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# The free energy of the metastable supersaturated vapor via restricted ensemble simulations. II. Effects of constraints and comparison with molecular dynamics simulations

Chu Nie, <sup>1,a)</sup> Jun Geng, <sup>2</sup> and W. H. Marlow <sup>2</sup> <sup>1</sup>School of Physics and Information Engineering, Jianghan University, Wuhan 430056,

People's Republic of China

Department of Nuclear Engineering, Texas A&M University, College Station, Texas 77843, USA

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Extensive restricted canonical ensemble Monte Carlo simulations [D. S. Corti and P. Debenedetti, Chem. Eng. Sci. 49, 2717 (1994)] were performed. Pressure, excess chemical potential, and excess free energy with respect to ideal gas data were obtained at different densities of the supersaturated Lennard-Jones (LJ) vapor at reduced temperatures from 0.7 to 1.0. Among different constraints imposed on the system studied, the one with the local minimum of the excess free energy was taken to be the approximated equilibrium state of the metastable LJ vapor. Also, a comparison of our results with molecular dynamic simulations [A. Linhart et al., J. Chem. Phys. 122, 144506 (2005)] was made. © 2008 American Institute of Physics. [DOI: 10.1063/1.2937893]

#### I. INTRODUCTION

The thermal properties of the supersaturated vapor are interesting and important to know because of their scientific and engineering applications. Numerous research efforts contribute to this field with their special interests. <sup>1–19</sup> For example, some focus on the free energy barrier of the nucleus, critical size of the liquidlike drop, or homogeneous condensation of the supersaturated vapor, <sup>1-13</sup> some works discuss the failure of the integral equation with (Percus-Yevick) or hypernetted chain approximation when applied to metastable systems, 14,15 while little work has been reported dealing with the thermal properties of the supersaturated vapor itself. 16-22

Baidakov and Protesenko<sup>16</sup> pointed out that when dealing with a metastable system with Monte Carlo (MC) or molecular dynamics (MD) simulation methods, a long range cutoff of least 5 or 6 times  $\sigma$ , where  $\sigma$  is the hard sphere radius, is essential. Corti and Debenedetti<sup>18</sup> emphasized that the difference of equilibrium between the normal state and the metastable state lies in the number of configurations appearing in the two kinds of systems. For a normal state, the probability of the appearance of a configuration is governed by the Boltzmann factor and the higher the system energy is, the lower is its probability to appear. However, in the metastable system, there exists an energy barrier; once the system overcomes the barrier, a bifurction of the system elements occurs, i.e., it goes to phase separation. Therefore, the central task for a simulation in the metastable region is to prevent the system from going to phase separation. They divided the system into small three-dimensional cells and limit the number of particles in each cell to remain a certain number; thus, the phase separation is effectively prohibited. They also pointed out that the way to prevent phase separation is not

unique. Their method is something similar to global density constraint; however, some other technique might also work, for example, to limit the maximum density of the surroundings of a particle and so on.

Other researchers<sup>1,6–8</sup> imposed a different constraint on the system when studying the free energy barrier of homogeneous condensation. They defined clusters by identifying the mean-center distance between atoms and considered two atoms to be in the same cluster only if their mean-center distance is less than  $1.5\sigma$ . By limiting the maximum in an atomic cluster, the system could be maintained in its one phase state. This method is efficient when studying the free energy barrier but not suitable to study the thermal properties of supersaturated vapor. Linhart et al. 19 developed a MD simulation method to study the thermal properties of the supersaturated vapor up to spinodal density at different temperatures. They average the thermal properties before the system goes to phase separation. They used the instantaneous pressure, the numbers of atoms in the largest cluster, and the number of atoms not belonging to any cluster as the indication of phase separation. What they found is that the atomic cluster exists even near the spinodal density, as would be expected in the coexistence region. Hence, simply limiting the distance between two atoms is not sufficient to prevent phase separation because by doing so, too many configurations will be eliminated. However, two aspects of their method have shortcomings. First, their method requires varying the system size to maintain a longer induction period especially when density approaches spinodal. It is also very interesting to point out that no conclusion has been reached on the system size of metastable liquid simulation. For example, in Ref. 20, the authors found that the metastable phase can exist longer the smaller the box is. This is probably because small simulation systems suppress phase separation, in which large density fluctuations leading to phase separation are difficult to create. Thus, small simulation sys-

a) Electronic mail: niechu@hotmail.com.

tems allow reaching spinodal more closely. However, in our work, in order to eliminate the effects of periodic boundary conditions, we choose the system size relatively large. Second, their method provides no quantitative criteria for identifying the phase separation. Because the instantaneous pressure, maximum number in an atomic cluster, and the number of atoms not belonging to any cluster all vary continuously in the coexistence region and they do not evolve like step functions, one has to judge phase separation based on their experience and different persons may have different judgment. Therefore, uncertainty will be introduced in this stage.

Recently, we reported the pressure, excess chemical potential, and excess free energy, with respect to ideal gas data at different densities of the supersaturated LJ vapor at the reduced temperature of 0.7 with restricted canonical ensemble MC simulation method. At each density studied, we imposed different constraints on the system and found that those densities below the spinodal density would exhibit a free energy local minimum, as discussed in Ref. 18, which is close to the equilibrium state of the metastable system. In that work, we only performed simulations at reduced temperature  $T^*=0.7$ , where  $T^*=kT/\varepsilon$ , and in this work we extend the simulation from  $T^*=0.7$  to 1.0. With the data obtained, we may estimate the thermal properties, as well as the spinodal densities at different temperatures.

#### **II. METHODS AND SIMULATIONS**

Here, we recapitulate the methodology we used in our previous work.<sup>21</sup> In this work, the vapor is modeled by simulations with Lennard-Jones (LJ) model interactions for atoms.

$$u(r) = 4\varepsilon \left[ \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^{6} \right]. \tag{2.1}$$

First, divide the system into subcells based on the number of atoms and density of the system studied. The length of the cubic subcell is taken with the following relations:

$$l = \frac{L}{N^{(1/3)}},\tag{2.2}$$

where L is the length of an edge of the simulation box and N is the total number of particles during simulation. Therefore, the ideal homogenous case should correspond to one atom per cell. However, density fluctuations exist and limiting the maximum number of atoms in each subcell to  $d_{\rm max}$  may help to maintain the one phase state for the system being studied. For a given density, a series of runs starts from  $d_{\rm max}$ , when no number limitation is imposed onto the subcell, to  $d_2$ , when only two atoms are allowed to remain in one subcell; otherwise, the trial movement of a selected atom will be rejected. For each run, the excess chemical potential and pressure are averaged and recorded. With the obtained excess chemical potential and pressure, the excess free energy can be computed with the following formula:

$$f^{\text{ex}} = \mu^{\text{ex}} - P^{\text{ex}}/\rho, \tag{2.3}$$

where  $f^{\text{ex}}$  is the excess free energy per particle,  $\mu^{\text{ex}}$  is the

excess chemical potential, and  $P^{\text{ex}}$  is the excess pressure of the system with respect to the ideal case. In this work, initially, all the atoms were placed on a simple cubic lattice. For all simulation, the system size was set at least to  $40\sigma$ . The number of steps to reach equilibrium is  $10 \times 10^6$  moves, and another  $200 \times 10^6$  moves were used to sample the phase space. The cutoff distance is set to be  $8.0\sigma$ , and beyond the cut off distance, standard long range corrections were employed. The simulations are carried out at  $T^*=0.7, 0.75, 0.8$ , 0.9, and 1.0, where  $T^* = kT/\varepsilon$ . For  $T^* = 0.7$ , 0.75, and 0.8, the number of particles involved in the simulation is 3375, and for  $T^*=0.9$  and 1.0, the numbers of particles involved in the simulation are 5832 and 6859, respectively. The excess chemical potential is measured by performing the brute force sampling of  $\exp(-\beta u)$ , where it is the energy increase brought by the insertion of a test atom. To see the original idea of the atom insertion method, the reader may go to Ref. 23 and our program is written based on the classical textbooks on molecular simulation.<sup>24,25</sup>

#### **III. RESULT AND DISCUSSION**

Table I gives simulation results for the excess free energy per particle, pressure, and constraints imposed on the supersaturated vapor system. Actually, the excess free energy listed in Table I is defined by

$$f' = \mu^{\text{ex}} - (P^{\text{id}} + P^{\text{ex}})/\rho.$$
 (3.1)

It does not eliminate the ideal partial contribution to the free energy per particle; however, the inclusion of the ideal part of pressure does not affect the comparison of the excess free energy for the same density. Together with Table I in Ref. 21, the current work gives a relatively complete table of all our simulation result from  $T^*=0.7$  to 1.0. We note that for most of the simulations with the same density, as the constraint number  $d_{\text{max}}$  changes, the free energy changes and there is a local minimum of the free energy. As we pointed in Ref. 21, for a certain constraint imposed on the system, sometimes the system may very closely approach the equilibrium configuration; hence the excess free energy per particle exhibits a local minimum. With the loosening of the constraint, the system gradually starts phase separation, until, finally, full phase separation occurs when the system has a larger excess free energy compared with that of gas phase. This is the reason why the excess free energy of the system finally decreases. The state with the local minimum of free energy should be very close to the equilibrium state and the thermal properties obtained under that constraint can be viewed as the approximate thermal properties of that state. Thus, most states in Table I can be easily determined and only some exceptions exist, for example,  $T^* = 0.75$  and  $\rho \sigma^3 = 0.035$ . At a first glance, the series of data has no local minimum, but this does not mean the failure of our method. Because we only imposed coarse constraints on the system studied based on the number of particles and system size, we perhaps missed the local minimum due to this coarse selection. Observing the data for 0.035, they are smooth and we may apply the polynomial interpolation to find the local minimum if we do not want to introduce the extra labor to locate the local mini-

TABLE I. Simulation results for the excess free energy [actually, in the table,  $f' = \mu^{\rm ex} - (P^{\rm td} + P^{\rm ex})/\rho$  is listed] and constraints imposed on the system of supersaturated vapor. The cutoff distance was  $r_c = 8.0\sigma$ .

TABLE I. (Continued.)

				$ ho\sigma^3$	$p\sigma^3/\varepsilon$	$f'/\varepsilon$	$d_{\mathrm{max}}$
$ ho\sigma^3$	$p\sigma^3/\varepsilon$	f'/arepsilon	$d_{ m max}$		0.015 86	-1.1952	8
				0.05	0.024 30	-1.2491	4
	$T^*=0$				0.023 47	-1.2650	5
0.030	0.016 51	-1.0651	4		0.022 46	-1.2761	6
	0.016 45	-1.0762	5		0.017 01	-1.2513	7
	0.016 32	-1.0824	6	0.055	0.024.06	1 2022	4
	0.016 20	-1.0852	7	0.055	0.024 86	-1.2932	4
	0.016 16	-1.0870		8	0.023 46	-1.3111	5
	0.015 97	-1.0889			0.019 75	-1.3101	6
	0.013 71	-1.0853	10		0.009 05	-1.2592	7
0.035	0.018 20	-1.1171	4	0.06	0.024 94	-1.3387	4
	0.017 91	-1.1320	5		0.022 44	-1.3615	5
	0.017 58	-1.1405	6		0.012 24	-1.3090	6
	0.017 30	-1.1450	7		0.000 82	-1.2555	7
	0.015 85	-1.1457	8		T*=0	0.00	
	0.013 13	-1.1320	9	0.02	0.021 97		4
	0.010 14	-1.1917	10	0.03	0.021 97	-1.1010	4
0.04	0.010.05	4.4000	_			-1.1069	5
0.04	0.018 87	-1.1888	5		0.021 85	-1.1097	6
	0.018 89	-1.2009	6		0.021 82	-1.1103	7
	0.016 34	-1.1982	7		0.021 81	-1.1105	8
0.045	0.020 22	-1.2218	4		0.021 81	-1.1107	9
0.015	0.019 21	-1.2441	5		0.021 81	-1.1104	10
	0.017 35	-1.2516	6	0.04	0.027 03	-1.1676	4
	0.017 33	-1.2277	7		0.026 88	-1.1757	5
	0.011 70	-1.2277	,		0.026 80	-1.1793	6
0.05	0.020 53	-1.2765	4		0.026 74	-1.1796	7
	0.018 76	-1.3009	5		0.026 74	-1.1814	8
	0.013 94	-1.3010	6		0.026 75	-1.1819	9
	0.004 80	-1.2727	7		0.026 75	-1.1813	10
	T*=0	0.80		0.05	0.030 94	-1.2340	4
0.03	0.018 46	-1.0691	4	0.03	0.030 64	-1.2436	5
	0.018 33	-1.0783	5		0.030 50	-1.2484	6
	0.018 24	-1.0825	6		0.030 38	-1.2513	7
	0.018 21	-1.0843	7		0.030 36	-1.2525	8
	0.018 20	-1.0852	8		0.030 30	-1.2525 -1.2519	9
	0.018 16	-1.0857	9				10
	0.018 17	-1.0860	10		0.027 34	-1.2338	10
0.025	0.000.44	4.4440	,	0.06	0.033 67	-1.3007	4
0.035	0.020 44	-1.1143	4		0.033 08	-1.3116	5
	0.020 24	-1.1248	5		0.032 70	-1.3188	6
	0.020 09	-1.1302	6		0.031 20	-1.3181	7
	0.020 02	-1.1328	7		0.023 43	-1.2791	8
	0.019 94	-1.1340	8		0.016 56	-1.2528	9
	0.019 84	-1.1351	9		0.012 98	-1.2858	10
	0.018 81	-1.1286	10	0.07	0.035 01	-1.3642	4
0.04	0.022 09	-1.1590	4	0.07	0.034 07	-1.3810	5
	0.021 76	-1.1775	5		0.029 14	-1.3717	6
	0.021 54	-1.1801	6		0.027 14	-1.3272	7
	0.021 29	-1.1824	7		0.017 31	1.3212	,
	0.020 68	-1.1828	8	0.08	0.034 88	-1.4299	4
	0.017 27	-1.1630	9		0.029 80	-1.4316	5
	0.011 51	-1.1347	10		0.012 44	-1.3600	6
0.45					0.005 03	-1.4032	7
0.45	0.023 37	-1.2044	4	0.085	0.034 38	-1.4638	4
	0.022 90 0.022 43	-1.1291 -1.2276	5 6	0.003	0.034 56	-1.4351	5
		-1 ///6	n		0.045 50	1.7331	J

$ ho\sigma^3$	$p\sigma^3/arepsilon$	f'/arepsilon	$d_{\mathrm{max}}$			
	0.000 60	-1.4619	7			
	T*=1.00					
0.04	0.031 74	-1.2025	4			
	0.031 66	-1.2081	5			
	0.031 62	-1.2110	6			
	0.031 61	-1.2114	7			
0.05	0.037 12	-1.2533	4			
	0.036 99	-1.2595	5			
	0.036 92	-1.2619	6			
	0.036 91	-1.2632	7			
0.06	0.041 46	-1.3030	4			
	0.412 7	-1.3107	5			
	0.411 9	-1.3142	6			
	0.411 3	-1.3152	7			
0.07	0.447 6	-1.3534	4			
	0.044 50	-1.3609	5			
	0.044 36	-1.3669	6			
	0.044 22	-1.3676	7			
0.08	0.046 98	-1.4022	4			
	0.046 55	-1.4130	5			
	0.046 16	-1.4167	6			
	0.044 73	-1.4145	7			
0.09	0.048 14	-1.4517	4			
	0.047 41	-1.4658	5			
	0.042 33	-1.4568	6			
	0.033 72	-1.4536	7			
0.1	0.047 97	-1.5002	4			
	0.041 28	-1.4943	5			
	0.029 36	-1.4802	6			
	0.028 86	-1.5368	7			

mum. We perform the polynomial interpolation for the 0.035 case and the result is displayed in Fig. 1, which clearly tells us that the location of the free energy minimum is between  $d_{\rm max}$ =7 and  $d_{\rm max}$ =8. Thus, we may conclude that the pressure for 0.035 under  $T^*$ =0.75 is between 0.017 30 and 0.017 58. The other exception is for  $T^*$ =1.0 and  $\rho\sigma^3$ =0.1 and  $T^*$ =1.0 and  $T^*$ =1

Figures 2 and 3 show the comparisons of pressure from our simulation and those from Ref. 19 at  $T^*$ =0.75 and  $T^*$ =1.0, respectively. It is obvious that at higher temperature, both results correspond very well but at low temperature the discrepancy is large. At this stage the reason why the huge discrepancy arises at low temperature is not clear; it may be ascribed to the uncertainties in identifying the beginning and ending of the induction period in Ref. 19 or the constraints imposed on the system in our simulation.

We would also like to estimate the spinodal density for a given temperature with the simulated data. Here, we omit

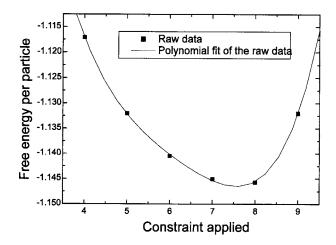


FIG. 1. Polynomial interpolation on the free energies for different constraints imposed on to  $T^*=0.75$  and  $\rho\sigma^3=0.035$ .

this stage because our data do not span a sufficient range to calculate this density. Our purpose here is to test the robustness of our method for a large range of densities and temperatures.

#### IV. SUMMARY

In this work we extend our simulation of free energies of the supersaturated LJ vapor from  $T^*=0.7$  to 1.0 and compared our data with those reported in Ref. 19. As concluding remarks, several things have to be pointed out. First, compared with the method used in Ref. 19, our method provides an effective way to identify the configurations which belong to the phase separation or close to the equilibrium of the metastable state, and it requires no judgment to determine the beginning of phase separation; second, our method is not difficult to apply. On average, four or five runs are enough to determine the thermal properties at given densities because the polynomial interpolation could be very useful when it is hard to determine the local minimum of the free energy due to the smoothness of the free energy data. Third, the system size remains unchanged during simulations for a certain temperature. In our method, what is required is to lower the  $d_{\text{max}}$ value; thus, there is no need to increase the system size in

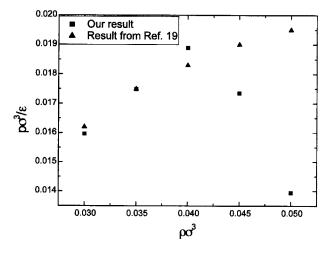


FIG. 2. Pressure-density data at  $T^*=0.75$ .

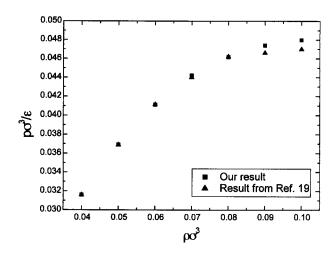


FIG. 3. Pressure-density data at  $T^*=1.0$ .

order to obtain longer induction time. Finally, our method is easy to extend to more complex molecular systems such as the water case.

We also need to point out that the discrepancies arising in the low temperature cases are not fully resolved. Further investigations are needed to explore the exact reasons for this disagreement with prior work.

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