

Instantiation of QMMMInputs

Get_linkcorrlist?

Class MMParams Instatiation of MMParams readInput.py → Reads information from mm.dat generate_pcf_from_top.py read_mmparams Instatiation of QMParams Class QMParams prepare_pcf_for_shift.py → Reads information from qm.dat read_qmparams _helper.py Instatiation of PathParams Class PathParams generate_charge_shift.py read_pathparams → Reads information from path.dat sum_pcf_tm.py Instatiation of QMMMParams Class QMMMParams generate_top.py read_qmmmparams → Reads information from qmmm.dat Read molecules from topology readmols Read charges of all atoms in all molecules readcharges checkformol Read xyz for all atoms readg96 Read number of atoms from structure file read_numatoms Read connection of atoms from topology read_conn_list_from_top get_curr_top checkformol get_mollength_direct get_connlist Read qm atoms read_qmatom_list If inout: make inner/outer list read inner list read_outer_list If structure file is gro: write to g96 write_highprec stepper Make xyzq make_xyzq Get m1 and m2 atoms and shift charges prepare_pfc_for_shit_fieldsonly identify_m1 get_bondpartners identify_m2 get_bondpartners get_qmcoords eliminate_and_shift_to_m1 Get vector from qm atom to mm atom if bond is cut get_linkatoms_ang

get_linkcorrlist

```
Make new point charge field
                                                          generate_charge_shift_fieldsonly
Calculate distance vectors * charge for m/q atoms
                                                                 sum_pcf_tm_nofile
                                                                 get_m2vec_fieldsonly
                                                                 create_corr_charges
                                                                        write_disp_charges
                                                                               uvec
                                                                 make_new_field
                                                                 sum_pcf_tm_nofile
While loop until maximum displacement below threshold
                                                                 create_corr_charges
                                                                        write_disp_charges
                                                                        uvec
                                                                 make_new_field
                                                                 sum_pcf_tm_nofile
                                                                 write_new_field_to_disk_listsonly
Write new topology
                                                                 generate_top_listsonly
                                                                        read_qmatom_list
                                                                        <u>readmols</u>
                                                                        getincludelist
                                                                        get_molfindlist
                                                                               molfinder
                                                                        make_exclude_index
                                                                        make_large_top
                                                                               get_mollength
                                                                                      molfinder
                                                                               find_ffnonbonded
                                                                               clean_exclusions
                                                                               clean_exclusions
                                                                 read qmatom list
```