Usage of the code for NN-supported GRASP computations

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1 Preparation for computations

1.1 Get the code

The code can be cloned from the GitHub repo: https://github.com/pavlobilous/neural_grasp

The main file here is the file neural_grasp.py containing the class NeuralGraspManager responsible for NN-supported computations. The user is supposed to write his/her own Python scripts importing and using this class. There is an auxiliary script process_grasp_inp.py for creating a NumPy array for the CSF set to be sorted out (see below). The user is supposed to write also a Python script employing process_grasp_inp.py. In this demo example, we consider computations of Re ground state. The exemplary user files are called Re187_....

1.2 Rebuild GRASP2018

The code works with GRASP2018, but GRASP has to be rebuild after the file src/appl/rmcdhf90_mpi/csfwgt.f90 is replaced with the attached file csfwgt_rmcdhf_mpi.f90. With this, the weights of all CSFs will be written in the file csfwgt_rmcdhf_mpi.dat each time the rmcdhf_mpi program is run (this file will be placed in the folder tmp_mpi/000). The order of the CSFs in this file is the same as in the input rcsf file.

1.3 GRASP input

Attached are the isodata file and the radial wave function file rwfn.prim.inp. Place them in the folder with the future NN-supported computations. The rcsf files should be created in the following way.

First, create rcsf-files from the attached CSF-files gendata1-2 and gendata1 by feeding them into the rcsfgenerate program. The CSF set originating from gendata1-2 (let's call the corresponding file rcsf1-2.inp) is the "big" CSF set which we cannot computed directly (in this case we can, but only because this is a demo version). The CSF set originating from gendata1 (let's call the corresponding file rcsf1.inp) is the so called *primary subset* of the "big" set which will be always included in the GRASP runs and won't be exposed to the NN.

The "big" rcsf-file has to be now restructured such that the primary CSFs come first. Use the rcsfzerofirst program on the files rcsf1-2.inp and rcsf1.inp to achieve this. Rename the obtained file to rcsf.full and the primary set file called before rcsf1.inp to rcsf.head.

1.4 Creating NumPy file with the "big" set

Run the Python script Re187_gen_rcsf.py. The input files are rcsf.full and rcsf.head, the output is the file rcsf.npy needed for the NN-supported computations. Note that in this Python script there is additional input $2J_{\text{total}} = 5$, which should be changed correspondingly for computations other than the Re ground state.

2 NN-supported computations

2.1 Initial iteration on random CSFs

Since at the beginning the NN is untrained and thus useless, the first iteration is performed by picking randomly CSFs from the "big" set (or "pool") to be sorted out (of course, excluding the primary CSFs). Run the script Re187_init_on_random.py which will pick randomly 1% of CSFs from the pool and write them to an rcsf.inp file on top of the primary subset. At the same time, it creates a blanc NN already at this stage, which is saved in the folder save_init_random together with some NumPy arrays needed for further computations. Note that in this script the NN is not trained, since the CSF weights are not known yet.

As soon as the script is finished, GRASP should be run by the user on the formed rcsf.inp file. After that, copy the csfwgt_rmcdhf_mpi.dat file from the folder tmp_mpi/000 to the folder with the computations — this is required for running the next Python script. Store the obtained energy value.

Note that in this demo example, the Python code never runs GRASP by itself (though it can, but this is good only for small computations). It forms only rcsf.inp files, and the user is expected to run GRASP "manually" on it. The idea behind this splitting is that for bigger computations, the NN training can be performed on a GPU, whereas GRASP requires many CPU nodes. The user would then submit two separate jobs requesting different kinds of resources.

2.2 Performing the first NN-supported iteration

Now we do the first NN-supported iteration by running the script Re187_iter_on_cutlog.py. This script has the cutoff value $10^{-8.6}$ (see the variable cutlog) which determined which CSFs are *important* and which are *unimportant*. First, the NN and the internal NumPy arrays are read from the folder save_init_random. The NN trains on the CSFs randomly picked in the previous iteration, for which the weights (and thus the binary classification as important / unimportant) are already known. Then the trained NN is applied to the whole pool and the next rcsf.inp file is formed. The trained NN and the internal NumPy arrays is saved in the folder save_cutlog_-8.6 for the next iteration.

Now run GRASP, copy the csfwgt_rmcdhf_mpi.dat file from the folder tmp_mpi/000 to the folder with the computations, and store the obtained energy value.

2.3 Performing subsequent NN-supported iteration

In the same manner we can run further NN-supported iterations. For this, the script Re187_iter_on_cutlog.py needs to be changed at two places

• The input folder should be now not save_init_random but save_cutlog_-8.6 (or the last folder saved). Edit the line ngm.load_state("save_init_random",fit_params) correspondingly.

• The cutoff value has to be changed. Edit the line cutlog=-8.6 for the values -9.2, -9.8, -10.4, -11.0 for the subsequent iterations.

As usually, after this Python script is finished, run GRASP, copy the <code>csfwgt_rmcdhf_mpi.dat</code> file from the folder <code>tmp_mpi/000</code> to the folder with the computations, and store the obtained energy value. Make sure that the obtained energies converge to the "true" value on the full CSF set.