

Users' guide of MakePES

Kiyoshi Yagi
kiyoshi.yagi@riken.jp

Theoretical Molecular Science Laboratory
RIKEN Cluster for Pioneering Research

2019/05/14

Contents of Sample Files

Sample files are found in `sindo-4.0/doc/MakePES/sample_MakePES`

0.harmonic_h2co

1.qff_h2co: Quartic force field for H₂CO

1-1.single

1-2.parallel

1-3.dryrun

1-4.generic

2.grid_h2co: Grid PES for H₂CO

2-1.1MR

2-2.2MR

2-3.3MR

2-4.1MR_generic

3.mrpes_h2co: Multi-resolution PES for H₂CO

4.water-hexamer

NOTE

MakePES is a command line based program. This manual assumes that you are familiar with basic commands in UNIX. Shell scripts are given for Bourne Shell (bash).

This manual also assumes that an alias is set to invoke RunMakePES,

```
sindo_jar=/path/to/sindo-4.0/jar  
alias RunMakePES='java -cp "$sindo_jar/*" RunMakePES'
```

Thus, the command “RunMakePES” in this document is the same as the java command above.

For a theoretical background on PES generation, see Lecture Notes #2 (koug2.pdf).

0.harmonic_h2co

Proceed to 0.harmonic_h2co and find an input file to perform harmonic vibrational analysis for formaldehyde using Gaussian.

```
> cd 0.harmonic_h2co
> ls
h2co-b3lyp-dz.inp    log/
```

- “log” folder contains sample output files.

h2co-b3lyp-dz.inp is the input file. Run Gaussian by the following command:

```
> runGaussian.sh ./ h2co-b3lyp-dz.inp
```

- review the installation if you cannot run Gaussian with this command.
- the two arguments are (working folder) and (input file), respectively.

You will obtain the following output files when the job ends.

```
> ls h2co-b3lyp-dz.*
h2co-b3lyp-dz.chk    h2co-b3lyp-dz.fchk    h2co-b3lyp-dz.inp
h2co-b3lyp-dz.out
```

.fchk is a formatted check point file, which archives the result of quantum chemistry calculations. Let us convert the fchk file to a minfo file,

```
> java -cp "/path/to/sindo-4.0/jar/*" Fchk2Minfo h2co-b3lyp-dz
```

- The argument after Fchk2Minfo is the basename of output files.

You will find a minfo file,

```
> ls h2co-b3lyp-dz.*  
h2co-b3lyp-dz.chk    h2co-b3lyp-dz.fchk  h2co-b3lyp-dz.inp  
h2co-b3lyp-dz.out   h2co-b3lyp-dz.minfo
```

Minfo file includes the equilibrium geometry, harmonic frequencies, and vibrational displacement vectors.

The same result can be obtained by JSindo; refer to the usage manual of JSindo. You can find details about the format of minfo file therein, too.

1.qff_h2co

1-1.single

Proceed to 1.qff_h2co/1-1.single to find input files to generate quartic force field (QFF) for formaldehyde,

```
> cd 1.qff_h2co/1-1.single
> ls
GaussianTemplate      makePES.xml          resources.info       log/
```

makePES.xml is the main file, which is structured using tags in xml format. It is divided into sections by, <makePES> ... </makePES>, <qchem> ... </qchem>, and <qff> ... </qff>. The options are specified in each section by <key value="value" />. The value is case insensitive except for filenames. Comment out is possible as usual by <!-- ... -->.

makePES

```
<makePES>
  <minfoFile value="../../0.harmonic_h2co/h2co-b3lyp-dz.minfo" />
  <MR      value="3" />
  read a minfo file of H2CO
  the order of the mode coupling expansion.
```

makePES

```
<qchem>
  <program value="gaussian" />   use Gaussian
  <dryrun value="false"/>       run Gaussian
  <removefiles value="true" />   remove the output of Gaussian
  <title value="B3LYP/cc-pVDZ" /> set the title
  <template value="GaussianTemplate" /> the name of template file to
  generate Gaussian input
</qchem>
<qff>
  <stepsize value="0.5" />      step size of numerical differentiations
  <ndiffstype value="hess"/>    numerical differentiations using Hessian
  <mopfile value="prop_no_1.mop" /> name of a mop file
</qff>
</makePES>
```

resources.info provides hostname of nodes to run Gaussian. When we run on a single node, it is not important (but still, it should exist). We will later discuss this file in detail for parallel calculations in 1.2-parallel.

GaussianTemplate is a template file to generate input files for Gaussian specified by <template> in <qchem> section. It is the same as the usual input file for Gaussian, except for the red colored text.

```
                                GaussianTemplate

%chk=#basename#.chk
%NprocShared=2
%mem=1GB
#P B3LYP/CC-PVDZ FREQ NOSYMMETRY MAXDISK=240GB

Frequency at B3LYP/cc-pVDZ

0 1
#coordinate#
```

MakePES replaces #basename# and #coordinate# by the filename and the coordinates, respectively, to create input files.

"FREQ" keyword is necessary because we use numerical differentiations of the Hessian matrix (i.e., ndiffdtype = hess).

MakePES is invoked by the following command

```
> RunMakePES >& makePES.out
```

The main input file is set to makePES.xml by default. You can specify a different input file with `-f` option,

```
> RunMakePES -f makePES.xml >& makePES.out
```

In the output file, the options are first printed, and then the status of electronic structure calculations is printed,

makePES.out

Execute electronic structure calculations.

Thread0> Running minfo.files/mkqff-eq.inp on kyagi-mac3.local at ...

Thread0> Running minfo.files/mkqff0-0.inp on kyagi-mac3.local at ...

Thread0> Running minfo.files/mkqff0-1.inp on kyagi-mac3.local at ...

During this step, Gaussian jobs are carried out in a folder minfo.files. Because we've set `<removefiles>` to true, the input and output files are removed leaving only minfo files in the folder.

When this step is done, you will see an output like this,

```
makePES.out

End of electronic structure calculations.
Storing electronic structure data in tempfile ... Done!
Generating prop_no_1.mop... Done!
Removing the tempfiles ... Done!
End of QFF generation.
```

The QFF coefficients are written to <mopfile>, i.e., prop_no_1.mop.

```
1.1832573615027308000000e-15 1
2.7040635655127780000000e-03 1 1
1.1590249883181242000000e-17 1 1 1
2.2875576159959352000000e-05 1 1 1 1
-2.1030452203654645000000e-16 2
```

one-body terms:
ci, cii, ciii, ciiii

```
-4.189735728085525000000e-18 1 2
-6.3982329963326570000000e-14 1 2 2
1.8221212840606143000000e-13 1 2 2 2
-1.9301098360709593000000e-15 1 1 2
2.9510063534107640000000e-05 1 1 2 2
3.2272058841394823000000e-15 1 1 1 2
9.6870281798527820000000e-19 1 3
3.1255300969821536000000e-13 1 3 3
```

two-body terms:
cij, cijj, cijjj, ciiij, ciiijj, ciiiij

```
1.443395067341624000000e-13 1 2 3
-1.6315521638821910000000e-13 1 2 3 3
-3.0819651776043526000000e-13 1 2 2 3
6.1712325565851670000000e-14 1 1 2 3
1.8082830483224921000000e-13 1 2 4
```

three-body terms:
cijk, cijkk, cijjk, ciijk

Note that four-body terms (cijkl) are missing because <MR> was set to 3.

1-2.parallel

The electronic structure calculations are an exclusive bottleneck for generating the PES. In the case of H₂CO, FREQ calculations at 13 grid points are required and they were carried out one by one in one node in the previous section.

MakePES can distribute the grid points to multiple nodes, and process the FREQ calculations in parallel. This function substantially speeds up the calculation. It requires that the nodes have shared disks (via NFS), where the input files as well as `sindo/gaussian` are located, and are inter-connectable with SSH without being asked for a password.

Proceed to `1-2.parallel` to find the same set of input files.

```
> cd 1.qff_h2co/1-1.parallel
> ls
GaussianTemplate      makePES.xml           resources.info        log/
```

Assuming that we use 16 core x 2 nodes, we make the following modification to `resources.info` and `GaussianTemplate`

resources.info

```
beluga01  
beluga01  
beluga02  
beluga02
```



The hostname of 2 nodes.

We will run 2 processes in each nodes.

GaussianTemplate

```
%chk=#basename#.chk
```

```
%NprocShared=8
```

Each Gaussian process uses 8 cores

```
%mem=1GB
```

```
...
```

makePES.xml is the same as before. Set an environment variable `SINDO_RSH=ssh`, and then invoke `MakePES` as before,

```
> export SINDO_RSH=ssh  
> RunMakePES >& makePES.out
```

You can see in the output that the grid points are distributed to `beluga01` and `02`, each in 2 processes.

makePES.out

Execute electronic structure calculations.

Thread2@beluga02> Running minfo.files/mkqff0-1.inp on beluga02 at ...

Thread3@beluga02> Running minfo.files/mkqff1-0.inp on beluga02 at ...

Thread0@beluga01> Running minfo.files/mkqff-eq.inp on beluga01 at ...

Thread1@beluga01> Running minfo.files/mkqff0-0.inp on beluga01 at ...

When you want to stop the job, a safe way to terminate all Gaussian jobs is to create a file with a name “terminate”.

```
> touch terminate
```

Then, the job stops after Gaussian jobs that are currently running are all finished.

If you want to immediately stop the job, you have to kill the main process, i.e., the java process responsible for RunMakePES. In that case, however, check carefully that all Gaussian child processes are killed as well.

1-3.dryrun

The <dryrun> option generates input files for grid points, and then stops the program without executing Gaussian. Proceed to 1-3.dryrun,

```
> cd 1.qff_h2co/1-3.dryrun
> ls
GaussianTemplate   log1_dryrun_true/   log2_dryrun_false/
makePES.xml        resources.info
```

The only difference is the value of <dryrun> in makePES.xml

```
makePES.xml
<qchem>
  <program value="gaussian" />
  <dryrun value="true"/> stop after generating the input files
```

Running the program creates the input files for Gaussian in the minfo.files folder,

```
> RunMakePES >& makePES.out
> ls minfo.files/
mkqff-eq.inp  mkqff0-1.inp  mkqff1-1.inp  mkqff2-1.inp
mkqff3-1.inp  mkqff4-1.inp  ..
```

You may transfer these input files to other computer systems and carry out Gaussian there. Then, convert the formatted checkpoint files to minfo format using Fchk2Minfo.

```
> java -cp "/path/to/sindo-4.0/jar/*" Fchk2Minfo mkqffx-x
```

Bring back the minfo files and locate them in the minfo.files folder,

```
> ls minfo.files/  
mkqff-eq.inp      mkqff-eq.minfo  mkqff0-0.inp    mkqff0-0.minfo  
mkqff0-1.inp     mkqff0-1.minfo  ...
```

Change dryrun to false and run the program again.

```
makePES.xml  
  
<qchem>  
  <program value="gaussian" />  
  <dryrun value="false"/>  don't stop after generating the input files
```

```
> RunMakePES >& makePES.out  
> ls  
GaussianTemplate  makePES.out      makePES.xml      minfo.files/  
prop_no_1.mop    resources.info
```

You will see that the program immediately produces the mopfile.

The two log folders contains the files for the first step (log1_dryrun_true), and the files for the second step (log2_dryrun_false).

Note that, in general, MakePES looks into minfo.files folder for minfo files before starting Gaussian jobs. The job is skipped if a minfo file is found, and starts from the grid point where it ended before. In this example, we provided all minfo files, and thus the electronic structure calculations were all skipped.

1-4.generic

Setting <program> to generic prints the coordinates to a file in xyz format. You have to generate the input files, carry out the electronic structure calculations, and return the information in minfo format by yourself. Nevertheless, this may be useful for users who wish to use programs other than Gaussian.

Proceed to 1-4.generic to find makePES.xml.

```
> cd 1.qff_h2co/1-4.generic
> ls
log1_genxyz/ log2_genmop/ makePES.xml
```

The file is different only in <qchem> section,

```
makePES

<qchem>
  <program value="generic" />      "generic" means no specific program
  <title   value="B3LYP/cc-pVDZ" />
  <xyzfile value="makeQFF" />     set the name of xyz file
</qchem>
```

Running the program creates makeQFF.xyz,

```
> RunMakePES >& makePES.out  
> ls  
makePES.out  makePES.xml  makeQFF.xyz
```

makeQFF.xyz is written in the usual xyz format,

```
                                makeQFF  
4                                The number of atoms  
mkqff-eq                        name of the first point  
C      0.0000000000      0.0000000000      -0.6014736819  
O      0.0000000000      -0.0000000000      0.6027247362  
H      0.0000000000      0.9459644267      -1.2020174143  
H     -0.0000000000     -0.9459644267      -1.2020174143  
4  
mkqff0-0                        name of the second point  
C     -0.0125897498     -0.0000000000     -0.6014736819  
O      0.0031430124      0.0000000000      0.6027247362
```

The name, colored in red, is the ID of each grid points. We assume you carry out the electronic structure calculations by yourself, and collect the data in a minfo file with a name, ID.minfo.

Locate the minfo files in a minfo.files folder,

```
> ls minfo.files/  
mkqff-eq.minfo  mkqff0-1.minfo  mkqff1-1.minfo  mkqff2-1.minfo  
mkqff3-1.minfo  mkqff4-1.minfo  mkqff5-1.minfo  mkqff0-0.minfo  
mkqff1-0.minfo  mkqff2-0.minfo  mkqff3-0.minfo  mkqff4-0.minfo  
mkqff5-0.minfo
```

Run the program again (no need to change anything in makePES.xml).

```
> RunMakePES >& makePES.out  
> ls  
makePES.out  makePES.xml  makeQFF.xyz  minfo.files/  
prop_no_1.mop
```

The program produces the mopfile.

The two log folders contains the files for the first step (log1_genxyz), and the files for the second step (log2_genmop).

2.grid_h2co

2-1.1MR

Proceed to 2.grid_h2co/2-1.1MR to find input files to generate grid PES for formaldehyde,

```
> cd 2.grid_h2co/2-1.1MR
> ls
GaussianTemplate      makePES.xml           resources.info        log/
```

makePES.xml has the same <qchem> section as before. A new section <grid> replaces <qff>.

```
makePES
<makePES>
  <minfoFile value="../../0.harmonic_h2co/h2co-b3lyp-dz.minfo" />
  <MR value="1" />      MR=1 for 1MR-PES
  <dipole value="true" /> calculate dipole moment surface
  ...
  <grid>
    <ngrid value="11" /> number of grid points for each coordinates
    <fullmc value="true"/> calculate all modes
  </grid>
</makePES>
```

The generation of a gridPES requires only the energy at grid points, so that `FREQ` is no longer needed in GaussianTemplate. **Don't forget to remove `FREQ`** if you copy the file from QFF. Note that MakePES still works as long as the energy is printed in the output; however, it would be an enormous waste of time! `SCF=TIGHT` is recommended for HF/DFT calculations.

GaussianTemplate

```
%chk=#basename#.chk
%NprocShared=8   Each Gaussian process uses 8 cores
%mem=1GB
#P B3LYP/CC-PVDZ SCF=TIGHT NOSYMMETRY MAXDISK=240GB
...              Don't put a FREQ keyword!
```

Modify `resource.info` and `%NprocShared` for your system. In this sample, we run 8 processes of Gaussian with 8 cores (64 cores in total).

resources.info

```
beluga01 }
beluga01 } 4 nodes x 2 = 8 processes
...      }
beluga04 }
beluga04 }
```

Set an environment variable `SINDO_RSH=ssh`, and then run `MakePES`,

```
> export SINDO_RSH=ssh
> RunMakePES >& makePES.out
```

You will find in the output,

```
makePES.out

Setup MakeGrid module

o ngrid = 11
o 1MR Grid:
  1  2  3  4  5  6 } 6 modes x 11 grid = 66 grid points

Enter GridPES generation:

Execute electronic structure calculations.

Thread2@beluga02.local> Running minfo.files/mkg-q1-11-1.inp on beluga02 at ...
Thread3@beluga02.local> Running minfo.files/mkg-q1-11-2.inp on beluga02 at ...
Thread0@beluga03.local> Running minfo.files/mkg-eq.inp on beluga03 at ...
```

After the energy calculations are done, pot/dipole files are created.

makePES.out

Generating pot files.

```
o q1.pot [OK]
o q1.dipole [OK]
...
o q6.dipole [OK]
```

} pot and dipole files are created.

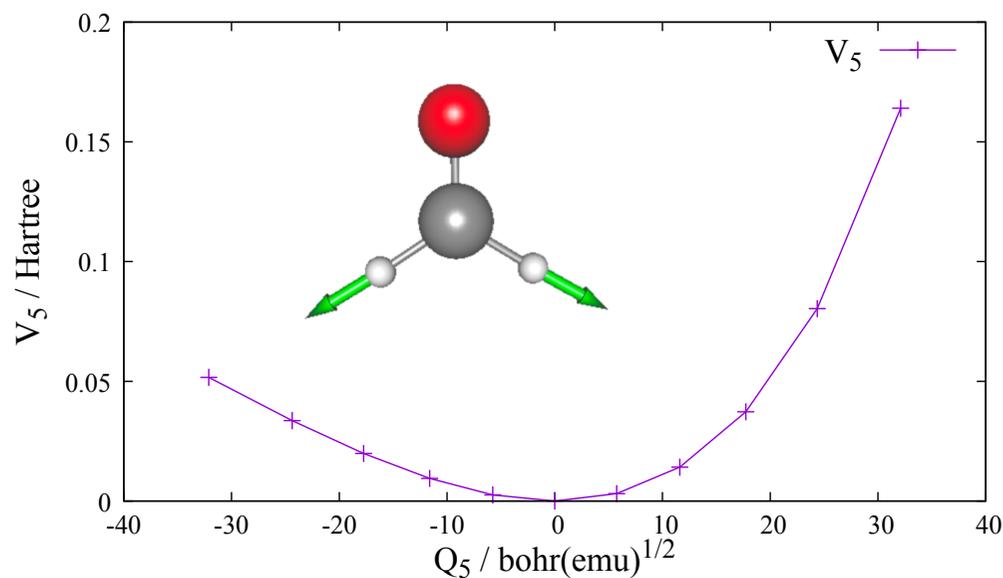
End of GridPES generation:

```
> ls q*
q1.dipole  q2.dipole  q3.dipole  q4.dipole  q5.dipole  q6.dipole
q1.pot     q2.pot     q3.pot     q4.pot     q5.pot     q6.pot
```

$qN.pot$ and $qN.dipole$ contain the change of the potential energy and dipole moment, respectively, with respect to the equilibrium geometry along mode N . The values at the equilibrium geometry are written in $eq.pot$ and $eq.dipole$.

grid points along Q_5

B3LYP/cc-pVDZ				B3LYP/cc-pVDZ				
# Number of grids and data				# Number of grids and data				
11	1			11	3			
#	q5	Energy V_5		#	q5	X dx	Y dy	Z dz
-32.11897636		5.1637111962e-02		-32.11897636		6.5354419100e-16	-2.1581328800e-14	2.2264673800e-01
-24.36885358		3.3576631399e-02		-24.36885358		1.5282778610e-15	-3.2503038080e-13	1.6380302600e-01
-17.73801106		1.9823126834e-02		-17.73801106		-3.5132508090e-15	2.9134816400e-15	1.1545683200e-01
-11.61455479		9.4296641468e-03		-11.61455479		6.1520087100e-16	-2.6271289280e-13	7.2992832000e-02
-5.75063885		2.5713497506e-03		-5.75063885		6.2180322100e-16	9.1610204700e-14	3.4762361000e-02
-0.00000000		0.0000000000e+00		-0.00000000		0.0000000000e+00	0.0000000000e+00	0.0000000000e+00
5.75063885		3.0938756174e-03		5.75063885		-5.2036891900e-16	-7.6053530000e-15	-3.1664072000e-02
11.61455479		1.4162827543e-02		11.61455479		-4.5814996000e-17	-2.2601766400e-14	-6.0408552000e-02
17.73801106		3.7256435616e-02		17.73801106		2.2404410010e-15	1.0871512036e-14	-8.6260919000e-02
24.36885358		8.0356295312e-02		24.36885358		3.7191305100e-16	1.0853080599e-14	-1.0913078900e-01
32.11897636		1.6406728197e-01		32.11897636		-1.4172511000e-16	1.1229081913e-14	-1.2867327600e-01
q5.pot (END)				q5.dipole (END)				



Plots of V_5 at grid points of Q_5 . Q_5 (symmetric CH stretching mode) is visualized in the inset. Note that the arrow corresponds to the negative direction of Q_5 , that is, the potential becomes flat as the CH bond extends.

2-2.2MR

In this section, we generate 2MR-grid PES for (Q1, Q2) and (Q5, Q6). Proceed to 2.grid_h2co/2-2.2MR,

```
> cd 2.grid_h2co/2-2.2MR
> ls *pot *dipole
eq.dipole  q1.dipole  q2.dipole  q5.dipole  q6.dipole
eq.pot     q1.pot     q2.pot     q5.pot     q6.pot
```

The pot and dipole files obtained in Sec. 2-1 for the equilibrium geometry and along Q1, Q2, Q5, Q6 are placed in the same folder. These files provide the information along the coordinates, and thus reduce the cost.

makePES

```
<makePES>
  <minfoFile value="../0.harmonic_h2co/h2co-b3lyp-dz.minfo" />
  <MR value="2" />      MR=2 for 2MR-PES
  ...
  <grid>
    <ngrid value="9" />      number of grid points is reduced to 9
    <mc2 value="1,2, 5,6"/> (Q1, Q2) and (Q5, Q6)
  </grid>
</makePES>
```

The two-mode terms are specified by <mc2> to (Q1,Q2) and (Q5,Q6). See the appendix on the details of format of mc2.

Calculating these terms with ngrid = 9 would require $9 \times 9 \times 2 = 162$ grid points. In this case, the information along the axis is provided by the files, so that the number of grid points is reduced to $8 \times 8 \times 2 = 128$.

GaussianTemplate and resources.info are the same as before. Run MakePES,

```
> export SINDO_RSH=ssh
> RunMakePES >& makePES.out
```

You may check the settings in the output,

```
makePES.out

Setup MakeGrid module

o ngrid = 9          number of grid points
o 1MR Grid:
  1 2 5 6          1-mode terms
o 2MR Grid:
  (1,2) (5,6)      2-mode terms

Enter GridPES generation:
```

pot/dipole files are created at the end of the calculation.

```
makePES.out

Generating pot files.

o q2q1.pot [OK]
o q2q1.dipole [OK]
o q6q5.pot [OK]
o q6q5.dipole [OK]
} 2-mode terms (9x9 grid points) are created.
End of GridPES generation:
```

In addition to the original files with 11 grid points (q[1-6].pot/dipole), new files for 2MR-PES with 9 grid points (q2q1.pot/dipole and q6q5.pot/dipole) are created.

```
> ls q*
q1.dipole    q2.pot      q5.dipole   q6.pot
q1.pot      q2q1.dipole q5.pot      q6q5.dipole
q2.dipole   q2q1.pot    q6.dipole   q6q5.pot
```

2-3.3MR

In this section, we generate 3MR-grid PES for (Q4, Q5, Q6). Proceed to 2.grid_h2co/2-3.3MR,

```
> cd 2.grid_h2co/2-3.3MR
> ls *pot *dipole
eq.dipole      q4.dipole      q5.dipole      q6.dipole      q6q5.dipole
eq.pot         q4.pot         q5.pot         q6.pot         q6q5.pot
```

Again, we use the existing information (q4, q5, q6, and q6q5) to reduce the cost. By placing these files in the same folder, the grid points are reduced from 729 (=9 x 9 x 9) points to 640 points.

```
                                makePES
<makePES>
  <minfoFile value="../0.harmonic_h2co/h2co-b3lyp-dz.minfo" />
  <MR      value="3" />      MR=3 for 3MR-PES
  ...
  <grid>
    <ngrid value="9" />
    <mc3 value="4,5,6"/>    (Q4, Q5, Q6)
  </grid>
</makePES>
```

GaussianTemplate and resources.info are the same as before. Run MakePES,

```
> export SINDO_RSH=ssh
> RunMakePES >& makePES.out
```

You will find in the output that the 2-mode terms, (Q4, Q5) and (Q4, Q6), are automatically added,

```
makePES.out

Setup MakeGrid module

o ngrid = 9
o 1MR Grid:
  4 5 6
o 2MR Grid:
  (4,5) (4,6) (5,6)
o 3MR Grid:
  (4,5,6)

Enter GridPES generation:
```

1- and 2-mode terms that are subsets of the 3-mode terms are automatically added by the program.

the 3-mode term specified in the input

After iteration over the grid points, we obtain pot and dipole files for (Q4, Q5), (Q4, Q6), and (Q4, Q5, Q6).

```
makePES.out

Generating pot files.

o q5q4.pot [OK]
o q5q4.dipole [OK]
o q6q4.pot [OK]
o q6q4.dipole [OK]
o q6q5q4.pot [OK]
o q6q5q4.dipole [OK]
End of GridPES generation:
```

Automatically added 2-mode terms

The target 3-mode term

```
> ls q*
q4.dipole      q5q4.dipole    q6q4.dipole    q6q5q4.dipole
q4.pot        q5q4.pot       q6q4.pot       q6q5q4.pot
q5.dipole     q6.dipole      q6q5.dipole
q5.pot        q6.pot         q6q5.pot
```

2-4.1MR_generic

In this section, we illustrate the generic mode for grid PES. Proceed to 2-4.1MR_generic,

```
> cd 2.grid_h2co/2-4.1MR_generic
> ls
log1_genxyz/ log2_genpot/ makePES.xml
```

makePES.xml has <qchem> section as follow,

```
makePES

<makePES>
...
<qchem>
  <program value="generic" />    "generic" means no specific program
  <title   value="B3LYP/cc-pVDZ" />
  <xyzfile value="makeGrid" />   set the name of xyz file
</qchem>
</makePES>
```

Running the program creates makeGrid.xyz,

```
> RunMakePES >& makePES.out
> ls
makePES.out  makePES.xml  makeGrid.xyz
```


3.mrpes_h2co

In this section, we calculate a multi-resolution PES, which is a combination of QFF and grid PES. All 1-mode terms are obtained by the grid method using 11 points. Mode coupling strength (MCS) is derived from QFF coefficients, and coupling terms with $MCS > 10.0$ are generated by the grid method using 9 points.

Proceed to 3.mrpes_h2co to find input files,

```
> cd 3.mrpes_h2co
> ls
GaussianTemplate1  GaussianTemplate2  log/
makePES.xml       resources.info
```

GaussianTemplate1 and GaussianTemplate2 are the template files to calculate a Hessian matrix (FREQ) and only the energy, respectively.

In makePES.xml, two `<qchem>` sections are given which specifies GaussianTemplate1 and GaussianTemplate2 attributed to `id = "freq"` and `"ene"`, respectively. The ID is associated with QCID of `<qff>` and `<grid>`.

makePES.xml

```
<qchem id="freq">
  <program value="gaussian" />
  ...
  <template value="GaussianTemplate1" />
</qchem>
<qchem id="ene">
  <program value="gaussian" />
  ...
  <template value="GaussianTemplate2" />
</qchem>
<qff>
  <QCID value="freq" />
  <mopfile value="prop_no_1.mop" />
  ...
</qff>
<grid>
  <QCID value="ene" />
  <ngrid value="11" />
  <mc1 value="1-6"/>
</grid>
<grid>
  <QCID value="ene" />
  <ngrid value="9" />
  <mcsstrength value="10"/>
  <mopfile value="prop_no_1.mop"/>
</grid>
```

The diagram illustrates the XML structure and connections between elements. An orange bracket groups the first two `<qchem>` blocks. A blue bracket groups the second `<qchem>` block and the `<qff>` block. An orange arrow points from the `<QCID value="freq" />` element to the `<QCID value="ene" />` element in the first `<grid>` block. A blue arrow points from the `<QCID value="ene" />` element in the second `<grid>` block to the `<QCID value="ene" />` element in the first `<grid>` block. A green arrow points from the `<mopfile value="prop_no_1.mop" />` element in the `<qff>` block to the `<mopfile value="prop_no_1.mop"/>` element in the second `<grid>` block.

<qchem> sections are followed by one <qff> and two <grid> sections. MakePES processes these sections **in the order they appear** in the input file. In the present example,

1. Generate QFF
prop_no_1.mop is created.
2. Generate 1MR-grid PES with ngrid = 11
q1 - q6.pot / dipole files are created.
3. Generate 3MR-grid PES with ngrid = 9
pot/dipole files with MCS > 10 are created.

Step 1 (QFF) must precede Step 3, because prop_no.1.mop is needed to calculate MCS. Step 2 generates 1MR-grid PES, q1 - q6.pot. These files are used in Step 3.

Run the program by typing,

```
> export SINDO_RSH=ssh  
> RunMakePES >& makePES.out
```

Find in the makePES.out that the calculation is performed in the above order.

makePES.out

```
...  
Enter QFF generation:  
...  
Generating prop_no_1.mop... Done!  
Removing the tempfiles ... Done!  
End of QFF generation.  
  
Setup MakeGrid module  
o ngrid = 11  
o 1MR Grid:  
  1 2 3 4 5 6  
Enter GridPES generation:  
...  
Generating pot files.  
o q1.pot [OK]  
o q1.dipole [OK]  
...  
o q6.pot [OK]  
o q6.dipole [OK]...  
End of GridPES generation:
```

} QFF generation

} 1MR-grid PES generation

makePES.out

Setup MakeGrid module

o Setup MCS: Read QFF Data via prop_no_1.mop ... [OK]

o ngrid = 9

o 1MR Grid:

5 1 2 3 6 4

Reads the output of QFF calc.

o 2MR Grid:

(1,5) (2,5) (3,5) (1,6) (2,6) (3,6) (5,6) (2,3) (2,4) (4,6)

o 3MR Grid:

(2,3,6) (2,4,6)

Coupling terms with MCS > 10.0

Enter GridPES generation:

...

Generating pot files.

o q5q1.pot [OK]

o q5q1.dipole [OK]

...

o q6q4q2.pot [OK]

o q6q4q2.dipole [OK]

} 2, 3MR-grid PES generation

End of 3MR-GridPES generation:

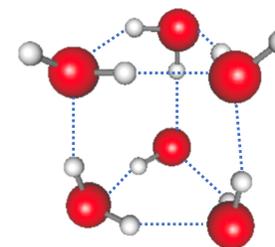
The program detects 10 two-mode terms and 2 three-mode terms where $MCS > 10.0$. When the calculation is done, you will find the following pot files in the working directory.

```
> ls *pot
eq.pot      q2.pot      q3q2.pot    q4q2.pot    q5q1.pot
q5q3.pot    q6q1.pot    q6q3.pot    q6q4.pot    q6q5.pot
q1.pot      q3.pot      q4.pot      q5.pot      q5q2.pot
q6.pot      q6q2.pot    q6q3q2.pot  q6q4q2.pot
```

4.water-hexamer

4-1.intra

In this section, we generate the PES of water hexamer using the intramolecular modes, i.e., OH stretching and HOH bending modes. There are 18 such modes, and corresponds to mode 31 – 48.



water hexamer
(prism)

Proceed to 4.water-haxamer/4-1.intra to find input files,

```
> cd 4.water-hexamer/4-1.intra
> ls
GaussianTemplate    h2o_6-mp2dz.minfo  log/
makePES.xml         resources.info
```

h2o_6-mp2dz.minfo is a minfo file that contains the equilibrium geometry and vibrational modes of water hexamer (prism shape) obtained at the MP2/c—pVDZ level of theory.

GaussianTemplate is a template file to perform FREQ calculations at the MP2/cc-pVDZ level of theory.

In makePES.xml, the active modes are specified by <activemode> to modes 31 – 48. Other sections, <qchem> and <qff>, are the same as before.

makePES.xml

```
<makePES>
  <minfoFile value="h2o_6-mp2dz.minfo" />
  <MR      value="3" />
  <activemode value="31-48" />
  <qchem>
    <program value="gaussian" />
    <dryrun value="false"/>
    <removefiles value="true" />
    <title value="MP2/cc-pVDZ" />
    <template value="GaussianTemplate" />
  </qchem>
  ...
```

active modes are set to the intramolecular modes, mode 31 – 48.

Run the program by,

```
> export SINDO_RSH=ssh
> RunMakePES >& makePES.out
```

Find in the output that mode 31-48 are active ,

```
makePES.out

o Input options read via makePES.xml ... [OK]
  - Molecular info via h2o_6-mp2dz.minfo ... [OK]
  - InterDomain = false
  - ActiveModes:
    * Domain 1
      31 32 33 34 35 36 37 38 39 40
      41 42 43 44 45 46 47 48
    - MR      = 3
    - Dipole = false
  } 31 – 48 are active, and 1
    } – 30 are inactive.
```

When the calculation ends, you will find that prop_no_1.mop has entry only for modes 31 – 48.

```
DALTON_FOR_MIDAS MP2/cc-pVDZ
1.2693093572610456000000e-07 31
3.8331230612195770000000e-03 31 31
5.2999094588990350000000e-05 31 31 31
-5.3384572915176750000000e-06 31 31 31 31
-3.8778975455244765000000e-06 32
3.8794005068453160000000e-03 32 32
1.1584964110841597000000e-04 32 32 32
-6.2778785001575310000000e-06 32 32 32 32
-2.5417319397503540000000e-06 33
3.8953636442193990000000e-03 33 33
```

Note that it is not easy to tell which modes should be set to active or inactive. The modes can be set to inactive later in the vibrational calculations using `sindo`. Therefore, I recommend to include as many modes as possible during the PES generation step.

Nevertheless, there are increasing demand to select vibrational modes as the size of system becomes large. For example, a solute in solvent, ligands in a protein, etc. A model that separates the inter- and intra-molecular modes, illustrated in this example, is often used for cluster systems. `<activemode>` is made to achieve such purposes.

Appendix: List of all keys

General keys

- minfoFile: file name
The name of minfo file containing the vibrational data. The value is case sensitive.
- MR: 1/2/3
The order of mode coupling expansion. Can take 1, 2, or 3. (default = 3)
- activemode: string of mode index
Specifies active modes for PES generation. All modes are active by default. The mode numbers should be separated by comma or space. A hyphen can be used for a sequence of mode number. For example,

```
<entry key="activemode"> 1,2,3,5 </entry>
```

is equivalent to,

```
<entry key="activemode"> 1-3 5 </entry>
```

which means Q_1, Q_2, Q_3 , and Q_5 are active, and Q_4 isn't.

- dipole: true/false
Generates the dipole moment surface in addition to the PES, when true. (default = false)

QCHEM section

ID: string

- program: string
 - gaussian ... Interface with Gaussian
 - generic ... Print the coordinates to a xyz file
- title: string
 - A title line that will be printed to PES files.

options for non-generic (=Gaussian)

- removefiles: true/false
 - Removes the input/output files of the quantum chemistry program, when true. (default = false)
- dryrun: true/false
 - Generates the input files for the quantum chemistry program and exits without execution, when true. (default = false)
- template: file name
 - The name of a template file to generate the input files for quantum chemistry jobs.

options for generic

- xyzfile: filename
 - Name of a xyz file, where the coordinates are written. (default = makeQFF for QFF and makeGrid for GRID)

QFF section

- QCID: string
The ID of associated <qchem>
- stepsize: real number
The step size for numerical differentiations in dimensionless unit ($\sqrt{\omega/\hbar} * Q$). (default = 0.5)
- ndifftype: grad or hess
The type of numerical differentiations.
grad : Numerical 3rd-order diff. of gradient.
hess (default) : Numerical 2nd-order diff. of hessian.
- mopfile: file name
The name of mop file, in which the QFF coefficients are written. (default = prop_no_1.mop)
This format is compatible with the MIDAS software developed by Christiansen and coworkers.
- interdomain_hc: true/false
Prints the harmonic coupling, when true. (default = true)

- gradient and hessian: input/current
Specifies where the gradient and Hessian are retrieved.
input (default) : From the input minfo file.
current : From the current calculation. (mkqff-eq.minfo)

“input” is useful for combining accurate geometry, gradient, and Hessian, read from the input minfo file, with lower-level cubic and quartic terms, which are calculated by MakePES module.

On the other hand, one might think of another strategy, where the geometry and coordinates are derived from a low-level of theory, and the QFF at a higher-level of theory. In that case, this option should be set to “current”, which incorporates the gradient and Hessian obtained from the current calculation.

- genhs: true/false
Generate the 001.hs file, when true. (default = false)
001.hs is a file which contains the QFF coefficients in the old format; however, this format is deprecated and not recommended to use unless for a debugging purpose to compare the result with the previous version of SINDO.

GRID section

- QCID: string
The ID of associated `<qchem>`
- ngrid: integer number
The number of grid points along each coordinates. (default = 11)
- fullmc: true/false
All the mode coupling up to the MR-th order is generated, when true. (default = false)
- mc1, mc2, mc3: string of mode index
The 1, 2, or 3MR terms separated by comma or space.

Examples:

- `<mc1 value="1,2,3,5" />` or `<mc1 value="1-3 5" />`

generates grid points for Q1,Q2,Q3, and Q5.

- `<mc2 value="1,2, 1,4, 2,4, 3,4" />` or `<mc2 value="1,2, 1-3,4" />`

generates the grid points for (Q2, Q1),(Q4, Q1),(Q4, Q2), and (Q4, Q3).

- `<mc3 value="1,2,3, 1,2,4" />`

generates the grid points for (Q3, Q2, Q1) and (Q4, Q2, Q1).

- **mcsstrength:** real number
The threshold value (in cm^{-1}) to select the mode coupling term for generating the grid potential.
The coupling terms with MCS larger than this value are generated.
- **mopfile:** file name
The name of mop file to obtain MCS.

NOTE: One of fullmc, mc1, mc2, mc3, or mcsstrength must be present to specify the coupling terms.