

Simulation of aggregation phenomena in solution

Leonardo Lo Presti
leonardo.lopresti@unimi.it



Università degli Studi di Milano



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Outline

- (i) Motivation
- (ii) Method
- (iii) Applications and Results
- (iv) Conclusions

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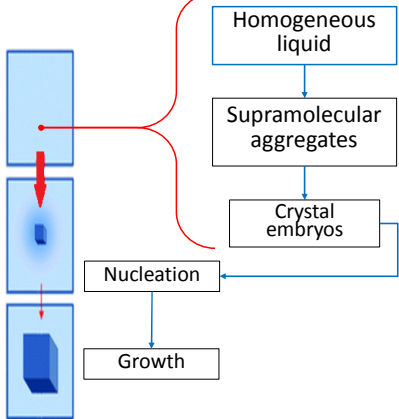
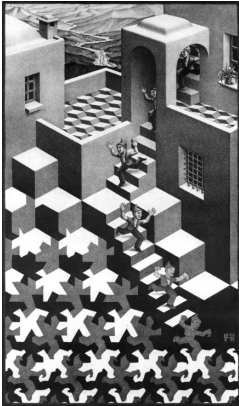
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Crystallization



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Chasing nucleation

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Capturing the Moment of Emergence of Crystal Nucleus from Disorder

Takayuki Nakamura, Masaya Sakakibara, Hiroki Nada, Koji Harano, and Eiichi Nakamura*

Cite this: *J. Am. Chem. Soc.* 2021, 143, 4, 1763–1767
 Publication Date: January 21, 2021
<https://doi.org/10.1021/jacs.0c12100>
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Molecular Dynamics Simulations of Crystal Nucleation from Solution at Constant Chemical Potential

Tarak Karmakar, Pablo M. Piaggi, and Michele Parrinello*

Cite this: *J. Chem. Theory Comput.* 2019, 15, 12, 6923–6930
 Publication Date: October 28, 2019
<https://doi.org/10.1021/acs.jctc.9b00795>
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The Kinetic-Biased algorithm

$$T_i^b = g \cdot T_i$$

$$\Delta T_i = T_i^b - T_i = T_i \cdot (g - 1)$$

$$\Delta T_{i,a} = \frac{\Delta T_i}{N_i} = T_{i,a}^{new} - T_{i,a} < 0$$

$$T_i^b = \sum_a^{N_i} T_{i,a}^b = \sum_a^{N_i} \frac{1}{2} m_{i,a} v_{i,a,b}^2 = g \cdot T_i$$

$$\sum_a^{N_i} \frac{1}{2} m_{i,a} v_{i,a,b}^2 = g \cdot \sum_a^{N_i} \frac{1}{2} m_{i,a} v_{i,a}^2 = \sum_a^{N_i} \frac{1}{2} m_{i,a} g (v_{i,a}^2)$$

$$\pm v_{i,a,s}^b = \pm \sqrt{g} \cdot v_{i,a,s}, s = x, y, z$$

$$\Delta T_k = -\Delta T_i / N_k > 0$$

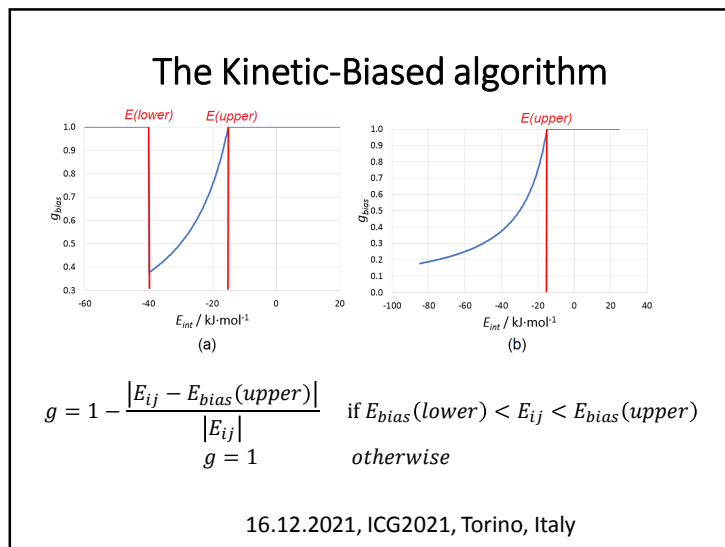
$$T_k^b = T_k + \Delta T_k = B \cdot T_k$$

$$B = 1 + \frac{\Delta T_k}{T_k}$$

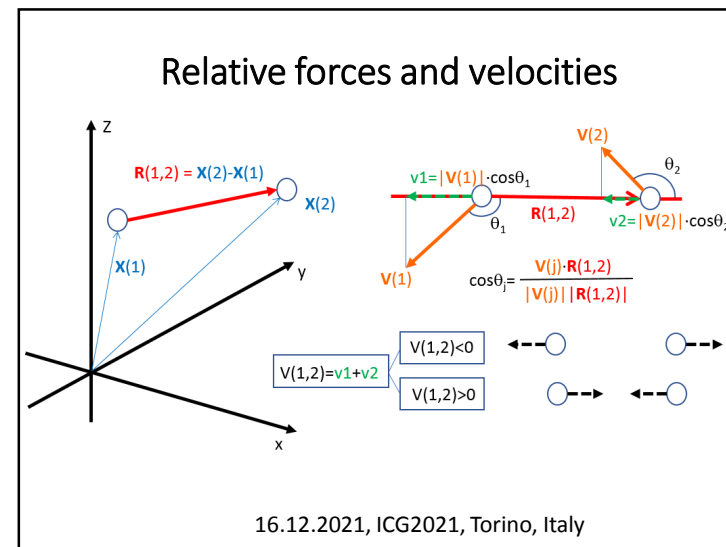
$$\pm v_{k,a,s}^{new} = \pm \sqrt{B} \cdot v_{k,a,s}, s = x, y, z$$

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MiCMoS

https://sites.unimi.it/xtal_chem_group/index.php

- *New J. Chem.* 2019, 43, 2077–2084
- *J. Appl. Cryst.* 2019, 52, 1253–1263
- *Cryst. Growth Des.* 2020, 20, 11, 7421–7428

Free!

Service programs

MD

Molecular topology and simulation box

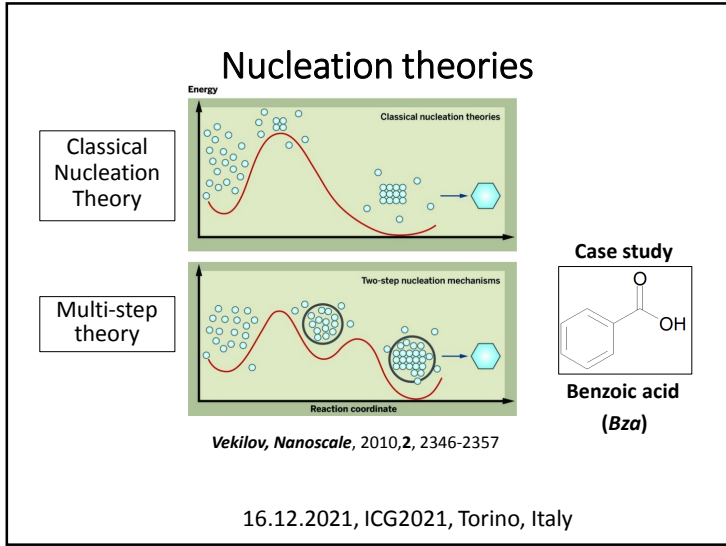
.oeh ASCII file: lattice, coordinates, charges and symmetry

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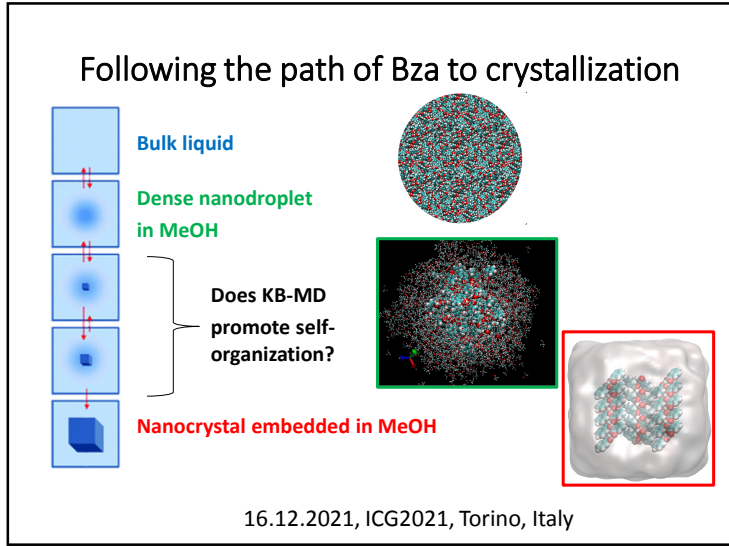
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- ### Outline
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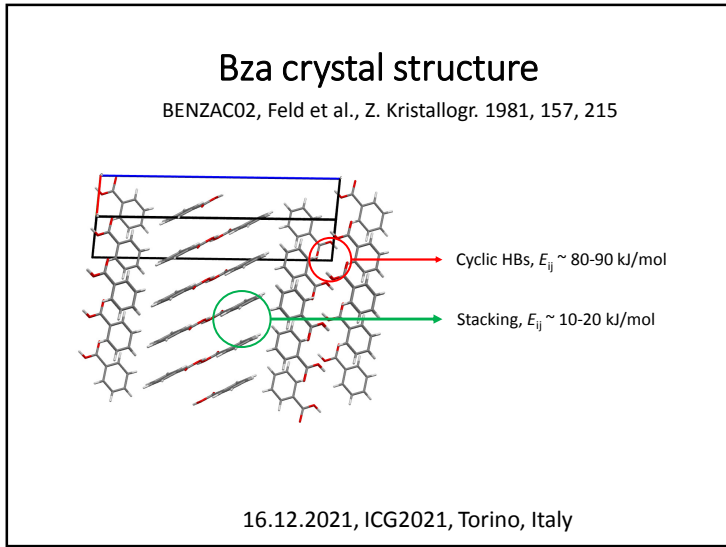
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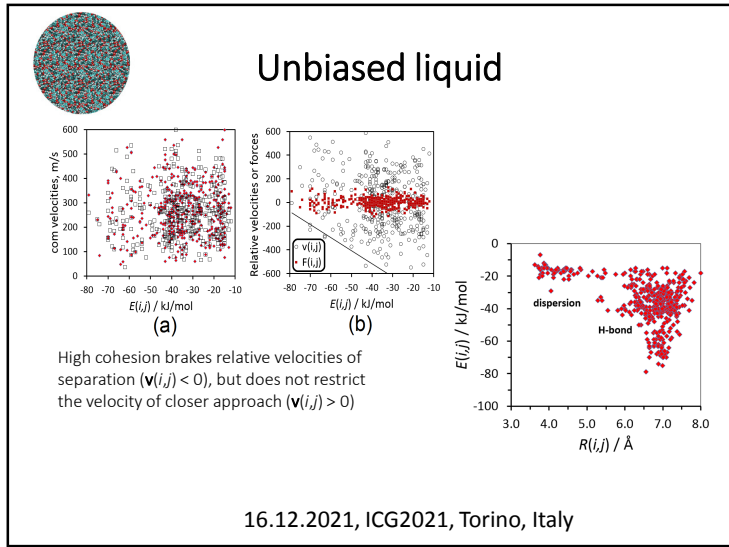
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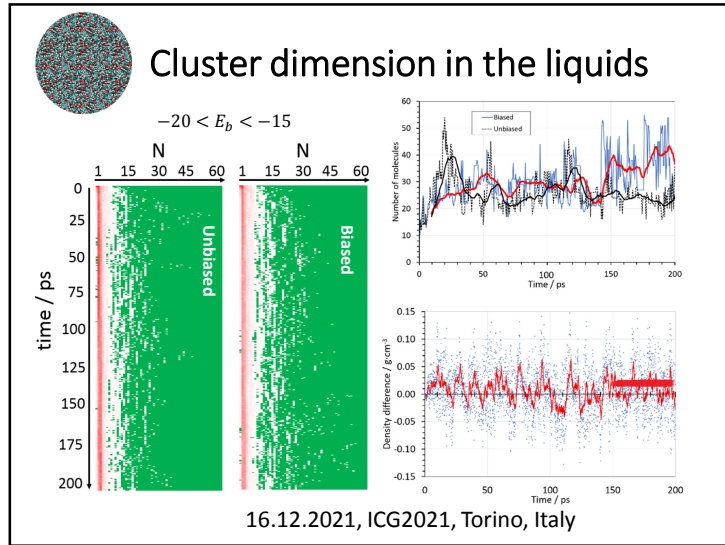
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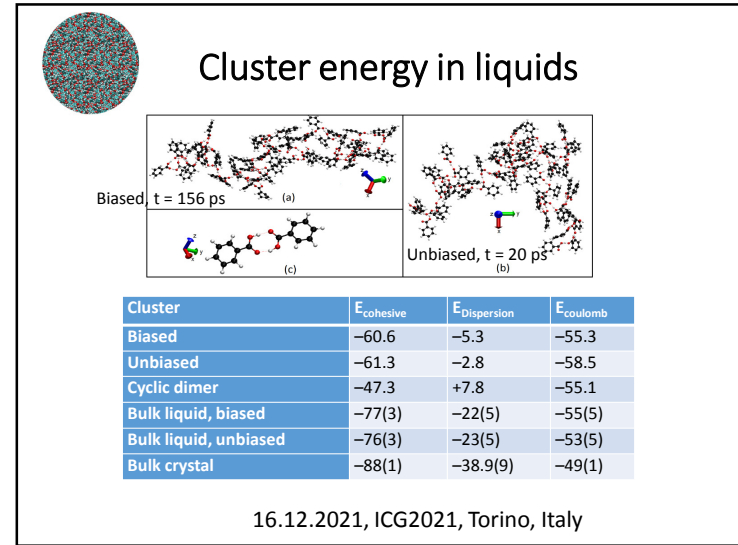
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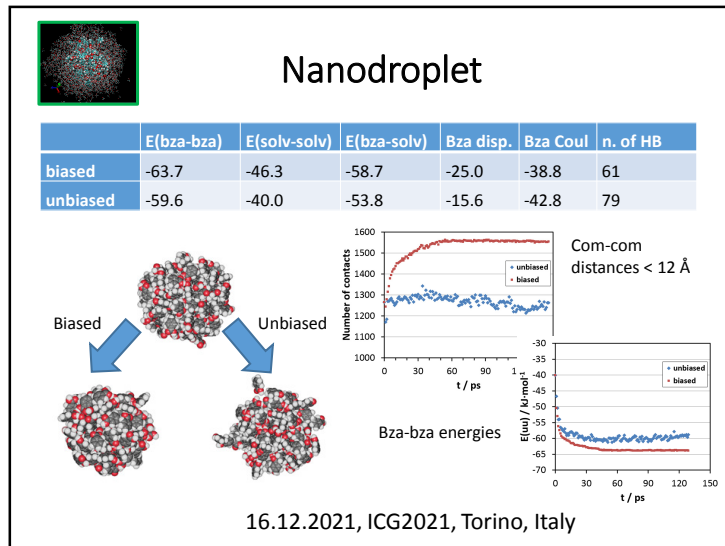
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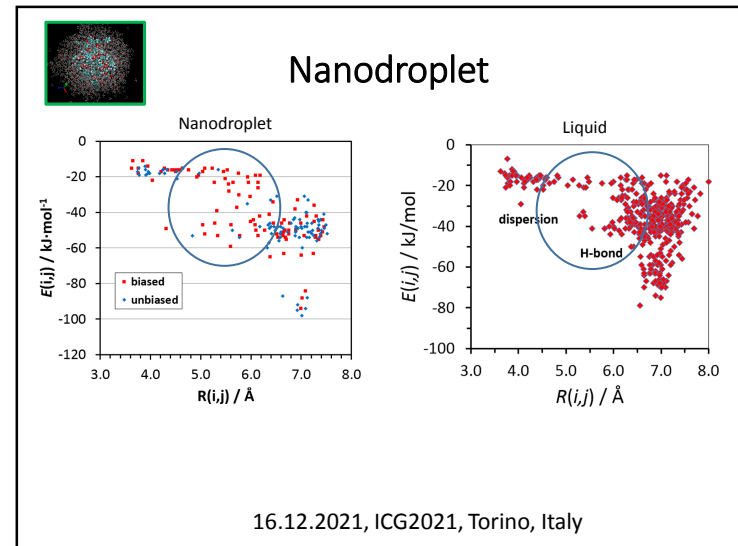
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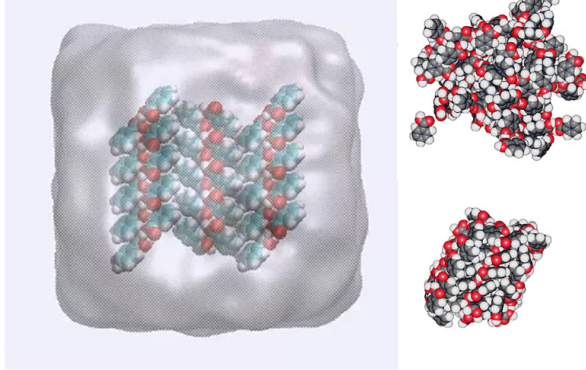


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Nanocrystal



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Conclusions

Under the biasing restrictions, pure liquid benzoic acid condenses by increasing its density and forming hydrogen-bonded clusters of irregular shape and multiform size

A solvated liquid droplet freezes to an amorphous semi-solid, into a compact form held together mainly by the dispersive potentials.

A microcrystal slab preserves its symmetric stability under biased dynamic conditions, while an unbiased simulation quickly runs into a liquid drop.

This last point strongly suggests (one could say demonstrates) **that isolated clusters cannot be crystalline**, contrary to hypotheses embedded into classical nucleation theory

The KB algorithm drives the system toward aggregation, not symmetrization (at least at these very short time scales)

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Thank you for your kind
attention

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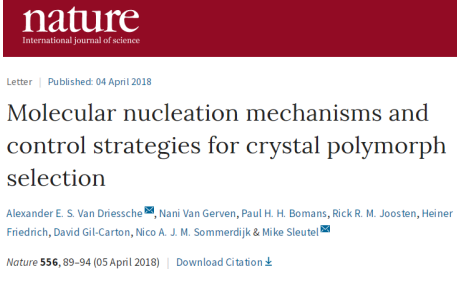
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Support slides

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Chasing nucleation



Letter | Published: 04 April 2018

Alexander E. S. Van Driessche, Nani Van Gerven, Paul H. H. Bomans, Rick R. M. Joosten, Heiner Friedrich, David Gil-Carton, Nico A. J. M. Sommerdijk & Mike Sleutel

Nature 556, 89–94 (05 April 2018) | Download Citation

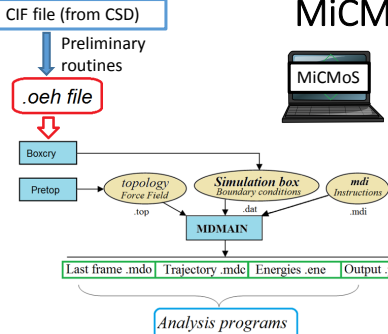


Van Driessche, Sleutel et al., Nature, 2018, 556, 89-94

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- For **bulk crystals**, periodicity is determined by repetition of the crystallographic unit cell. **Bulk liquids** use cubic simulation boxes. Also **non-periodic clusters** (nanocrystals and droplets) can be treated.
- Leap-Frog and velocity-Verlet integrator.**
- Berendsen (T, p) (anisotropic) rescaling allowed (NpT ensemble).**
- Heterogeneous mixtures (liquids+solids) are allowed.**

$$\lambda(t) = \left[1 + f \left(\frac{T_{set} - T}{T} \right) \right]^{1/2}$$

$$P_{ii} = \frac{2}{3V} (E_{kin,ii} - W_{ii})$$

$$\mu_i = [1 - c(P_{set} - P_{ii})]^{1/3}$$

$$V(t + 1/2\Delta t) = V(t - 1/2\Delta t) + \frac{\Delta t}{M} F(t)$$

$$r(t + \Delta t) = r(t) + \Delta t \cdot V(t + 1/2\Delta t)$$

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The method

Lattice Energy, Packing Energy

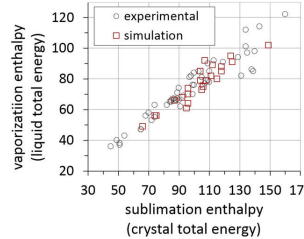
$$E_{latt} = PE = U(\text{crystal}) = \frac{1}{2} PPE$$

$\Delta H = -PE - 2RT$

Ignoring zero-point and molecular relaxation corrections

Parametrization

- (i) Enthalpies of sublimation (solids) and of evaporation (liquids);
- (ii) Observed lattice parameters (solids) and specific volumes (liquids);
- (iii) $U(i,j)$ from MP2 calculations



Gavezzotti & Lo Presti, *Cryst. Growth Des.* **2015**, *15*, 3792–3803

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Intramolecular force field

$$E(\text{str}) = 1/2 \cdot k_{\text{str}} \cdot (R - R^0)^2$$

$$E(\text{bend}) = 1/2 \cdot k_{\text{bend}} \cdot (\cos\theta - \cos\theta^0)^2$$

$$E(\text{tors}) = k_{\text{tors}} [1 + f \cdot \cos(m\varphi)]$$

k_{str} and k_{bend} from MP2/6-31G** ab initio calculations on the corresponding deformation of prototypic compounds (Gavezzotti, New J. Chem. **2016**, *40*, 6848)

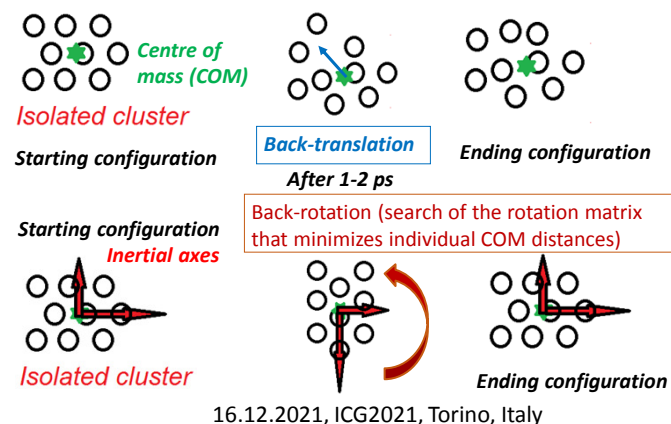
k_{tors} calibrated by fitting against MP2/6-31G** potential curves of ~ 50 organic compounds *in vacuo* (Gavezzotti & Lo Presti, J. Appl. Cryst. **2019**, submitted)

Intramolecular potentials here serve only as restraints to avoid undue molecular bond and angle distortions

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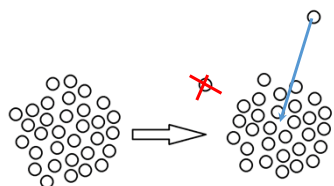
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Isolated clusters: suppression of net momenta



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Isolated clusters: Evaporation



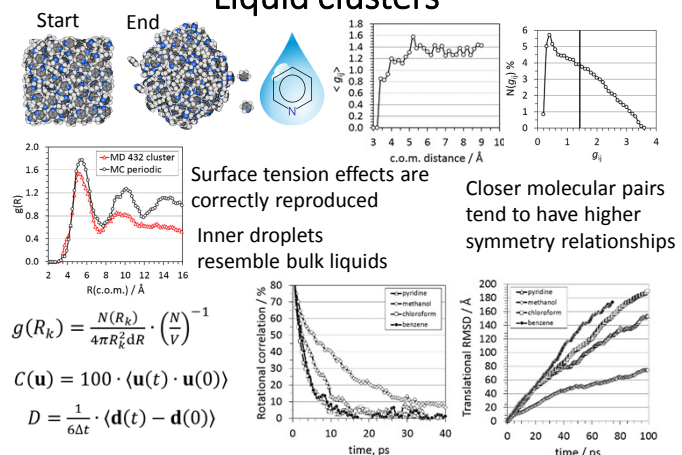
(1) Tethering: if the COM of a give molecule is more distant than a given threshold (for example $\sim 2 \cdot \text{cluster radius}$) from the cluster COM, the distance is reduced by a user-defined factor $0 < F_{\text{ev}} < 1$, usually 0.9.

(2) Deletion: if the COM of a give molecule is more distant than a given threshold (for example $\sim 2 \cdot \text{cluster radius}$) from the cluster COM, the molecule is deleted.

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Liquid clusters



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