
4  <project id="c4q" series="0"/>
5  <!--Link to the location of the Atomic Coordinates and the location of the Wavefunction.-->
6  <include href="c4q.structure.xml"/>
7  <include href="c4q.wfj.xml"/>
8  <!--Hamiltonian of the system.-->
9  <hamiltonian name="h0" type="generic" target="e">
10    <pairpot name="ElecElec" type="coulomb" source="e" target="e" physical="true"/>
11    <pairpot name="IonIon" type="coulomb" source="ion0" target="ion0"/>
12    <pairpot name="IonElec" type="coulomb" source="ion0" target="e"/>
13  </hamiltonian>
14  <!--Example initial VMC to measure initial energy and variance -->
15  <qmc method="vmc" move="pbyp" checkpoint="-1">
16    <estimator name="LocalEnergy" hdf5="no"/>
17    <parameter name="warmupSteps">100</parameter>
18    <parameter name="blocks">20</parameter>
19    <parameter name="steps">50</parameter>
20    <parameter name="substeps">8</parameter>
21    <parameter name="timestep">0.5</parameter>
22    <parameter name="usedrift">no</parameter>

```



## Adding symmetric portion of wave function

---

$$\Psi_T(R) = J(R)\Psi_{AS}(R)$$

$$J(R) = e^{J_1+J_2+\dots}$$

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$$J_1 = \int_I \sum_i^e u_{ab}(|r_i - R_I|)$$

$$J_2 = \sum_i^e \sum_{j<i}^e u_{ab}(|r_i - r_j|)$$

- We will optimize the  $J_1$  and  $J_2$

```
1 # optimize 2-body Jastrow
2 optJ2 = generate_qmcpack(
3     identifier      = 'opt',
4     path            = 'Be2/' + y + '/' + x + '/optJ2
5     job             = job(cores=cores),
6     system          = system,
7     J2              = True,           # 2-body
8     J1_rcut         = 6.0,           # 6 Bohr
9     J2_rcut         = 8.0,           # 8 Bohr
10    seed            = 42,             # Fix the
11    qmc              = 'opt',         # Wavefun
12    minmethod        = 'oneshift',    # Energy
13    init_cycles      = 4,             # 4 itera
14    cycles           = 8,             # 8 produ
15    warmupsteps      = 10,
16    blocks           = 20,
17    steps            = 3,
18    timestep         = 0.1,
19    init_minwalkers  = 0.1,
20    minwalkers       = 0.5,
21    samples          = 25600,         # VMC sam
22    dependencies     = orbdeps,
```

## Adding symmetric portion of wave function

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## Adding symmetric portion of wave function

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23 )
```

## Adding symmetric portion of wave function

The three-body Jastrow form:

$$u_{\sigma\sigma'I}(r_{\sigma I}, r_{\sigma'I}, r_{\sigma\sigma'}) = \sum_{\ell=0}^{M_{eI}} \sum_{m=0}^{M_{eI}} \sum_{n=0}^{M_{ee}} \gamma_{\ell mn} r_{\sigma I}^{\ell} r_{\sigma'I}^m r_{\sigma\sigma'}^n \\ \times \left( r_{\sigma I} - \frac{r_c}{2} \right)^3 \Theta \left( r_{\sigma I} - \frac{r_c}{2} \right) \\ \times \left( r_{\sigma'I} - \frac{r_c}{2} \right)^3 \Theta \left( r_{\sigma'I} - \frac{r_c}{2} \right)$$

- correlation is only a function of the interparticle distances
- correlations are set to zero beyond a distance
- J3 will impact J1 and J2, so optimizing them all together is important

```
1 # optimize 3-body Jastrow
2 optJ3 = generate_qmcpack(
3     identifier      = 'opt',
4     path            = 'Be2/'+y+'/' + x + '/optJ3
5     job             = job(cores=cores),
6     system          = system,
7     J3              = True,           # 3-body
8     seed            = 42,            # Fix the
9     qmc              = 'opt',        # Wavefun
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14    blocks            = 20,
15    steps             = 5,
16    timestep          = 0.1,
17    init_minwalkers   = 0.1,
18    minwalkers        = 0.5,
19    samples           = 25600,        # VMC sam
20    dependencies      = orbdeps+[(optJ2, 'jastro
21 )
```

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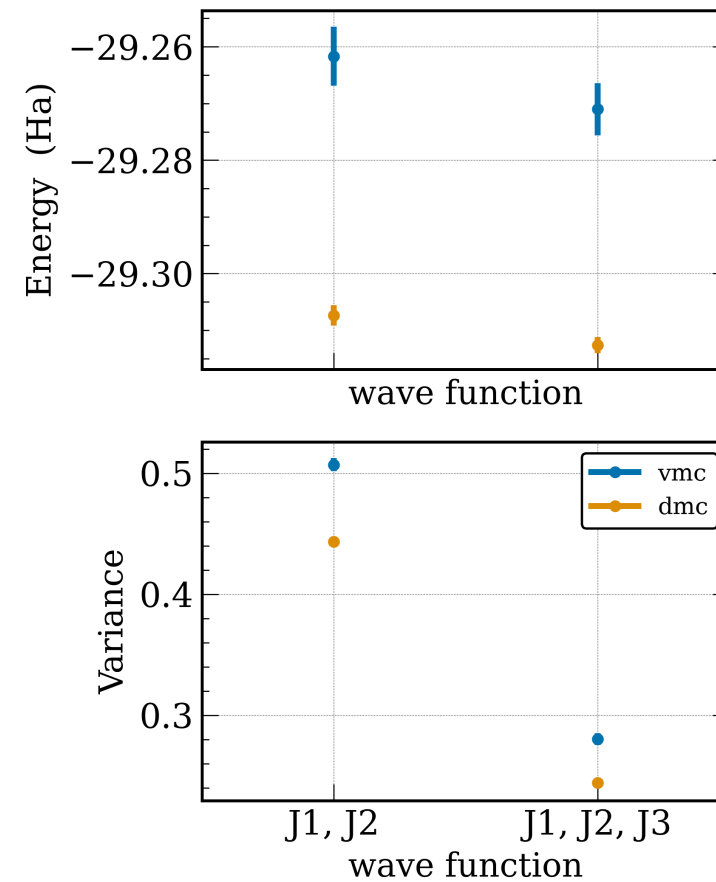
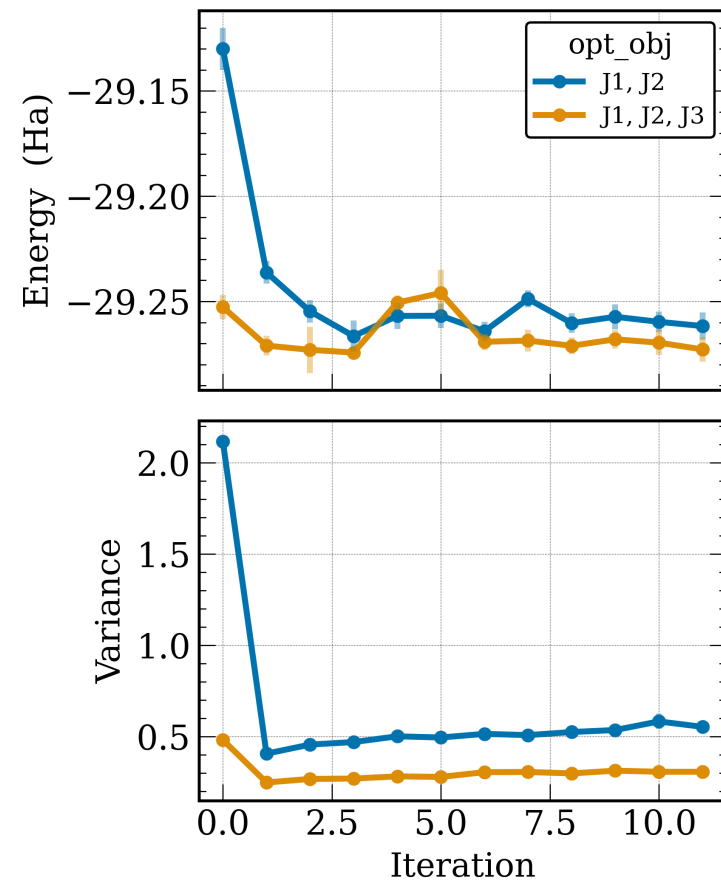
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```

## Analyzing a wave function optimization run

```
$ qmca -qev *.scalar.dat
```

		LocalEnergy	Variance	ratio
opt	series 0	-29.130060 +/- 0.010435	2.117672 +/- 0.018035	0.0727
opt	series 1	-29.236196 +/- 0.005300	0.406483 +/- 0.006288	0.0139
opt	series 2	-29.254653 +/- 0.005361	0.456244 +/- 0.004171	0.0156
opt	series 3	-29.266499 +/- 0.007434	0.470493 +/- 0.010077	0.0161
opt	series 4	-29.256925 +/- 0.006055	0.501486 +/- 0.005856	0.0171
opt	series 5	-29.256848 +/- 0.005754	0.494502 +/- 0.023800	0.0169
opt	series 6	-29.264160 +/- 0.004457	0.514852 +/- 0.006401	0.0176
opt	series 7	-29.248796 +/- 0.004104	0.508019 +/- 0.005036	0.0174
opt	series 8	-29.260340 +/- 0.004567	0.524970 +/- 0.006886	0.0179
opt	series 9	-29.257258 +/- 0.005917	0.535870 +/- 0.006333	0.0183
opt	series 10	-29.259730 +/- 0.004787	0.583399 +/- 0.041853	0.0199
opt	series 11	-29.261702 +/- 0.006473	0.553353 +/- 0.004991	0.0189

# Results of wave function optimization



Including three-body terms decreases energy and variance for VMC

Jastrow function improves significantly the variance, but eventually DMC will recover the missing correlation

## Diffusion Monte Carlo



Considerations	Files to explore
timestep error	<code>session4_molecules/01_Be2_dimer/run1_dmc_timestep.py</code>
reducing error bars	<code>session4_molecules/01_Be2_dimer/run1_dmc_errorbars.py</code>
population bias	<code>session4_molecules/01_Be2_dimer/run4_dmc_population.py</code>
production run	<code>session4_molecules/01_Be2_dimer/run5_dmc_production.py</code>

## DMC: timestep



run2\_dmc\_timestep.py

```
1 x="PBE"
2 y="cc-pvtz"
3 orbdeps = [(c4q, 'particles'), # pyscf changes
4             (c4q, 'orbitals'),
5             (cc, 'cuspcorr')]
6
7 qmc = generate_qmcpack(
8     identifier      = 'dmc',
9     seed            = 42,
10    driver           = 'batched',
11    path             = 'Be2/'+y+'/'+x+'/dmc_tst',
12    job              = qmc_job,
13    system           = system,
14    jastrows          = [],
15    qmc              = 'dmc',
16    warmupsteps      = 50,
17    vmc_blocks       = 200,
18    vmc_steps        = 20,
19    vmc_timestep     = 0.3,
20    timestep         = 0.01,
21    timestep_factor  = 0.5,
22    ntimesteps       = 4,
```

## DMC: timestep



run2\_dmc\_timestep.py

```
5         (cc, 'cuspcorr')]  
6  
7 qmc = generate_qmcpack(  
8     identifier      = 'dmc',  
9     seed            = 42,  
10    driver           = 'batched',  
11    path             = 'Be2/'+y+'/' +x+' /dmc_tst  
12    job              = qmc_job,  
13    system           = system,  
14    jastrows         = [],  
15    qmc              = 'dmc',  
16    warmupsteps      = 50,  
17    vmc_blocks        = 200,  
18    vmc_steps         = 20,  
19    vmc_timestep      = 0.3,  
20    timestep         = 0.01,  
21    timestep_factor  = 0.5,  
22    ntimesteps        = 4,  
23    total_walkers     = 512,  
24    blocks            = 400,  
25    dependencies      = orbdeps+[(optJ3, 'jastrow  
26    )
```

## DMC: timestep



run2\_dmc\_timestep.py

```
6
7 qmc = generate_qmcpack(
8     identifier      = 'dmc',
9     seed            = 42,
10    driver           = 'batched',
11    path             = 'Be2/'+y+'/'+'x+'/'dmc_tst
12    job              = qmc_job,
13    system           = system,
14    jastrows         = [],
15    qmc              = 'dmc',
16    warmupsteps      = 50,
17    vmc_blocks       = 200,
18    vmc_steps        = 20,
19    vmc_timestep      = 0.3,
20    timestep         = 0.01,
21    timestep_factor  = 0.5,
22    ntimesteps       = 4,
23    total_walkers    = 512,
24    blocks           = 400,
25    dependencies     = orbdeps+[(optJ3, 'jastrow
26 )
27
```

## DMC: timestep



run2\_dmc\_timestep.py

```
6
7 qmc = generate_qmcpack(
8     identifier      = 'dmc',
9     seed            = 42,
10    driver           = 'batched',
11    path             = 'Be2/'+y+'/'+'x+'/'dmc_tst
12    job              = qmc_job,
13    system           = system,
14    jastrows         = [],
15    qmc              = 'dmc',
16    warmupsteps      = 50,
17    vmc_blocks       = 200,
18    vmc_steps        = 20,
19    vmc_timestep     = 0.3,
20    timestep         = 0.01,
21    timestep_factor  = 0.5,
22    ntimesteps       = 4,
23    total_walkers    = 512,
24    blocks           = 400,
25    dependencies     = orbdeps+[(optJ3, 'jastrow
26 )
27
```

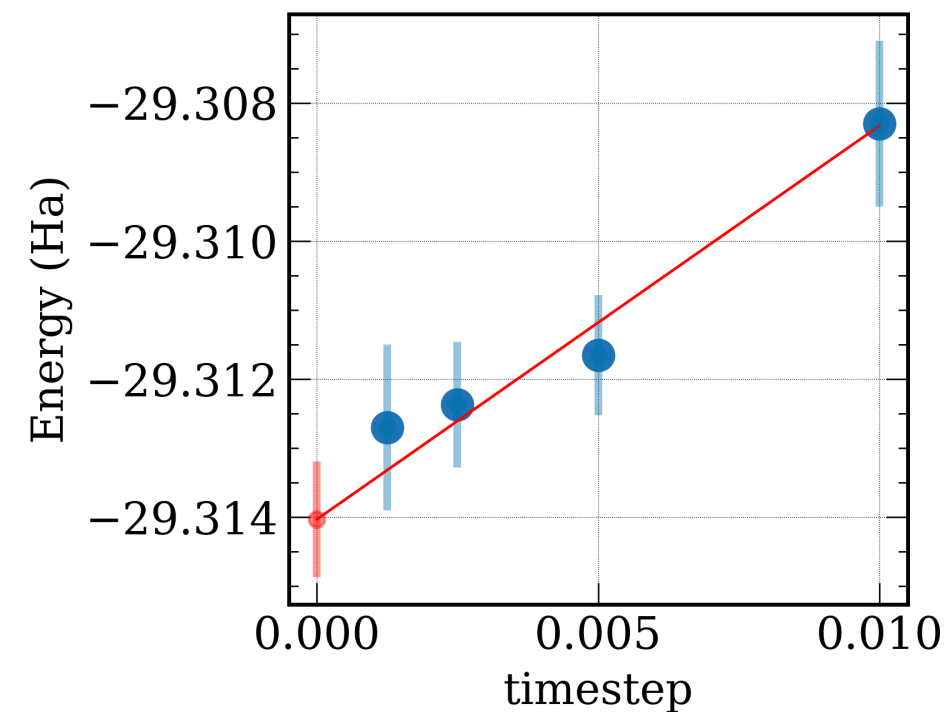


## DMC: timestep

run2\_dmc\_timestep.py

```
1 x="PBE"
2 y="cc-pvtz"
3 orbdeps = [(c4q,'particles'), # pyscf changes
4             (c4q,'orbitals'),
5             (cc, 'cuspcorr')]
6
7 qmc = generate_qmcpack(
8     identifier      = 'dmc',
9     seed            = 42,
10    driver           = 'batched',
11    path             = 'Be2/'+y+'/'+x+'/'dmc_tst
12    job              = qmc_job,
13    system            = system,
14    jastrows          = [],
15    qmc               = 'dmc',
16    warmupsteps       = 50,
17    vmc_blocks        = 200,
18    vmc_steps         = 20,
19    vmc_timestep       = 0.3,
20    timestep          = 0.01,
21    timestep_factor    = 0.5,
22    ntimesteps        = 4,
```

```
qmc-fit ts -e 20 '2 4 4' -t '0.01 0.005 0.0025'
*s00{1,2,3}.scalar.dat
```



## DMC: reducing error bars

To reduce the error bar by a factor  $N$ , multiply the population or the number of blocks by  $N^2$

run3\_dmc\_errorbars.py

```
1 init_blocks=100
2 for i in range(1,6):
3     myblocks=init_blocks*i*i
4     qmc = generate_qmcpack(
5         identifier      = 'dmc_error'+str(i),
6         seed            = 42,
7         path            = 'Be2/'+y+'/'+x+'/dmc_error',
8         job             = job(cores=cores),
9         system          = system,
10        jastrows         = [],
11        qmc              = 'dmc',
12        vmc_samples      = 1024,
13        warmupsteps      = 50,
14        vmc_blocks       = 100,
15        vmc_steps        = 10,
16        vmc_timestep     = 0.1,
17        timestep         = 0.00250,
18        steps            = 80,
19        blocks           = myblocks,
20        dependencies     = orbdeps+[(optJ3, 'jastrow
21    )
```

## DMC: reducing error bars

To reduce the error bar by a factor  $N$ , multiply the population or the number of blocks by  $N^2$

run3\_dmc\_errorbars.py

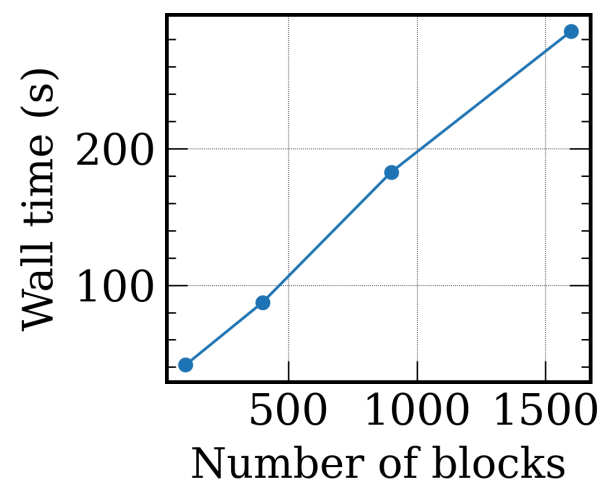
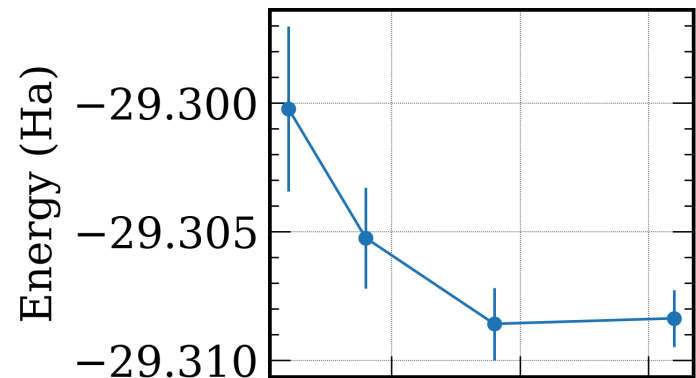
```
1 init_blocks=100
2 for i in range(1,6):
3     myblocks=init_blocks*i*i
4     qmc = generate_qmcpack(
5         identifier      = 'dmc_error'+str(i),
6         seed            = 42,
7         path            = 'Be2/'+y+'/'+x+'/dmc_error',
8         job             = job(cores=cores),
9         system          = system,
10        jastrows         = [],
11        qmc              = 'dmc',
12        vmc_samples      = 1024,
13        warmupsteps      = 50,
14        vmc_blocks       = 100,
15        vmc_steps        = 10,
16        vmc_timestep     = 0.1,
17        timestep         = 0.00250,
18        steps            = 80,
19        blocks           = myblocks,
20        dependencies     = orbdeps+[(optJ3, 'jastrow'),
21    )
```

## DMC: reducing error bars

To reduce the error bar by a factor  $N$ , multiply the population or the number of blocks by  $N^2$

run3\_dmc\_errorbars.py

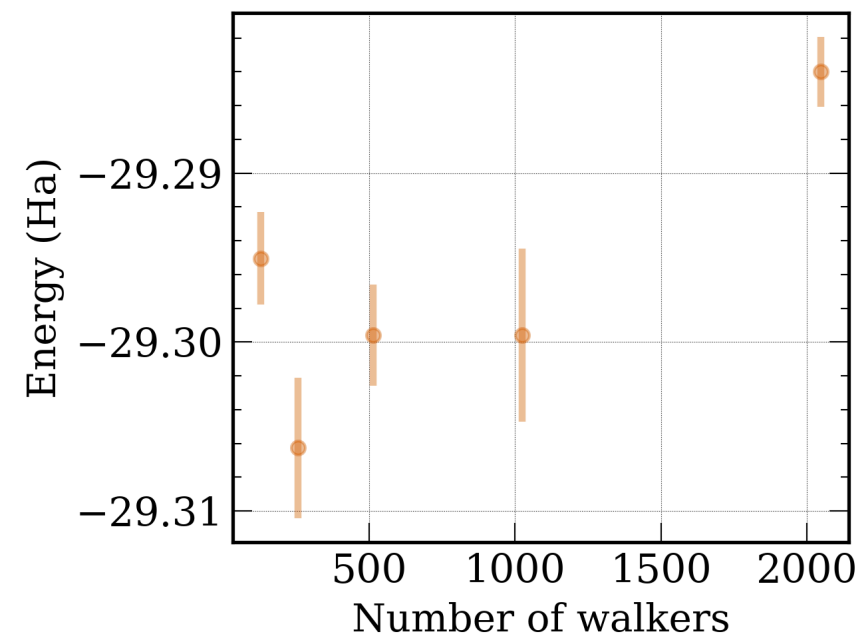
```
1 init_blocks=100
2 for i in range(1,6):
3     myblocks=init_blocks*i*i
4     qmc = generate_qmcpack(
5         identifier    = 'dmc_error'+str(i),
6         seed          = 42,
7         path          = 'Be2/'+y+'/' +x+' /dmc_err
8         job           = job(cores=cores),
9         system        = system,
10        jastrows       = [],
11        qmc            = 'dmc',
12        vmc_samples    = 1024,
13        warmupsteps    = 50,
14        vmc_blocks     = 100,
15        vmc_steps      = 10,
16        vmc_timestep   = 0.1,
17        timestep       = 0.00250,
18        steps          = 80,
19        blocks         = myblocks,
20        dependencies   = orbdeps+[(optJ3, 'jastrow
21    )
```



## DMC: reducing population bias

run4\_dmc\_population.py

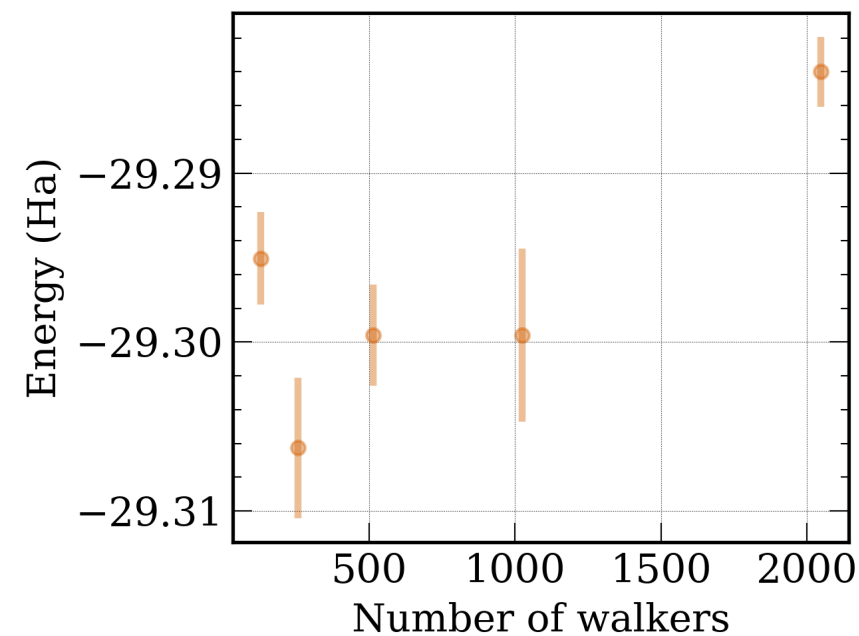
```
1 pop=64
2 while pop<2049:
3     qmc = generate_qmcpack(
4         identifier    = 'dmc_pop'+str(pop),
5         seed          = 42,
6         path          = 'Be2/'+y+'/' +x+' /dmc_',
7         job           = qmc_job,
8         system        = system,
9         jastrows       = [],
10        qmc            = 'dmc',
11        total_walkers  = pop,
12        warmupsteps    = 0,
13        vmc_blocks     = 10,
14        vmc_steps      = 1,
15        vmc_timestep   = 0.3,
16        timestep       = 0.01,
17        steps          = 20480//pop,
18        blocks         = 10,
19        dependencies   = orbdeps+[(optJ3, 'jastrows')]
20    )
21    pop=pop*2
```



## DMC: reducing population bias

run4\_dmc\_population.py

```
1 pop=64
2 while pop<2049:
3     qmc = generate_qmcpack(
4         identifier    = 'dmc_pop'+str(pop),
5         seed          = 42,
6         path          = 'Be2/'+y+'/' +x+' /dmc_',
7         job           = qmc_job,
8         system        = system,
9         jastrows       = [],
10        qmc            = 'dmc',
11        total_walkers  = pop,
12        warmupsteps    = 0,
13        vmc_blocks     = 10,
14        vmc_steps      = 1,
15        vmc_timestep   = 0.3,
16        timestep       = 0.01,
17        steps          = 20480//pop,
18        blocks         = 10,
19        dependencies   = orbdeps+[(optJ3, 'jastrows')]
20    )
21    pop=pop*2
```



# Oxygen dimer



# Oxygen dimer: using pseudopotentials

## Perform SCF with Pseudopotential

```
1 system = generate_physical_system(  
2     units      = 'A',  
3     elem       = ['O', 'O'],  
4     pos        = [[0.000000, 0.000000, 0.000000],  
5                  [0.000000, 0.000000, 1.208]],  
6     O=6,  
7 )  
8 settings(  
9     machine     = 'ws4',  
10    pseudo_dir  = './pseudos/')
```





# Oxygen dimer: using pseudopotentials

## Perform SCF with Pseudopotential

```
1 system = generate_physical_system(  
2     units      = 'A',  
3     elem       = ['O', 'O'],  
4     pos        = [[0.000000, 0.000000, 0.000000],  
5                   [0.000000, 0.000000, 1.208]],  
6     O=6,  
7 )  
8 settings(  
9     machine     = 'ws4',  
10    pseudo_dir  = './pseudos/')  

```

## Create pseudopotentials set

```
1 ppset(  
2     label      = 'ccecp',  
3     qmcpack    = ['O.ccECP.xml'],  
4 )  

```



# Oxygen dimer: using pseudopotentials

## Perform SCF with Pseudopotential

```
1 system = generate_physical_system(  
2     units      = 'A',  
3     elem       = ['O', 'O'],  
4     pos        = [[0.000000, 0.000000, 0.000000],  
5                   [0.000000, 0.000000, 1.208]],  
6     O=6,  
7 )  
8 settings(  
9     machine     = 'ws4',  
10    pseudo_dir  = './pseudos/')
```

## Create pseudopotentials set

```
1 ppset(  
2     label      = 'ccecp',  
3     qmcpack    = ['O.ccECP.xml'],  
4 )
```

## Point programs towards pseudopotentials

```
1 scf = generate_pyscf(  
2     ...  
3     mole      = obj(  
4         basis  = 'ccecp-ccpvtz',  
5         ecp    = 'ccecp',
```

```
1 qmc = generate_qmcpack(  
2     identifier = 'dmc',  
3     ...  
4     system     = system,  
5     pseudos    = 'ccecp',
```

# Oxygen dimer: workflow

## Calibration runs([run1\\_calibration.py](#))

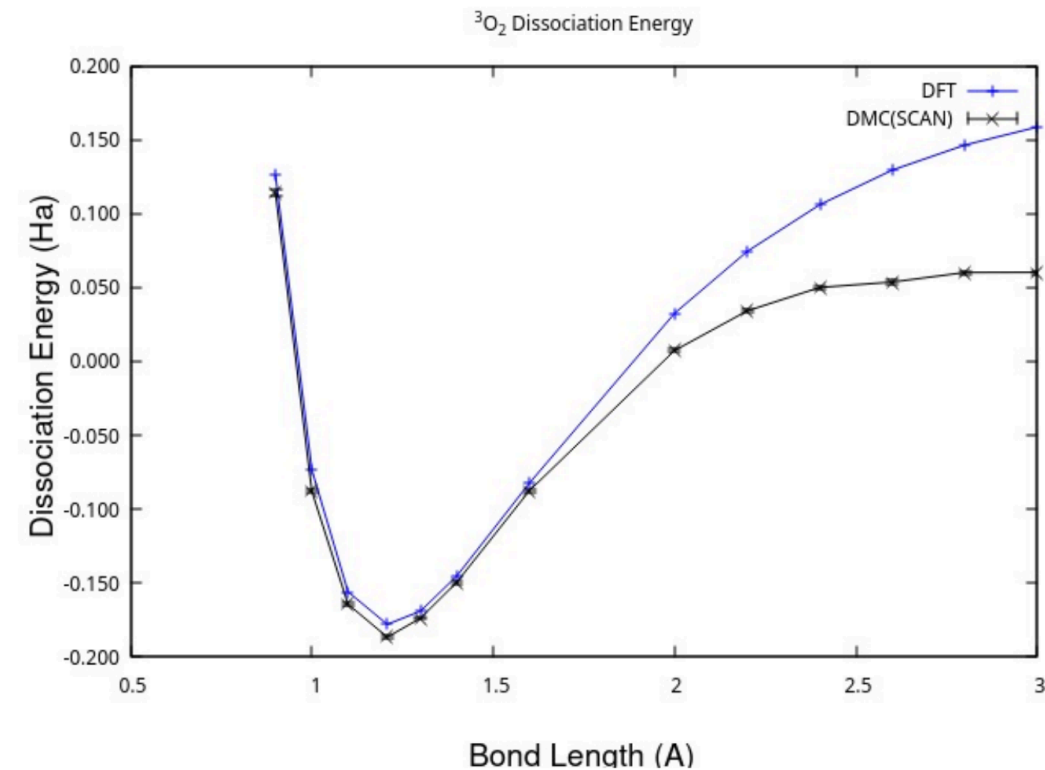
- determine appropriate timestep, walker count, acceptable error

## Production ([run2\\_dissociation\\_curve.py](#))

run2\_dissociation\_curve.py

```

1 oo_dists=[0.9, 1.0, 1.1, 1.208, 1.3, 1.4, 1.6]
2 for i in oo_dists:
3     system = generate_physical_system(
4         units      = 'A',
5         elem       = ['O','O'],
6         pos        = [[0.000000, 0.000000, 0.000000],
7                       [0.000000, 0.000000, i]],
8         O=6,
9     )
10    scf = generate_pyscf(...)
11    c4q = generate_convert4qmc(...)
12
13    orbdeps = [(c4q, 'particles'),
14              (c4q, 'orbitals' )]
15
16    optJ2 = generate_qmcpack(...)
17    optJ3 = generate_qmcpack(...)
18    qmc = generate_qmcpack(...)
```





## Additional Considerations

---

### How to start your own system

The materials of this workshop are a great starting place.

- while the scale of these problem are small, the workflow files give a great framework to start from
- Tune parameters to make the most of your computational resources
- We've provided the start of a water molecule example in the materials for you to try and set up your own workflow.  
[session4\\_molecules/03\\_water\\_molecule/](#)

## Additional Considerations

### Choices of trial wave functions

Choice of DFT functional

Multideterminant expansions

- link: previous workshop discussing selected CI
- link: qmcpack manual

Orbital optimization

- link: previous workshop with examples

TABLE I. Total energies of Be and Be<sub>2</sub> and the Be<sub>2</sub> dissociation energy computed with DMC using various trial functions.

Trial function <sup>a</sup>	Total energy (a.u.)		D <sub>e</sub> (cm <sup>-1</sup> )
	Be <sup>b</sup>	Be <sub>2</sub>	
HF/QZ- <i>g</i>	-14.657 30(4)	-29.317 89(6)	724(21)
LDA/QZ- <i>g</i>	-14.657 21(4)	-29.319 77(7)	1174(25)
PBE/QZ- <i>g</i>	-14.657 31(5)	-29.319 60(8)	1094(26)
BLYP/QZ- <i>g</i>	-14.657 25(4)	-29.319 56(8)	1113(26)
B3LYP/QZ- <i>g</i>	-14.657 27(3)	-29.319 46(8)	1079(23)
PBE0/QZ- <i>g</i>	-14.657 28(3)	-29.319 07(8)	992(21)
BH&HLYP/QZ- <i>g</i>	-14.657 26(5)	-29.318 91(7)	966(26)
BD/QZ- <i>g</i>	-14.657 18(4)	-29.318 72(7)	955(24)
CAS(4,8)/QZ- <i>fg</i> <sup>c</sup>	-14.667 23(1)	-29.337 07(3)	573(8)
CAS(4,16)/QZ- <i>fg</i> <sup>c</sup>	-14.667 30(1)	-29.338 32(3)	819(8)
Ext. CAS(4,16)/QZ- <i>fg</i>	-14.667 30(1)	-29.338 41(2)	838(7)
CAS(4,16)/QZ- <i>g</i> <sup>c</sup>	-14.667 27(2)	-29.338 38(3)	845(8)
Ext. CAS(4,16)/QZ- <i>g</i>	-14.667 27(2)	-29.338 45(2)	857(9)
CI/QZ- <i>g</i> <sup>c</sup>	-14.667 25(1)	-29.338 48(2)	873(6)
Ext. CI/QZ- <i>g</i>	-14.667 25(1)	-29.338 64(2)	908(6)
Experimental <sup>d</sup>	-14.667 356	-29.338 97	934.9(4)



## Summary

---

### The goals were to learn:

How to drive the calculations using Nexus

- examples of beryllium dimer, and oxygen binding curve

How to define a trial wave function

- convert4qmc, defining a jastrow factor

How to optimize  $\Psi_T$

- convergence in energy and variance and the ratio between the two
- nexus does have a way to guide this decision

Considerations for our DMC calculations

- timestep, desired error bars, population bias

Next session: Solid-State calculations Tuesday, July 15th 11AM EST US time