**Phase Transitions and the Liquid State**

Physical Chemistry

Fall 2017

* 1. 1 Introduction

Much — if not most — chemistry of industrial and biological significance takes place in solution, rather than in the gas phase. In this lab, we investigate condensation of gas to liquid and the general properties of the liquid state. Our simulation will compress argon gas at 100 K under 10 bar of pressure, slowly cooling it to 85 K. This causes a transition from gas to liquid. Along the way, we will make passing observations of the general properties of phase transitions, and we will finally calculate the heat of vaporization of argon.

* 1. 2 Procedure

Follow the instructions of your professor to start a terminal and copy the files you need into an appropriate place. Then execute the following commands. This sets up and runs the simulation,[[1]](#footnote-1) which should take up to 5 minutes to complete.

gmx grompp -f equil.mdp -c initial.gro -p argon.top -o equil.tpr  
gmx mdrun -nt 1 -nb cpu -s equil.tpr -c equil.gro -e equil.edr \   
 -g equil.log -x equil.xtc -cpo equil.cpt  
gmx grompp -f md.mdp -c equil.gro -t equil.cpt -p argon.top -o md.tpr   
gmx mdrun -nt 1 -nb cpu -s md.tpr -c md.gro -e md.edr -g md.log -x md.xtc

If you are typing these commands each on one line, omit the backslash in the second command. The first two commands bring our system of 100 argon atoms to a temperature of 140 K, and the second two commands perform the simulated compression.

View the evolution of this simulation in vmd. Start vmd, then in the “VMD Main” window, select “File”, then “New Molecule”. Click the “Browse” button, navigate to the directory you performed the simulation in, then choose 100.gro. After returning to the “Molecule File Browser” window, click “Load”. This loads the first frame (initial positions) of the simulation. Click “Browse” again, then choose md.xtc. When in the “Molecule File Browser” window, choose “Load all at once”, then click “Load”. This loads the remainder of the simulation. Close the “Molecule File Browser” window.

To see the motion of the atoms clearly, the display representation must be changed. In the “VMD Main” window, click on the “Graphics” menu, then “Representations”. Change the “Drawing Method” drop-down to “VDW”.

In the “VMD Main” window, rewind the movie, set the “Step” counter to 10 and the representation to “VDW”, then play the trajectory movie. In your notebook, carefully describe what you see, noting the simulation time that particularly striking changes occur.[[2]](#footnote-2)

Describe the qualitative behavior of the liquid state. Using the last quarter or so of the movie, zoom in so that you can see the entire box. Select a single atom and change its color.[[3]](#footnote-3) Describe the motion of the atom as time progresses. Is it smooth? Does it make steady progress through the bulk liquid, or does it make halting progress?

Quit vmd.

Prepare data for plots of temperature, volume, density, and energy by typing the following in the terminal:

echo 7 | gmx energy -f md.edr -s md.tpr -o temperature.xvg -skip 10  
echo 13 | gmx energy -f md.edr -s md.tpr -o volume.xvg -skip 10   
echo 14 | gmx energy -f md.edr -s md.tpr -o density.xvg -skip 10  
echo 4 | gmx energy -f md.edr -s md.tpr -o pot\_energy.xvg -skip 10   
echo 5 | gmx energy -f md.edr -s md.tpr -o kin\_energy.xvg -skip 10  
echo 6 | gmx energy -f md.edr -s md.tpr -o tot\_energy.xvg -skip 10

This produces the files temperature.xvg, volume.xvg, density.xvg, etc. in the current directory.[[4]](#footnote-4)

Prepare three plots in your plotting tool of choice. Using the xvg files you just produced, plot each of volume, density, kinetic energy, potential energy, and total energy (in separate plots), along with temperature on a second vertical axis for each. From these plots, identify (by simulation time) the following four periods of the simulation: (1) compression, (2) cooling, (3) condensation, and (4) liquid. Describe what happens to volume, density, kinetic energy, potential energy, and total energy in each of these four regions. If you observe any unusual features in your graphs other than the phase transition, view the corresponding portion of the simulation trajectory in vmd to help you explain them. Explain how the first law is preserved despite the total system energy changing.

Calculate the density of argon (1) after compression but before condensation, and (2) after condensation using the van der Waals equation and compare to the density at the corresponding points in your simulation. How well do the van der Waals values and the simulated values agree? If agreement is poor, use a more advanced equation of state[[5]](#footnote-5) to calculate the density. Comment on which (if either) equation of state is appropriate for modeling argon under each of the two conditions.

Identify the phase transition temperature (or temperature range).

Use your plots to determine the molar energy ( ) and enthalpy ( ) of vaporization of argon. If there is any ambiguity about how to determine either value, devise a way to turn this ambiguity into error bars. Find an experimental value for for the vaporization of argon. How well does your value compare? What are possible explanations for any discrepancies between your measured value and the experimental value of ?

* 1. 3 Report

Write a narrative report containing the observations you made and answers[[6]](#footnote-6) to the questions posed in the above procedure. Include a brief discussion (probably at the beginning) of why cooling argon under pressure results in condensation. Discuss the phase change you observed, emphasizing atomic interactions and how they give rise to changes in volume, density, and energy (all three kinds considered) during each of the four main portions of the simulation (compression, cooling, phase transition, and post-transition). Explain in detail how you calculated . Present the phase transition temperature and you calculated for argon and the corresponding experimental values and discuss (i.e. attempt to explain) any discrepancy between your results and the experimental results. Include all graphs produced and scans of your notebook pages. Submit your report electronically.

1. a 5 ns simulation in steps of 1 fs, with atomic positions (the “movie”) recorded every 0.1 ps [↑](#footnote-ref-1)
2. The simulation time can be obtained from the frame number, which is in the box between the “rewind” button and the trajectory time slider. This box only displays 4 digits, but you need 5; you can click and drag in the frame number box either to read the number better or to select and copy the number out. You can then use this frame number and the time between frames to calculate the time at which changes occur. [↑](#footnote-ref-2)
3. For example, go to the “Representations” dialog as before, select “Create Rep”, then use index 0 for the atom selection and “ColorID” for the coloring method. [↑](#footnote-ref-3)
4. The -skip 10 parameter requests that data be saved every 10 frames (1 ps), which ensures that Excel is capable of handling these files. If you intend to use Python or another similarly efficient system to analyze your data, you can (and probably should) omit the -skip 10 flag so that you have full-resolution data available for analysis. [↑](#footnote-ref-4)
5. such as the Redlich-Kwong equation [↑](#footnote-ref-5)
6. in prose [↑](#footnote-ref-6)