

# **Gaussian Basis AIMD**





#### Gaussian Basis AIMD: QMD Module



- Finite Cluster Molecular Dynamics
- Compatible with all Gaussian basis function based electronic structure methods in NWChem
  - DFT, LR-TDDFT, MP2
  - Will switch to numerical gradients if analytical gradients are absent
- Velocity Verlet
- Constant Energy & Constant Temperature Ensembles
  - Berendsen, Langevin, and Stochastic Velocity Rescaling

#### **Documentation:**

https://github.com/nwchemgit/nwchem/wiki/Gaussian-Basis-AIMD

Details: Journal of Physical Chemistry B, 120(8), 1429 (2015)





### Sample Input



```
# water - SO HF/6-31G* geometry
# ground state HF/6-31G* MD
# time step = 10 \text{ a.u.}
# number of steps = 200
# SVR thermostat, default tau
# 200 K
# translations and rotations every 10 steps
# print trajectory every 5 steps
# random seed set to 12345
#
```

```
echo
start qmd_dft_h2o_svr
print low
geometry noautosym noautoz
O 0.00000000
                -0.01681748
                             0.11334792
   0.00000000
                0.81325914
                            -0.34310308
H 0.00000000
               -0.67863597
                            -0.56441201
end
basis
 * library 6-31G*
end
dft
xc hfexch 1.0
end
amd
nstep nucl 200
dt_nucl 10.d0
targ_temp 200.d0
com step 10
rand seed 12345
thermostat svr
print_xyz 5
end
task dft amd
```

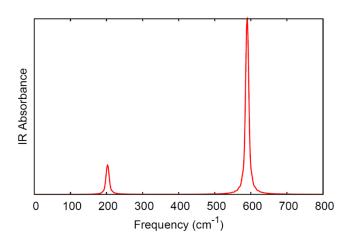


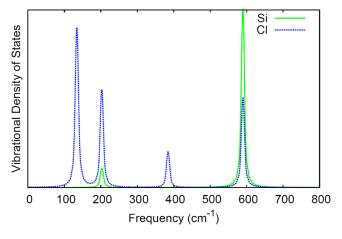


# Detailed Example: SiCl<sub>4</sub>



```
start SiCI4
echo
print low
geometry noautosym noautoz
Si
         -0.00007905
                       0.00044148
                                    0.0000001
CI
          0.71289590
                       1.00767685
                                     1.74385011
CI
         -2.13658008
                       -0.00149375
                                    -0.0000001
CI
          0.71086735
                      -2.01430142
                                    -0.0000001
CI
          0.71289588
                      1.00767684
                                    -1.74385011
end
basis
* library 6-31G
end
dft
xc hfexch 1.0
end
amd
nstep_nucl 20000
dt nucl
        10.0
targ_temp 20.0
com step
rand seed 12345
thermostat none
end
task dft amd
```









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## **Combining QMD with Properties**



```
echo
start amd_props
amd
nstep_nucl 200
dt nucl
        10.d0
targ_temp 200.d0
com_step
           10
rand seed 12345
thermostat berendsen
# calculate properties as defined in
# the properties block every 5 steps
property 5
end
property
dipole
aoresponse 1.0911267060
velocity
damping 0.007
shielding 1 1
end
```

task dft amd

NOTE: The choice of property can slow down the QMD depending on how oftern you want to calculate it





## **Combining QMD with TDDFT**



```
echo
start amd_tddft
amd
nstep_nucl 200
dt nucl
         10.d0
targ_temp 200.d0
com step
           10
rand seed 12345
thermostat berendsen
# calculate TDDFT every 3 steps
tddft 3
end
tddft
nroots 2
notriplet
end
task dft amd
```

NOTE: The choice of property/response can slow down the QMD depending on how oftern you want to calculate it



