

UNIVERSITY OF YORK
DEPARTMENT OF PHYSICS
Third Year Examinations 2016
MOLECULAR SIMULATION
THEORETICAL PHYSICS STUDENTS ONLY

Issued at: 12:00 25 February 2016

Submission deadline: 16:00 Thursday 28 April 2016

Feedback by: Thursday 26 May 2016

**Please identify yourself by your examination number and not your name
on each sheet of paper you use.**

**See the Undergraduate Handbook for the regulations regarding academic
misconduct and plagiarism.**

Molecular Simulation - PHY000016H

- 1 Using the molecular dynamics program with which you have been provided, generate trajectories for the Lennard-Jones fluid with density $\rho^* = 0.7$, using $N = 256$ particles over a range of initial temperatures, T^* , from 1.0 to 2.5. A time-step $\Delta t^* \simeq 0.004$ should be satisfactory for these simulations. (All quantities are quoted in MD units). Hence:
 - (a) Using the method of block averages, estimate the errors in the simulated values of internal configuration energy $\langle U^* \rangle$ and temperature $\langle T^* \rangle$. Ensure that you have used sufficient steps in your trajectories to give one standard deviation from the mean of U^* which is less than 0.002.
 - (b) Plot $\langle U^* \rangle$ v. $\langle T^* \rangle$ and obtain the best fit cubic polynomial relating these quantities. Hence find an expression for the constant volume thermal capacity, C_v^* as a function of temperature.
 - (c) For each trajectory find C_v^* from the fluctuation formula:

$$C_v^* = \frac{3}{2} \left[1 - \frac{2}{3NT^{*2}} \langle (\delta E_k^*)^2 \rangle \right]^{-1}$$

Where total kinetic energy $E_K^* = \frac{3}{2}NT^*$ and

$$\langle (\delta E_k^*)^2 \rangle = \langle (E_k^*)^2 \rangle - \langle E_k^* \rangle^2.$$

Estimate the error in each of these measurements.

- (d) Plot your estimates of C_v^* from the fluctuation formula and the polynomial expression on a graph.

Present your work in the form of a Laboratory Formal Report. Your report should be no more than 10 sides of A4, using 12 point font, including figures, tables and references. In addition to this you should include a listing of your own code (not the molecular dynamics code unless modified by you) in an appendix.

End of paper