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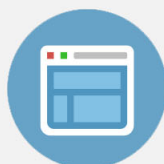
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Finite-size scaling investigation of the liquid-liquid critical point in ST2 water and its stability with respect to crystallization

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The liquid-liquid critical point scenario of water hypothesizes the existence of two metastable liquid phases—low-density liquid (LDL) and high-density liquid (HDL)—deep within the supercooled region. The hypothesis originates from computer simulations of the ST2 water model, but the stability of the LDL phase with respect to the crystal is still being debated. We simulate supercooled ST2 water at constant pressure, constant temperature, and constant number of molecules N for $N \leq 729$ and times up to $1 \mu\text{s}$. We observe clear differences between the two liquids, both structural and dynamical. Using several methods, including finite-size scaling, we confirm the presence of a liquid-liquid phase transition ending in a critical point. We find that the LDL is stable with respect to the crystal in 98% of our runs (we perform 372 runs for LDL or LDL-like states), and in 100% of our runs for the two largest system sizes ($N = 512$ and 729 , for which we perform 136 runs for LDL or LDL-like states). In all these runs, tiny crystallites grow and then melt within $1 \mu\text{s}$. Only for $N \leq 343$ we observe six events (over 236 runs for LDL or LDL-like states) of spontaneous crystallization after crystallites reach an estimated critical size of about 70 ± 10 molecules. © 2013 AIP Publishing LLC. [<http://dx.doi.org/10.1063/1.4808355>]

I. INTRODUCTION

For many centuries, water and its anomalies have been of much interest to scientists. A particular rise of interest occurred in the late 1970s after experiments done by Angell and Speedy^{1–4} seemed to imply some kind of critical phenomenon in supercooled liquid water at very low temperatures. Even though liquid water experiments are limited by spontaneous crystallization below the homogeneous nucleation temperature ($T_H \approx 233 \text{ K}$ at 1 bar), it is possible to further explore the phase diagram by quenching water to far lower temperatures.^{5–7} The result of these experiments is an amorphous solid, i.e., a glassy ice, corresponding to an out-of-equilibrium state that is very stable with respect to the equilibrium crystalline ice phase. The type of amorphous solid depends on the applied pressure: at low pressure, below $\approx 0.2 \text{ GPa}$, the low density amorphous ice (LDA) is formed, while at higher pressure the high density amorphous ice (HDA) is observed.⁸ It has been shown by Mishima *et al.*^{9–12} that these two amorphous ices are separated by a reversible abrupt change in density that resembles in all its respects an equilibrium first order phase transition.

Raising the temperature of either LDA or HDA does not turn the sample into a liquid, but leads once again to spontaneous crystallization (around $T_X \approx 150 \text{ K}$). In fact, between T_H and T_X , often called the “no man’s land” of bulk water, crystallization occurs at a time scale that is too short for current experimental methods, although a new technique is

possibly succeeding in the task of measuring the metastable liquid phase.¹³ Computer simulations of water, however, involve time scales small enough to witness spontaneous crystallization and are therefore able to explore liquid water in the “no man’s land.” In 1992, Poole *et al.*¹⁴ performed a series of molecular dynamics simulations using the ST2 water model,¹⁵ using the reaction field method for the long-range interactions (ST2-RF), and discovered a liquid-liquid phase transition (LLPT) ending in a critical point, separating a low density liquid (LDL) and a high density liquid (HDL). These two liquids can be considered to be the liquid counterparts of the LDA and HDA, respectively.

The hypothesis of the critical point also allows one to understand X-ray spectroscopy results,^{16–19} explains the increasing correlation length in bulk water upon cooling as found experimentally,²⁰ the hysteresis effects,²¹ and the dynamic behavior of protein hydration water.^{22–24} It would be consistent with a range of thermodynamical and dynamical anomalies^{25–33} and experiments.^{34–37}

Many more computer simulations investigating the phenomenology of the liquid-liquid critical point (LLCP) have been performed since then.^{38–55} Detailed studies using ST2-RF have been made by Poole *et al.*⁵⁶ using molecular dynamics, while Liu *et al.*^{57,58} simulated ST2 with Ewald summation (ST2-Ew) for the electrostatic long-range potential using Monte Carlo. Also in other water models the LLPT and its LLCP are believed to be found, for example, by

Yamada *et al.*⁵⁹ in the TIP5P model, by Paschek *et al.*⁶⁰ in the TIP4P-Ew model, and in TIP4P/2005 by Abascal and Vega.^{61,62}

Recently, Limmer and Chandler⁶³ used Monte Carlo umbrella sampling to investigate the ST2-Ew model, but claimed to have found only one liquid metastable phase (HDL) rather than two. They therefore concluded that LDL does not exist because it is unstable with respect to either the crystal or the HDL phase. The emphasis in their work is about the difference between a metastable phase, i.e., separated from the stable phase by a finite free-energy barrier, and an unstable state, where the free-energy barrier is absent and the state does not belong to a different phase.

Shortly after, Poole *et al.*⁶⁴ and Kesselring *et al.*⁶⁵ presented results using standard molecular dynamics for ST2-RF showing the occurrence of the LLCP with both HDL and LDL phases metastable with respect to the crystal, but with the LDL not unstable with respect to either the crystal or the HDL. This result was confirmed, using the same method as Limmer and Chandler,⁶³ by Sciortino *et al.*⁶⁶ and Poole *et al.*⁶⁷ in ST2-RF water and by Liu *et al.*⁶⁸ in ST2-Ew water.

The aim of this paper is to investigate the possible existence of a liquid-liquid critical point in simulated water in the thermodynamic limit using finite-size scaling techniques, and confirm that LDL is a bona fide metastable liquid. We use the ST2-RF model because it has been well-studied in the supercooled region, making it easier to compare and verify our data. In the supercooled phase, it has a relatively large self-diffusion compared to other water models, therefore, suffers less from the slowing down of the dynamics at extremely low temperatures. We explore a large region of the phase diagram of supercooled liquid ST2-RF water (Fig. 1) using molecular dynamics simulations with four different system sizes by keeping constant the number N of molecules, the pressure P , and the temperature T (NPT ensemble).

Within the explored region we find both LDL and HDL, separated at high pressures by a LLPT, ending in a LLCP estimated at $P_C \approx 208$ MPa and $T_C \approx 246$ K. This phase transition is particularly clear in Fig. 2 where one can see from the density how the system continuously flips between the two states. However, due to finite-size effects this phase flipping also occurs below the critical point along the Widom line (the locus of correlation length maxima).^{69,70} For this reason, it is necessary to apply finite-size scaling methods to establish the exact location of the critical point.

For six state points and for small system size $N \leq 343$ we observe, in only one over the (on average) seven simulations we performed for each state point, irreversible crystal growth, indicated as full red circles in Fig. 1. Each of these crystallization events occurred within the LDL (or LDL-like) region. Analysis of these crystals revealed them to have a diamond cubic crystal structure. As we will discuss later, because these events disappear for larger systems, we ascribe these crystallization to finite-size effects.

We start in Sec. II with a description of the model and the procedures that we use. In Sec. III, we discuss the use of the intermediate scattering function to analyze the structure of the liquid, and in Sec. IV its use in defining the correlation time. The analysis of the liquid structure is continued

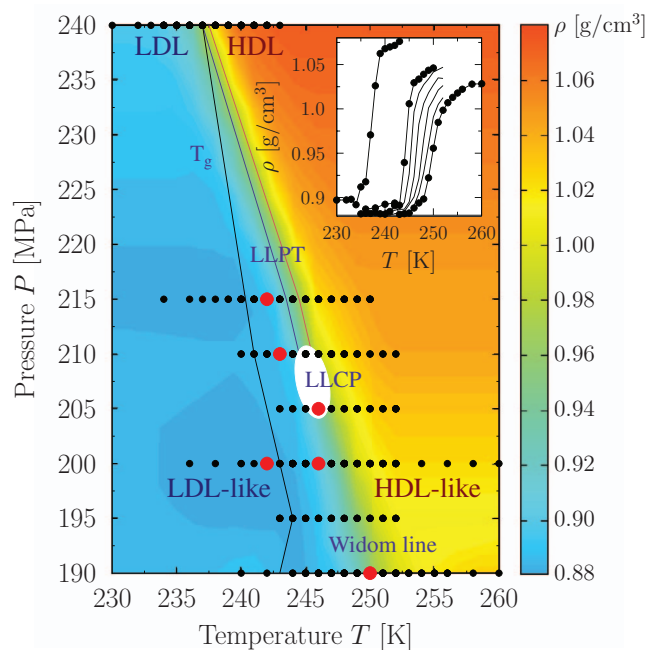


FIG. 1. Overview of the state points at which simulations have been performed. Colors away from the simulated points (full black circles) are a linear interpolation of ρ for the sake of presentation. At high temperatures, we observe a high-density liquid state (HDL, shaded in orange), while at lower temperatures we find a low-density liquid (LDL, in blue). These are separated by a region where the system is continuously flipping between the two states, as seen in Fig. 2. This transition region (in green) is identified as the liquid-liquid phase transition (LLPT) line at high pressures, and the Widom line at low pressures. These lines join at the liquid-liquid critical point (LLCP) estimated at $P_C = 208 \pm 3$ MPa and $T_C = 246 \pm 1$ K (see Sec. VII). At low temperatures, the LDL (or LDL-like) region is bound by the glass transition temperature T_g , below which we can no longer fully equilibrate the system within 100 ns, and consider the liquid to have become a glass (see Sec. IV). For small sizes ($N \leq 343$), we observe spontaneous crystallization within 1 ns-long simulations at six state points (indicated by the red circles), all of them within the LDL (or LDL-like) region. We never observe crystallization for sizes $N = 512$ and 729 for simulations of comparable duration. Because the probability of crystallization should increase with N , these results suggest that our crystallization events are a finite-size effect that becomes negligible for large sizes. Crystallization events are discussed in Sec. VI. (Inset) Average $\rho(T)$ for pressures $P = 240$ MPa (left-most), 215, 210, 205, 200, 195, and 190 (right-most). Here, the average is taken over all N ; excluding $N \leq 343$ shows a much sharper transition at 240 MPa.

in Sec. V where we define and compare a selection of structural parameters. The parameter d_3 is found to be particularly well-suited to distinguish between the liquid and the crystal state, and this fact is subsequently used in Sec. VI where we discuss the growth and melting of crystals within the LDL liquid. In Sec. VII, by defining the appropriate order parameter, we show that the LLCP in ST2-RF belongs to the same universality class as the 3D Ising model. We accurately determine where the LLCP is located in the phase diagram in the thermodynamic limit by applying finite-size scaling on the Challa-Landau-Binder parameter. We discuss our results and present our conclusions in Sec. VIII.

II. SIMULATION DETAILS

In the ST2 model,¹⁵ each water molecule is represented by a rigid tetrahedral structure of five particles. The central

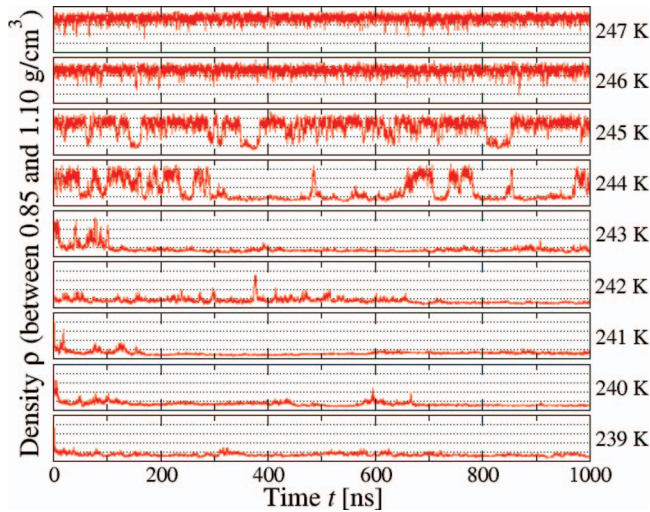


FIG. 2. Phase flipping near the phase transition line ($P = 215$ MPa, with $N = 343$ molecules). At high T , the system is in the HDL phase (with a density $\rho \simeq 1.03$ g/cm³), while at low T the system is in the LDL phase (density $\rho \simeq 0.88$ g/cm³). However, near the phase transition line (at $T \simeq 244.5$ K for this pressure) the system is flipping between the two phases.

particle carries no charge and represents the oxygen atom of water. It interacts with all other oxygen atoms via a Lennard-Jones (LJ) potential, $U_{\text{LJ}}(r_{ij}) \equiv 4\epsilon[(\sigma/r_{ij})^{12} - (\sigma/r_{ij})^6]$ with $\epsilon \equiv 0.31694$ kJ/mol and $\sigma \equiv 3.10$ Å. Two of the outer particles represent the hydrogen atoms. Each of them carries a charge of $+0.2357$ e, and is located a distance 1 Å away from the central oxygen atom. The two remaining particles carry a negative charge of -0.2357 e, are positioned 0.8 Å from the oxygen, and represent the lone pairs of a water molecule.

The electrostatic potential in ST2 is treated in a special way. To prevent charges a and b from overlapping, the Coulomb potential is reduced to zero at small distances

$$U_{\text{el}}(r_{ab}) \equiv S(r_{ij}) \frac{1}{4\pi\epsilon_0} \frac{q_a q_b}{r_{ab}}, \quad (1)$$

where $S(r_{ij})$ is a function that smoothly changes from one to zero as the distance between the molecules decreases,

$$S(r_{ij}) \equiv \begin{cases} 0 & (r_{ij} \leq R_L) \\ \frac{(r_{ij}-R_L)^2(3R_U-R_L-2r_{ij})}{(R_U-R_L)^3} & (R_L \leq r_{ij} \leq R_U) \\ 1 & (r_{ij} \geq R_U) \end{cases} \quad (2)$$

with $R_L \equiv 2.0160$ Å, $R_U \equiv 3.1287$ Å, and where r_{ij} is the distance between the oxygen atoms of the interacting molecules. In the original model, a simple cutoff was used for the electrostatic interactions. In this paper, however, we apply the reaction field method⁷¹ which changes the ST2 Coulomb potential to

$$U_{\text{el}}(r_{ab}) \equiv S(r_{ij})T(r_{ij}) \frac{q_a q_b}{4\pi\epsilon_0} \left(\frac{1}{r_{ab}} + \frac{r_{ab}^2}{2R_c^3} \right), \quad (3)$$

where $T(r_{ij})$ is another smoothing function

$$T(r_{ij}) \equiv \begin{cases} 1 & (r_{ij} \leq R_T) \\ 1 - \frac{(r_{ij}-R_T)^2(3R_c-R_T-2r_{ij})}{(R_c-R_T)^3} & (R_T \leq r_{ij} \leq R_c) \\ 0 & (r_{ij} \geq R_c) \end{cases} \quad (4)$$

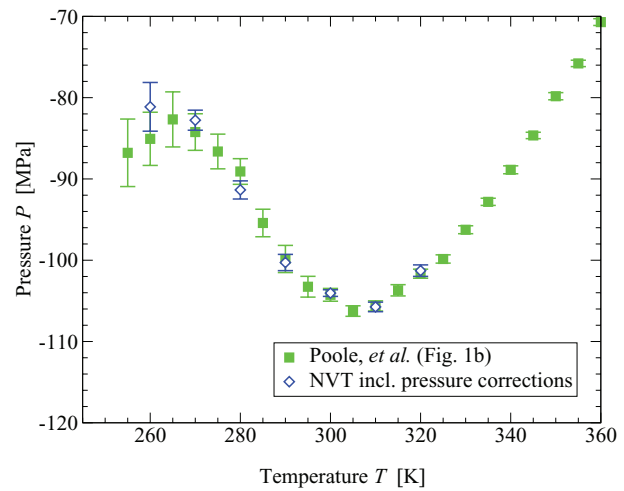


FIG. 3. To validate our code, we compare our simulation results with those from Poole *et al.*⁵⁶ at density $\rho = 0.83$ g/cm³ and for $N = 216$ molecules. We performed simulations in the NVT ensemble applying pressure corrections and find the same results as Ref. 56 within the error bars. At this density, the pressure correction due to the LJ cutoff (proportional to ρ^2) is equal to -12.66 MPa. The variation of P with T along this isochore shows the occurrence of both a density maximum at 305 K and a density minimum near 265 K, as at these state points $(\partial\rho/\partial T)_P = -\rho K_T(\partial P/\partial T)_V = 0$ with $K_T > 0$ the isothermal compressibility.

We use a reaction field cutoff $R_c \equiv 7.8$ Å together with $R_T \equiv 0.95R_c$. These parameters define our ST2-RF water model and are the same that were used in previous ST2-RF simulations.

For the LJ interaction, we use a simple cutoff at the same distance of 7.8 Å. We do not adjust the pressure to correct for the effects of the LJ cutoff,^{72,73} since these adjustments come from mean field calculations which become increasingly unreliable as one approaches a critical point. In order to facilitate comparing results with and without this correction, we denote that for LDL ($\rho \approx 0.90$ g/cm³) the correction is $P \approx -15$ MPa, and for HDL ($\rho \approx 1.05$ g/cm³) the correction is $P \approx -20$ MPa.

We use the SHAKE algorithm⁷⁴ to keep the relative position of each particle within a ST2 molecule fixed. The temperature and pressure are held constant using a Nosé-Hoover thermostat^{73,75,76} together with a Berendsen barostat.⁷⁷ In all simulations, periodic boundary conditions are applied.

Our code is validated by simulating the same state points as those published by Poole *et al.*, see Fig. 1(b) in Ref. 56, where pressure corrections for the LJ cutoff were applied in the NVT (constant N , T , and volume V) ensemble. Averaging at each state point over 10 simulations with different initial conditions allows us to estimate the error bars. In Fig. 3, we compare our results for $N = 216$ molecules and density 0.83 g/cm³, and find that our data, after pressure correction, match that of Ref. 56 well.

For each of the simulations done in the NPT ensemble, we use the following protocol. We first create a box of N molecules at n different initial densities (with n up to 21) ranging from 0.85 to 1.05 g/cm³. We then perform a 1 ns NVT simulation at $T = 300$ K. In this way, we obtain n independent configurations all at $T = 300$ K in the prefixed range of densities. Next, we use these independent configurations as

starting points for NPT simulations at $T = 265$ K and different pressures ranging from 190 to 240 MPa, and continue the simulation for an additional 1 ns. This results in n independent configurations at $T = 265$ K and the given pressure. For all pressures considered here, this will lead the system into the HDL phase. Finally, the system is quenched to the desired temperature at the given pressure, followed by 100–200 ns of equilibration time. In Sec. IV, it will be shown that this provides enough time for the system to reach equilibrium for the state points above the line marked with the label T_g in Fig. 1.

III. INTERMEDIATE SCATTERING FUNCTION

The intermediate scattering function $S(\mathbf{k}, t)$ plays an essential role in the analysis of liquid structure, since it is frequently measured in experiments as well as easily calculated from simulation data. It describes the time evolution of the spatial correlation at the wave vector \mathbf{k} , and can be used to distinguish between phases of different structure, such as LDL and HDL or crystal. It is defined as

$$S(\mathbf{k}, t) \equiv \frac{1}{N} \left\langle \sum_{\ell, m} e^{i\mathbf{k} \cdot [\mathbf{r}_\ell(t') - \mathbf{r}_m(t' + t)]} \right\rangle_{t'},$$

where $\langle \dots \rangle_{t'}$ denotes averaging over simulation time t' , and $\mathbf{r}_\ell(t')$ the position of particle ℓ at time t' . For simplicity, we only apply the intermediate scattering function to the oxygen atoms, which we denote as $S_{OO}(k, t)$.

Since the system has periodic boundary conditions, the components of \mathbf{k} have discrete values $2\pi n/L$, where L is the length of the simulation box and $n = 1, 2, 3, \dots$. We define $S_{OO}(k, t) \equiv \langle S_{OO}(\mathbf{k}, t) \rangle_n$ where the average is taken over all vectors \mathbf{k} with magnitude k belonging to the n th spherical bin $\pi(n - \frac{1}{2})/L \leq k < \pi(n + \frac{1}{2})/L$ for $n = 2, 3, \dots, 300$. Similarly, we define the structure factor $S_{OO}(k) \equiv \langle S_{OO}(k, t) \rangle_t$ as the time-averaged intermediate scattering function, with (unless indicated otherwise) the average taken over the whole duration of the run.

We study $S_{OO}(k)$ above and below our estimate for the LLCP pressure. At $P = 210$ MPa $> P_C$ (Figs. 4(a) and 4(b)), we observe a discontinuous change in the first two peaks of $S_{OO}(k)$ as T changes between 245 and 246 K, and a continuous change above and below these temperatures. This is the expected behavior for a first order phase transition occurring at $245 \text{ K} \lesssim T \lesssim 246 \text{ K}$ and $P = 210$ MPa between two phases with different structure, consistent with our results in Fig. 1. The fact that for both phases $S_{OO}(k) \sim O(1)$ for all k shows that both phases are fluid. Indeed, for a crystal-like configuration, with a long-range order, there would be at least one wave vector such that $S_{OO}(k) \sim O(N)$.⁷⁸ Furthermore, the fact that at lower T the first peak increases and the other peaks only have minor changes indicates that the lower- T liquid has a smaller density than the higher- T liquid. Therefore, this result shows a first-order phase transition between the LDL at lower- T and HDL at higher- T . This transition occurs at the same temperature at which we observe the phase flipping in density (Fig. 2) and corresponds to the yellow/green region at $P > P_C$ in Fig. 1.

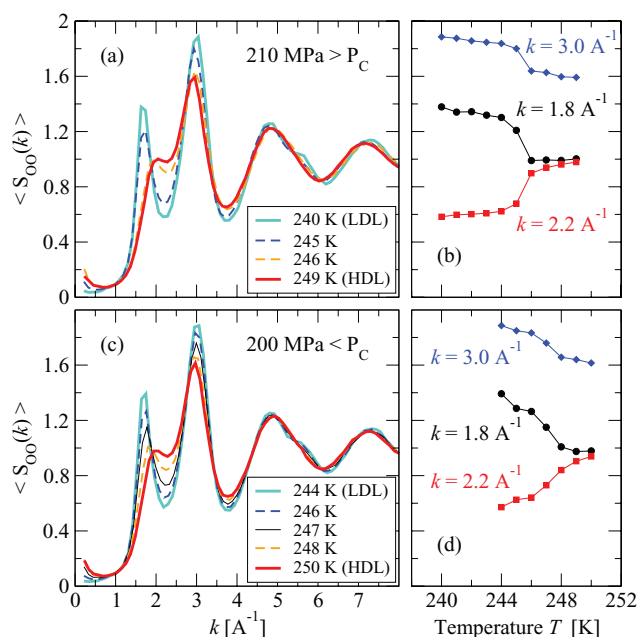


FIG. 4. The structure factor $S_{OO}(k)$ for a range of temperatures at (a) 210 MPa and (c) 200 MPa for $N = 729$. (a) For $P > P_C$, the structure has a large change between $T = 245$ and 246 K, corresponding to the LDL-HDL first-order phase transition. (b) The value of S_{OO} for k corresponding to the first maximum, the first minimum, and the second maximum as a function of T for $P = 210$ MPa as in panel (a). (c) For $P < P_C$, the structure changes in a way that is smoother than the case in panel (a), with the more evident change occurring between $T = 249$ and 250 K, corresponding to the crossing of the Widom line, as marked by the value of S_{OO} at first maxima and minima in panel (d).

The fact that the peaks of $S_{OO}(k)$ are sharper in LDL than HDL is an indication that the LDL phase is more structured. We can also observe that the major structural changes in $S_{OO}(k)$ between LDL and HDL are for $k \simeq 1.8$ and 2.8 Å^{-1} , corresponding to $r = 4\pi/k \simeq 7$ and 4.5 Å , respectively, i.e., the third and the second neighbor water molecules. This change in the structure is consistent with a marked shift inwards of the second shell of water with increased density, and almost no change in the first shell (at $k \simeq 4.6 \text{ Å}^{-1}$ and $r \simeq 2.75 \text{ Å}$), as seen in structural experimental data for supercooled heavy water interpreted with reverse Monte Carlo method.⁷⁹ These changes are visible also in the OO radial distribution function $g_{OO}(r)$.

For $P < P_C$ (Figs. 4(c) and 4(d)) by increasing T , we observe that the first peak of $S_{OO}(k)$ merges with the second, transforming continuously into a shoulder (the same qualitative behavior can be observed for $g_{OO}(r)$). These quantities also show us that the lower- T structure is LDL-like, while the higher- T structure is HDL-like. However, the absence of any discontinuous change in the structure implies the absence of a first-order phase transition in the structure of the liquid. This is consistent with the occurrence of a LLCP at the end of the first-order phase transition somewhere between 200 and 210 MPa, at a temperature between 245 and 250 K. In Sec. VII, we shall apply a different method to locate the LLCP with more precision.

At $P < P_C$, in the one-phase region, we expect to find the Widom line emanating from the LLCP. The Widom line

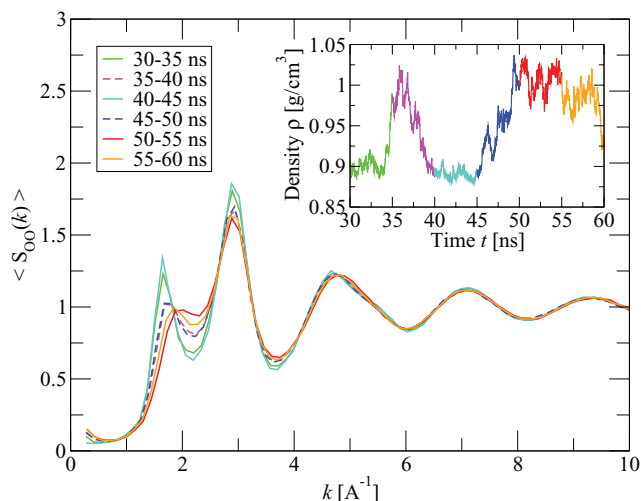


FIG. 5. As the density changes from $\rho(\text{LDL})$ to $\rho(\text{HDL})$, also the structure changes. The inset shows how the density is changing with time for six consecutive time intervals of 10 ns, with the corresponding $S_{OO}(k)$ shown in the main plot ($N = 343$ at 200 MPa and 248 K).

is by definition the locus of maxima of the correlation length, therefore, from general thermodynamic considerations⁷⁰ near the LLC, it must be also the locus of maxima of the response functions. In particular, it must be the locus where the isobaric heat capacity $C_P \equiv T(\partial S/\partial T)_P$, where S is the entropy of the system, has its maximum along a constant- P path. This maximum occurs where the entropy variation with T is a maximum, and is expected where the structural variation of the liquid is a maximum, i.e., where the derivatives of the values of $S_{OO}(k)$ (Fig. 4(d)) and $g_{OO}(r)$ with T are at a maximum. The interval of temperatures for each P where this occurs corresponds to the green region at $P < P_C$ in Fig. 1, indicated as the Widom line.

It is actually possible to follow the structural changes during the simulation. An example is given in Fig. 5 where we focus on a 30 ns time period of a simulation at 200 MPa and 248 K. We divide this time period into six 5 ns intervals and for each interval we calculate the intermediate scattering function, time-averaged over those 5 ns. We observe that the liquid is LDL-like for the first and third interval, having low density and LDL-like $S_{OO}(k)$ (first peak near 2 \AA^{-1} , separated from the second). On the contrary, for the fifth and sixth intervals the density is high and $S_{OO}(k)$ is HDL-like (the first peak is merely a shoulder of the second peak), indicating that the liquid is HDL-like. For the second and fourth intervals, the liquid has an intermediate values of density and $S_{OO}(k)$, indicating that it is a mix of LDL-like and HDL-like structures.

IV. CORRELATION TIME

Apart from its use in structure analysis, the intermediate scattering function $S_{OO}(k, t)$ can also be used to define a correlation time τ , i.e., the time it takes for a system to lose most of its memory about its initial configuration.^{80,81}

In Fig. 6, we show how $S_{OO}(k, t)$ decays with time for a fixed value of k . Its decay is characterized by two relaxation times, the α -relaxation time τ_α and the β -relaxation time τ_β .

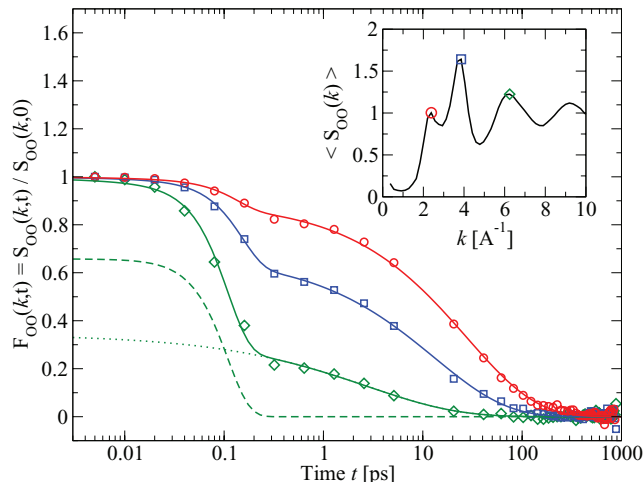


FIG. 6. Decay of $S_{OO}(k, t)$ with time, for $P = 210 \text{ MPa}$, $T = 250 \text{ K}$, and $N = 343$. Symbols indicate $F_{OO}(k_i, t)$ for three different values of k : the first maximum of $S_{OO}(k)$ at k_1 (red circles), the second maximum at k_2 (blue squares), and the third maximum k_3 (green diamonds). Solid lines are fits according to Eq. (5). The two components of Eq. (5) are explicitly shown for $F_{OO}(k_3, t)$: the green dashed line represents the β -relaxation and is given by $[1 - A(k)]\exp[-(t/\tau_\beta)^2]$, the green dotted line represents the α -relaxation and satisfies $A(k)\exp[-(t/\tau_\alpha)^b]$. The solid green line going through $F_{OO}(k_3, t)$ is the sum of both.

On very short time scales, the molecules do not move around much and each molecule is essentially stuck in a cage formed by its neighbors. This is represented by the β -relaxation time τ_β which is on the order of picoseconds. On longer time scales, the molecule can escape from its cage and diffuse away from its initial position. The time τ_α is the relaxation time of this structural process.

Mode-coupling theory of supercooled simple liquids predicts that⁸²

$$F_{OO}(k, t) \equiv S_{OO}(k, t)/S_{OO}(k, 0) \\ = [1 - A(k)]e^{-(t/\tau_\beta)^2} + A(k)e^{-(t/\tau_\alpha)^b}. \quad (5)$$

The factor $A(k)$ is the Debye-Waller factor arising from the cage effect, which is independent of the temperature and follows $A(k) = \exp(-a^2k^2/3)$ with a the radius of the cage. We are able to fit Eq. (5) remarkably well to all our data, as, for example, in Fig. 6.

Data in Fig. 6 were collected every 10 fs for simulations of 1 ns. This rate of sampling results in a large amounts of data and is unfeasible for our runs up to 1000 ns. Therefore, for the 1000 ns runs we collect data at 10 ps intervals. At this rate of sampling, it is no longer possible to estimate τ_β or the cage size a , but it is still possible to determine τ_α accurately, utilizing the fact that $S_{OO}(k, t)$ reaches a plateau near $t \approx \tau_\beta$. One can therefore define

$$C_{OO}(k, t) \equiv S_{OO}(k, t)/S_{OO}(k, \tau_\beta), \quad (6)$$

which is $S(k, t)$ normalized by its value at the plateau. A good estimate of τ_α is then the time for which $C_{OO}(k, \tau_\alpha) = 1/e \approx 0.37$.

From the shorter 1 ns runs (which were mostly done in the HDL regime), we find that the cage radius is $a = 0.35 \pm 0.09 \text{ \AA}$ with a stretching exponent of $b = 0.63 \pm 0.09$. Both

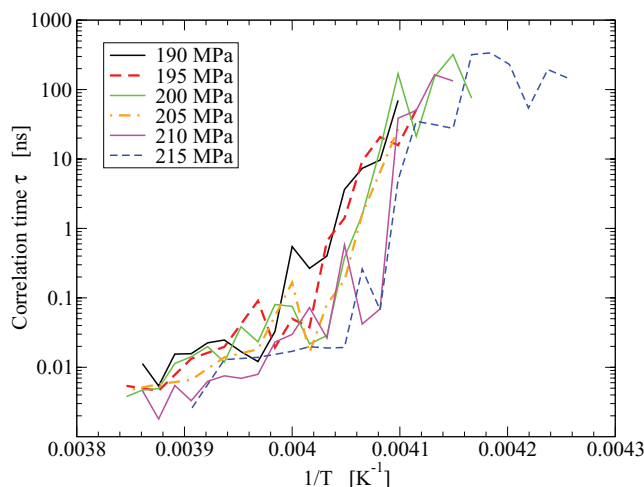


FIG. 7. Arrhenius plot of the correlation time τ for different pressures. Errors on our estimates are of the order of the discontinuities along the curves. At high temperatures (the HDL regime), the correlation time is of the order of 10–100 ps, which jumps several orders up as we pass the phase transition line and enter the LDL regime. To obtain this plot, we dismissed the simulations that had a significant increase in τ because of crystal growth (see Sec. VI).

parameters a and b do not show a significant dependence on the state point within the studied range of temperatures and pressures.

As shown in Fig. 6, different k result in slightly different values for τ_α . We use as the correlation time τ the largest value of τ_α which is usually found at $k = k_1$, the first maximum in $\langle S_{OO}(k) \rangle$ (inset Fig. 6).

As is to be expected, the correlation time does not seem to depend on the box size. It does however depend strongly on the phase, which is evident from Fig. 7.

At high temperatures, the system is in the HDL phase, and has a correlation time τ on the order of 10–100 ps. As we decrease the temperature at fixed pressure, the value of τ has a large increase when we cross the phase transition line or the Widom line, depending on if P is above or below P_C , respectively. Apparently, the LDL states evolve nearly four orders of magnitude slower than HDL states, with correlation times in the nanosecond range.

If we lower the temperature further, the correlation time slowly increases until the system becomes a glass rather than a liquid, and we are no longer able to fully equilibrate the system. As we can only run simulations up to 1000 ns, we consider the state points with a correlation time above 100 ns to be beyond our reach. We therefore designate the effective glass transition temperature T_g as the temperature for which $\tau > 100$ ns (see Fig. 1).

V. STRUCTURAL PARAMETERS

Apart from the intermediate scattering function, there are other ways to quantify the structure of a liquid. In this section, we shall examine several structural parameters, and determine which of those are the most effective in distinguishing between LDL, HDL, and the crystal. Apart from considering several global parameters, we shall mainly focus on local parameters which can be used to analyze the local environment

of a molecule. In addition, we study here not only the parameters based on the nearest neighbors, but also those based on the next-nearest neighbors (the second coordination shell). Finally, we determine the best local parameter to use for identifying tiny crystals in the liquid, an important result that will be used in Sec. VI.

The structural parameters are designed to distinguish between different phases by analyzing the geometrical structure. This is typically done by evaluating the spherical harmonics $Y_\ell^m(\varphi, \vartheta)$ for a particular set of neighboring atoms, with φ and ϑ the polar angles between each pair of oxygen atoms in that set. In this paper, we consider two different sets: we define the first coordination shell $n_1(i)$ to be the four nearest neighbors of molecule i , and define the second coordination shell $n_2(i)$ as the fifth to 16th nearest neighbors (the 16th nearest neighbors minus those in the first shell). For simplicity, we approximate the center of mass of a water molecule with the center of its oxygen atom.

Different values of ℓ are sensitive to different symmetries. The spherical harmonics with $\ell = 3$, for example, are sensitive to a diamond structure. Those with $\ell = 6$ are more sensitive to the hexagonal closest packing (hcp) structure. Since we expect the liquid and crystal structures to be hcp, diamond, or a mix of these, we focus primarily on $\ell = 3$ and $\ell = 6$.

A. Parameters q_3 and q_6

All parameters defined in this section are based on $q_{\ell,m}^{(s)}(i)$ which quantifies the local symmetry around molecule i . It is defined as

$$q_{\ell,m}^{(s)}(i) \equiv \frac{1}{N_s} \sum_{j \in n_s(i)} Y_\ell^m(\varphi_{ij}, \vartheta_{ij}) \quad -\ell \leq m \leq \ell, \quad (7)$$

where ℓ and m are integers, $s = 1, 2$ indicates the shell we are considering, and with N_s the number of molecules within that shell (i.e., $N_1 \equiv 4$ for the first coordination shell, and $N_2 \equiv 12$ for the second). Y_ℓ^m is normalized according to $\int |Y_\ell^m|^2 \sin(\vartheta) d\varphi d\vartheta = 1$. We can consider $q_{\ell,m}^{(s)}(i)$ as a vector $\mathbf{q}_\ell^{(s)}(i)$ in a $(4\ell + 2)$ -dimensional Euclidean space having components $\text{Re}(q_{\ell,m}^{(s)}(i))$ and $\text{Im}(q_{\ell,m}^{(s)}(i))$. This means that we can define an inner product

$$\mathbf{q}_\ell^{(s)}(i) \cdot \mathbf{q}_\ell^{(s)}(j) \equiv \sum_{m=-\ell}^{\ell} [\text{Re}(q_{\ell,m}^{(s)}(i)) \text{Re}(q_{\ell,m}^{(s)}(j)) + \text{Im}(q_{\ell,m}^{(s)}(i)) \text{Im}(q_{\ell,m}^{(s)}(j))] \quad (8)$$

and a magnitude

$$q_\ell^{(s)}(i) \equiv \sqrt{\mathbf{q}_\ell^{(s)}(i) \cdot \mathbf{q}_\ell^{(s)}(i)}. \quad (9)$$

The local parameter $q_\ell^{(s)}(i)$ is one way to distinguish between different structures, and can be used to label individual molecules as LDL-like or HDL-like. We can convert it into a global parameter by averaging over all molecules,

$$q_\ell^{(s)} \equiv \frac{1}{N} \sum_{i=1}^N q_\ell^{(s)}(i), \quad (10)$$

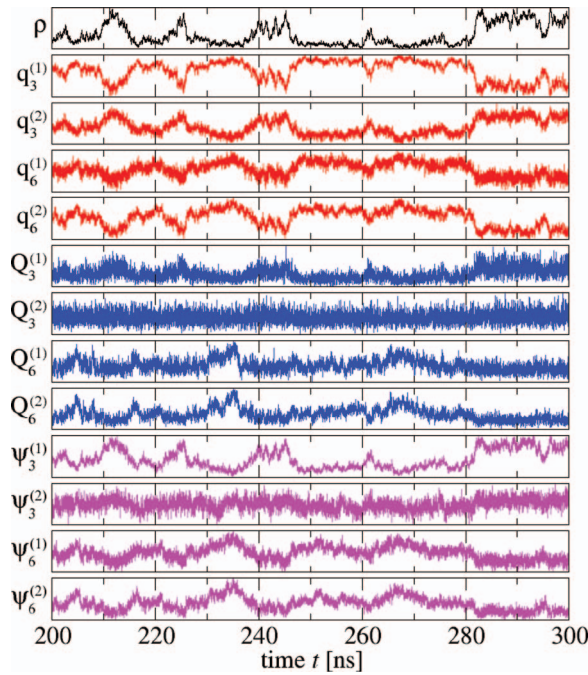


FIG. 8. Fluctuations of the density and the global structural parameters as a function of time. The parameters are shown for one run using 343 molecules at 200 MPa and 248 K, the same as in Fig. 5. Parameters $q_3^{(1)}$, $q_6^{(2)}$, and $\psi_3^{(1)}$ (defined in the text) are as sensitive as ρ to the difference between LDL-like and HDL-like structures, while the others are more noisy, being $Q_3^{(2)}$ and $\psi_3^{(2)}$ much less sensitive than all the others. $Q_6^{(s)}$ and $\psi_6^{(s)}$, for both $s = 1$ and 2, have similar behaviors that might be related to the temporary appearance of crystal-like structures.

where N is the total number of molecules. In Fig. 8, we see that all global $q_\ell^{(s)}$ are sensitive to the difference between LDL and HDL, especially $q_3^{(1)}$ and $q_6^{(2)}$. We conclude that the structural difference is visible in both the first and second shell, and that LDL and HDL differ mostly in the amount of diamond structure of the first shell and the amount of hcp structure in the second shell. This is confirmed by the histograms in Fig. 9, in which the largest difference between LDL and HDL is seen in $q_3^{(1)}$ and, next, in $q_6^{(2)}$. The latter is the parameter that better discriminates with respect to the crystal structure.

B. Global parameters Q_3 and Q_6

An alternative approach, as used by Steinhardt *et al.*,⁸³ is to first average $q_{\ell,m}^{(s)}(i)$ over all molecules, defining $Q_{\ell,m} \equiv \sum_{i=1}^N q_{\ell,m}^{(s)}(i)$, and then calculate the magnitude

$$Q_\ell^{(s)} \equiv \frac{1}{N} \left(\sum_{m=-\ell}^{\ell} Q_{\ell,m} Q_{\ell,m}^* \right)^{1/2}. \quad (11)$$

Our calculations show that the parameters $Q_3^{(s)}$ and $Q_6^{(s)}$, with $s = 1, 2$, are not efficient in discriminating between LDL and HDL (Fig. 8), although $Q_6 \equiv Q_6^{(1)}$ has been proposed recently as a good parameter to this goal⁶³ and consequently has been used by several authors.^{66–68} In particular, we observe that there is not much correlation between the fluctuations of $Q_\ell^{(s)}$ and those of the density, except for $Q_3^{(1)}$.

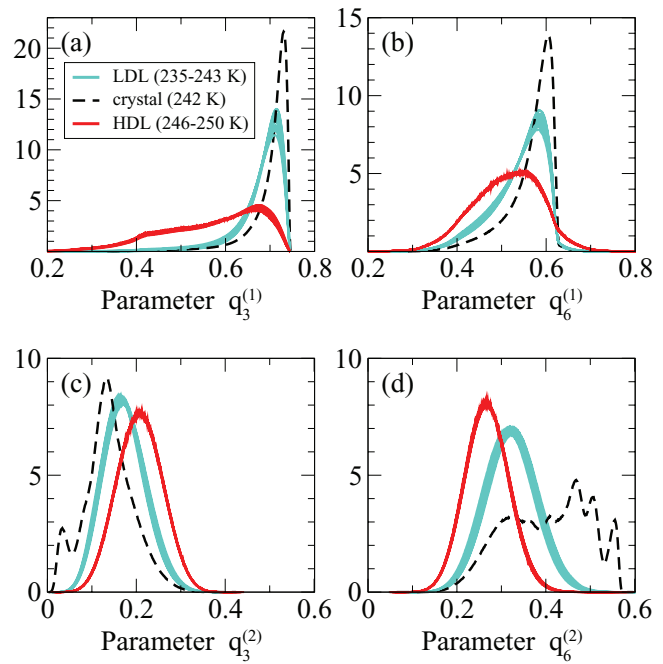


FIG. 9. Histograms of $q_\ell^{(s)}$ for $\ell = 3, 6$ and coordination shells $s = 1, 2$ at 215 MPa with $N = 343$ molecules. The solid red (dark) curves correspond to HDL structures, and the solid blue (light) curves to LDL structures. The dashed black curve corresponds to the crystal structure found in run C described in Sec. VI. The parameter $q_3^{(1)}$ (a) discriminates better between HDL and LDL structures, while the parameter $q_6^{(2)}$ (d) discriminates better between liquid-like and crystal-like structures. Parameters in (b) and (c) are much less sensitive to structural changes.

However, we confirm that $Q_6^{(1)}$ and $Q_6^{(2)}$ are excellent parameters to distinguish between the liquids (LDL and HDL) and the crystal, with the value of $Q_6^{(s)}$ approximately 10 times larger for the crystal than it is for the liquids (Fig. 10). This large increase of $Q_6^{(s)}$ for crystal-like structures might be related to the few instances in Fig. 8 where an increase in $Q_6^{(s)}$ corresponds to a decrease of density (such as within interval $t = 230$ – 237 ns), consistent with the observation that the crystal-like structures have a density comparable to the LDL structure and smaller than the HDL structure.

To confirm that LDL remains a liquid in the thermodynamic limit, we look at how Q_6 changes with the system size. For liquids, Q_6 scales like $N^{-1/2}$ while for crystals the value Q_6 remains finite as $N \rightarrow \infty$. We find that the probability distribution functions of $Q_6 N^{1/2}$ for $N = 216, 343, 512$, and 729 overlap, which means that $Q_6 N^{1/2}$ is independent of the system size, therefore, $Q_6 \sim N^{-1/2}$ (Fig. 11). We conclude that the metastable LDL is not transforming into the stable crystal in the thermodynamic limit. This implies that the LDL and the crystal phase are separated by a free-energy barrier that is higher than $k_B T$ at the temperatures we consider here, and that the system equilibrates to the stable (crystal) phase only on a time scale that is infinite with respect to our simulation time (1000 ns), as occur in experiments for metastable phases. Therefore, the LDL is a bona fide metastable state. Our conclusion is consistent with recent calculations by other authors.^{66–68}

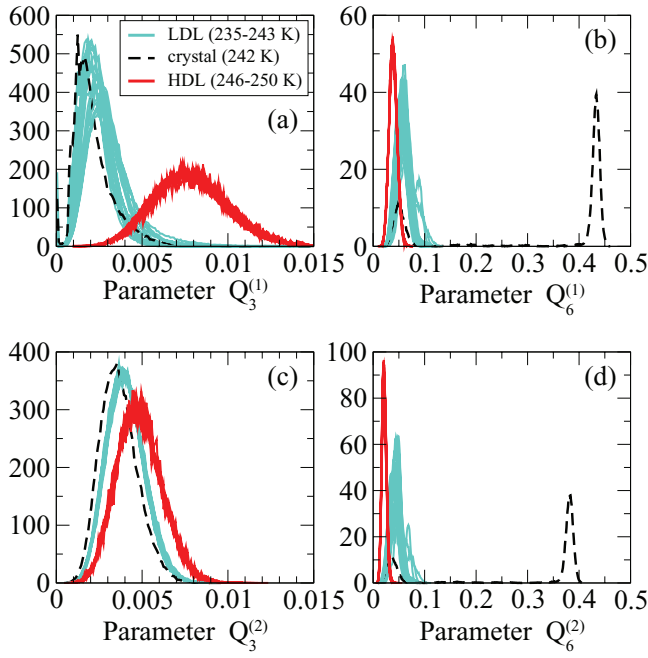


FIG. 10. Histograms of $Q_\ell^{(s)}$ for $\ell = 3, 6$ and coordination shells $s = 1, 2$ at 215 MPa with $N = 343$ molecules. The symbols are as in Fig. 9. The parameter $Q_6^{(s)}$, for the first shell in (b) and the second in (d), shows a clear difference between the liquid-like structures and the crystal-like structure, but not between the two liquids. Note that scales on x-axis in panels (a) and (c) are one order of magnitude smaller than those in panels (b) and (d). As a consequence, $Q_3^{(s)}$, for the first shell in (a) and the second in (c), is much less sensitive to structural changes than $Q_6^{(s)}$.

C. Bond parameters d_3 and ψ_3

We define the bond order parameter $d_\ell^{(s)}$ similar to that defined by Ghiringhelli *et al.* in Ref. 84, where the quantity $d_3^{(1)}(i, j)$ characterizes the bond between molecules i and j , and is designed to distinguish between a fluid and a diamond structure. The local parameter $d_\ell^{(s)}(i, j)$ is defined as the co-

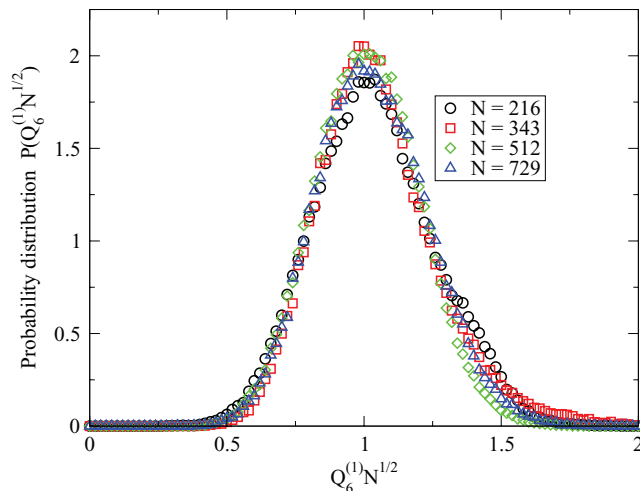


FIG. 11. Finite size scaling of parameter Q_6 in the LDL phase (210 MPa, 243 K). The probability distribution function of $Q_6 N^{1/2}$ is independent of the system size N , which means LDL scales like a liquid in the thermodynamic limit: $Q_6 \sim N^{-1/2}$.

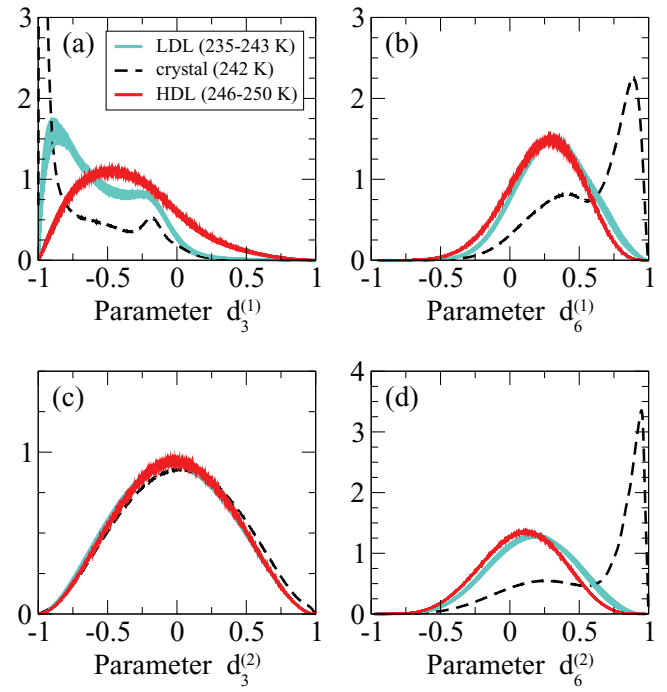


FIG. 12. Histograms of $d_\ell^{(s)}$ for $\ell = 3, 6$ and coordination shells $s = 1, 2$ at 215 MPa with $N = 343$ molecules. The symbols are as in Fig. 9. Apart from $d_3^{(1)}$ in (a), these parameters do not distinguish well between the two different liquid-like structures, but $d_3^{(1)}$ and $d_6^{(s)}$ for the first shell (b) and the second (d) are suitable to distinguish between the crystal and the liquids. The parameter $d_3^{(2)}$ in (c) is remarkably the same for the three structures.

sine of the angle between the vectors $\mathbf{q}_\ell^{(s)}(i)$ and $\mathbf{q}_\ell^{(s)}(j)$

$$d_\ell^{(s)}(i, j) \equiv \frac{\mathbf{q}_\ell^{(s)}(i) \cdot \mathbf{q}_\ell^{(s)}(j)}{|\mathbf{q}_\ell^{(s)}(i)| |\mathbf{q}_\ell^{(s)}(j)|} \quad (12)$$

with the inner product and magnitude as defined in Eqs. (8) and (10).

A crystal with a perfect diamond structure has $d_3^{(1)}(i, j) = -1$ for all bonds. For a graphite crystal, only the bonds within the same layer (three out of four) have $d_3^{(1)}(i, j) = -1$, while the bonds connecting atoms in different layers (one out of four) have $d_3^{(1)}(i, j) = -1/9$.

We find that the parameters $d_\ell^{(s)}$ for $\ell = 3, 6$ and $s = 1, 2$ do not distinguish well between the two different liquid-like structures, but that $d_3^{(1)}$ and $d_6^{(s)}$ for both $s = 1$ and 2 are suitable to discriminate between the crystal-like structure and the liquids (Fig. 12). In particular, for the crystal, most molecules have $d_3^{(1)} < -0.87$, and we therefore consider a molecule to be part of a crystal if at least three out of its four bonds with its nearest neighbors have $d_3^{(1)} < -0.87$. This is the same cutoff used by Ghiringhelli *et al.* in Ref. 84.

The global parameter associated with $d_\ell^{(s)}(i, j)$ is defined as

$$\psi_\ell^{(s)} \equiv \frac{1}{N} \sum_{i=1}^N \psi_\ell^{(s)}(i), \quad (13)$$

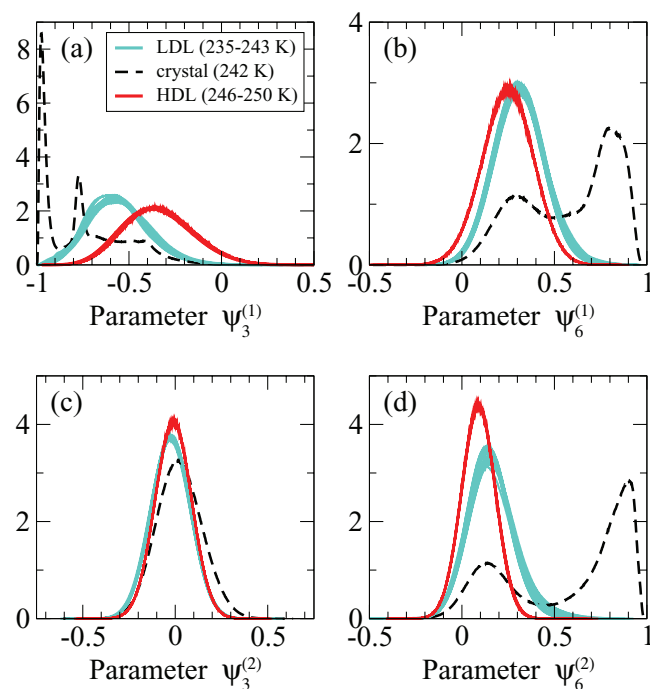


FIG. 13. (a)–(d) Histograms of $\psi_\ell^{(s)}$ for $\ell = 3, 6$ and coordination shells $s = 1, 2$ at 215 MPa with $N = 343$ molecules. The symbols are as in Fig. 9. Each $\psi_\ell^{(s)}(i)$ has similar features as the corresponding $d_\ell^{(s)}(i, j)$ in Fig. 12.

where

$$\psi_\ell^{(s)}(i) \equiv \frac{1}{4} \sum_{j=1}^4 d_\ell^{(s)}(i, j) \quad (14)$$

is the average of $d_\ell^{(s)}(i, j)$ over the first four nearest neighbors of the molecule i . We observe that each $\psi_\ell^{(s)}(i)$ has the same features of the corresponding $d_\ell^{(s)}(i, j)$, with $\psi_3^{(1)}(i)$ discriminating well between the crystal-like and the liquids-like structures (Fig. 13). We observe that $\psi_3^{(1)}$ discriminates well between LDL-like and HDL-like structures (Fig. 8), while $\psi_6^{(s)}$ for $s = 1$ and 2 might be able to emphasize the temporary appearance of crystal-like structures, as noted for $Q_6^{(s)}$.

In Fig. 14, we see that there is no significant difference between the low- T curves at $P = 240$ MPa, 215 MPa (Fig. 13(a)), and 195 MPa, and no significant difference between the high- T curves. We therefore conclude that at the Widom line the low density liquid (at low T) is LDL-like, and similarly that the high density liquid (at high T) is HDL-like.

From the figure, we can also see that the difference between the LDL- and HDL-structure decreases when we decrease P , indicating that far from the LLPT the distinction between LDL-like and HDL-like disappears as is to be expected in the one-phase region.

VI. GROWTH AND MELTING OF CRYSTAL NUCLEI

In a small percentage of our simulations, the system was found to spontaneously crystallize. These are interesting events because spontaneous crystallization of water in molecular dynamics is extremely rare; only recently Matsumoto *et al.*⁸⁵ were the first to successfully simulate the

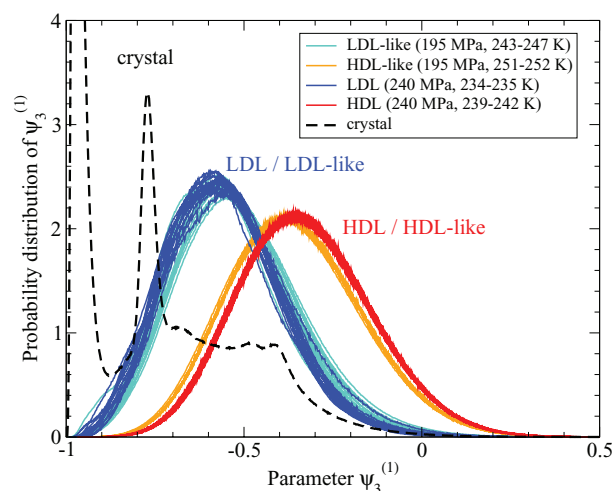


FIG. 14. Probability distribution function of $\psi_3^{(1)}$ for both $P = 195$ MPa and 240 MPa, with $N = 343$ molecules (see Fig. 13(a) for $P = 215$ MPa). At the Widom line (195 MPa), the structure of the low-density liquid is similar to that of LDL at 240 MPa, and the structure of the high-density liquid near the Widom line is practically the same as HDL. This demonstrates that the LDL-like and HDL-like phases are indeed structurally similar to LDL and HDL. Furthermore, the structural difference between the LDL-like and HDL-like phases becomes smaller as we move away from the phase transition line to lower and lower pressures.

freezing of water on a computer. Crystallization events in supercooled ST2 water are particularly important to study, as it has been proposed that LDL is unstable with respect to crystallization.⁶³

Following the discussion in Sec. V, we define a crystal as a cluster of molecules which has three out of four bonds with $d_3^{(1)} < -0.87$ and belong to the first coordination shell of each other. In this section, we shall study the growth and melting of these crystal nuclei, and estimate the critical nucleus size needed to overcome the free energy barrier. The existence of this barrier allows us to conclude that LDL is in fact a bona fide metastable state with respect to the crystal.

In Fig. 15, we show the density evolution for 11 different configurations, each with 343 molecules and at 205 MPa and 246 K. Each of these runs started at a different initial density (between 0.85 and 0.95 g/cm³) and was subsequently equilibrated to the final temperature and pressure using the procedure described in Sec. II. Because this state point lies close to the LLPT, we see phase flipping in all of them. However, the two configurations C and F display a sudden jump to a stable low density plateau. This is a hallmark of crystallization. We confirm this by calculating the size of the largest crystal as a function of time (Fig. 16). During most runs the largest crystal continuously grows and shrinks, but never reaches a size larger than 30 molecules. On the other hand, configurations C and F show a jump in crystal size exactly matching the jump in density. Run F ends up partially crystallized, while for C we find that over 90% of the box is crystallized in a diamond structure with a density of about 0.92 g/cm³ (Fig. 17).

The correlation time increases dramatically if crystals appear with a size comparable to the system size, as is evident from Fig. 18. The correlation functions of C and F decay very

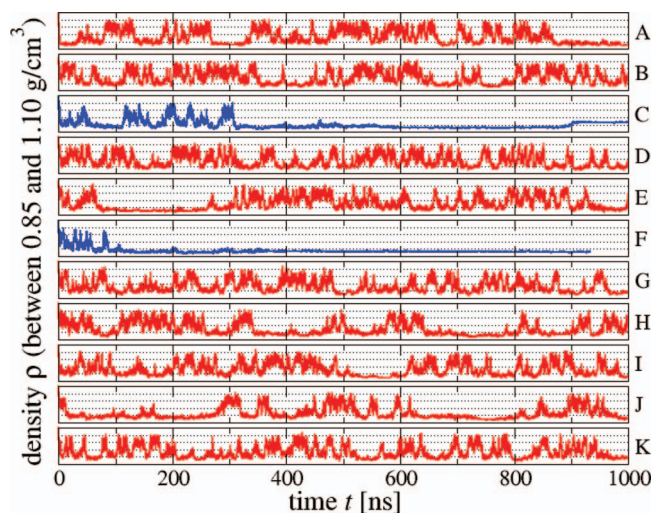


FIG. 15. Density vs. time near the phase transition line at $P = 205$ MPa and $T = 246$ K for several different configurations of $N = 343$ molecules. This state point lies near the phase transition, and therefore phase flipping is seen to occur. Runs C and F (partially) crystallize and, at that moment, cease to phase flip and remain stable at a low density.

slowly, leading to correlation times of 200–400 ns, while the other configurations have a correlation time of less than 4 ns.

For spontaneous crystallization to occur, a sufficiently large crystal nucleus needs to form within the liquid. According to classical nucleation theory, this nucleus needs to reach a minimum size to prevent it from melting. We observed in many simulations that a small nucleus grows and melts, and a few runs in which the nucleus grows further or remains stable. Therefore, we can make an estimate of the critical nucleus size.

The two largest crystals that formed and subsequently melted, both reached a size of about 50–60 molecules (Figs. 19(a) and 19(b)). The smallest crystal that formed and remained stable, had a size of about 50–80 molecules

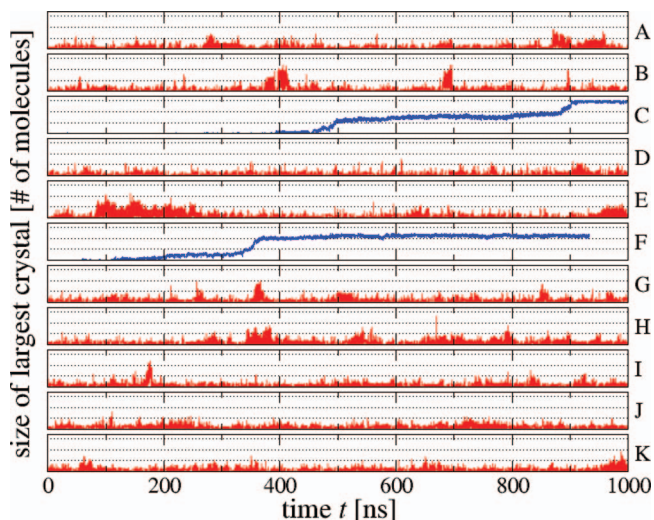


FIG. 16. Evolution of crystal size with time for the same configurations as in Fig. 15. The y-axis goes from 0 to 34, except for configurations C and F which go up to 343. The system spontaneously crystallizes in both C and F, while the largest crystals in the remaining configurations never reach a size larger than 30 molecules.

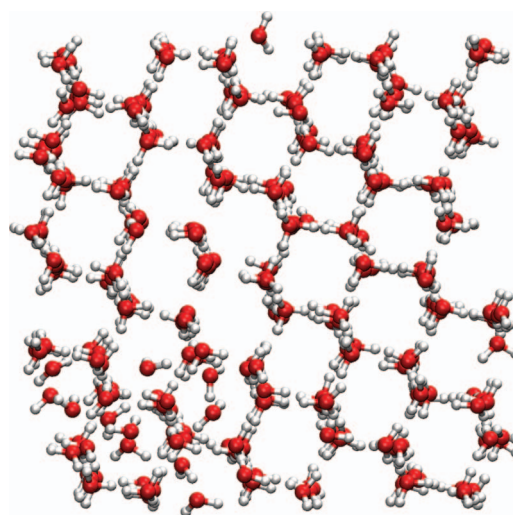


FIG. 17. A snapshot (at $t = 1000$ ns) of the diamond cubic crystal produced by run C of Figs. 15 and 16. Shown here are all $N = 343$ molecules, with a small part still in the liquid state (bottom-left corner), and a crystal defect in the center. Note that the defect only affects the position of the hydrogen atoms, and not that of the oxygen.

(Fig. 19(c)). We therefore conclude that the critical nucleus size is approximately 70 ± 10 molecules. A similar value of $\simeq 85$ molecules was found by Reinhardt and Doye⁸⁶ for ice nucleation in the monatomic water model.⁸⁷

For a more accurate estimate, it is necessary to run longer simulations, as the crystal nuclei can survive for hundreds of nanoseconds (e.g., Fig. 19(d) in which a small crystal lasts for 700 ns).

VII. LOCATING THE CRITICAL POINT

In Sec. III, we used the intermediate scattering function $S_{OO}(k)$ to estimate the position of the liquid-liquid critical point, and found it to lie near 200–210 MPa and 244–247 K. It is commonly believed that the LLCP falls in the same

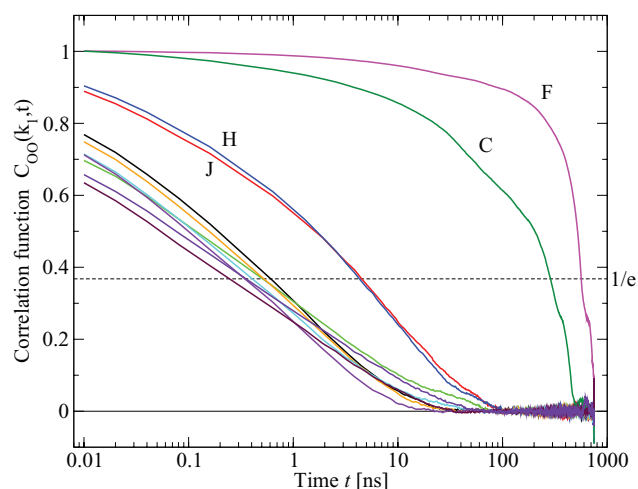


FIG. 18. The correlation time increases dramatically if crystals of a size comparable to the system size appear (i.e., runs C and F of Figs. 15 and 16). The correlation time of two other runs (H and J) are slightly larger than average because these runs spend more time in the LDL phase (see Fig. 15).

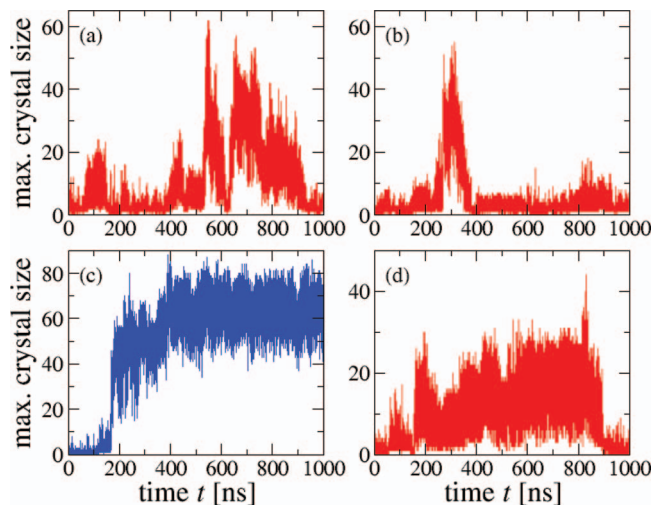


FIG. 19. Growth and melting of crystal nuclei. (a) The largest nucleus that melted reached a size of 62 molecules during a simulation of 512 molecules at 210 MPa and 244 K. (b) The second-largest nucleus was 55 molecules during a simulation of 343 molecules at 210 MPa and 243 K. (c) A few runs lead to irreversible crystallization ($N = 216$ at 195 MPa and 245 K). (d) Some crystal nuclei survive for hundreds of nanoseconds ($N = 343$ at 195 MPa and 246 K) before disappearing.

universality class as the three-dimensional Ising model.⁵⁷ At the critical point, the order parameter distribution function (OPDF) of a system has the same bimodal shape as all other systems that belong to the same universality class. Therefore, one can locate the LLCP accurately by fitting our data to the OPDF of the 3D Ising model, as was done in Ref. 65. Based on our fit, we locate the LLCP to be at $P_C = 206 \pm 3$ MPa and $T_C = 246 \pm 1$ K.

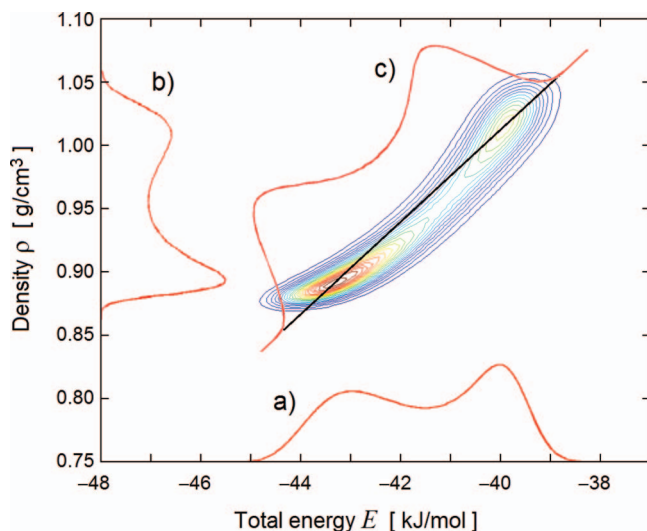


FIG. 20. 2D histogram (contour plot) of the density and the total energy for a system at 247.5 K and 200 MPa (on the Widom line), obtained via histogram reweighting, together with its projection in arbitrary units along the horizontal axis (curve (a)), the vertical axis (curve (b)) and the diagonal black-line (curve (c)). The histogram of the energy (curve (a)) seems to indicate that the system is mostly in the LDL state, while the histogram of the density (curve (b)) indicates the HDL state is more predominant. For liquids, the order parameter $M \equiv \rho + sE$ is actually a linear combination of the density ρ and the energy E (curve (c)), with $s = 0.0362$ (g/cm³)/(kJ/mol). By fitting the OPDF (curve (c)) to the critical OPDF of the 3D Ising model, it is possible to accurately locate the critical point.⁶⁵

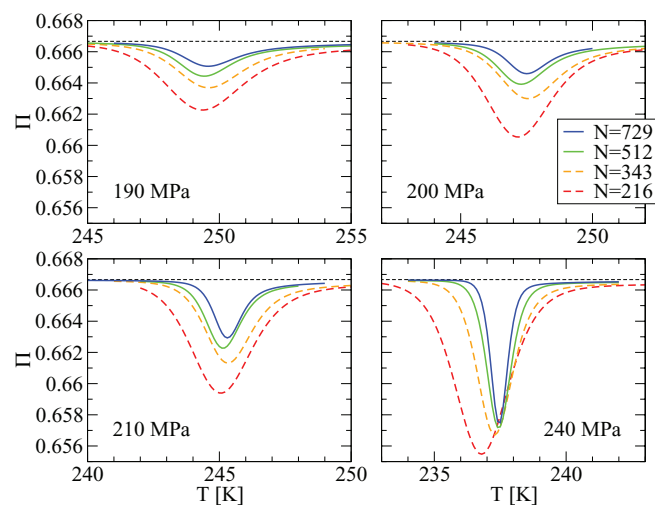


FIG. 21. The Challa-Landau-Binder parameter Π as a function of temperature and system size N , for four different pressures. For finite system, size Π shows a minimum at the LLPT and the Widom line, while $\Pi \approx 2/3$ (thin dashed line) at temperatures where $\mathcal{D}(\rho)$ is given by a single Gaussian. The finite-size scaling of the minimum of Π , indicates that the critical point exists in the thermodynamic limit (Fig. 22).

To establish that the LLPT does not vanish in the thermodynamic limit $N \rightarrow \infty$, we consider the finite-size scaling of the Challa-Landau-Binder parameter.^{88–93} Near the critical point the density distribution function $\mathcal{D}(\rho)$ has a bimodal shape that can be approximated by the superposition of two Gaussians. The Challa-Landau-Binder parameter Π is a measure of the bimodality of $\mathcal{D}(\rho)$ and is defined as

$$\Pi \equiv 1 - \frac{\langle \rho^4 \rangle}{3\langle \rho^2 \rangle^2}. \quad (15)$$

When there is only one phase, $\mathcal{D}(\rho)$ is unimodal and $\Pi = 2/3$. But in a two-phase region, with two phases that have different densities, the shape of $\mathcal{D}(\rho)$ is bimodal (Fig. 20) and $\Pi < 2/3$. For a finite system, $\mathcal{D}(\rho)$ is always bimodal at both the Widom line and the LLPT, but in the thermodynamic limit there exists only one phase at the Widom line, while there remain two at the phase transition line. Therefore, $\Pi \rightarrow 2/3$ at the Widom line, while $\Pi < 2/3$ at the LLPT even in the limit $N \rightarrow \infty$. Hence, the finite-size scaling of Π allows us to distinguish whether an isobar crosses the LLPT or the Widom line, and is yet another method of estimating the location of the critical point.

We study Π versus temperature T and system size N for different pressures, finding minima Π_{\min} at a specific temperature for each pressure (Fig. 21). The finite-size dependence of $\Pi_{\min}(P)$ reveals if $P < P_C$ or $P > P_C$ (Fig. 22).

For $P < P_C$, the minimum Π_{\min} approaches $2/3$ linearly with $1/N$, while for $P \leq P_C$ it approaches the limit⁸⁸

$$\Pi_{\min} \rightarrow \frac{2}{3} - \frac{1}{3} \frac{(\rho_H^2 - \rho_L^2)^2}{(\rho_H^2 + \rho_L^2)^2}. \quad (16)$$

This limiting value is also approached linearly with $1/N$. Here, $\rho_H \equiv \rho_H(P)$ and $\rho_L \equiv \rho_L(P)$ are the densities of the two phases LDL and HDL.⁹⁰ Above the critical pressure, the limiting value of Π_{\min} decreases as P increases, i.e., the two peaks

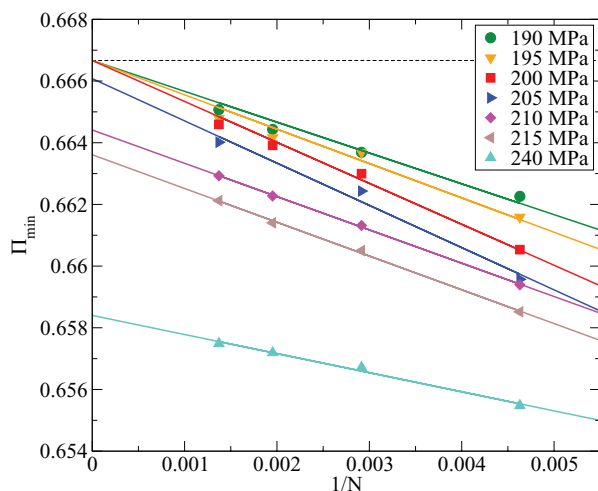


FIG. 22. Minima of the Challa-Landau-Binder parameter Π as a function of system size N for different pressures. The minimum Π_{\min} occurs at the pressures and temperatures of the LLPT and the Widom line, and is always less than $2/3$ for a finite system because of the bimodality of the density histogram. As $N \rightarrow \infty$, the bimodality disappears in the one-phase region but remains at the LLPT, and therefore $\Pi_{\min} \rightarrow 2/3$ at the Widom line while $\Pi_{\min} < 2/3$ on the LLPT, even in the thermodynamic limit. We conclude that the critical point survives in the thermodynamic limit, and that it is located between $P = 200$ and 210 MPa (in agreement with previous results of this paper).

of the bimodal $\mathcal{D}(\rho)$ move further apart. This happens because $\rho_H - \rho_L$ increases at coexistence as $(P - P_C)^\beta$ where $\beta \approx 0.3$ is the critical exponent of the 3D Ising universality class.^{94,95}

From this analysis (Fig. 22), we conclude that our results agree with theory and that the critical pressure $P_C \approx 190$ – 210 MPa, in agreement with the estimate of Sec. III. Furthermore, as Π remains less than $2/3$ for $P > P_C$ even in the limit $N \rightarrow \infty$, we conclude that the LLPT does not vanish in the thermodynamic limit.

VIII. CONCLUSIONS

We performed molecular dynamic simulations in the NPT ensemble for ST2-RF water in the supercooled region of the phase diagram for different system sizes with simulation times of up to 1000 ns. Using several different techniques, we confirmed the existence of two liquid phases, LDL and HDL, separated by a liquid-liquid phase transition line. Near the LLPT line, the system continuously flips between the two phases. Because of finite-size effects this phenomenon also occurs near the Widom line, but by fitting the order parameter distribution function to that of the 3D Ising model we were able to accurately determine the location of the liquid-liquid critical point (at $T_C = 246 \pm 1$ K, $P_C = 208 \pm 3$ MPa). These results agree exactly with ($T_C = 246$ K, $P_C \approx 188$ MPa) as found by Poole *et al.*,⁵⁶ with the discrepancy in pressure attributed to the LJ pressure correction of $\simeq -20$ MPa (see Sec. II). Finite size scaling of the Challa-Landau-Binder parameter indicates that the critical point does not disappear in the thermodynamic limit.

Both phases have been confirmed to be bona fide metastable liquids that differ substantially in structural as well as dynamical properties. It is found that the LDL phase is a

more “structured” liquid, and that it has a correlation time of almost four orders of magnitude larger than that of HDL, with LDL correlation time of the order of 100–1000 ns. We show that Q_6 structural parameter is not able to discriminate between HDL and LDL, but can discriminate well between liquids and crystal. Finite size scaling of the Q_6 parameter confirms that LDL scales as a liquid and not as a crystal.

The different structures of LDL and HDL are better discriminated by structural parameters such as $q_3^{(1)}$ and $q_6^{(2)}$. These parameters show that LDL and HDL differ mostly in the amount of diamond structure of the first shell and the amount of hcp structure in the second shell.

For small box sizes ($N = 343$), there were a few simulation runs that resulted in spontaneous crystallization, always within the LDL region of the phase diagram. Further analysis revealed that during all simulations small crystals grow and melt within the liquid, a clear indication that LDL is metastable with respect to the crystal. From the few crystallization events that occurred, we were able to conclude that the critical nucleus size is approximately 70 ± 10 molecules.

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