

Workflow

Identify Possible
Substrates

Reaxys®

Acids: 8,676

Export SMILES to Excel



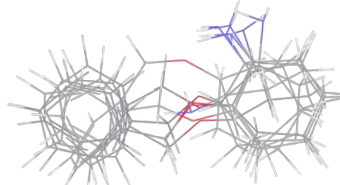
SMILES to .sdf Files

SMILES_to_sdf.py

Open Babel



MacroModel
Conformational
Search



prep_MM_npsh.bash
mm_reference.com



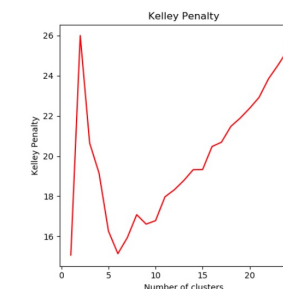
MacroModel
Clustering +
.maegz to .sdf Files

cluster based on
Kelley penalty value
if >20 confs.

MM_clustering_and_
converting.bash



Confirm clustering



check_manual_
clustering.bash



Generate job files

one job per
conformer

Group script: Convert
Gaussian



Gaussian
Geometry
Optimization +
Single Point

Opt: B3LYP/6-31G(d,p)
SP: M062X/def2tzvp



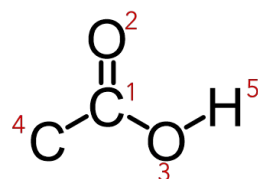
Check .log Files

normal job
termination
+ no imaginary
frequencies

post_processing.bash
log_check_for_
processing.py



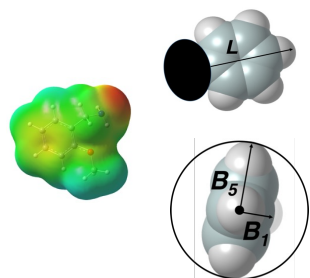
Atom Number
Collection



get_properties_
notebook.ipynb



Get Properties



get_properties_
notebook.ipynb

Automated Get_Properties_Notebook

Implemented Descriptors

- energies (goodvibes)
- nbo
- nmr
- angle
- dihedral angle
- distance
- plane angle
- total time
- frontier molecular orbitals
- Volume
- polarizability
- dipole
- Sterimol (morfeus)
- buried Sterimol (morfeus)
- buried volume (morfeus)
- buried volume scan (morfeus)
- pyramidalization (morfeus)
- SASA & sphericity (morfeus)
- Sterimol (dbstep)
- Sterimol2Vec (dbstep)
- Hirshfeld charges
- ChelpG

Partially Implemented / Not Generalizable

- IR stretching frequency - *works for one stretch in the input range*
- Ask Melissa or Brittany for a sample if you want to modify these to your needs:
 - Buried Volume: hemispheres, quadrants, and octants (morfeus)
 - Nborbs - *can be customized depending the types of orbital occupancy/energies you want out*

Not Implemented

- e, thermos, efg, tz_e - *use goodvibes instead*
- imaginary frequencies - *should be done in log check*
- qpole
- method
- route lines
- nmr tensors
- dorbs
- buried volume (dbstep) - *takes significantly longer than morfeus*
- Vol2Vec (dbstep) - *takes significantly longer than morfeus Vbur scan*
- cavity (*volume & surface area from solvation model*)