Tutorial: MOL

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3 Oligoacetylenes: DMRG, DMRG-SCF, and AC0



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Connection to the GPU cluster in Prague

- Connect using **ssh** to gpu.jh-inst.cas.cz with -X option
- User: student1
- Password: TREX23@Heyr

```
ssh student1@gpu.jh-inst.cas.cz -X
```

• Connect to g24, g30, g32, or g33 node (dependeng on to which group you belong)

ssh g24 ... for group 1ssh g30 ... for group 2ssh g32 ... for group 3ssh g33 ... for group 4

 \bullet Connect to $\mathbf{g33}$

ssh g33 -X

• Go to the /data/student1/student_X directory (where X is your number)

cd /data/student1/student_X

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Structure of MOLMPS input

- MOLMPS is not yet public on Github, but should be soon.
- Structure of input:



• You will run it locally using

molmps_run dmrg_input ncores > dmrg_output

- Go to subdirectory naphthalene
- We will compare DMRG performance in canonical HF orbitals and split-localized orbitals



- Go to subdirectory canonical
- Take a look at FCIDUMP and dmrg.inp files
- Run DMRG: molmps_run dmrg.inp 4 > dmrg.out
- Analyze the results with mmps_ext dmrg.out and mmps_ext dmrg.out swp
- Do the same in the subdirectory split_localized, record the final energies
- Probe the effect of ordering optimization, record the energy

```
$chain
optimize_ordering 1
optimize_bond_dims [50 50 50]
end
```

Try moltools correls_5.moltools, imrove manually the ordering and re-run DMRG

• Try very cheap optimization based on the exchange integrals $\langle ij||ji
angle$

```
$chain
optimize_ordering 2
end
```

Take-home message

Split-localized basis is the best suitable for DMRG (exploits locality).

Ordering optimization based on the exchange integrals is in most cases good-enough.

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```
$chain
optimize_ordering 2
end
```

Take-home message

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- **②** Ordering optimization based on the exchange integrals is in most cases good-enough.

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- For pre-DMRG stuff, we will use Orca
- Orca counts orbitals from 0!
- We will compute singlet-triplet (S-T) gaps for the following three examples:



• Go to directory /data/student1/student_X/oligoacetylenes

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Oligoacetylenes (n = 4): DMRG, DMRG-SCF, AC0

- Go to subdirectory units_4
- According to instructions run HF calculation
- Select which orbitals to rotate, copy the rhf.gbw file to 2_rot/inputOrbs.gbw and rotate them
- Split-localize the active space (in 3_loc directory)
- Dump the FCIDUMP file (in 4_dump directory)
- Run the singlet and triplet state DMRG calculations (in 5_dmrg/singlet and 5_dmrg/triplet)
- Use dump.gbw as an initial guess for DMRG-SCF, i.e. copy it to inputOrbs.gbw
- Run the DMRG-SCF singlet and triplet state calculations
- In order to compute DMRG-SCF-AC0, copy the RDMs (G1.bin and G2.bin) and orbitals from the next-to-last iteration (rename them to inputOrbs.gbw) to 7_dmrgscf_ac0
- Compute also DMRG-AC0, try to sort out what files do you need.

- Go to subdirectory units_7
- Not to waste your time, MO integrals have already been dumped, check the orbitals with Molden
- Compute the singlet and triplet states by means of DMRG and DMRG-SCF, notice that this time DMRG-SCF employes the resolution of identity for integral transformation
- Compute DMRG-AC0 for the singlet state (group 1), DMRG-AC0 for the triplet state (group 2), DMRG-SCF-AC0 for the singlet state (group 3), and DMRG-SCF-AC0 for the triplet state (group 4), it will take a while.

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- Go to subdirectory units_9
- Not to waste your time, MO integrals have already been dumped, no need to check the orbitals with Molden
- Run the DMRG and possibly also DMRG-SCF calculations of the singlet and triplet states
- Analyze all the S-T gaps of oligoacetylenes obtained by different computational methods

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- Go to directory /data/student1/student_X/f2
- Check the DMRG input file f2.inp several DMRG blocks with increasing bond dimensions
- Run the DMRG calculation
- Use a linear function to extrapolate the energy w.r.t. the truncation error. You can find both values below the lines Final energy: in the output file

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