

Tutorial : influence of environment on the uv-visible spectra of a small peptide : histidine amino-acid.

Prerequisite

- Gaussian
- Amber (sander, tleap)
- [Amber 12 manual](#)
- Any text editor (vi, emacs, gedit, geany or any you prefer)
- Visualisation programs : VMD, avogadro

1) Create a directory where you will put all your data. Go to this directory in a terminal

2) Creation of the small molecule histidine aminoacid using tleap

a) write done with an editor a script for tleap in the file tleap.in

\$ vi tleap.in

```
source leaprc.ff14SB #Source leaprc file for ff14SB protein force field
source leaprc.gaff #Source leaprc file for gaff
source leaprc.tip3p #Source leaprc file for TIP3P water model
his=sequence {ACE HIS NME} #create the peptide and store in variable named his
desc his #check what you have created
savepdb his his-ini.pdb #save the initial structure in a pdb file named his-
ini.pdb
saveamberparm his his-ini.prmtop his-ini.inpcrd #Save AMBER topology and
coordinate files for the structure in vacuo
solvatebox his TIP3PBOX 8 #Solvate the molecule with a cubic water box
addions mol Cl- 0 #Add Cl- ions to neutralize the system
savepdb his his-solv.pdb #Save a pdb file of the coord of peptide + water
saveamberparm his his-solv.prmtop his-solv.inpcrd #Save AMBER topology and
coordinate files for the structure in a water box
quit #Quit tleap program
```

b) run tleap and check that in the output file tleap.log, there is no warning or problems.

\$ tleap < tleap.in > tleap.log

you should now have created 6 files : his-ini.pdb his-ini.prmtop his-ini.inpcrd
his-solv.pdb his-solv.prmtop his-solv.inpcrd

c) open with vmd the files his-solv.pdb and his-ini.pdb.

Check which protonation state of the histidine was chosen by tleap.

As you can see a structure is far from the ideal one (two Hydrogen atoms are too close).

We will first relax the structure before doing any calculations.

3) optimization of the structure using sander

a) create input file for sander program to perform a minimization (remove the comments in pink):

\$ vi mini-ini.in

```
Minimize in vacuo
&cntrl
imin=1, maxcyc=200, (Perform minimization for 200 steps)
ntb=0, (Non-periodic simulation)
cut=20., (20 angstrom classical non-bond cut off)
&end
```

```
$ vi mini-solv.in
```

```
Minimize in Water
&cntrl
  imin=1,
  ntx=1,
  irect=0,
  maxcyc=200,
  ntpr=10,
  ntwx=0,
  cut=8.0,
/
```

b) run minimisation:

For structure in vacuo:

```
$ $AMBERHOME/bin/sander -O -i mini-ini.in -o his-iniMin.out -p his-
ini.prmtop -c his-ini.inpcrd -r his-iniMin.rst -inf his-iniMin.mdinfo &
```

sander uses a consistent syntax for each step of MD simulation. Here is a summary of the command line options of **sander**:

-O	Overwrite the output files if they already exist
-i 01_Min.in	Choose input file (default mdin)
-o 01_Min.out	Write output file (default mdout)
-p prmtop	Choose parameter and topology file prmtop
-c inpcrd	Choose coordinate file inpcrd
-r 01_Min.rst	Write output restart file with coordinates and velocities (default restrt)
-inf 01_Min.mdinfo	Write MD info file with simulation status (default mdinfo)
&	To make the calculation in background and be able to still use the terminal

Do the same for structure in water:

```
$ $AMBERHOME/bin/sander -O -i mini-solv.in -o his-solvMin.out -p his-
solv.prmtop -c his-solv.inpcrd -r his-solvMin.rst -inf his-solvMin.mdinfo &
```

check the .out files to see if the minimization is done (open with vi)

c) check that the obtained structures are indeed optimised (no more H atoms too close)

To do this, you need to creation of files .pdb using software ambpdb. Ambpdb need to have the correct topology file in name prmtop so the first command do a copy of the prmtopfile into prmtop and third command remove it after the use of ambpdb :

```
$ cp his-ini.prmtop prmtop
$ ambpdb < his-iniMin.rst > his-iniMin.pdb
$ rm prmtop

$ cp his-solv.prmtop prmtop
$ ambpdb < his-solvMin.rst > his-solvMin.pdb
$ rm prmtop
```

Open the .pdb files with vmd and check the structures.

d) If the structures do not seem to be optimized enough, rerun the minimization by increasing the number of cycles.

4) Calculation of the electronic transitions for the molecule in vacuo using Gaussian With Avogadro software, open his-*iniMin.pdb*
 Create a Gaussian input file in ground state of the structure, with the basis set 6-31G(d), functional B3LYP, and modify the file with an editor to have the heading for the calculation of the 3 first singlet states. Charge is equal to zero, spin should be singlet. The file should look like this :

`$vi his-iniMin-TDDFT.com`

```
%chk=his-iniMin-TDDFT.chk
# td=(singlets,nstates=3) b3lyp/6-31g(d)

TDDFT in vacuo

0 1
H      2.20900000  1.03600000  0.46300000
C      2.12900000  2.10000000  0.23100000
H      1.48800000  2.59300000  0.96300000
H      1.71300000  2.22700000 -0.76900000
C      3.50800000  2.71800000  0.28300000
O      4.45500000  2.08100000  0.72700000
N      3.61100000  3.96400000 -0.17600000
H      2.77200000  4.44300000 -0.46800000
C      4.85500000  4.74200000 -0.21200000
H      5.43800000  4.49500000  0.67700000
C      5.67800000  4.33500000 -1.45300000
H      5.71400000  3.24500000 -1.51300000
H      5.16600000  4.69400000 -2.35200000
C      7.10500000  4.82900000 -1.46000000
N      7.76300000  5.33600000 -2.58400000
C      9.01500000  5.60400000 -2.18800000
H      9.79900000  5.95200000 -2.85300000
N      9.17400000  5.30400000 -0.88000000
H     10.03300000  5.40100000 -0.35600000
C     7.98500000  4.79200000 -0.41400000
H     7.79600000  4.39200000  0.57400000
C     4.56000000  6.25900000 -0.17600000
O     3.40100000  6.66800000 -0.24300000
N     5.60800000  7.09400000 -0.06000000
H     6.52400000  6.67600000 -0.08400000
C     5.53300000  8.55600000 -0.00100000
H     4.49400000  8.88800000  0.01500000
H     6.03300000  8.91600000  0.89800000
H     6.02800000  8.98700000 -0.87300000
```

Check the transition energy (see in Gaussian output Excited States energies and oscillators section)

Remarks: It would be better to do before an optimization in vacuo using the same functional and basis set but it takes around one hour to perform it on one processor.

5) Calculation of the electronic transition for the molecule in implicit Water using Gaussian (QM/MM)

Take the obtained structure to perform a TD-DFT in Gaussian using the same basis set and functional for the 3 first singlet states but now with implicit water (PCM).

6) Calculation of the electronic transition for the molecule in explicit Water using Gaussian-ONIOM

Take the his-solvMin.pdb, file from the minimization in water.

a) First, we need to select only the water molecules that are at 6 Å from the peptide.

With VMD:

- Open his-solvMin.pdb

- To save pdb file with the selection of all the atoms of the peptide and the water where at least one atom is less than 6 Å from one atom of the peptide: First, check the selection "same resid as within 6 of resname ACE HIE NME" in the Graphical representation.

Then, in VMD Main, right click on the molecule and save Coordinate with selected atoms "same resid as within 6 of resname ACE HIE NME"

Save in file his-solv6.pdb

b) Open his-solv6.pdb with Avogadro and create a Gaussian input file.

c) We will modify with an editor this input file to make a QM/MM input file. All the water molecules will be in the MM description and the peptide in the QM description. It is a 2-layer ONIOM calculation.

See <http://gaussian.com/oniom/> for more information.

Modify the input to get this info:

\$ vi his-QMMM-TDDFT-UFF.com

```
%chk=his-QMMM-TDDFT.chk
# td=(nstates=3,root=1) ONIOM(b3lyp/6-31g(d):UFF) geom=connectivity

QM/MM TD calculation

0 1 0 1 0 1
H 0 11.57400000 11.09600000 13.96400000 H
C 0 11.47300000 12.18600000 13.83500000 H
H 0 10.73700000 12.53500000 14.55400000 H
H 0 11.12000000 12.37900000 12.82200000 H
C 0 12.78400000 12.85600000 14.10100000 H
O 0 13.60600000 12.31800000 14.83300000 H
N 0 12.98500000 14.02900000 13.51500000 H
H 0 12.27700000 14.38600000 12.89300000 H
C 0 14.22500000 14.81700000 13.55800000 H
H 0 14.78000000 14.55000000 14.47700000 H
C 0 15.10100000 14.43300000 12.35900000 H
H 0 15.11100000 13.34600000 12.28500000 H
H 0 14.64100000 14.82000000 11.45700000 H
C 0 16.49800000 14.99900000 12.35800000 H
N 0 17.24300000 15.30400000 11.20000000 H
C 0 18.50300000 15.52300000 11.61900000 H
H 0 19.31000000 15.90700000 10.98900000 H
N 0 18.57400000 15.41900000 12.94400000 H
H 0 19.40400000 15.52700000 13.50900000 H
C 0 17.34000000 15.09800000 13.43000000 H
H 0 17.15100000 14.74700000 14.43100000 H
```

C	0	13.92800000	16.32700000	13.64000000	H
O	0	12.75800000	16.71000000	13.75400000	H
N	0	14.98100000	17.15400000	13.69600000	H
H	0	15.91400000	16.76000000	13.57700000	H
C	0	14.84900000	18.61200000	13.82000000	H
H	0	13.80000000	18.95900000	13.87700000	H
H	0	15.36600000	18.96700000	14.72000000	H
H	0	15.30700000	19.10900000	12.96000000	H
O	0	17.96200000	20.70200000	16.86200000	L
H	0	16.98600000	20.73300000	16.67400000	L
H	0	18.28900000	21.05400000	16.00400000	L

.....
0 1 0 1 0 1 give the Charge and spin for entire molecule (real system), then the charge and spin for the high lever model (peptide) and the charge and spin for the low-model (waters)
You need to add the letter H at the end of each atom that is described in the high level model (peptide) and a L at the end of each atom that is described in the low-model (water).

The force field used is UFF

Check the transition energy (see in Gaussian output Excited States energies and oscillators section)

d) Let do the same with amber force field.

We need to modify the input file with the types of atoms :

\$ vi his-QMMM-TDDFT-UFF.com

```
%chk=his-QMMM-TDDFT-amb.chk
# td=(nstates=3,root=1) ONIOM(b3lyp/6-31g(d):Amber) geom=connectivity
```

QM/MM TD calculation

```
0 1
H-HC      0  11.57400000  11.09600000  13.96400000 H
C-CT      0  11.47300000  12.18600000  13.83500000 H
H-HC      0  10.73700000  12.53500000  14.55400000 H
H-HC      0  11.12000000  12.37900000  12.82200000 H
C-C-      0  12.78400000  12.85600000  14.10100000 H
O-O-      0  13.60600000  12.31800000  14.83300000 H
N-N-      0  12.98500000  14.02900000  13.51500000 H
H-H-      0  12.27700000  14.38600000  12.89300000 H
C-CT      0  14.22500000  14.81700000  13.55800000 H
H-H1      0  14.78000000  14.55000000  14.47700000 H
C-CT      0  15.10100000  14.43300000  12.35900000 H
H-HC      0  15.11100000  13.34600000  12.28500000 H
H-HC      0  14.64100000  14.82000000  11.45700000 H
C-CC      0  16.49800000  14.99900000  12.35800000 H
N-NB      0  17.24300000  15.30400000  11.20000000 H
C-CR      0  18.50300000  15.52300000  11.61900000 H
H-H5      0  19.31000000  15.90700000  10.98900000 H
N-NA      0  18.57400000  15.41900000  12.94400000 H
H-H-      0  19.40400000  15.52700000  13.50900000 H
C-CW      0  17.34000000  15.09800000  13.43000000 H
H-H4      0  17.15100000  14.74700000  14.43100000 H
C-C-      0  13.92800000  16.32700000  13.64000000 H
O-O-      0  12.75800000  16.71000000  13.75400000 H
N-N-      0  14.98100000  17.15400000  13.69600000 H
H-H-      0  15.91400000  16.76000000  13.57700000 H
C-CT      0  14.84900000  18.61200000  13.82000000 H
H-H1      0  13.80000000  18.95900000  13.87700000 H
H-H1      0  15.36600000  18.96700000  14.72000000 H
H-H1      0  15.30700000  19.10900000  12.96000000 H
O-OW      0  17.96200000  20.70200000  16.86200000 L
```

H-HW	0	16.98600000	20.73300000	16.67400000	L
H-HW	0	18.28900000	21.05400000	16.00400000	L
O-OW	0	19.92600000	19.21100000	17.96600000	L
H-HW	0	20.21000000	19.96300000	18.53000000	L
H-HW	0	19.10300000	19.61300000	17.58200000	L
O-OW	0	17.88500000	23.13500000	18.31200000	L
H-HW	0	17.88200000	23.86000000	17.64100000	L
H-HW	0	17.80400000	22.36000000	17.71700000	L
O-OW	0	23.59500000	16.55900000	16.56200000	L
H-HW	0	22.87700000	16.50400000	17.22400000	L
H-HW	0	23.90300000	15.62800000	16.54800000	L
O-OW	0	16.58100000	18.66000000	20.07900000	L
H-HW	0	16.12700000	18.01300000	20.67100000	L
H-HW	0	16.79600000	18.09200000	19.32400000	L
O-OW	0	15.60900000	23.30900000	14.61600000	L
H-HW	0	16.57000000	23.37500000	14.74100000	L
H-HW	0	15.42300000	24.09200000	14.07100000	L
O-OW	0	19.23000000	14.87600000	19.17300000	L
H-HW	0	19.26700000	14.80500000	20.15900000	L
H-HW	0	18.28000000	14.64000000	19.04400000	L
O-OW	0	20.67600000	19.71000000	15.25200000	L
H-HW	0	21.62900000	19.91600000	15.34500000	L
H-HW	0	20.47200000	19.40000000	16.15500000	L
O-OW	0	18.58200000	22.47900000	14.67700000	L
H-HW	0	19.37500000	22.04200000	14.27200000	L
H-HW	0	18.93800000	23.39900000	14.73800000	L
O-OW	0	20.59100000	21.55900000	13.10700000	L
H-HW	0	20.64700000	20.63000000	13.39800000	L
H-HW	0	20.47100000	21.44600000	12.14000000	L
O-OW	0	22.96400000	17.38500000	14.03000000	L
H-HW	0	22.73900000	16.48600000	13.74600000	L
H-HW	0	23.17000000	17.23000000	14.98000000	L
O-OW	0	15.43800000	21.18600000	16.24000000	L
H-HW	0	14.64000000	21.34400000	16.78600000	L
H-HW	0	15.37300000	21.94600000	15.61500000	L
O-OW	0	21.36800000	16.93800000	18.41200000	L
H-HW	0	20.66900000	16.27700000	18.60100000	L
H-HW	0	20.79900000	17.71200000	18.17500000	L
O-OW	0	16.89600000	18.73900000	6.33900000	L
H-HW	0	16.64700000	17.83600000	6.61600000	L
H-HW	0	16.17000000	19.25600000	6.76200000	L
O-OW	0	20.04500000	16.54800000	8.03000000	L
H-HW	0	20.02300000	17.26100000	7.33800000	L
H-HW	0	19.16900000	16.14900000	7.87200000	L
O-OW	0	19.61000000	18.51800000	6.26600000	L
H-HW	0	19.86600000	19.46300000	6.17600000	L
H-HW	0	18.63000000	18.61000000	6.23500000	L
O-OW	0	18.30500000	19.41500000	9.61100000	L
H-HW	0	17.65300000	20.14200000	9.76000000	L
H-HW	0	17.72100000	18.71800000	9.27700000	L
O-OW	0	24.33300000	18.93200000	12.39600000	L
H-HW	0	23.73900000	18.40800000	13.00700000	L
H-HW	0	24.89900000	19.35100000	13.09000000	L
O-OW	0	17.16700000	23.82700000	8.70200000	L
H-HW	0	16.68800000	24.33000000	9.39800000	L
H-HW	0	17.04200000	22.91300000	9.05000000	L
O-OW	0	16.33500000	25.03400000	11.13200000	L
H-HW	0	16.83000000	24.29000000	11.55300000	L
H-HW	0	15.64500000	25.17800000	11.81700000	L
O-OW	0	17.79800000	22.94300000	12.04900000	L
H-HW	0	17.93400000	22.65400000	12.98100000	L
H-HW	0	18.73800000	23.05100000	11.77700000	L
O-OW	0	19.94100000	21.48700000	10.40600000	L

H-HW	0	19.47000000	20.74000000	9.98900000	L
H-HW	0	20.44400000	21.83600000	9.64400000	L
O-OW	0	20.84300000	18.38100000	9.95800000	L
H-HW	0	20.70000000	17.63800000	9.33100000	L
H-HW	0	19.93900000	18.77000000	9.95600000	L
O-OW	0	23.46500000	17.42400000	7.26300000	L
H-HW	0	23.10000000	17.51400000	6.34400000	L
H-HW	0	22.70300000	17.01800000	7.70700000	L
O-OW	0	22.96100000	20.10600000	10.35000000	L
H-HW	0	23.40900000	19.69500000	11.12600000	L
H-HW	0	22.18800000	19.49600000	10.26800000	L
O-OW	0	16.58700000	21.47000000	10.05400000	L
H-HW	0	15.70300000	21.22200000	10.43000000	L
H-HW	0	16.97300000	21.91200000	10.85000000	L
O-OW	0	17.07100000	15.60600000	4.54500000	L
H-HW	0	17.89500000	15.23400000	4.14100000	L
H-HW	0	17.40800000	15.85300000	5.42200000	L
O-OW	0	16.94800000	15.94500000	7.66300000	L
H-HW	0	16.95500000	15.78700000	8.62800000	L
H-HW	0	16.21100000	15.33200000	7.41000000	L
O-OW	0	24.73700000	16.68900000	10.88900000	L
H-HW	0	24.53700000	17.53600000	11.35400000	L
H-HW	0	25.62000000	16.91900000	10.51500000	L
O-OW	0	16.06500000	8.42700000	15.89000000	L
H-HW	0	16.39000000	8.92700000	15.11500000	L
H-HW	0	16.10800000	9.13700000	16.57700000	L
O-OW	0	16.56600000	9.53500000	20.23100000	L
H-HW	0	16.47400000	9.77900000	19.27900000	L
H-HW	0	17.51300000	9.32500000	20.27000000	L
O-OW	0	21.27000000	13.36500000	18.15700000	L
H-HW	0	21.06000000	12.46300000	18.46200000	L
H-HW	0	20.52200000	13.86800000	18.55000000	L
O-OW	0	22.31000000	11.37500000	15.27300000	L
H-HW	0	22.08300000	12.16900000	14.73900000	L
H-HW	0	21.40600000	11.05400000	15.48600000	L
O-OW	0	16.22300000	10.44500000	17.68200000	L
H-HW	0	15.32300000	10.84900000	17.64200000	L
H-HW	0	16.76400000	11.21400000	17.39600000	L
O-OW	0	21.90700000	13.50800000	13.64100000	L
H-HW	0	22.72500000	14.00100000	13.36000000	L
H-HW	0	21.80100000	12.94900000	12.82800000	L
O-OW	0	18.33500000	9.30900000	13.92200000	L
H-HW	0	18.87500000	9.86800000	14.52900000	L
H-HW	0	18.84900000	8.47000000	13.97200000	L
O-OW	0	16.70600000	13.77700000	19.08800000	L
H-HW	0	16.48200000	13.28300000	19.91100000	L
H-HW	0	15.79200000	13.93900000	18.75000000	L
O-OW	0	17.95800000	12.54400000	16.97400000	L
H-HW	0	18.50800000	13.22400000	16.53300000	L
H-HW	0	17.60800000	13.06700000	17.73000000	L
O-OW	0	20.15800000	14.19300000	15.66100000	L
H-HW	0	20.77700000	13.98400000	14.92600000	L
H-HW	0	20.73500000	14.00900000	16.42900000	L
O-OW	0	19.69600000	10.67700000	15.82700000	L
H-HW	0	19.10700000	11.29000000	16.31400000	L
H-HW	0	19.77600000	9.94900000	16.48100000	L
O-OW	0	21.90000000	11.96500000	11.40400000	L
H-HW	0	21.82300000	12.51400000	10.58900000	L
H-HW	0	21.55500000	11.11000000	11.06200000	L
O-OW	0	17.63900000	8.08500000	9.51100000	L
H-HW	0	17.12500000	7.52200000	10.13400000	L
H-HW	0	17.89600000	8.81500000	10.12200000	L
O-OW	0	21.74200000	13.49000000	9.12000000	L

H-HW	0	21.09900000	14.11000000	8.73900000	L
H-HW	0	21.89200000	12.89200000	8.34800000	L
O-OW	0	24.36600000	14.42800000	9.35100000	L
H-HW	0	24.33600000	15.28200000	9.83000000	L
H-HW	0	23.42500000	14.16600000	9.38500000	L
O-OW	0	15.99600000	13.10200000	4.98500000	L
H-HW	0	15.66400000	13.27800000	5.89100000	L
H-HW	0	16.34700000	13.99200000	4.75800000	L
O-OW	0	18.47800000	9.94800000	11.29500000	L
H-HW	0	18.37900000	9.73100000	12.25400000	L
H-HW	0	18.10400000	10.84400000	11.27800000	L
O-OW	0	24.06400000	14.76300000	12.69200000	L
H-HW	0	24.89900000	14.26900000	12.67300000	L
H-HW	0	24.26100000	15.47500000	12.04100000	L
O-OW	0	9.81600000	21.51300000	15.35600000	L
H-HW	0	10.58700000	22.08200000	15.11900000	L
H-HW	0	10.16100000	20.64000000	15.08200000	L
O-OW	0	11.78600000	23.22500000	14.53800000	L
H-HW	0	11.78900000	23.27000000	13.56600000	L
H-HW	0	12.71300000	23.40900000	14.75000000	L
O-OW	0	11.82800000	19.51300000	17.08000000	L
H-HW	0	11.44300000	19.21000000	16.23200000	L
H-HW	0	11.37600000	18.91900000	17.71100000	L
O-OW	0	9.38000000	16.47000000	15.24100000	L
H-HW	0	9.74800000	15.97800000	15.99000000	L
H-HW	0	8.62000000	16.92300000	15.68400000	L
O-OW	0	10.81300000	18.73900000	14.66000000	L
H-HW	0	11.53800000	18.25600000	14.20800000	L
H-HW	0	10.16500000	18.00300000	14.75600000	L
O-OW	0	13.75400000	24.12900000	18.20000000	L
H-HW	0	13.54700000	23.16400000	18.09800000	L
H-HW	0	12.83300000	24.47500000	18.27300000	L
O-OW	0	13.33500000	21.49300000	17.97600000	L
H-HW	0	13.65300000	20.95400000	18.74100000	L
H-HW	0	12.71800000	20.82000000	17.57600000	L
O-OW	0	14.25300000	20.17200000	20.17500000	L
H-HW	0	14.16500000	19.73200000	21.05200000	L
H-HW	0	15.14200000	19.82500000	19.92000000	L
O-OW	0	15.06100000	20.35000000	7.55400000	L
H-HW	0	14.43100000	21.06200000	7.26700000	L
H-HW	0	15.40300000	20.75900000	8.37300000	L
O-OW	0	14.05600000	20.83100000	10.60200000	L
H-HW	0	13.66400000	21.68600000	10.28400000	L
H-HW	0	13.25800000	20.26800000	10.55500000	L
O-OW	0	11.41100000	20.29700000	9.43400000	L
H-HW	0	10.55900000	20.22300000	9.91700000	L
H-HW	0	11.14300000	20.89800000	8.70600000	L
O-OW	0	11.90600000	14.94400000	10.68800000	L
H-HW	0	12.48500000	14.19900000	10.40700000	L
H-HW	0	11.92000000	15.46500000	9.84500000	L
O-OW	0	12.96200000	18.58400000	7.98000000	L
H-HW	0	12.40300000	19.18400000	8.52500000	L
H-HW	0	13.74000000	19.16000000	7.83500000	L
O-OW	0	11.04700000	23.30200000	11.47000000	L
H-HW	0	11.74800000	23.02500000	10.83900000	L
H-HW	0	10.71900000	24.10600000	11.00600000	L
O-OW	0	8.99200000	19.91900000	10.86300000	L
H-HW	0	8.41100000	19.23700000	10.46100000	L
H-HW	0	9.30000000	19.43700000	11.65200000	L
O-OW	0	13.05200000	23.04200000	9.50100000	L
H-HW	0	13.11400000	22.81700000	8.54700000	L
H-HW	0	13.21000000	24.02100000	9.44300000	L
O-OW	0	12.22200000	16.03200000	8.16500000	L

H-HW	0	12.65900000	16.90100000	7.99400000	L
H-HW	0	12.21700000	15.66900000	7.25000000	L
O-OW	0	8.25300000	17.22700000	10.42000000	L
H-HW	0	8.87300000	17.22600000	9.65600000	L
H-HW	0	8.78000000	16.68700000	11.06000000	L
O-OW	0	9.82100000	17.52900000	8.23200000	L
H-HW	0	9.57000000	16.70000000	7.73400000	L
H-HW	0	10.78900000	17.36100000	8.29600000	L
O-OW	0	9.64800000	15.89400000	12.28600000	L
H-HW	0	9.78300000	16.01500000	13.24000000	L
H-HW	0	10.53900000	15.64100000	11.98500000	L
O-OW	0	8.03900000	8.13500000	18.13300000	L
H-HW	0	7.61000000	8.48800000	17.31600000	L
H-HW	0	7.88100000	8.89600000	18.73400000	L
O-OW	0	11.68400000	9.62000000	18.17600000	L
H-HW	0	11.53600000	8.67200000	18.41700000	L
H-HW	0	11.22800000	10.05300000	18.92000000	L
O-OW	0	9.22900000	6.25100000	15.20800000	L
H-HW	0	9.69500000	5.42300000	15.39700000	L
H-HW	0	9.96100000	6.90200000	15.34300000	L
O-OW	0	7.20000000	10.63300000	19.28200000	L
H-HW	0	7.95700000	11.19500000	19.00300000	L
H-HW	0	6.84900000	11.21400000	20.00600000	L
O-OW	0	9.11400000	12.88200000	19.62500000	L
H-HW	0	9.71600000	12.26300000	20.10200000	L
H-HW	0	9.45100000	13.73700000	19.96300000	L
O-OW	0	6.46300000	9.20100000	16.12000000	L
H-HW	0	6.32300000	10.06700000	15.68200000	L
H-HW	0	5.63800000	9.12600000	16.63600000	L
O-OW	0	13.67700000	11.33600000	17.36100000	L
H-HW	0	12.95600000	10.70900000	17.60600000	L
H-HW	0	13.49900000	11.43500000	16.40100000	L
O-OW	0	11.32900000	7.88500000	15.70700000	L
H-HW	0	11.29100000	8.52800000	16.43700000	L
H-HW	0	12.20800000	7.48000000	15.86800000	L
O-OW	0	6.02500000	11.57000000	14.63900000	L
H-HW	0	6.38400000	11.17900000	13.81600000	L
H-HW	0	5.13400000	11.83800000	14.32700000	L
O-OW	0	7.23400000	7.55000000	13.97700000	L
H-HW	0	6.89100000	8.07600000	14.73500000	L
H-HW	0	7.95400000	7.05300000	14.44600000	L
O-OW	0	6.02300000	13.82000000	16.34300000	L
H-HW	0	6.88000000	14.11700000	16.00000000	L
H-HW	0	5.93600000	12.95400000	15.88300000	L
O-OW	0	13.84200000	6.81500000	15.97600000	L
H-HW	0	14.64600000	7.36900000	16.11300000	L
H-HW	0	13.88500000	6.26300000	16.79000000	L
O-OW	0	14.49000000	7.05700000	13.30100000	L
H-HW	0	13.97700000	7.77000000	12.88800000	L
H-HW	0	14.15400000	7.09900000	14.22400000	L
O-OW	0	12.37400000	14.07800000	20.09500000	L
H-HW	0	13.05600000	13.84600000	19.43100000	L
H-HW	0	12.49900000	13.34000000	20.73300000	L
O-OW	0	14.17800000	13.86000000	18.05400000	L
H-HW	0	14.00100000	12.92000000	17.79900000	L
H-HW	0	13.66300000	14.32600000	17.37900000	L
O-OW	0	11.21600000	12.72900000	7.66200000	L
H-HW	0	10.71900000	13.44900000	7.23900000	L
H-HW	0	10.49200000	12.16600000	8.00300000	L
O-OW	0	13.94000000	10.09300000	9.41500000	L
H-HW	0	14.68300000	9.95100000	8.77600000	L
H-HW	0	13.62100000	9.17700000	9.49900000	L
O-OW	0	7.72400000	13.37500000	8.29200000	L

H-HW	0	8.05400000	13.06500000	9.14800000	L
H-HW	0	6.91200000	13.86600000	8.58300000	L
O-OW	0	7.24000000	10.04600000	9.87500000	L
H-HW	0	7.09800000	9.07100000	9.88800000	L
H-HW	0	6.50800000	10.32500000	10.47200000	L
O-OW	0	9.70100000	10.80700000	9.01900000	L
H-HW	0	9.97700000	10.77700000	9.95200000	L
H-HW	0	8.75100000	10.54000000	9.12000000	L
O-OW	0	5.47000000	10.60500000	11.91100000	L
H-HW	0	5.05100000	9.88200000	12.44900000	L
H-HW	0	4.65400000	11.10400000	11.67300000	L
O-OW	0	8.71400000	9.26000000	12.26400000	L
H-HW	0	7.99500000	9.58000000	11.69800000	L
H-HW	0	8.22700000	8.66700000	12.87400000	L
O-OW	0	9.79300000	6.85300000	10.25000000	L
H-HW	0	10.25900000	7.61700000	10.63000000	L
H-HW	0	8.86000000	7.10300000	10.43600000	L
O-OW	0	13.36100000	12.82400000	9.35300000	L
H-HW	0	12.59800000	12.74200000	8.73500000	L
H-HW	0	13.58400000	11.86800000	9.46300000	L
O-OW	0	11.39000000	9.01200000	7.74400000	L
H-HW	0	10.82300000	9.68900000	8.18200000	L
H-HW	0	11.86500000	9.61100000	7.10400000	L
O-OW	0	14.82600000	14.16500000	7.29200000	L
H-HW	0	13.99300000	14.39600000	6.83400000	L
H-HW	0	14.46500000	13.72600000	8.09600000	L
O-OW	0	5.37200000	14.69300000	12.55700000	L
H-HW	0	6.27100000	14.32100000	12.57200000	L
H-HW	0	5.53500000	15.50500000	12.01300000	L

Check the transition energy (see in Gaussian output Excited States energies and oscillators section)