

Supporting Material of: Anomalous behavior in the nucleation of ice at negative pressures

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DIRECT COEXISTENCE SIMULATIONS

To determine the solid-liquid equilibrium line we place a block of ice Ih consisting of 4032 molecules in contact with an equal number of liquid molecules. We perform *NPT* simulations, with anisotropic P to relax the crystal structure, counting the number of solid molecules in time. Along an isobar, if T is larger than the coexistence temperature the ice melts, hence the number of solid particles decreases in time. On the other hand, for T below the coexistence temperature the number of solid molecules increases in time. The coexistence T is determined when, for independent trajectories, the ice shows equal probability of melting or growing.

SEEDING SIMULATIONS

The purpose of this technique is to find the critical nucleus size at a certain temperature T and pressure P . Given that the critical nucleus is situated at the top of the energy barrier, it is in unstable equilibrium and the probability for the nucleus to grow or to shrink is equally 50%. Therefore, by inserting a crystal seed of a given size N within the bulk liquid and tracking its evolution with time over multiple independent runs, we can estimate if N is larger than the critical size N_c (meaning that all the crystal seeds grow) or if it is smaller (in which case all the seeds melt). When $N \sim N_c$ we expect to observe the ice growing for half of the trajectories, and melting for the remaining ones.

Introduced in 2005 [1], the *seeding* technique's popularity has grown over the last few years being extensively applied in nucleation studies [2–16]. The *seeding* scheme can be used in two distinct ways: tracking several seeds differing in size at a given (T, P) or studying a seed with a fixed size at different (T, P) . We found that the first protocol is more suitable when the critical size does not change significantly with the thermodynamic state point (i.e. $\partial N_c / \partial P \sim 0$ or $\partial N_c / \partial T \sim 0$) and we adopted it to identify N_c at most of the negative pressures. On the other hand, the second protocol (fixed seed simulated at different (T, P)) works well when the critical size is sensitive to changes of T and/or P , as observed at positive pressures, and has been adopted in this regime.

To create the initial configuration for a single *seeding* run we first equilibrate separately bulk liquid water and bulk ice at the T and P of interest. Then, we create a spherical cavity of radius R within the bulk liquid water and insert in the empty space a spherical seed of a slightly smaller radius extracted from the bulk ice. The system is quickly equilibrated at constant volume for ~ 10 ps keeping the ice-molecules motionless, which allows us to equilibrate the interface. Finally, we simulate the system in the *NPT* ensemble, where the pressure is applied isotropically.

In the following we show the estimation of N_c according to our *seeding* simulations.

MISLABELING CALCULATION

In Fig. 1 we report the % of mislabeled solid and liquid molecules, according to the threshold $\bar{q}_{6,t}$ of the \bar{q}_6 order parameter. The value of $\bar{q}_{6,t}$ is determined from the crossing of the mislabeling curves of ice and liquid water in the bulk. For any P the optimal value of $\bar{q}_{6,t}$ is well fitted by the expression $\bar{q}_{6,t}(P) = a_{q_6}(T) - b_{q_6}(T) \exp(c_{q_6}(T)P)/[d_{q_6}(T) + \exp(c_{q_6}(T)P)]$ (see Fig. 2 of SM). The fitting parameters are reported in Tab. II.

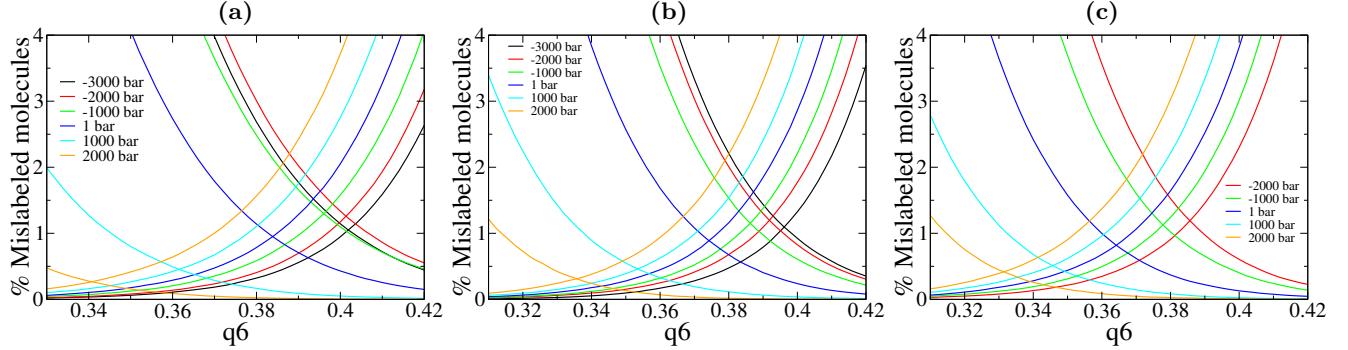


FIG. 1. Percentage of mislabeled molecules for each phase at $T = 230$ K (a), $T = 240$ K (b) and $T = 250$ K (c). Data are over equilibrium simulations of 3820 molecules in the liquid phase, and 4032 molecules in the solid Ih phase.

T [K]	$a_{q_6}(T)$	$b_{q_6}(T)$	$c_{q_6}(T)$	$d_{q_6}(T)$
230	0.401839	0.0765991	0.00127229	3.29737
240	0.396191	0.0772493	0.00106575	2.35767
250	0.392375	0.0776309	0.000938774	1.7289

TABLE II. Fitting parameters for the $\bar{q}_{6,t}$.

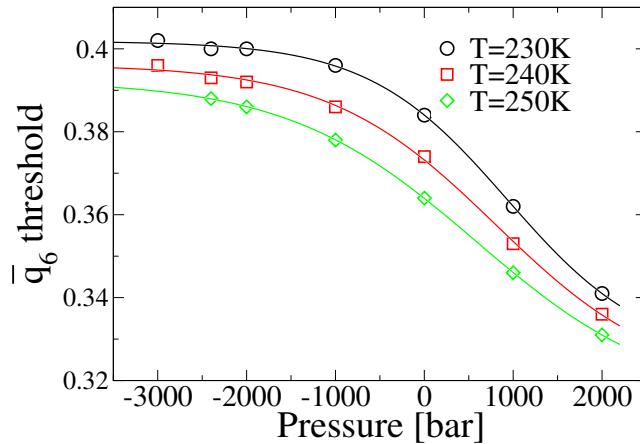


FIG. 2. Threshold $\bar{q}_{6,t}$ for the \bar{q}_6 order parameter as function of P , for different temperatures. Curves are fitted according to the expression reported in the SM.

ATTACHMENT RATE f^+

The attachment rate f^+ can be rigoursly computed from the expression proposed by Auer and Frenkel[17]

$$f_{AF}^+ = \frac{\langle (N(t) - N(0))^2 \rangle}{2t} \quad (2)$$

and approximated by the equation

$$f^+ = \frac{24DN_c^{2/3}}{\lambda^2} \quad (3)$$

where D is the diffusion coefficient of the supercooled liquid, N_c the size of the critical nucleus, and $\lambda \equiv 3.8\text{\AA}$ is a characteristic length of the order of a molecular diameter [4]. In a previous work, we have applied this approximation in nucleation of ice at positive pressures finding good agreement with the rigourous procedure [4]. Here, we test it at negative pressures, also finding good agreement as can be seen in Table VI. In particular, along the 240 K isotherm, we calculate f^+ for three different negative pressures: -300 bar, -1700 bar, and -2900 bar.

P (bar)	f_{AF}^+ (ns^{-1})	f^+ (ns^{-1})
-300	129.5	111.4
-1700	22.1	16.6
-2900	15.0	13.9

TABLE VI. Attachment rate computed according the rigourous Eq. 2 and approximated by the Eq. 3.

