

Steady-state simulation of homogeneous vapor-liquid nucleation “by the intervention of intelligent beings“

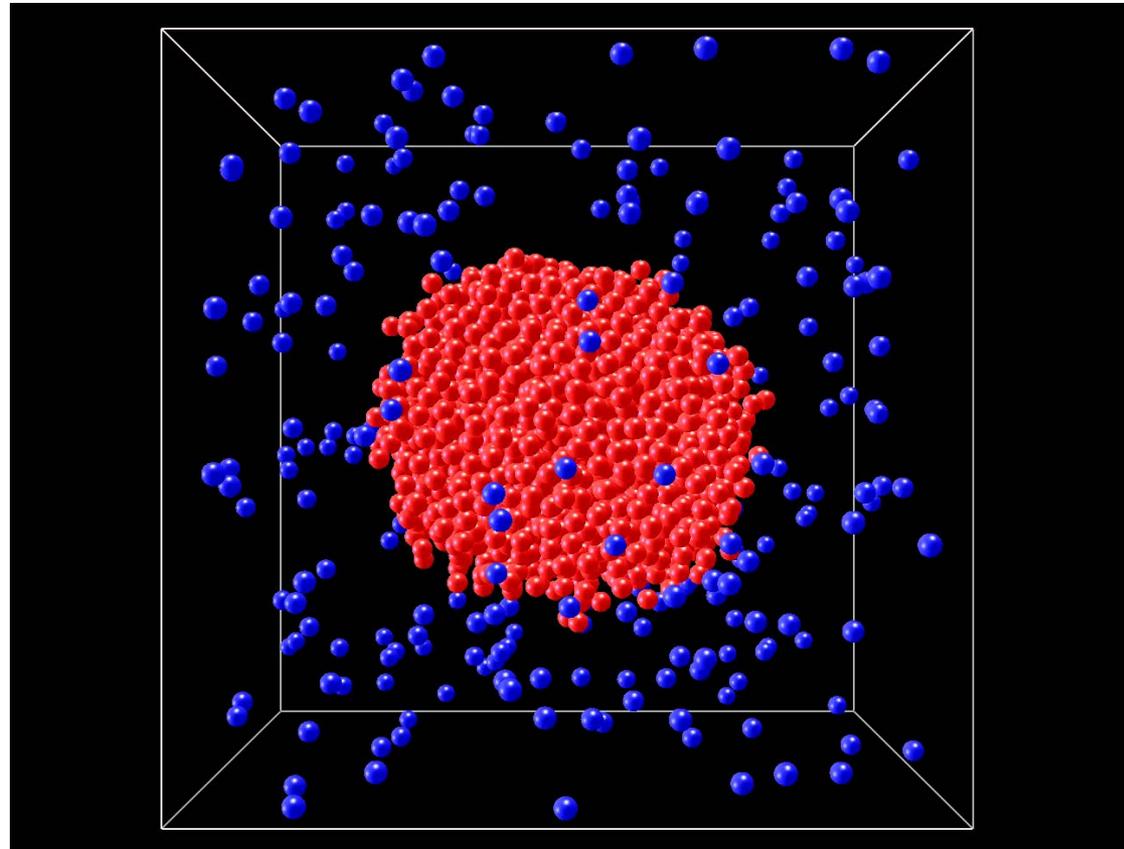
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International Conference on Nucleation & Atmospheric Aerosols

Prague, August 14, 2009

MD simulation of a single droplet

- Vapor and liquid are equilibrated separately
- A small ($j < 10000$) droplet is inserted into the vapor
- If the droplet cannot evaporate completely, an equilibrium is established within a few nanoseconds



t-s-LJ fluid ($r_c = 2.5 \sigma$)

Single droplet in non-equilibrium

t-s-LJ fluid

Vapor properties:

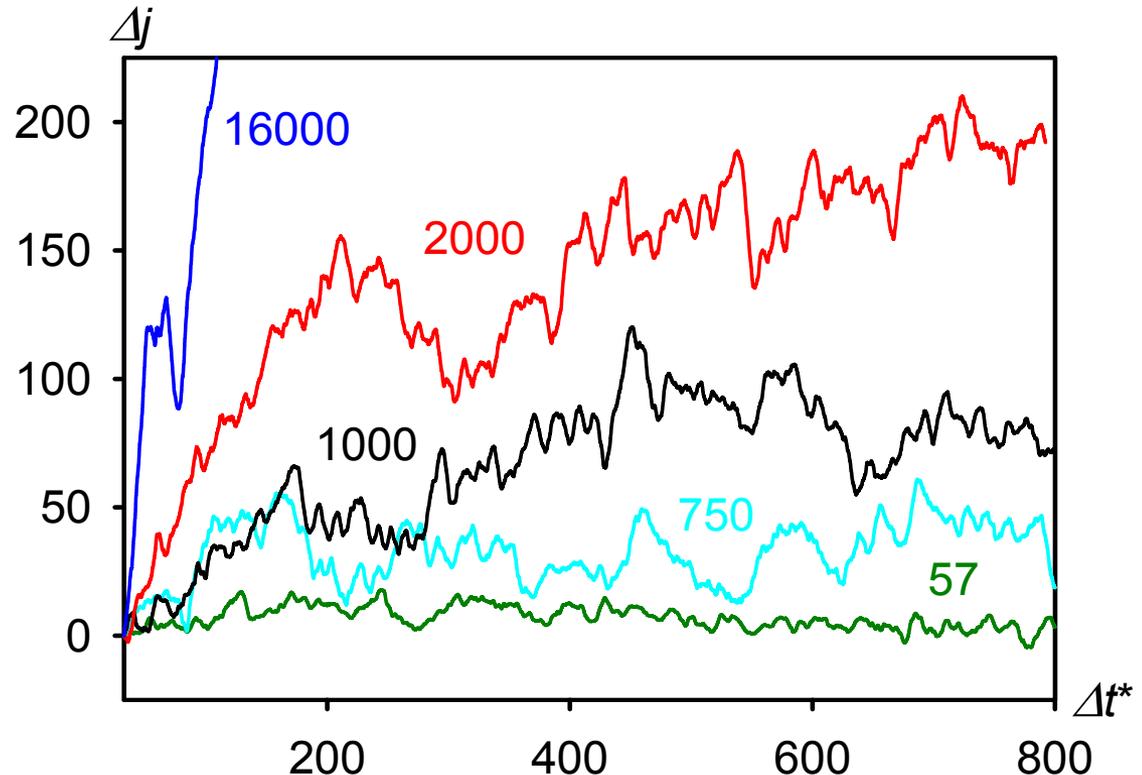
$$N = 130,000$$

$$\rho = 0.0268 / \sigma^3$$

$$T = 0.80 \epsilon/k$$

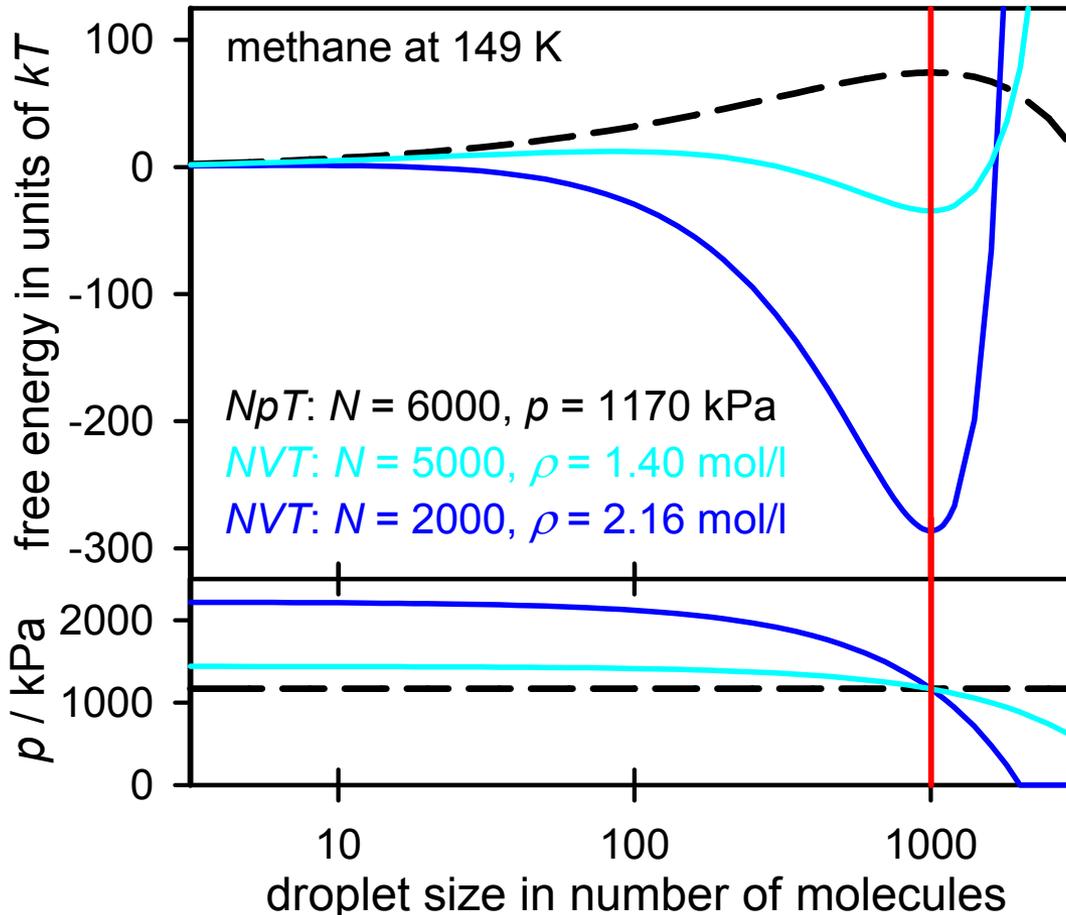
Time interval ≈ 1 ns

Classical theory: $j^* = 850$



The critical nucleus size cannot be clearly detected by non-equilibrium simulation

Equilibrium vapor pressure



Equilibrium condition for a droplet containing j molecules:

$$p = p(T, j)$$

ΔG at constant p and T :

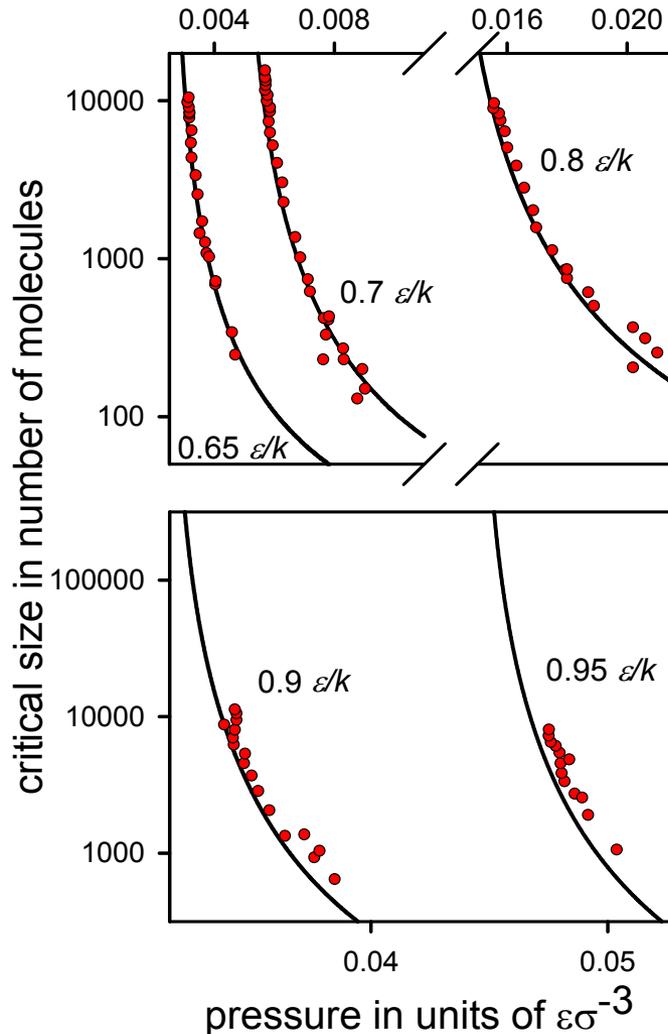
1 unstable equilibrium

ΔF at constant V and T :

1 unstable equilibrium

1 stable equilibrium

Critical nucleus from equilibrium MD simulation



Kelvin equation:

$$j^* = \left(\frac{2\gamma A_1}{3\Delta\mu} \right)^3$$

- present simulation data
- classical theory

- Good prediction of the critical size for low temperatures
- Above $0.8 T_c$, significant deviations are present

Nucleation from MD in the *NVT* ensemble

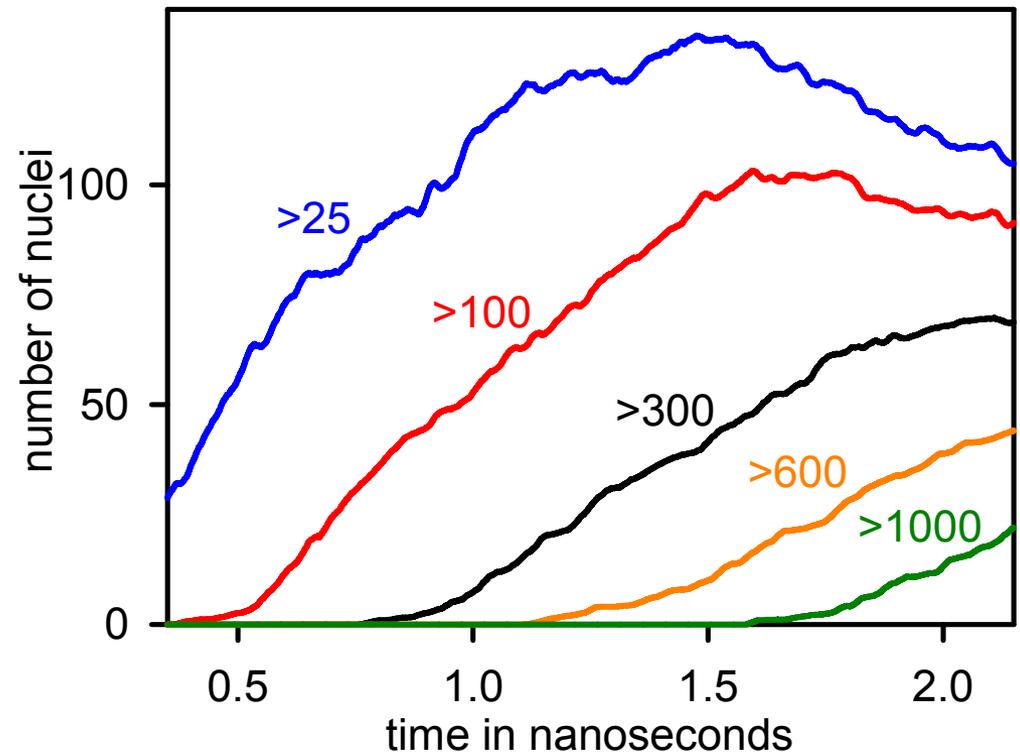
Yasuoka and Matsumoto (1998):

Number of emerging nuclei with $>j$ molecules per volume and time

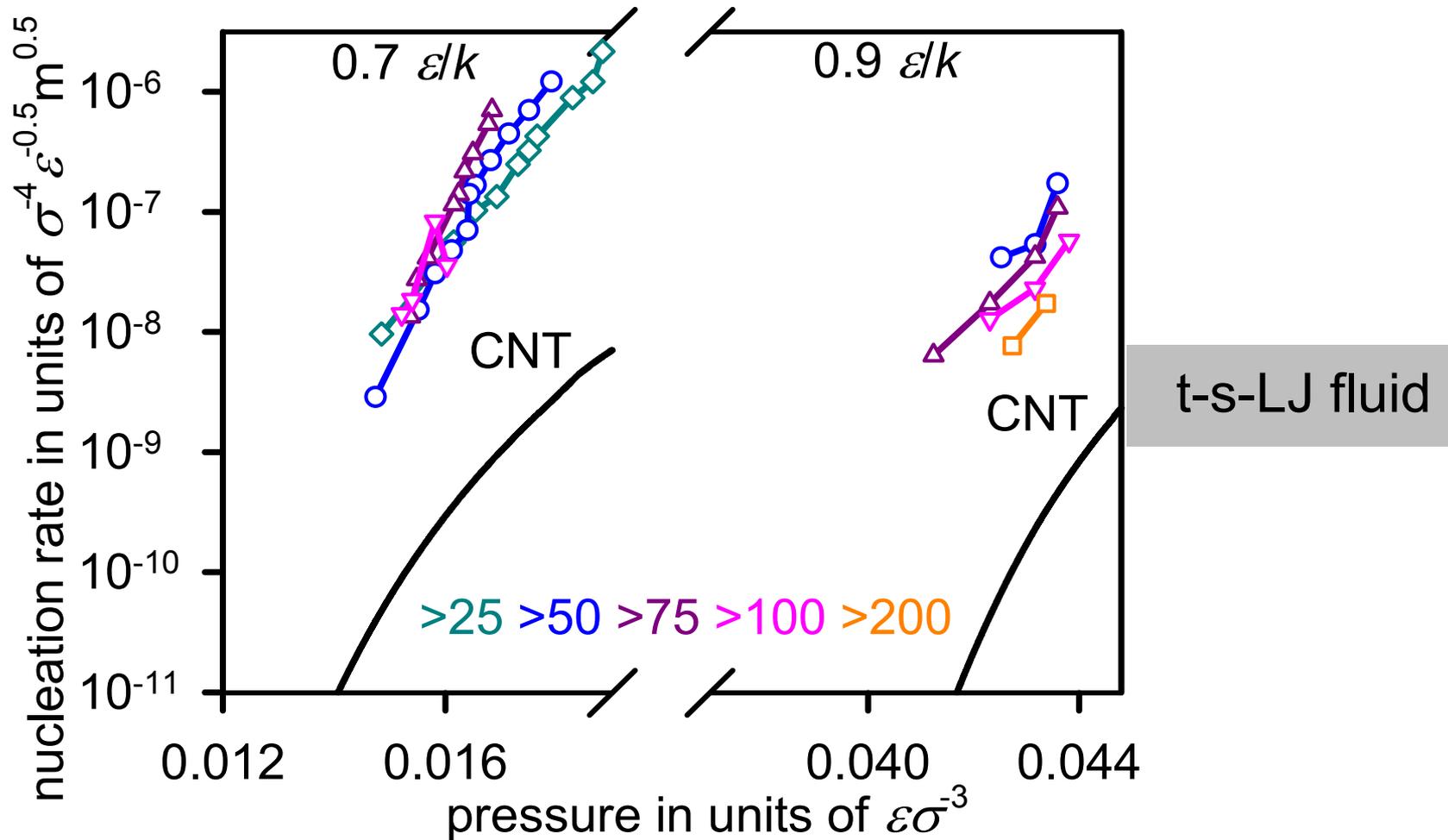
Typical approach:

determine the nucleation rate for various j values

250,000 methane molecules
at 130 K and 1.606 mol/l
(full LJ potential)



Nucleation rate following Yasuoka & Matsumoto



Grand canonical MD

Algorithm: MC insertion/deletion steps alternating with MD steps

- fixed values of μ , V und T
- test insertion of a molecule at a random position

$$P_{\text{ins}} = \min \left[1, \exp \left(\frac{\mu - \Delta U_{\text{pot}}}{kT} \right) \frac{V}{\Lambda^3 (N+1)} \right]$$

- test deletion of a random molecule

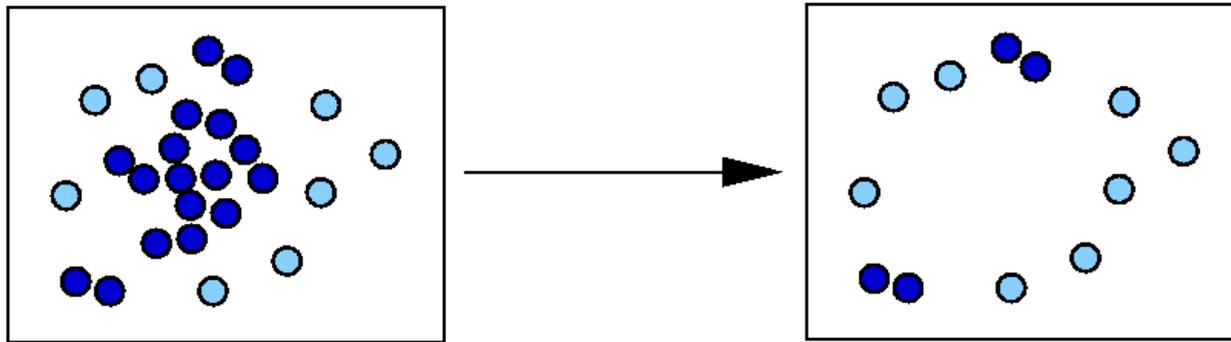
$$P_{\text{del}} = \min \left[1, \exp \left(\frac{-\mu - \Delta U_{\text{pot}}}{kT} \right) \frac{V}{\Lambda^3 N} \right]$$

- equal number of insertions and deletions ($10^{-6}N$ to $10^{-3}N$ / step)



Thermodynamic conditions of supersaturated state are maintained

McDonald's demon



Intervention rate as a function of the threshold size

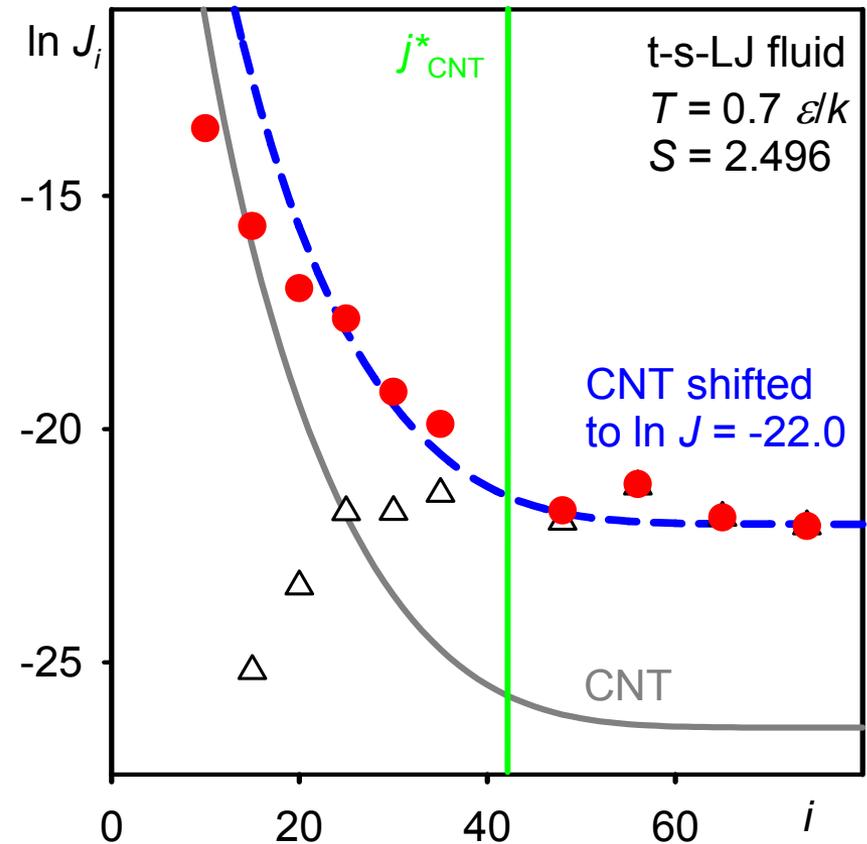
$$J_i = J/q_i$$

Probability for a nucleus of growing from size i to infinite size:

$$q_i = \frac{\int_1^i \exp(2\beta\Omega) dj}{\int_1^\infty \exp(2\beta\Omega) dj},$$

and in particular:

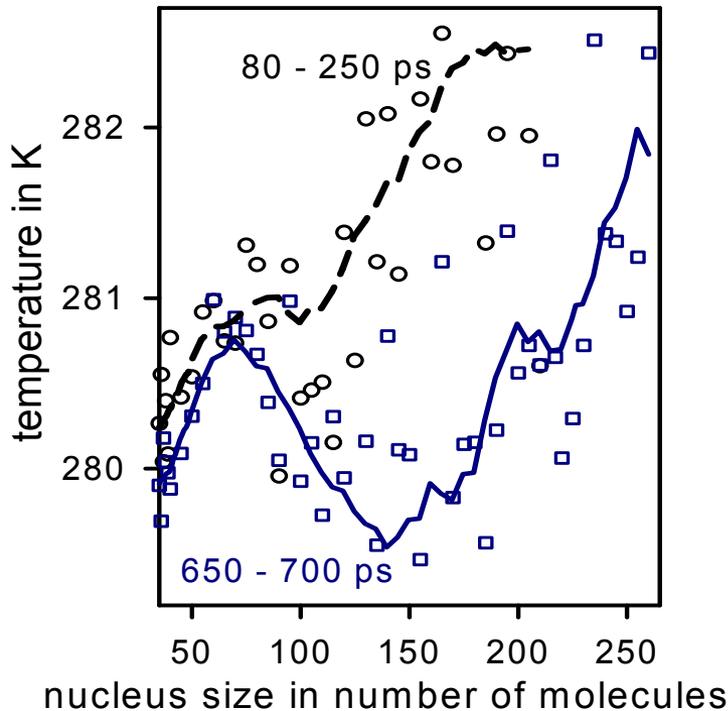
$$q_{j^*} \approx \frac{1}{2}$$



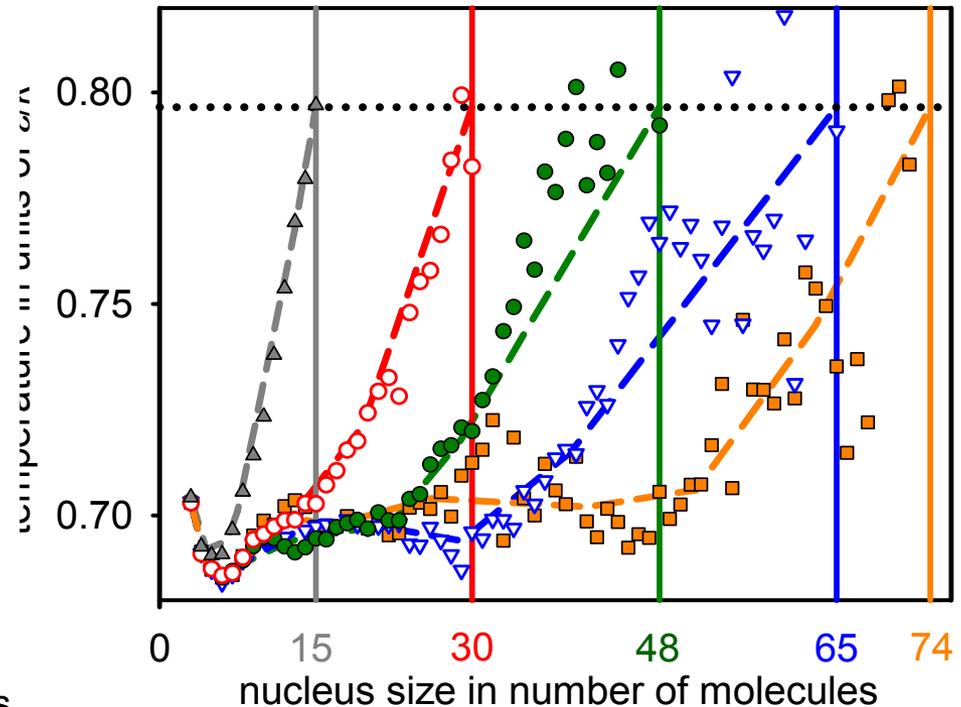
CNT predicts an acceptable value for j^* and underestimates J significantly

Maximal overheating of growing nuclei

Ethane: 280 K, 2.80 mol/l, *NVT*



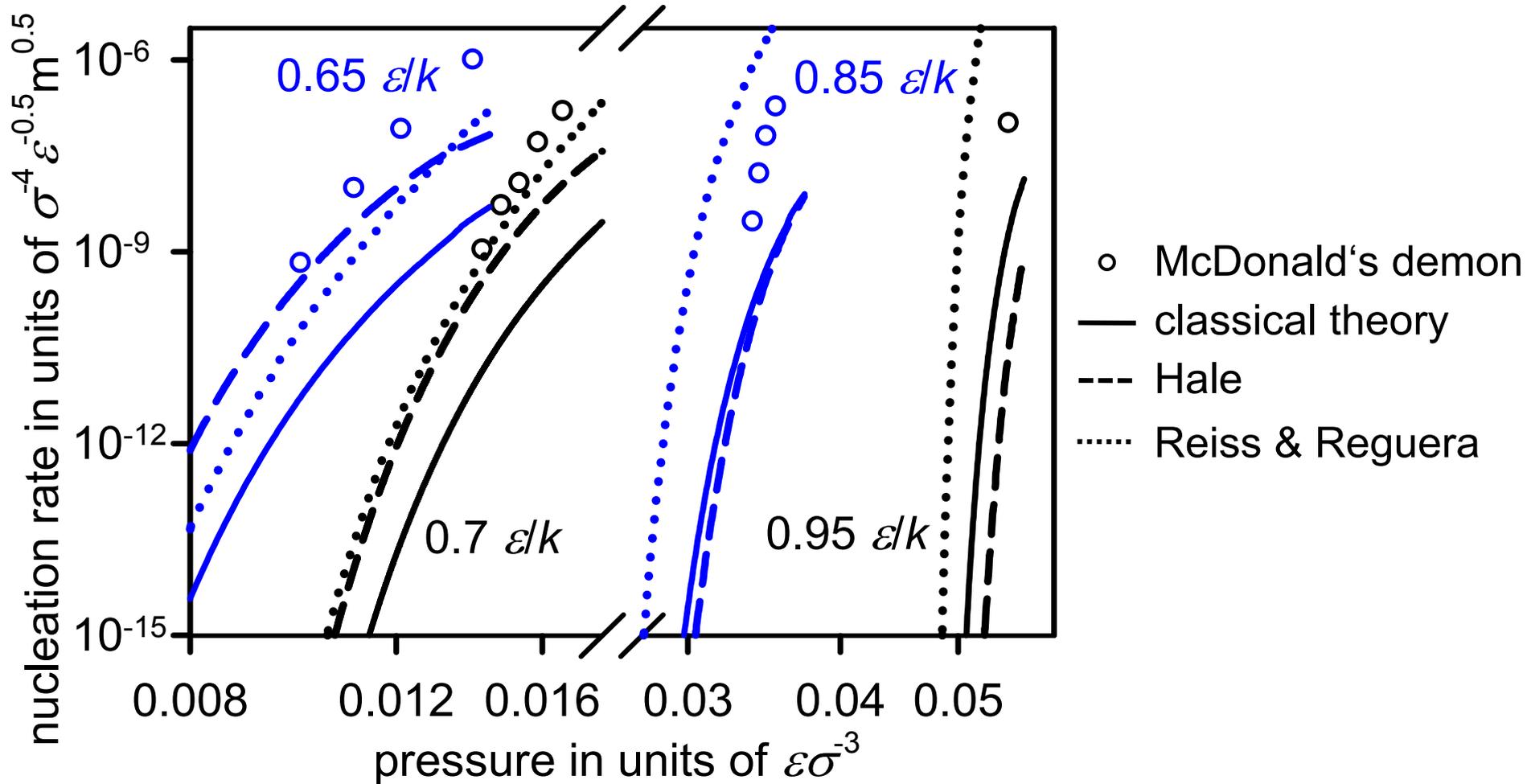
t-s-LJ fluid: $0.7 \epsilon/k$, μVT :
supercooled by $\Delta T = -0.0965 \epsilon/k$



$$(\Delta T)_{j=j^*=78}^{\text{CNT}} = \frac{2f_z}{\beta\Delta h^v} T = 1.4 \text{ K}$$

$$(\Delta T)_{j \rightarrow i} \gg (\Delta T)_{j=j^*=41}^{\text{CNT}} = 0.00608 \epsilon/k$$

GCMD simulation of nucleation: Results



Conclusion

- MD simulation of **steady states** allows sampling over an arbitrary time interval, eventually leading to the desired level of accuracy
- **Single droplets** can be stable in the **canonical** ensemble
- A **supersaturated vapor** near the spinodal line can be stabilized by grand canonical simulation with **McDonald's demon**
- **Classical nucleation theory** leads to acceptable results for the t-s-LJ fluid