

Calculate $g(r)$

You can calculate the second order radial distribution function, $g(r)$, using `gr.c`. Compile `gr.c` by invoking your c compiler on `gr.c`. To do this:

1. Open your terminal app
 2. Navigate to the directory that `gr.c` is in
 3. Figure out the name of your c compiler program (google this)
 4. Typically it will be called either `gcc` or `cc`, if it goes by a different name then replace `gcc` with that name in the following
 5. Run: `gcc -O3 gr.c -o gr`
 1. `-O3` (note that it is the capital letter O) turns on the highest levels of optimization by the compiler
 2. `-o gr` specifies that we want the binary to be named `gr`
 1. replace `gr` with whatever you want the binary to be named
 6. Then, you can run the $g(r)$ program by typing `./gr` (arguments)
- I recommend adding the directory `gr` is in to your path (just google this)

The arguments to `gr.c` are as follows (all are required)

input file -> a file in `.tr` format

output file name -> the name of the file that you want to store the (x,y) pairs of the $g(r)$ plot in

diameter -> the diameter of the smaller particle, or whatever particle size you want to correspond to $x = 1$ on the $g(r)$ plot

So, for example you could do:

`gr dump.tr outplot.dat 2.56` to run `gr` on the track file in `dump.tr` and store the results in `outplot.dat` with $x = 1$ corresponding to a diameter of 2.56

Note it will overwrite whatever was stored in `outplot.dat`

Convert `.xyz` to `.tr`

You can convert `lammps .xyz` files to track files using the `read_tr.py` script. To run it, make sure that you have python installed (it's tested with 2.7 but should work with others) and also have `numpy` installed. Then, run:

`python read_tr.py (name of .xyz file) (name of track file you want to create)` and it will save the track file in the specified location

Note that it will overwrite anything stored in (name of track file you want to create)

Calculate $s(k)$

You can calculate the 2D $s(k)$ and it's related functions using `sk.py`. To do so, open the python REPL, then do the following:

```
>>> import sk
```

```
>>> sof_k = sk.run(track file to run it on)
```

```
>>> sk_avg = sk.runskAverage(sof_k)
```

Calling `sk.plotSKAverage(sk_avg)` or `sk.plotSK(sof_k)` will produce the plots of the data in `sof_k` and `sk_avg`

Misc. useful scripts:

Get all tiffs from the cluster

Run: `getTiffs (file name)` where file name is the directory you want to put the tiffs in

Find differences in files in a more readable manner

Run: `diffD (file 1) (file 2)`

Plot from the command line (requires gnuplot)

Run: `plot (file to plot) (optional arguments)`

If no arguments are provided it will plot to the screen with lines connecting points

If it's called as `plot (file to plot) -nl` then it will plot without lines connection points

If it's called as `plot (file to plot) -l -o (output file)` then it will save the plot to the output file with lines connecting points

Todo:

This can certainly be made better through Latexing it or just using better formatting

Add $s(k)$ theta dependence part (it wasn't up when I made this)

Have someone who hasn't used the programs read over this to see if it actually made sense