

CHL Lab

## 11/01/17: Building Gromacs and SoleFP on Stampede2

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### Gromacs:

Added /opt/apps/intel17/impi17\_0/gromacs/5.1.2/share/gromacs/bin to \$PATH  
Exported GMXLIB=\$HOME/gmx/forcefields

To run gmx features: type gmx

To run mdrun with parallelization: type ibrun mdrun\_mpi

### SoleFP:

1. Added module load intel to ~/.bashrc

2. Installed python2.7.12 in \$HOME/programs/python2.7.  
./configure --prefix=\$HOME/programs/python2.7  
make  
make test  
make install

Added \$HOME/programs/python2.7/bin to \$PATH (to overwrite the existing python, add to before the existing paths!)

3. Installed pp:  
python setup.py install

4. Installed numpy linked with mkl libraries following globulion's instructions.

1. Note1: Add lapack\_libs = to site.cfg, because the lapack libraries are deprecated in the newer version of intel. So your site.cfg should look like:

```
[mkl]
library_dirs = /opt/intel/mkl/lib/intel64
include_dirs = /opt/intel/mkl/include
mkl_libs = mkl_rt
lapack_libs =
```

2. Note2: No need for the --prefix=\$HOME option in install; leave it out

3. Note3: The MKL library paths ( LD\_LIBRARY\_PATH ) are already set in stampede2

5. Installed scipy linked with mkl libraries following globulion's instructions.

1. Note1: No need for the --prefix=\$HOME option in install; leave it out

6. Installed matplotlib:  
python setup.py install

7. MDAnalysis:  
python setup.py install

8. Installed scitools:  
python setup.py install

9. Installed PyQuante:  
1. Added basis for Ca in PyQuante/Basis/p6311pp\_1d\_1p.py  
2. python setup.py install

10. Installed libbbg  
1. Modified the install script:  
1. Change prefix to \$HOME/programs/python2.7  
2. Change export LIBBBG\_LIB=\$PREFIX/lib/python2.7/dist-packages/ to  
export LIBBBG\_LIB=\$PREFIX/lib/python2.7/site-packages/  
2. Modified utilities.py:  
1. Change import\_matplotlib =  
int(os.environ['LIBBBG\_IMPORT\_MATPLOTLIB']) to  
import\_matplotlib=1  
3. bash install

11. Installed coulomb  
python setup.py install

12. Installed solvshift  
1. Added fragments for ca++ ad indole to slv-master/frg  
2. Modified solvshift/slvpar.py:  
1. Added 'ca++', 'indole' to solvshift.slvpar.params  
2. Added 20:'Ca' to solvshift.slvpar.\_\_atomic\_symbols  
3. Modified solvshift/setup.py:  
1. Added 'ca++', 'indole' to molecules  
4. Modified the install\_slv script:  
1. Change prefix to \$HOME/programs/python2.7  
2. Change export LIBBBG\_LIB=\$PREFIX/lib/python2.7/dist-packages/ to  
export LIBBBG\_LIB=\$PREFIX/lib/python2.7/site-packages/  
5. bash install\_slv  
6. Exported SLV\_DATA=\$HOME/programs/python2.7/lib/python2.7/site-packages/solvshift-dat  
7. Copied util as \$HOME/programs/slv\_util  
8. Added \$HOME/programs/slv\_util to path

9. uploaded solefp.files (with modified gmx.tc) to \$HOME/solefp

Success!