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Centre for Scientific Computing

Predictive modelling: a view from the atomic level

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- MidPlus
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Philosophy

- Complex materials
 - complete model is not possible
- Mechanistic information
 - “predictions” not reliable without the correct underlying physics/chemistry
- Dimensionality
 - identify key degrees of freedom
 - target analysis
 - drive or simplify modelling

Unexpected Mechanisms: old examples

- Decay of the Velocity Autocorrelation Function

B. J. Alder and T. E. Wainwright

Phys. Rev. A **1**, 18 – Published 1 January 1970

- Hyperdynamics: Accelerated Molecular Dynamics of Infrequent Events

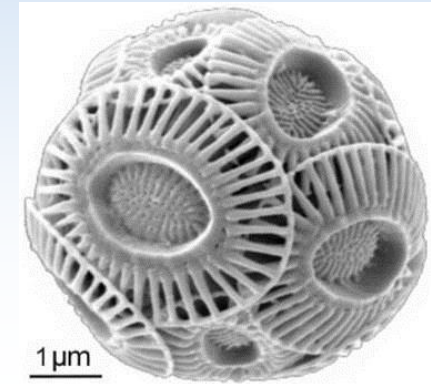
Arthur F. Voter

Phys. Rev. Lett. **78**, 3908 – Published 19 May 1997

Motivation: Biominerals

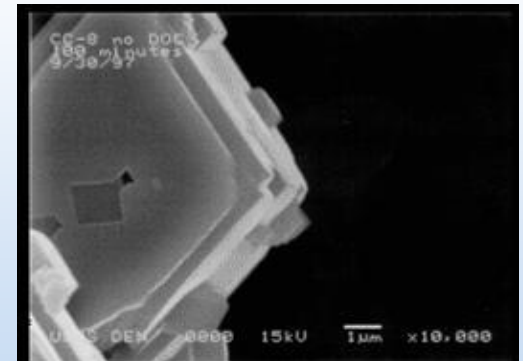
- Nature:
 - Exquisite control of crystal morphology & aggregation found in nature
 - Leads to very well “designed” functional materials
 - Ability to simulate implies ability to guide synthesis

Emiliania huxleyi
coccoliths



Henriksen, K., S. L. S. Stipp, et al.
American Mineralogist **89**, 1709-1716
(2004)

- Laboratory:



Reddy, M. M. and A. R. Hoch.
*Journal of Colloid and Interface
Science* **235**, 365-370 (2001)

Motivation: additives for inhibition

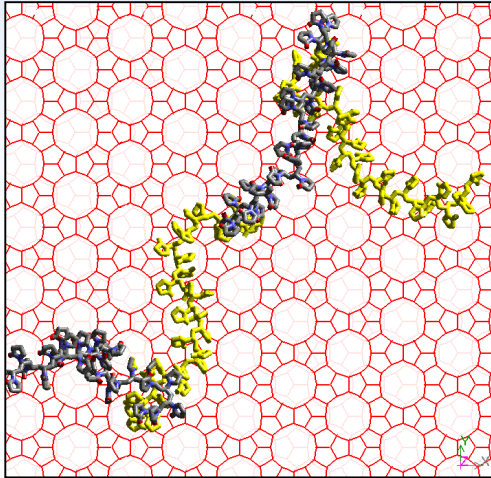
- Often need to suppress crystal growth
 - Scale, wax, hydrate
- “kinetic” inhibitors
 - *Delay* nucleation or *slow* growth
 - Active at low concentrations
 - Require *molecular* understanding of nucleation and growth



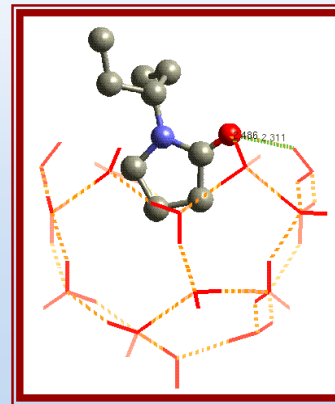
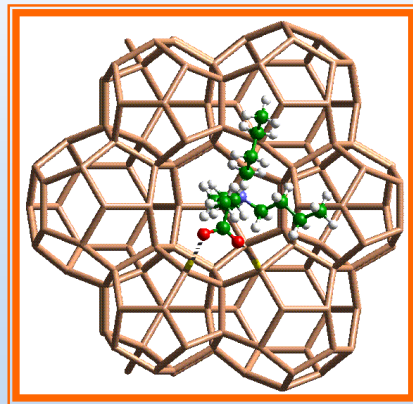
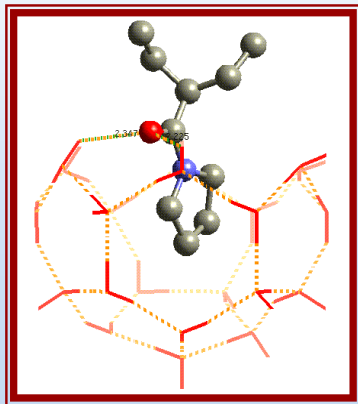
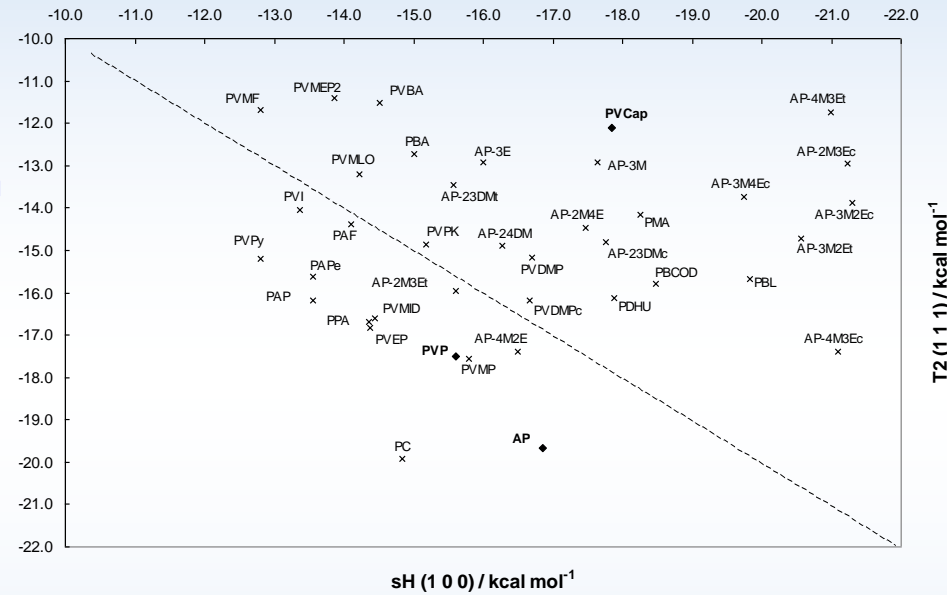
Mechanistically inspired prediction?

- Inhibition by surface adsorption

validate

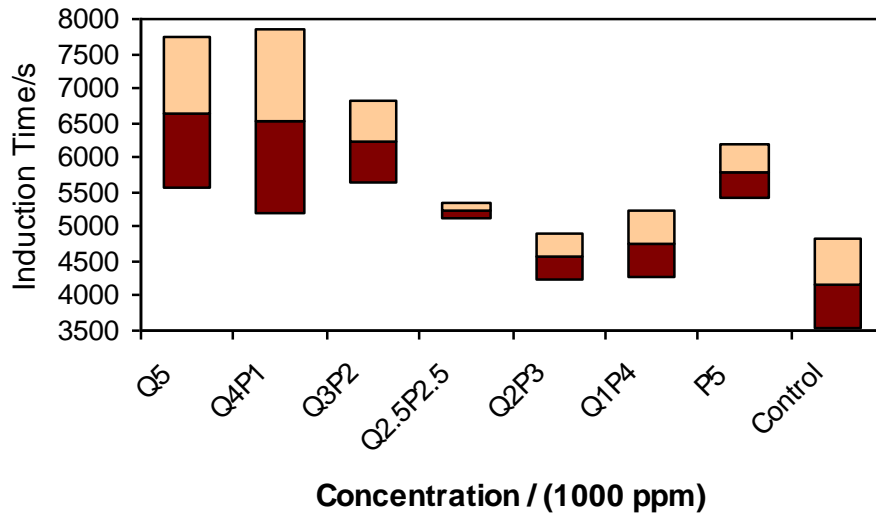


screen



new modes?

Synthesise and test



THF Hydrate Tests

induction times for 0.5% mixtures of tba3S (the “quat”, or Q) and PVP (P). The Control contains no inhibitor. Bars indicate \pm one standard deviation

	$t_{\text{induction}} / \text{s}$	$R_{\text{nucl}} / (\text{ml}^{-1} \text{min}^{-1})$	$R_{\text{growth}} / (\mu\text{m min}^{-1})$
Control	-90 (32)	1439 (1031)	1.56 (2.18)
tba3S (0.1 %)	1705 (422)	89 (-)	0.61 (0.23)
tba3S (0.5 %)	2678 (231)	116 (13)	0.86 (0.25)

Ethane Hydrate Tests

Induction times, nucleation rates (R_{nucl}) and growth rates (R_{growth}). Standard deviations are given in parentheses

2nd generation

Test No.	KI	Gas	Test P/T psia/C	Subcooling /C	Induction Time/ hrs
1	JI-C003:1	NG	1023 / 5.3	10.7	5
2	JI-C003:1	NG	1023 / 5.3	10.7	>67
3	JI-C003:1	NG	1015 / 5	11	>67
4	JI-C003:1	NG	1023 / 5	11	>150
5	JI-C003:1	Methane	1700 / 4.5	10	>150
	Commercial	NG		10	16
	Commercial	Methane		10	5
6	JI-C002:1	NG	1095 / 4.5	12.3	3
7	JI-C002:1	NG	1095 / 4	12.8	40
8	JI-C002:1	NG	1095 / 5	11.8	>42

- 2nd generation inhibitors perform about 4 times better than current commercial inhibitor

But!

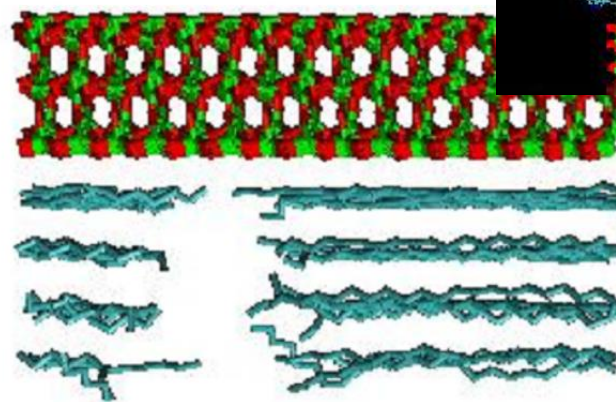
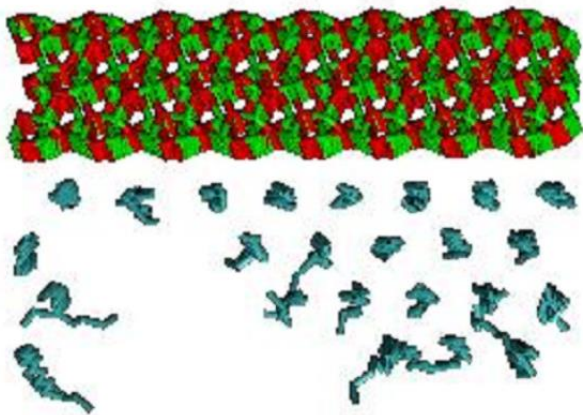
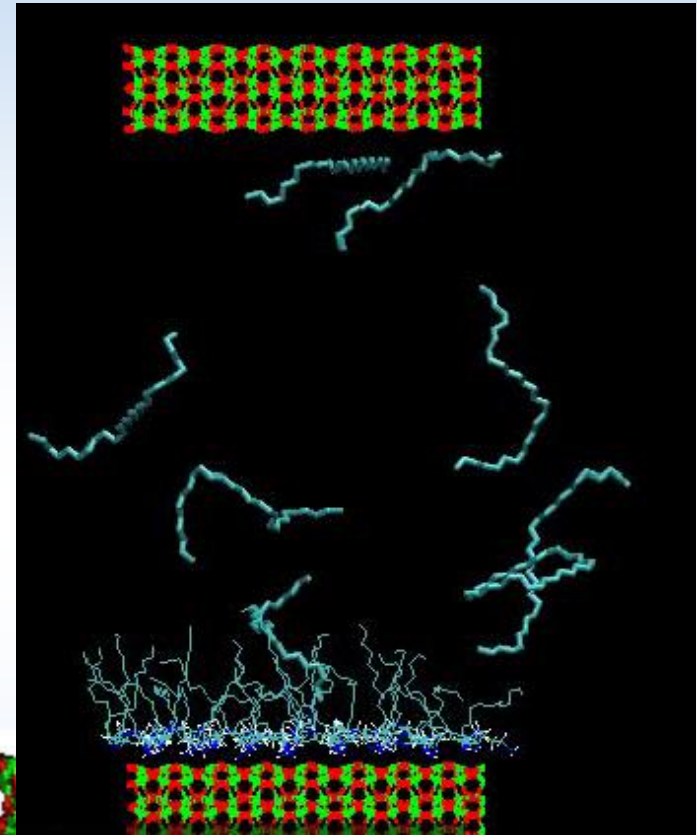
- Experiments not repeatable!
 - Oil companies can't repeat each other's screens
 - Heriot-Watt: activity changed with test-cell stirrer design
 - Scale of activity with new compound by test-site
 - Heriot-Watt > Toulouse > Halliburton > Heriot-Watt
- Similar story with wax inhibition (model oil using well characterised edible oil mixture)
- Experimentalists working with nanotoxicology want legislative standards based on computational tests because experiments are too irreproducible!
- Simplistic model: dynamic interface; nucleation; ...

Nucleation from MD?

- Homogeneous nucleation?
 - Experimental nucleation rates $< O(10^6)$ nuclei per cm^3 per second
 - MD simulation 10,000 molecules:
 - 1 nucleation event every 32 millenia!
- Heterogeneous nucleation?
 - (some) success with seeding
 - (some) success with interfaces

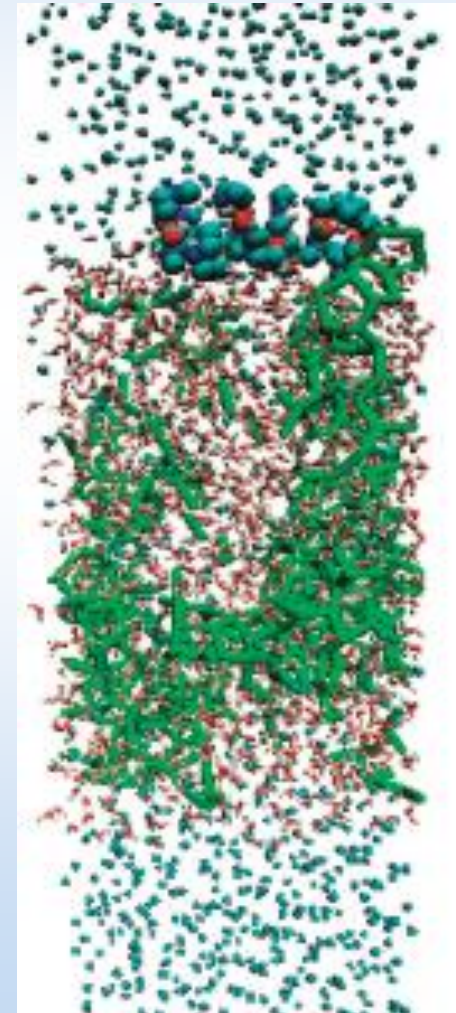
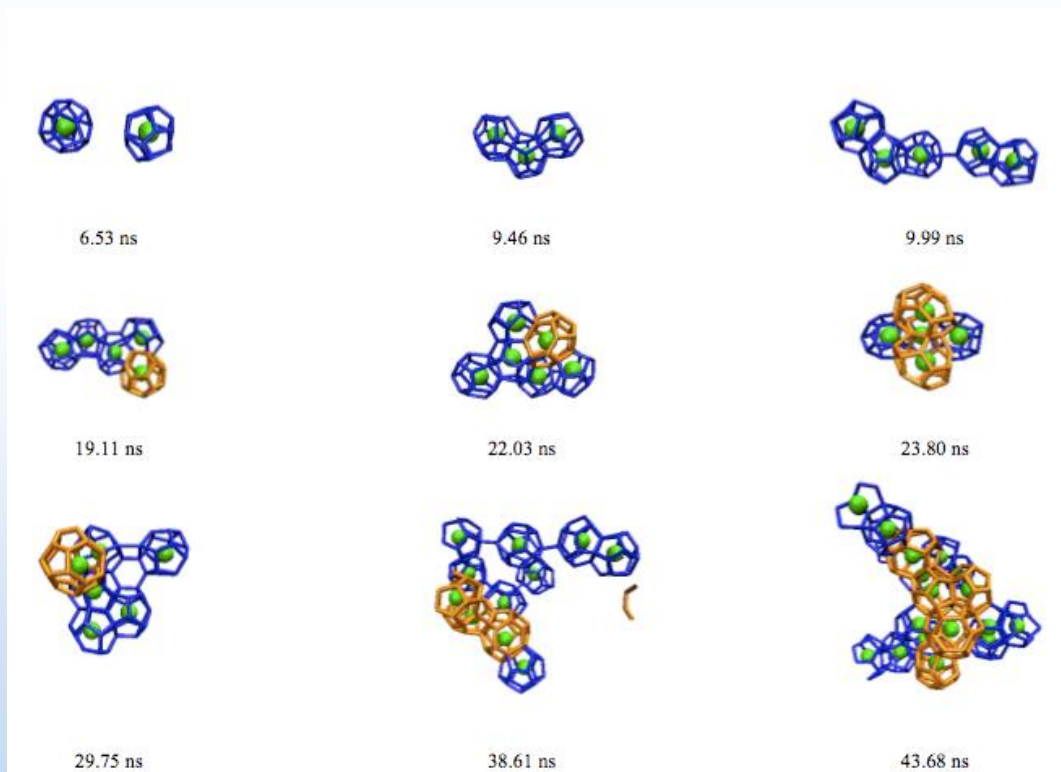
Standard MD: modelling surfaces

- Deposition of wax on Fe_2O_3
 - C_{28} from C_7



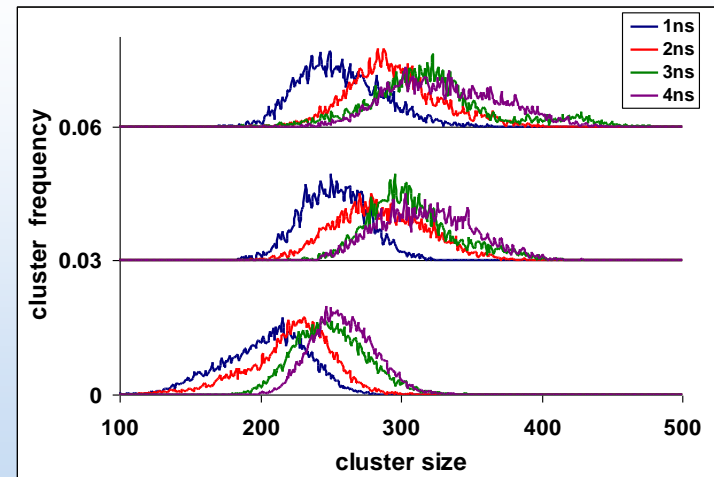
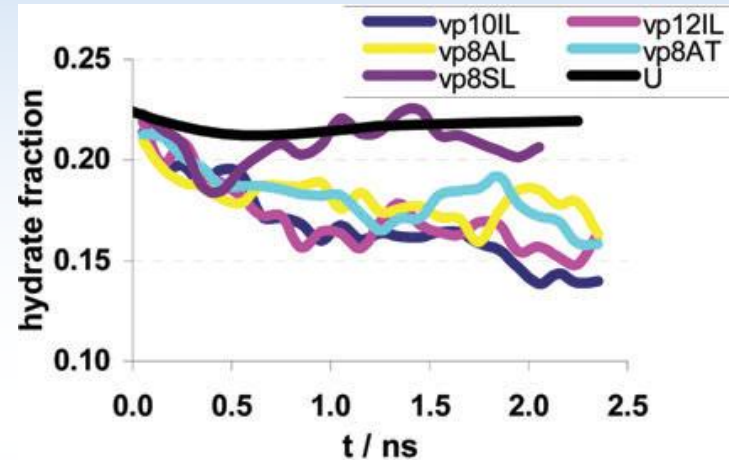
Standard MD: crystal nucleation

- heterogeneous and/or high supersaturation/subcooling



Polydispersity in inhibitors

- molecular weight
 - amount of hydrate present
 - PVP
- molecular shape
 - growth of hydrate clusters
 - pDMAEMA

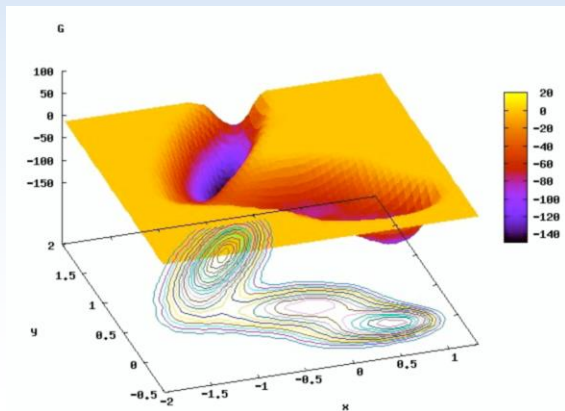


Nucleation more generally

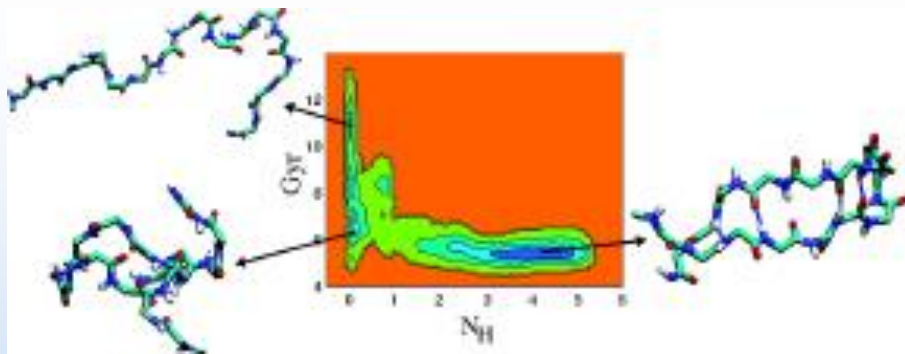
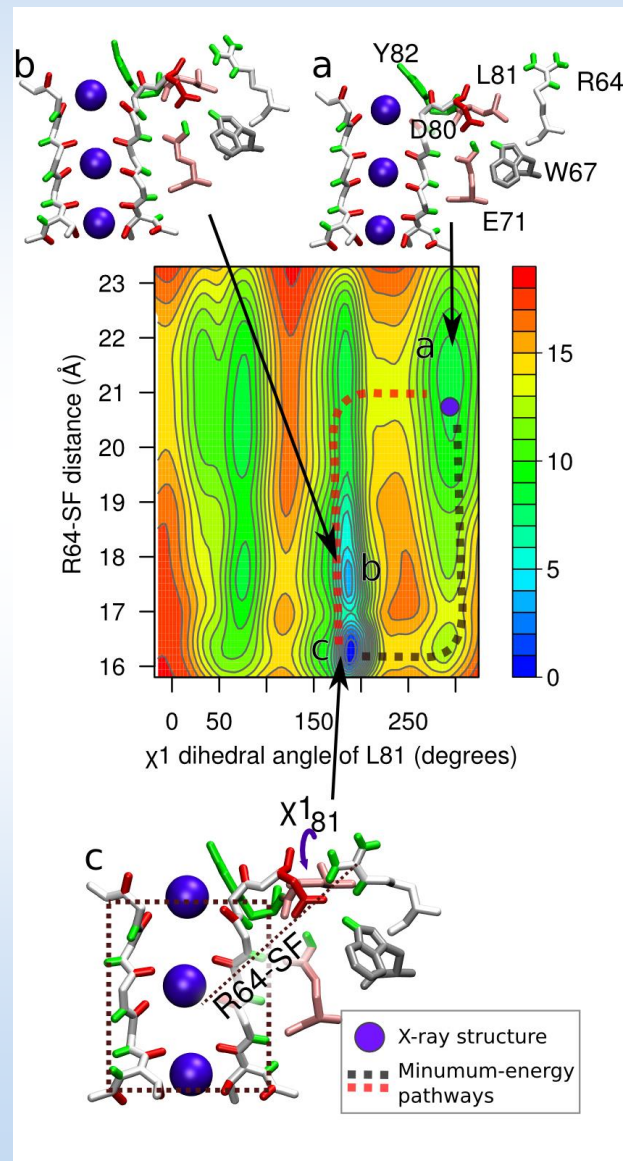
- Project onto “important” manifold
 - Define key dimensions (order parameters / reaction coordinates / collective variables)
 - Project analysis onto these dimensions
 - Bias simulations to explore these dimensions
 - constrain to portion of manifold (umbrella sampling)
 - trap fluctuations that explore the manifold (adaptive bias force)
 - disfavour current portion of manifold (Wang-Landau; metadynamics)

⇒ Free energy hypersurfaces

3-basin model
Quigley, PMR



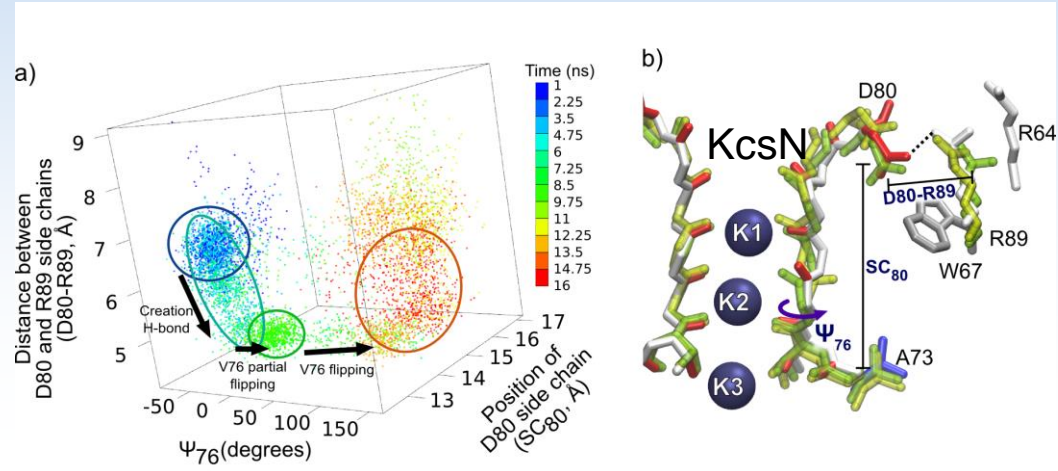
KcsN ion channel
Cosseddu, PMR, Khovanov



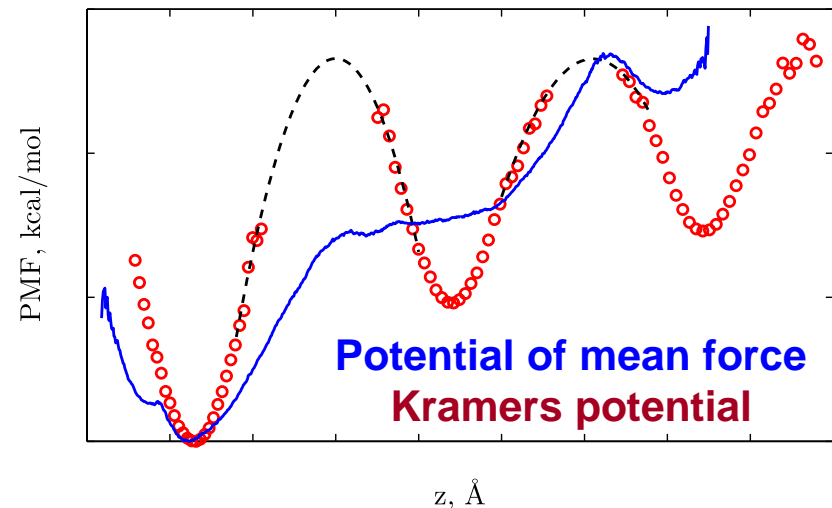
peptide folding
Bussi, ..., Parrinello

Limitations and Pitfalls: dimensionality!

- Equilibrium / comprehensive sampling
 - Typically 1–2D (≤ 6 ; can do ~ 100 for basin escape & rough sampling)
 - projection merges basins

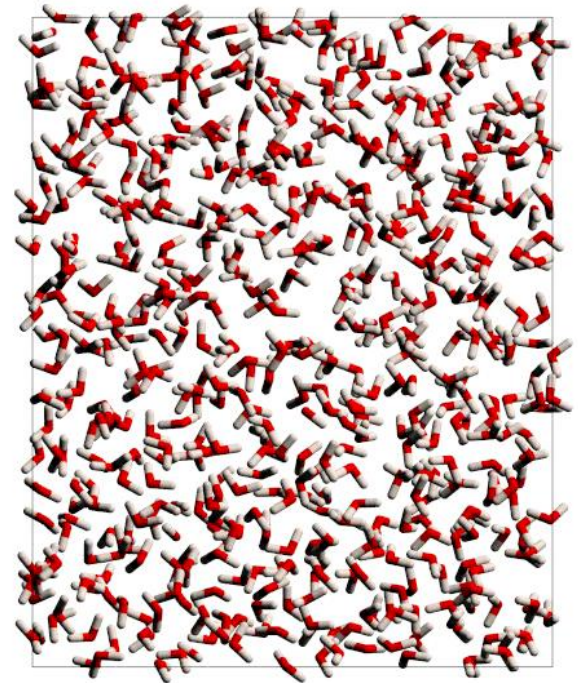


- Adiabatic Surfaces
 - dynamics defined by metastable substates?

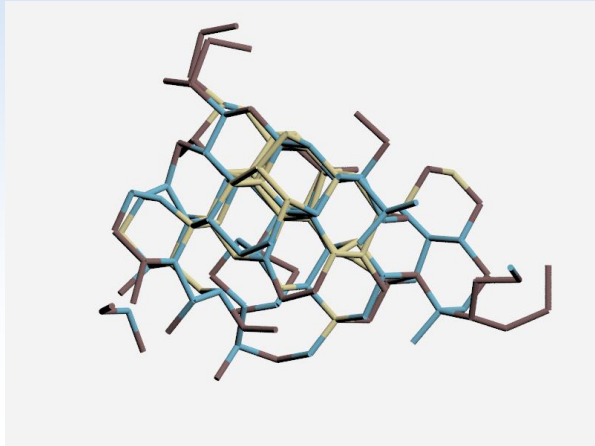


Nucleation with metadynamics: ice

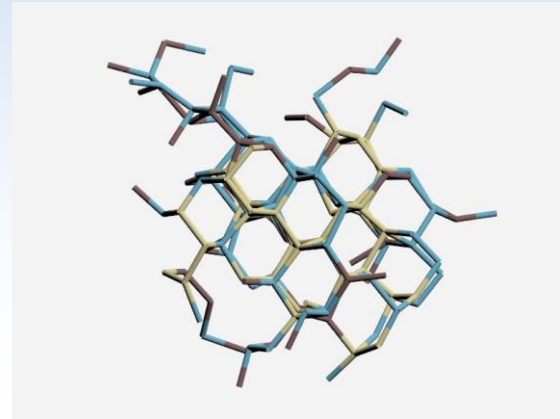
- 4D bias:
 - Q4, Q6, ζ and potential energy (simultaneous)
- *NPT* simulation
 - density change is spontaneous
- $T = 180$ K; *ca.* 11 “ns” simulation
- Results independent of periodic boundaries



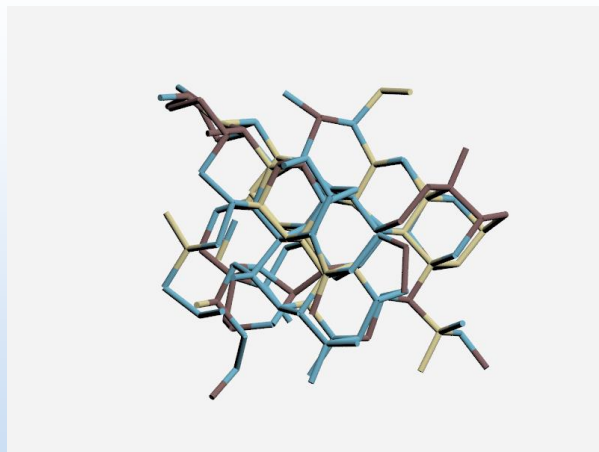
An ensemble of critical nuclei



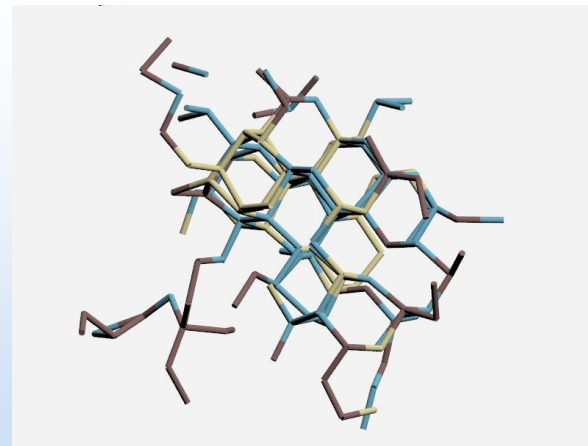
N = 147



N = 157



N = 138



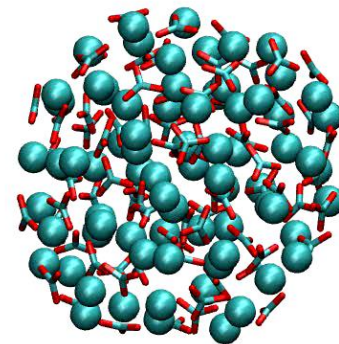
N = 161

ice I_c
ice I_h
both

Nucleation with metadynamics: CaCO_3

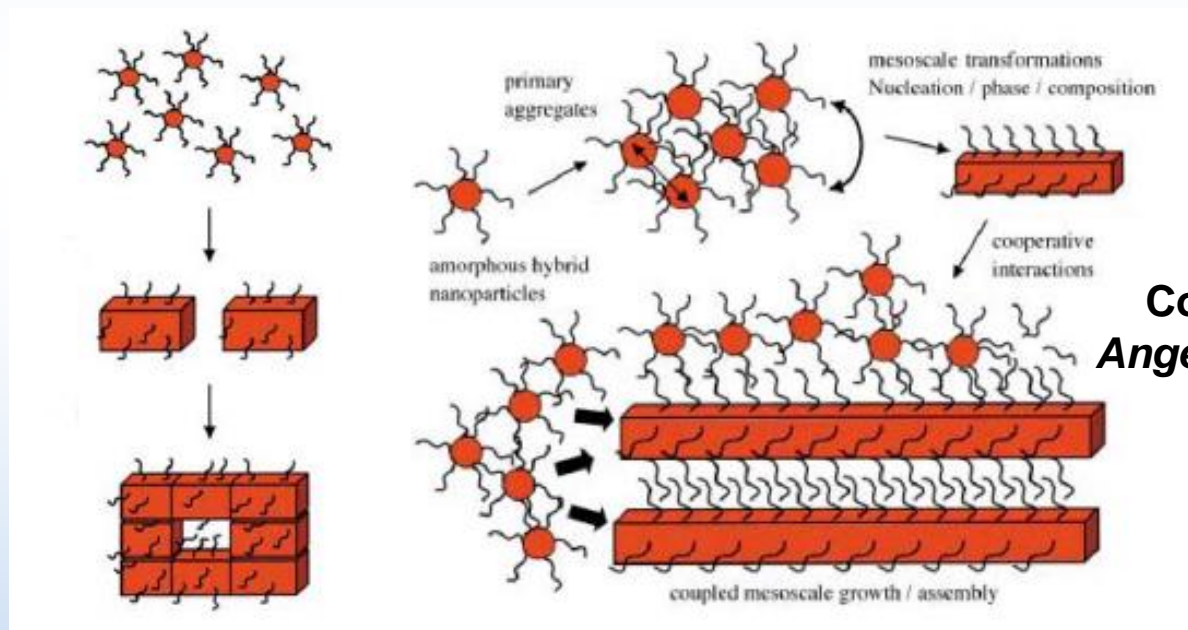
- 6 order parameters
 - Q_4 for: Ca-Ca, Ca-C, Ca-O, C-C, C-O
 - measures orientations of X about Y
 - energy associated with CaCO_3
- *ca.* 10 ns of MD

Exploring configurations
for 75 units (water not
shown)



Calcium Carbonate Biominerals

- Multi-stage hierarchical formation
- ο ρ γ α ν ι χ ϖ χ ο ν τ ρ ο λ



Colfen & Mann
Angew. Chem, 2003

Early stages of carbonate biomineralisation

ions & ion pairs → hACC → ACC → crystalline

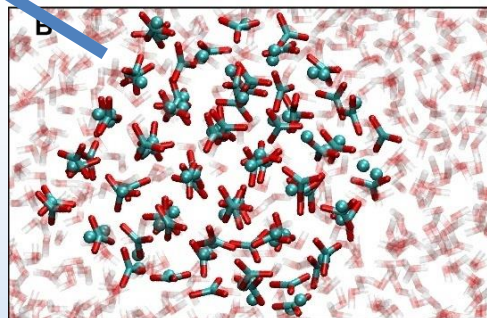
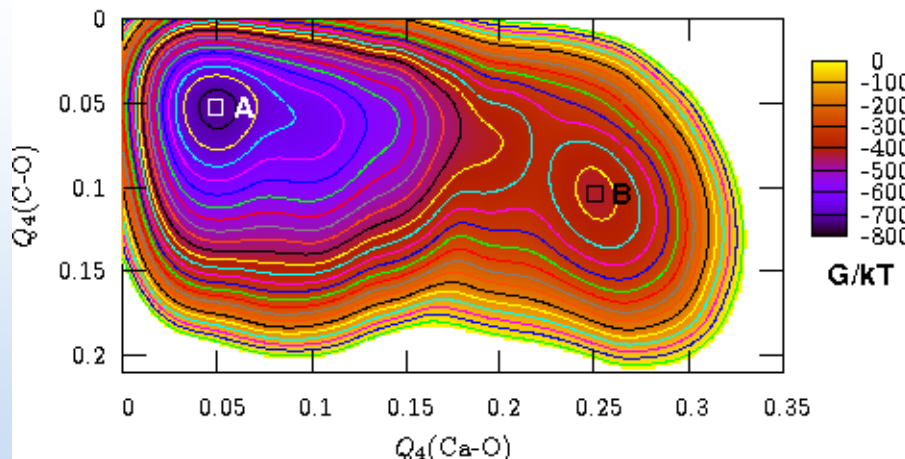
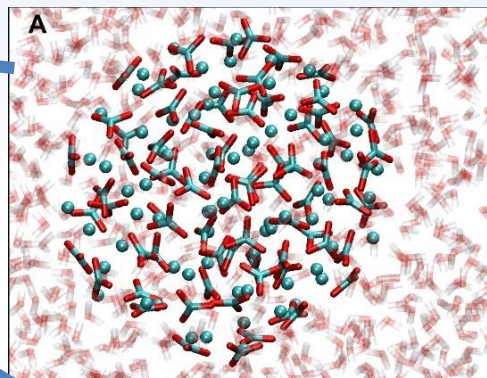
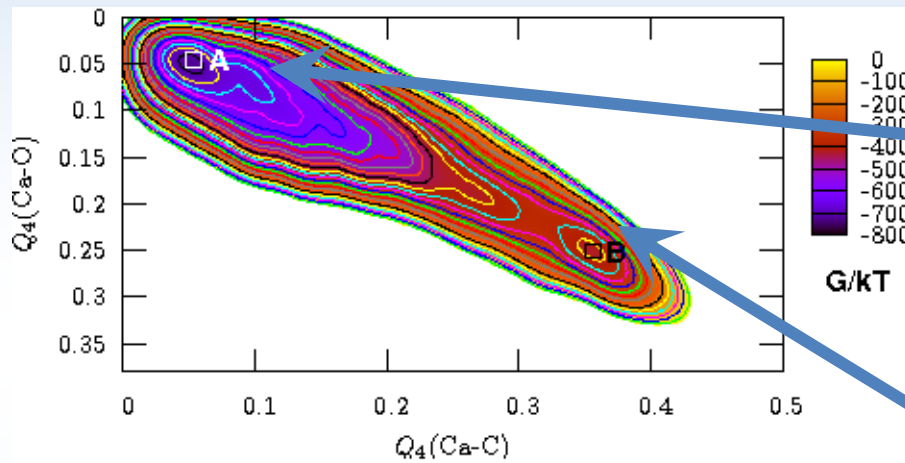
hACC: hydrated amorphous calcium carbonate
composition *ca.* $\text{CaCO}_3 \cdot \text{H}_2\text{O}$

ACC: (anhydrous) amorphous calcium carbonate

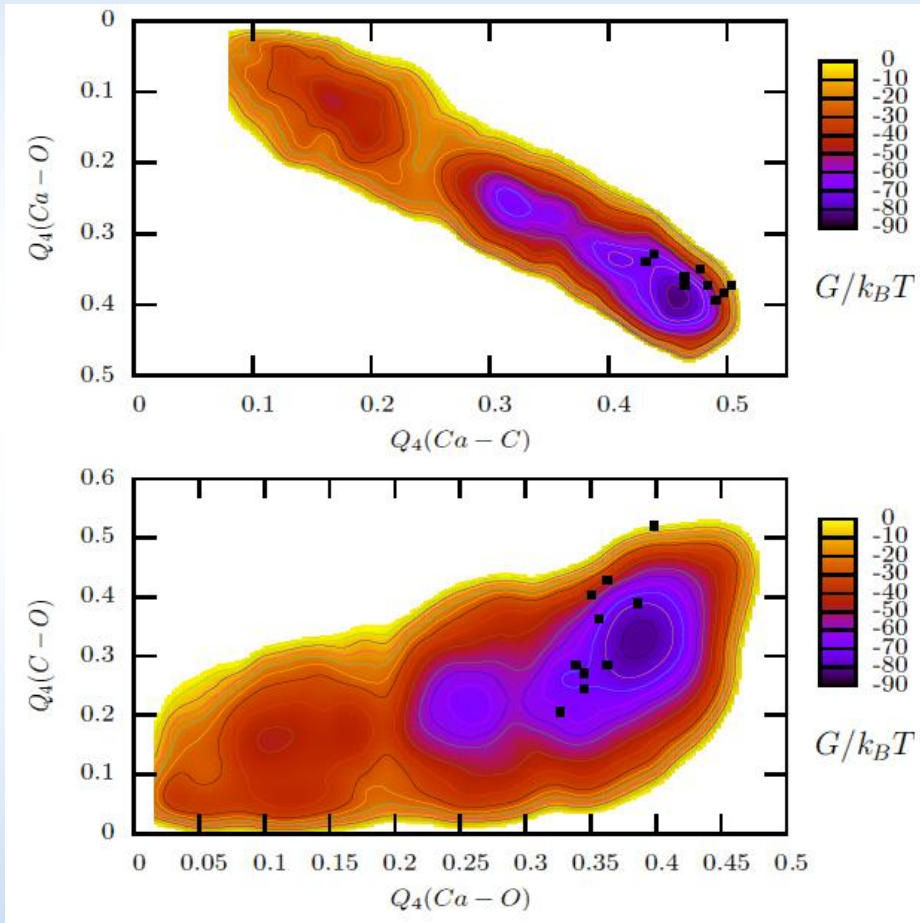
Crystalline: calcite, aragonite, vaterite

Free energy maps: 75 CaCO_3 units

- *NVT*
- favours amorphous

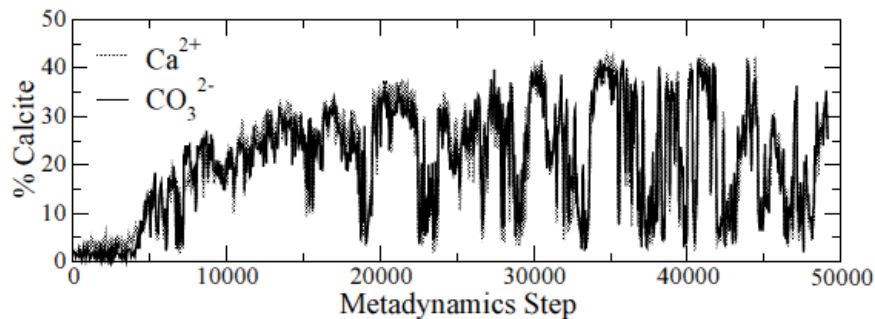
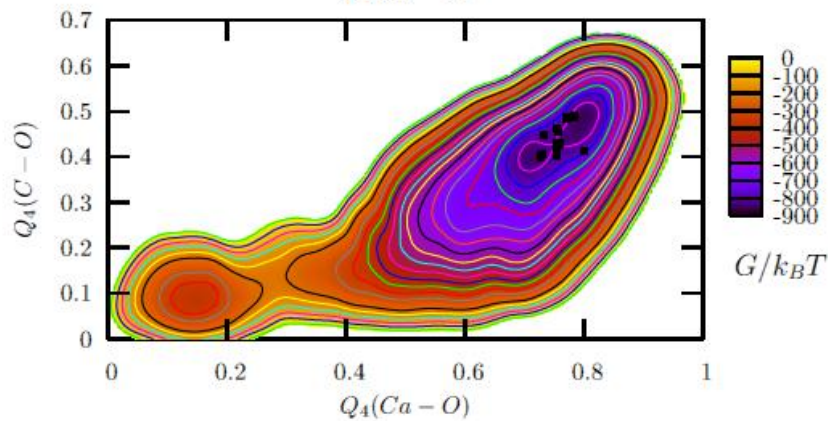
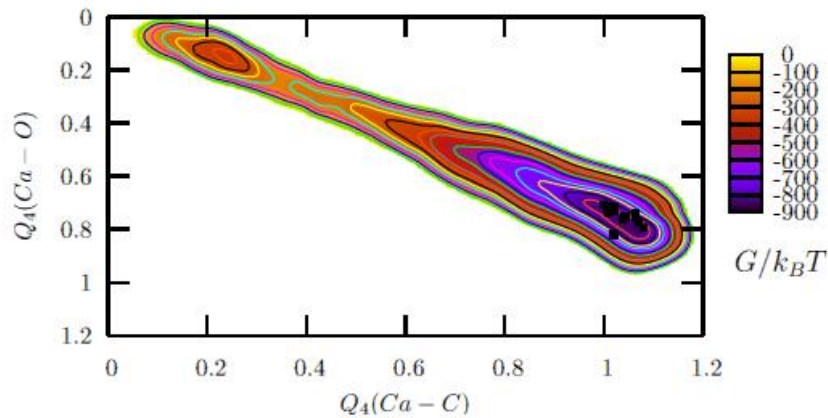


Free energy maps: 75 CaCO_3 units



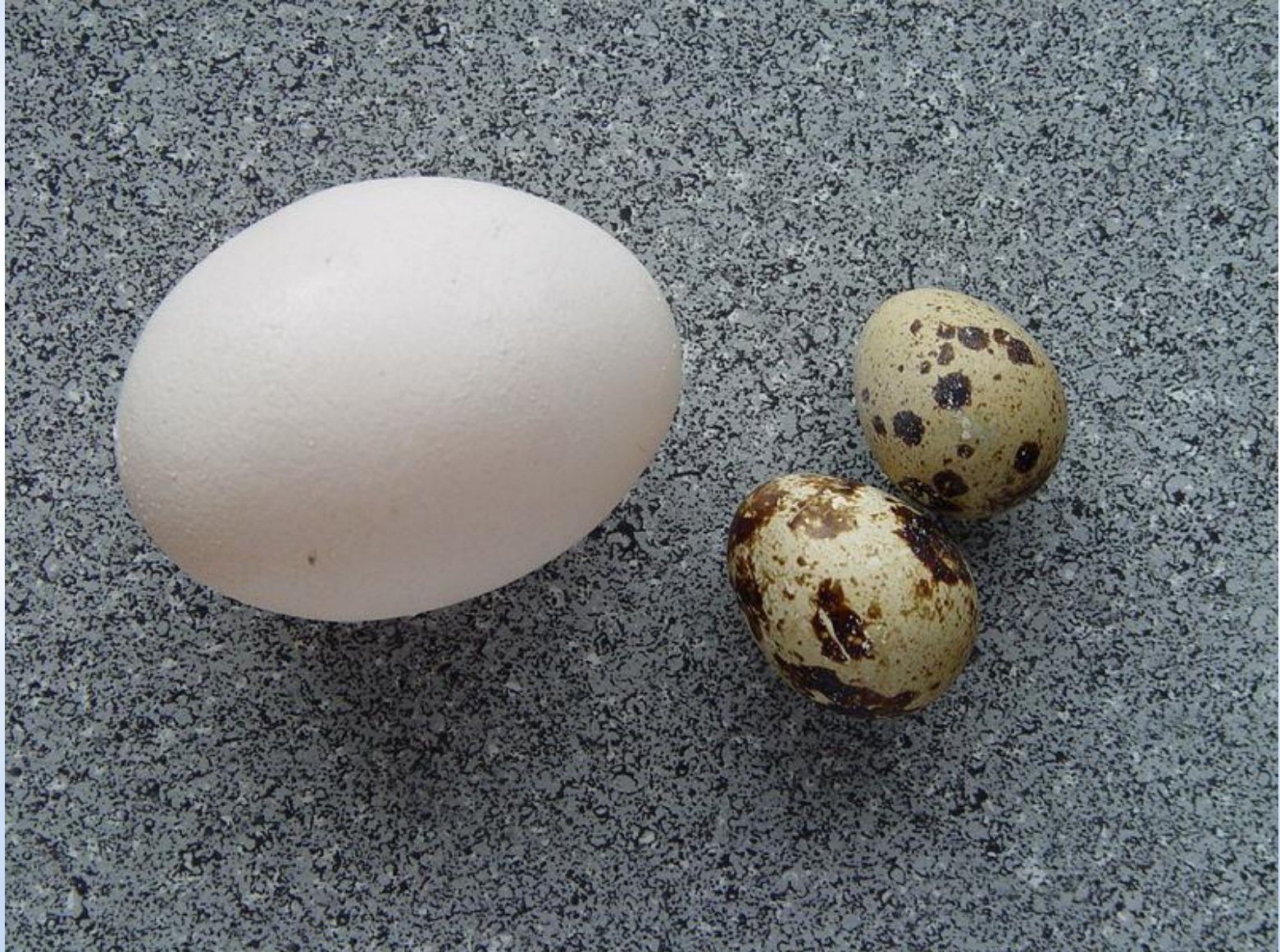
- *NPT*
- Dominated by calcite
 - very low barrier (10–20 kT)

300 CaCO₃ Units in water

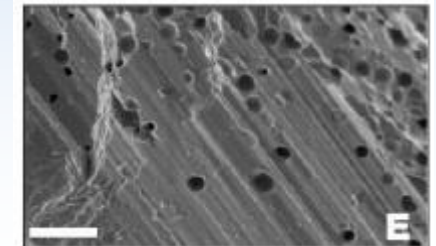
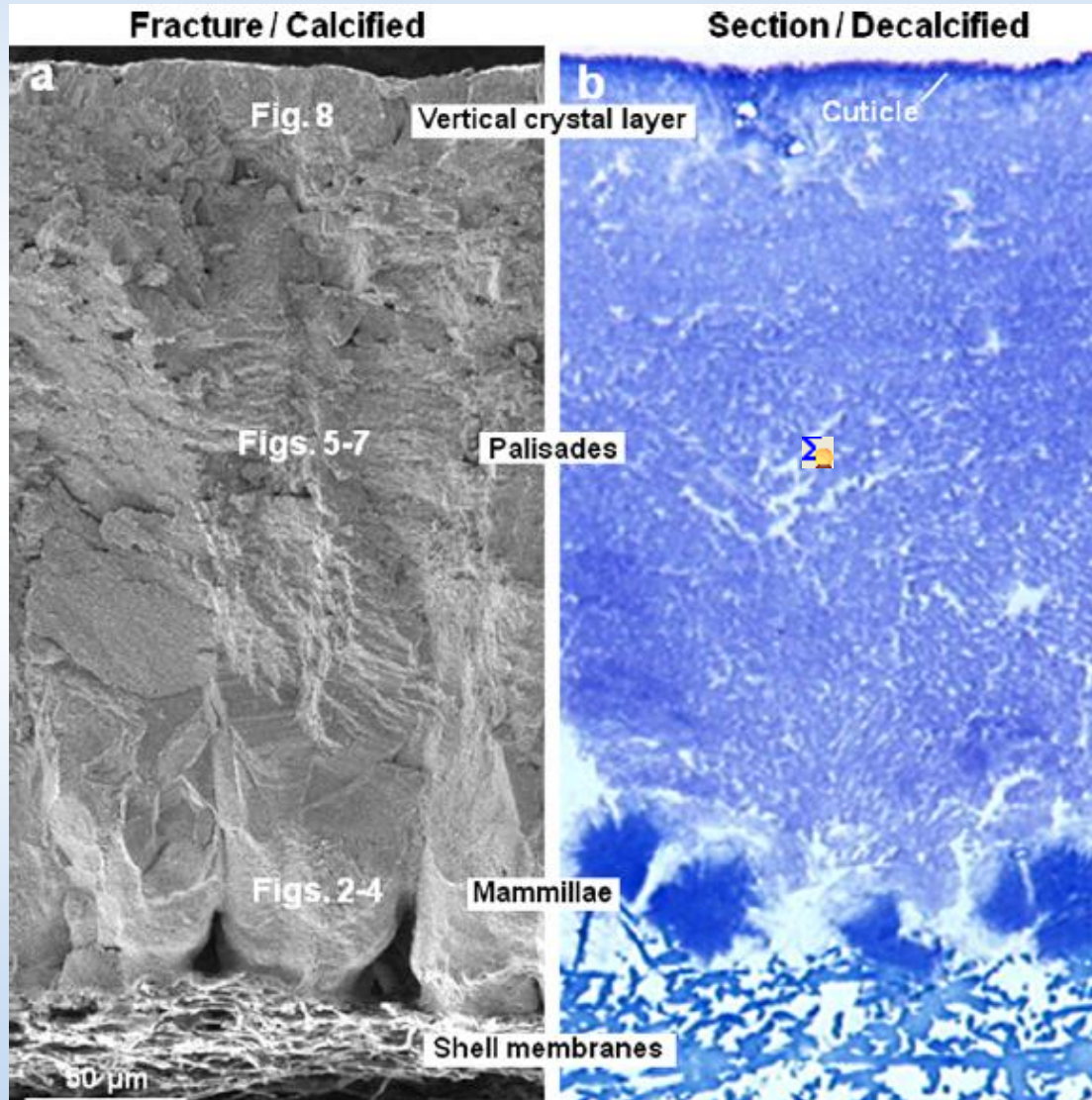


- two stable states
- calcite (dominant)
- amorphous
- energy barrier $> 10^2 kT$
- multiple transitions

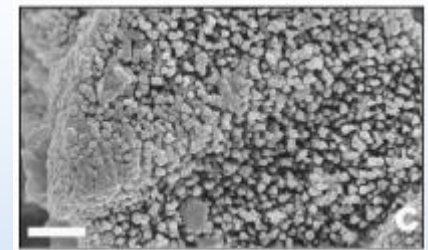
Biom mineralisation: egg shells



Structure of an eggshell



Biomacromolecules, Vol. 7, No. 11, 2006 3203
Lakshminarayanan et al.



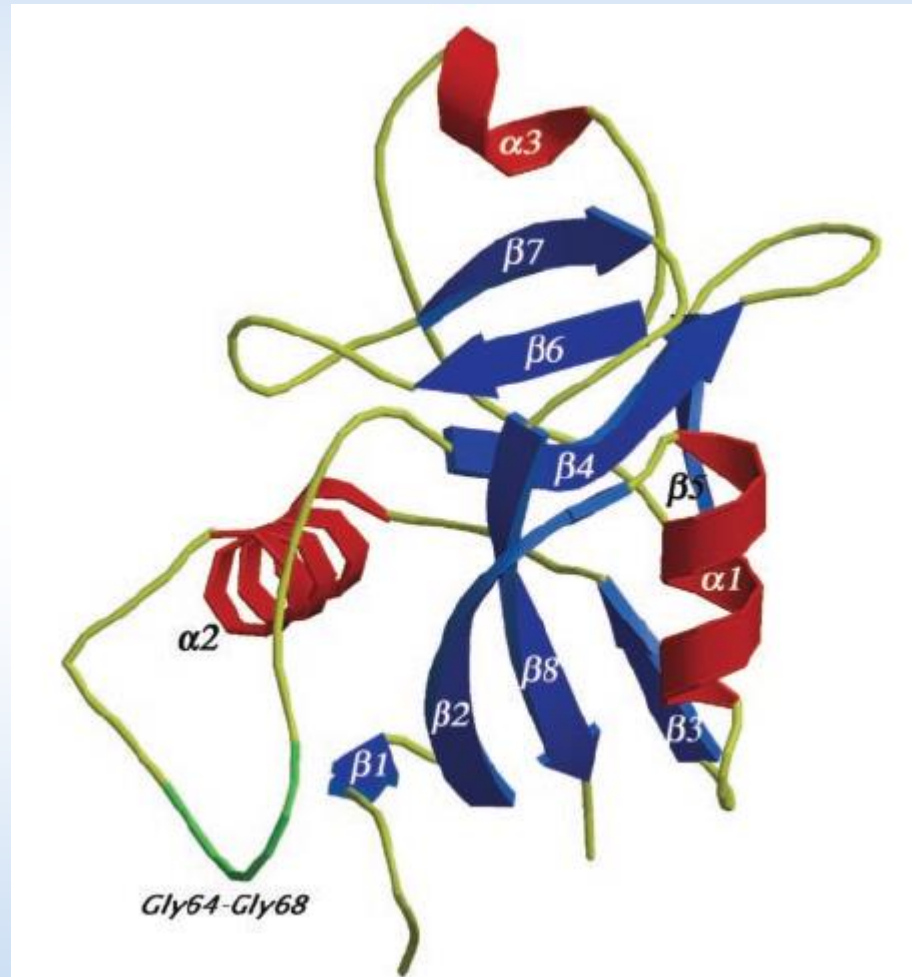
G-type lectin-type proteins

K. Mann, F. Siedler / Comparative Biochemistry and Physiology, Part B 143 (2006) 160-170

- found in
 - Rhea
 - Duck
 - Chicken
 - Emu
 - Ostrich

RCA-1	1	V	R	A	N	R	C	L	K	G	W	L	D	F	R	G	N	C	Y	G	Y	F	R	Q	E	L	P	W	R	K	A		
DCA-1	8	V	R	A	S	P	C	P	K	G	W	L	D	F	R	G	N	C	Y	G	Y	F	R	H	E	L	P	W	R	K	A		
SCA-1	1				D	K	C	P	K	G	W	L	D	F	R	G	N	C	Y	G	Y	F	R	Y	E	L	P	W	R	K	A		
ANCA	1				N	K	C	P	K	G	W	L	D	F	R	G	N	C	Y	G	Y	F	R	Q	E	L	P	W	R	K	A		
RCA-2	1	D	E	Q	E	I	G	C	A	S	G	W	V	P	F	D	G	R	C	F	G	F	P	Q	E	L	S	W	R	R	A		
DCA-2	1	E	E	E	I	G	C	T	S	G	W	V	P	F	D	G	R	C	Y	G	F	F	P	Q	E	L	S	W	R	R	A		
SCA-2	1	R	E	R	A	G	C	A	K	G	W	V	P	F	D	G	R	C	Y	G	F	F	P	Q	E	L	S	W	R	R	A		
OC-17	1	D	P	D	G	C	G	P	G	W	V	P	F	D	G	R	C	L	G	F	F	S	R	E	L	S	W	S	R	A			
Hu-PSP	1	I	S	C	P	E	G	T	N	A	Y	R	S	Y	C	Y	F	N	E	D	P	E	T	W	V	D	A						
RCA-1	31	E	A	W	C	R	V	V	R	G	G	C	H	L	A	S	I	H	T	S	E	E	H	R	A	V	A	K	F	I	-		
DCA-1	38	Q	A	W	C	R	A	L	R	D	G	C	H	L	A	S	I	H	S	A	E	E	H	R	A	I	A	R	F	V	-		
SCA-1	28	E	A	W	C	R	S	I	R	A	G	A	H	L	A	S	I	H	T	S	E	E	H	R	A	I	A	R	F	I	-		
ANCA	28	E	A	W	C	K	V	I	H	A	G	C	H	L	A	S	I	H	S	E	E	H	Q	A	I	A	V	A	R	F	I		
RCA-2	31	E	G	F	C	Q	R	L	G	A	R	T	H	L	A	S	I	H	S	E	E	H	Q	A	I	I	M	S	M	L	A		
DCA-2	31	E	S	F	C	Q	R	L	G	A	R	T	H	L	A	S	I	H	N	E	E	H	Q	A	I	I	S	M	L	A			
SCA-2	31	E	G	F	C	Q	R	L	G	A	R	T	H	L	A	S	I	H	S	E	E	H	Q	A	I	I	V	S	M	L	A		
OC-17	30	E	S	F	C	Q	R	L	G	A	R	T	H	L	A	S	I	H	S	E	E	H	Q	A	I	I	V	S	M	L	A		
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RCA-1	60	-	S	Q	C	R	R	G	-	E	E	G	-	-	-	D	D	V	W	I	G	L	Y	H	W	N	-	-	K	S	W		
DCA-1	67	-	S	Q	C	Q	R	G	-	E	E	E	-	-	-	E	N	V	W	I	G	L	R	Q	L	V	-	-	K	L	W		
SCA-1	57	-	S	Q	Y	H	H	G	-	E	E	E	-	-	-	E	D	V	W	I	G	L	F	R	W	N	-	-	S	V	W		
ANCA	58	K	F	Q	-	R	R	-	-	E	E	E	-	-	-	D	N	V	W	I	G	L	H	H	W	N	Q	A	R	-	-		
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DCA-2	61	S	S	Q	P	Y	S	D	S	E	E	E	E	V	A	A	N	G	D	V	W	I	G	L	R	L	S	L	-	R	R	L	W
SCA-2	61	S	S	Q	P	Y	S	D	S	E	E	E	E	V	A	G	E	E	V	W	I	G	L	H	R	P	L	G	R	R	N	W	
OC-17	60	A	S	R	G	G	D	G	S	G	E	G	-	-	-	A	D	G	R	V	W	I	G	L	H	R	P	A	G	S	R	W	
Hu-PSP	57	E	S	S	T	-	D	D	S	-	-	-	-	-	-	N	V	W	I	G	L	H	D	P	K	K	N	R	R				
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ANCA	80	V	W	I	D	G	S	K	K	R	Y	S	A	W	D	D	D	D	E	L	P	R	G	K	-	-	Y	C	T	V	L	E	
RCA-2	91	E	W	S	D	G	T	K	M	D	Y	S	S	W	Y	R	E	G	F	P	R	R	R	-	-	A	C	A	A	L	E		
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SCA-2	90	E	W	S	D	G	T	K	L	D	Y	G	S	W	Y	R	D	V	L	P	R	R	R	-	-	A	C	A	A	L	E		
OC-17	90	R	W	S	D	G	T	A	P	R	F	A	S	W	H	R	T	A	K	A	R	R	G	-	G	R	C	V	A	A	L	R	
Hu-PSP	79	H	W	S	S	G	S	L	V	S	Y	K	S	W	D	T	G	S	P	S	S	A	N	A	G	Y	C	A	S	L	T		
RCA-1	111	D	S	S	G	F	L	S	W	E	D	D	A	C	S	E	R	N	A	F	I	C	K	C	A	A	A	A	A				
DCA-1	118	D	S	S	G	F	L	S	W	E	D	D	S	C	G	E	R	N	A	F	I	C	K	Y	A	A	A	A					
SCA-1	108	E	S	S	G	F	L	S	W	D	N	D	S	C	G	E	R	N	A	F	I	C	K	C	T	A	A						
ANCA	108	G	S	S	G	F	M	S	W	E	D	N	A	C	S	E	R	N	P	F	V	C	K	Y	S	A	A						
RCA-2	119	D	S	T	D	F	A	S	W	D	T	E	L	C	S	D	R	K	P	F	I	C	E	Y	H	V	V						
DCA-2	118	D	T	A	D	F	A	S	W	D	V	E	L	C	S	D	R	K	P	F	I	C	E	Y	R	T							
SCA-2	118	D	T	T	D	F	A	T	W	D	V	E	L	C	S	D	R	K	P	F	I	C	E	Y	R	T							
OC-17	119	D	E	E	A	F	T	S	W	A	A	R	P	C	T	E	R	N	A	F	V	C	K	A	A	A							
Hu-PSP	108	S	C	S	G	F	K	K	W	K	D	E	S	C	E	K	K	F	S	F	V	C	K	F	K	N							

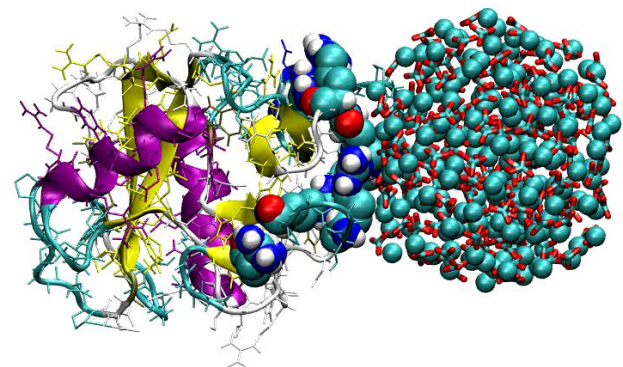
Ovocleidin-17



Reyes-Grajeda, Moreno, Romero;
J. Biol. Chem., 2004

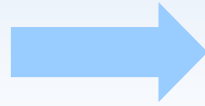
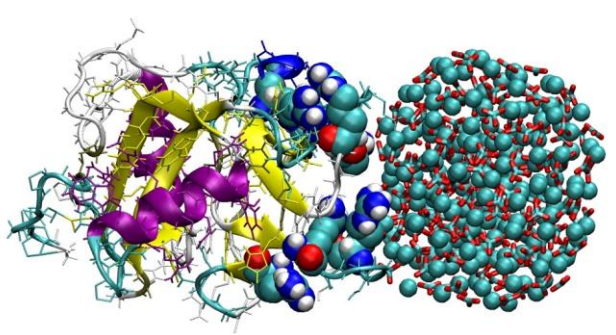
metaDynamics of OC17 + CaCO₃

- 192 & 300 unit nanoparticles
 - 22,000 water molecules
- 20 different protein/nanoparticle orientations
- Select “best” 4 for long meta-dynamics
- Potentials: due to Freeman *et al.*
 - Pavese CaCO₃ ; Amber protein; Tip3p water;
 - cross terms derived from crystals with scaled charges



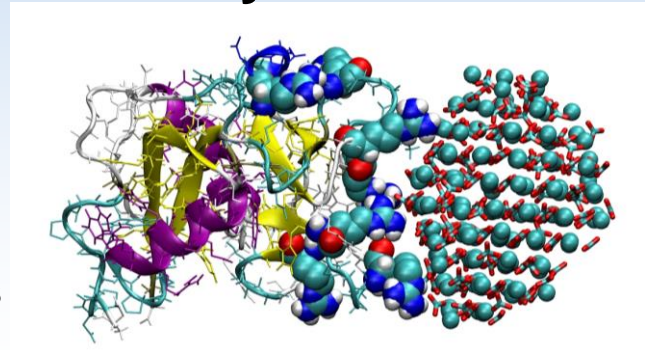
Free energy landscapes

Initial

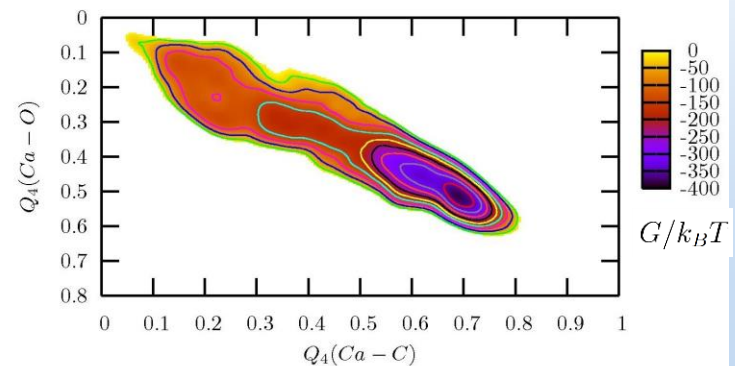
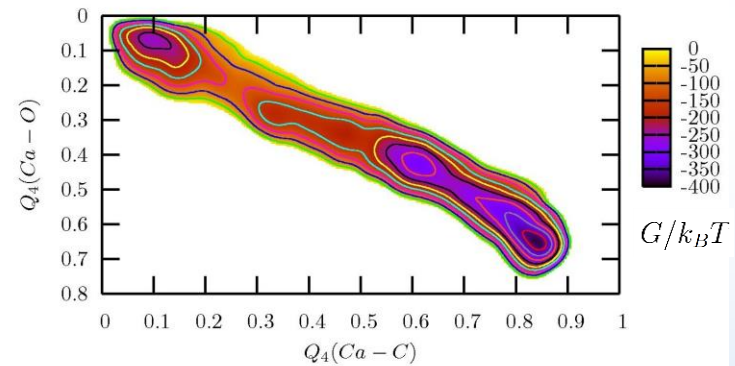


metadynamics

Crystallised

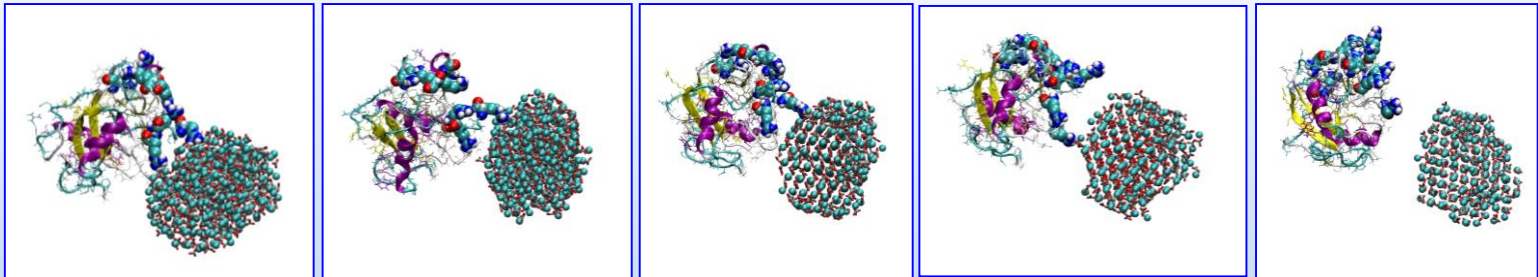


- 192 units of CaCO₃ in water
- 192 units of CaCO₃ bound to Ovocleidin-17 in water

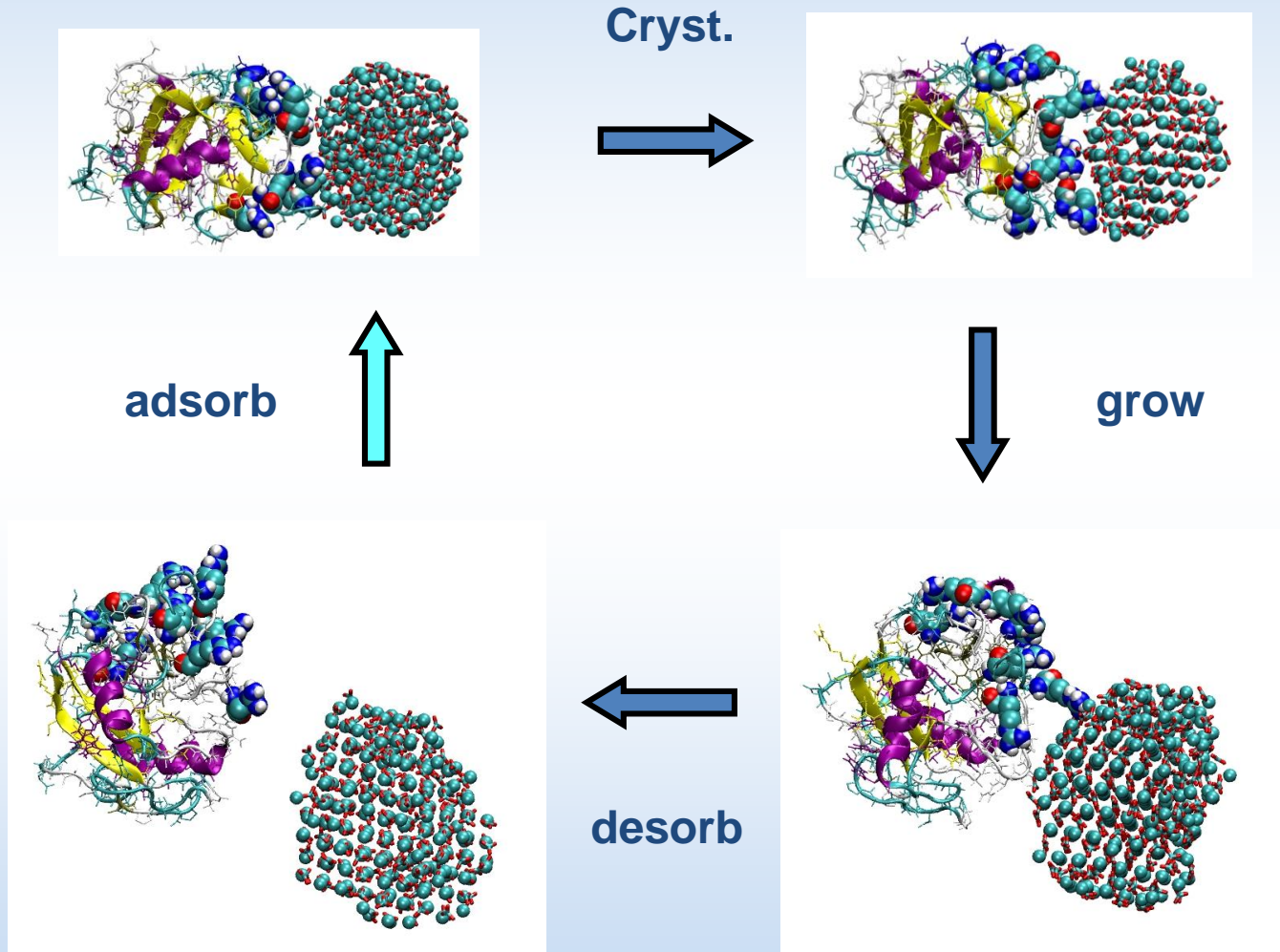


Larger nonaparticles?

- 300 CaCO_3 unit particles don't stay bound on crystallisation
- Mechanism for proteins
 - Bind to small nanoparticles
 - Facilitate transformation to calcite
 - Desorb as crystal grows
- Gives catalytic cycle for polycrystalline mamillary layer

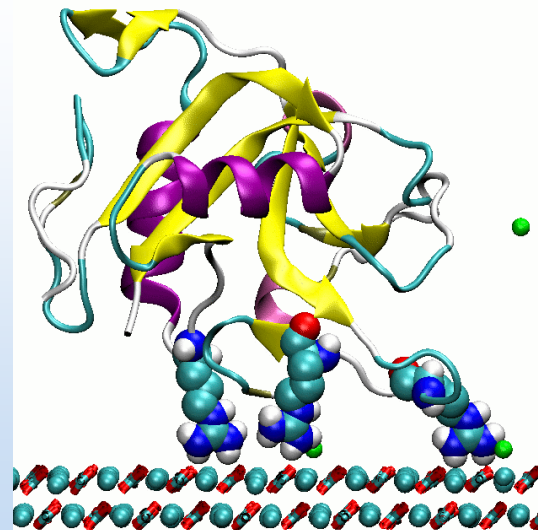
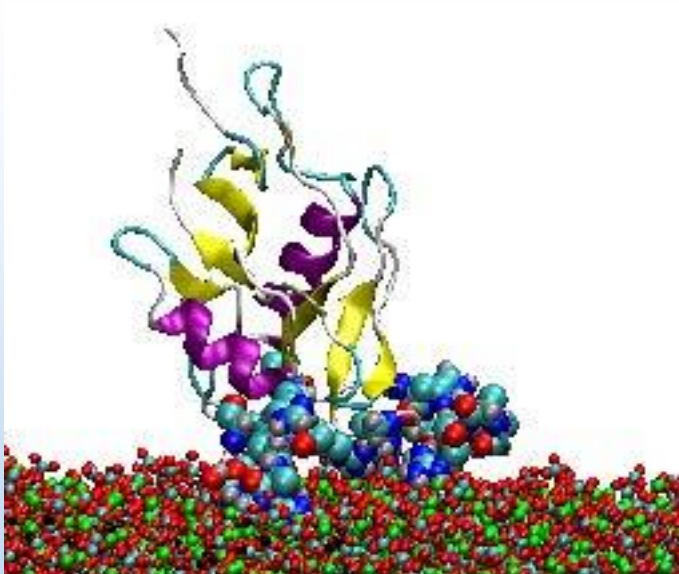


OC17: catalytic cycle?



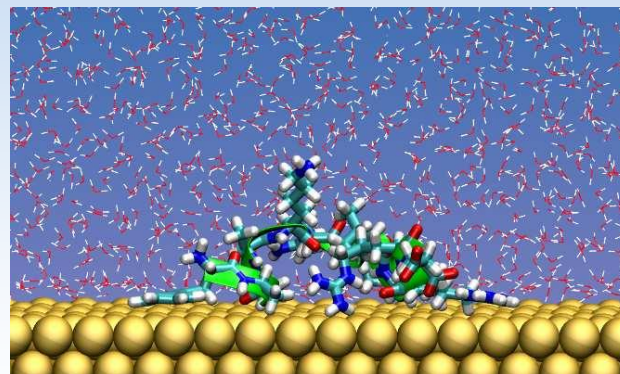
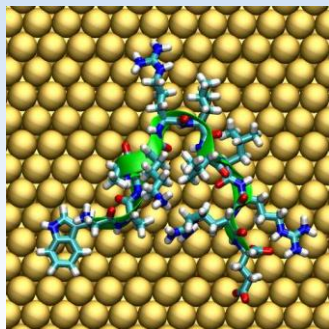
Surface binding & structured water

- Binding is seen to planar surfaces
 - Structured water dominates adsorption energy on large crystalline surfaces
 - Calcite nanoparticles and ACC do not give structured water layer; leads to “flat” adsorption

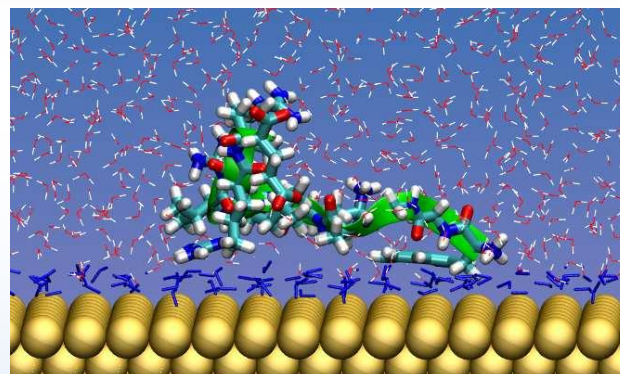
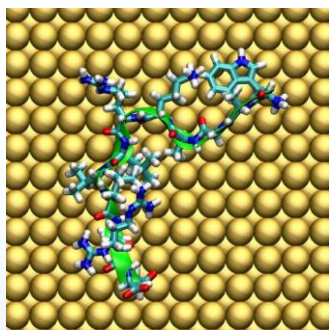


Also seen with AuBP1 on gold

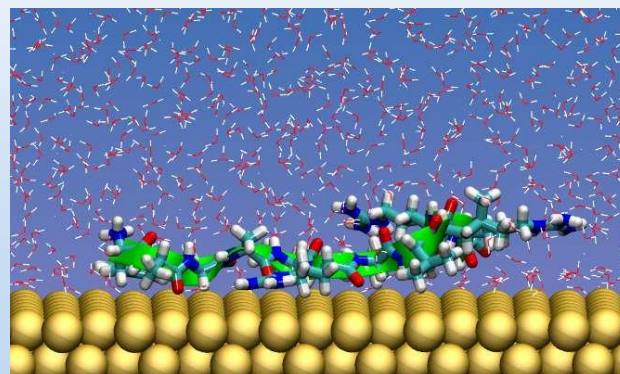
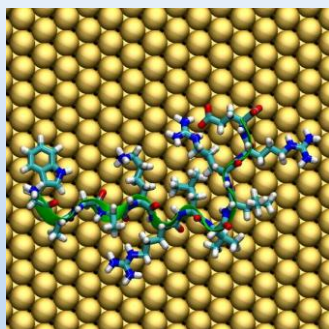
(100) 5x1



(100)



(111)

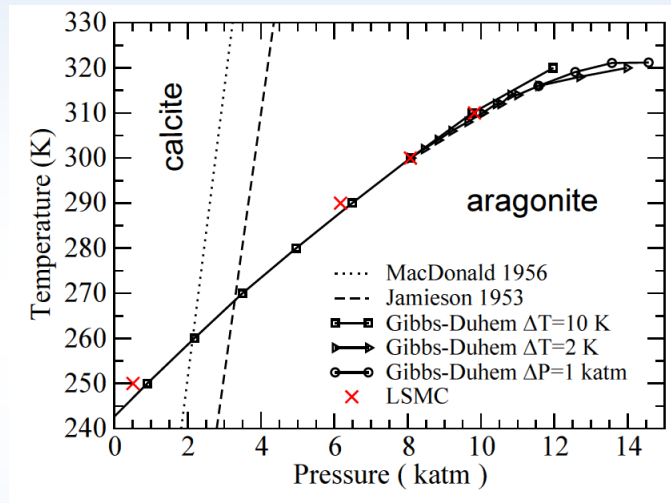


hACC nucleation

- Problem
 - intermolecular potentials
- Controversy

Aragonite?

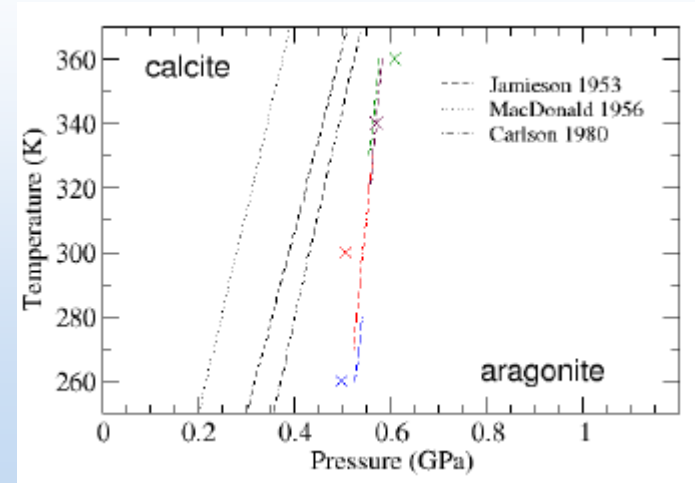
- needs a better potential (Gale & co workers)



Pavese, Catti, Parker, Wall, *Phys Chem. Miner.*, 1996

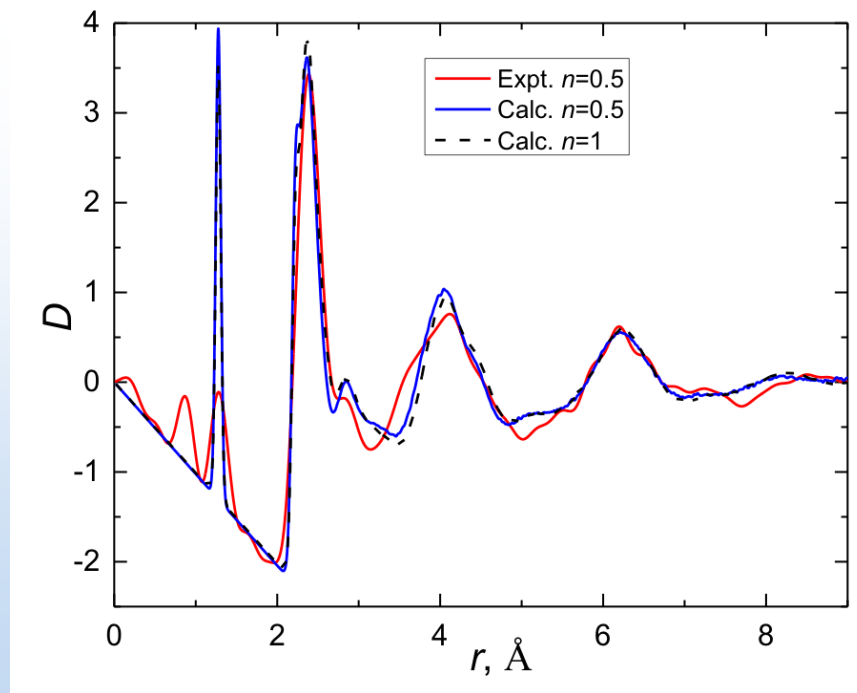


Raiteri, Gale, Quigley, Rodger, *J. Phys. Chem. C*, 2010



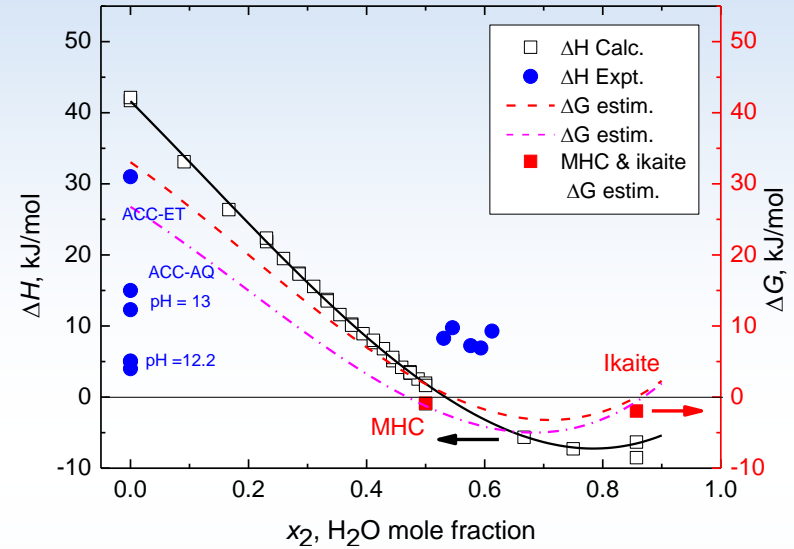
Structure of hACC

- create with several protocols
 - melt monohydrocalcite; melt and dehydrate ikaite; anneal random distribution ...
- Good agreement with expt structure factor

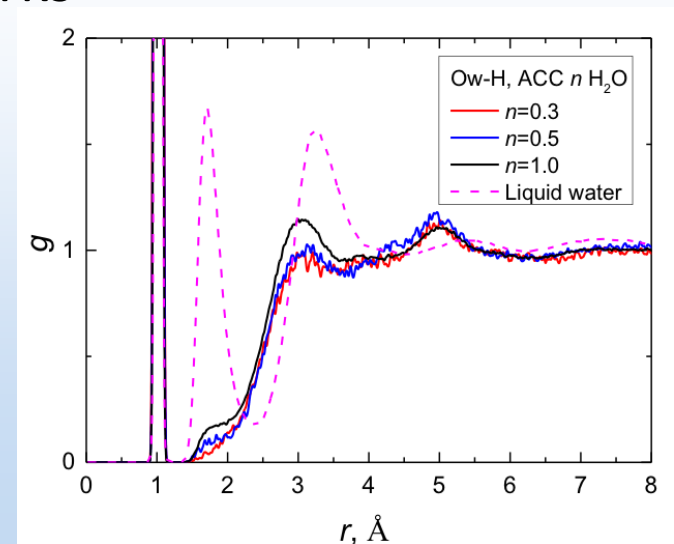
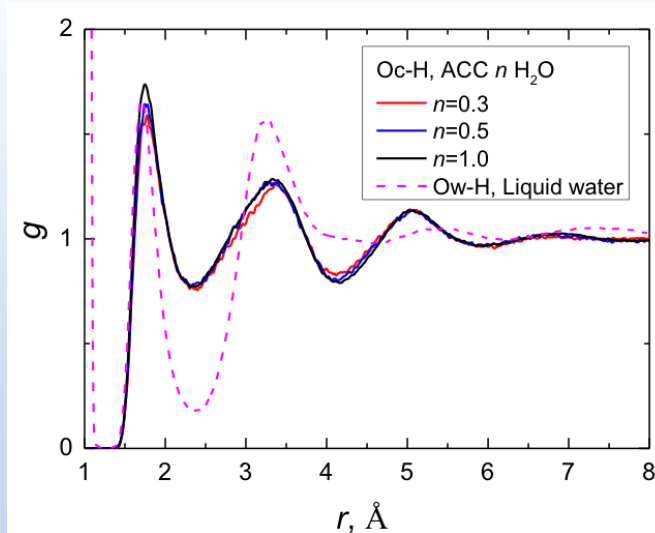


Water in hACC

- zones of stability for hACC

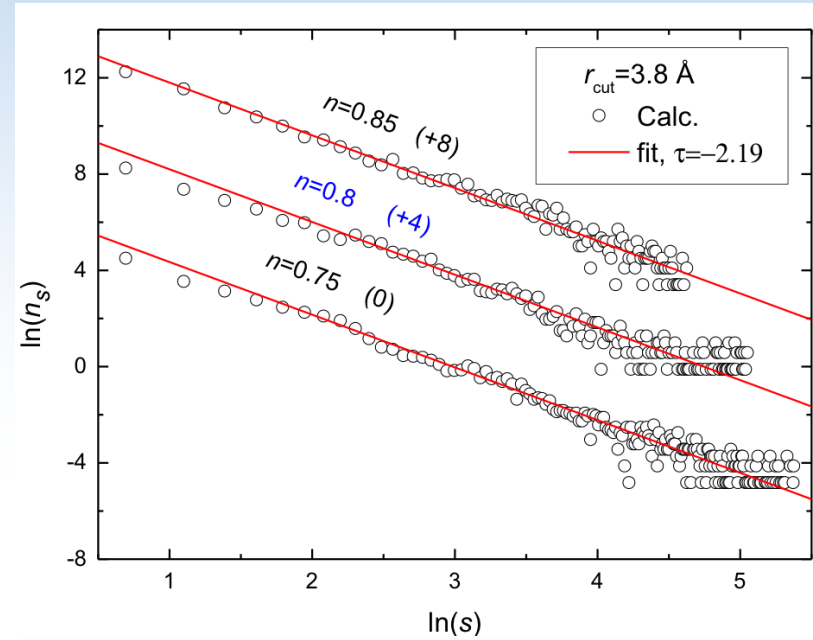
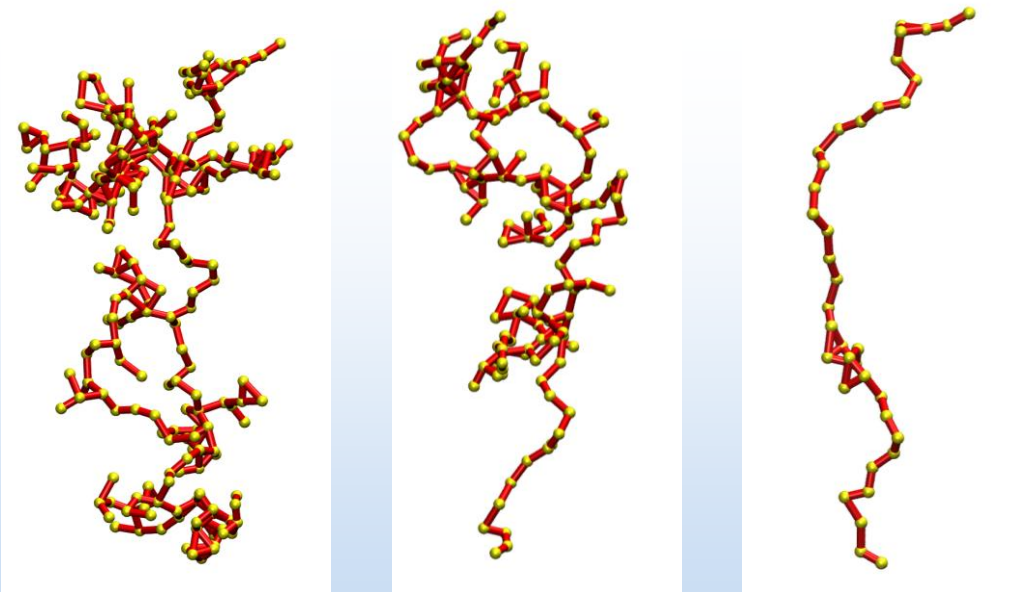


- carbonate-mediated H-bond networks



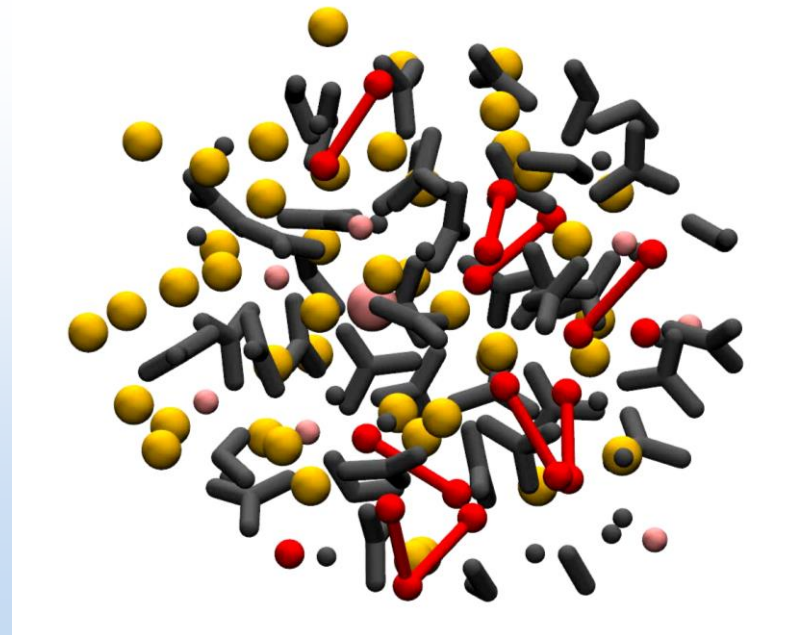
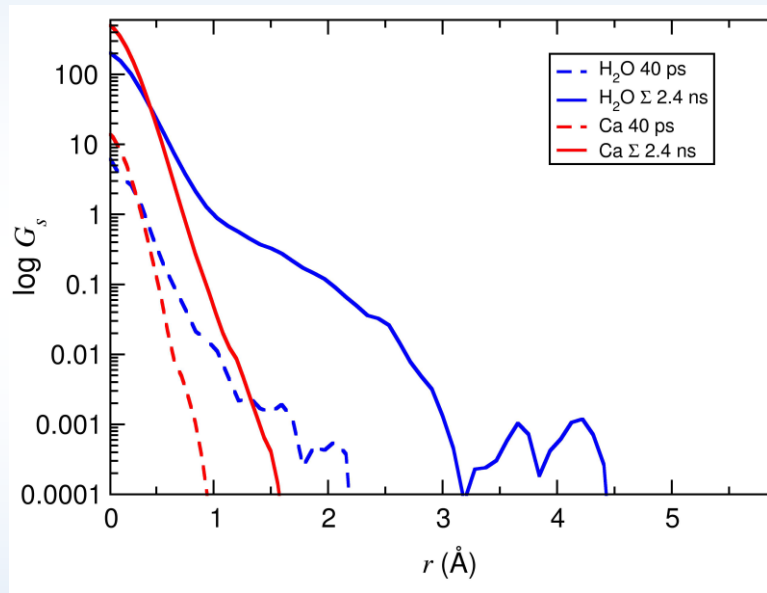
Percolating water clusters

- Universal scaling behaviour
- percolation threshold $\sim n = 0.9$
- underlying hexagonal lattice



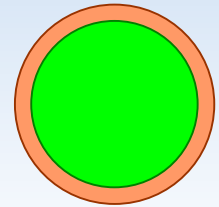
correlated water dynamics

- correlated large hops along chains of water



Classical nucleation

- Activated process:
 - Favourable “bulk” energy
 - Unfavourable interfacial energy
 - Critical “cluster” size

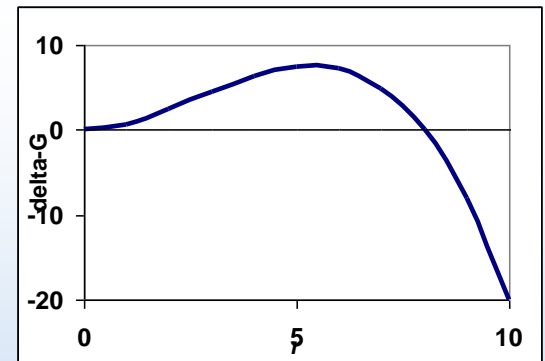


Favourable
Unfavourable

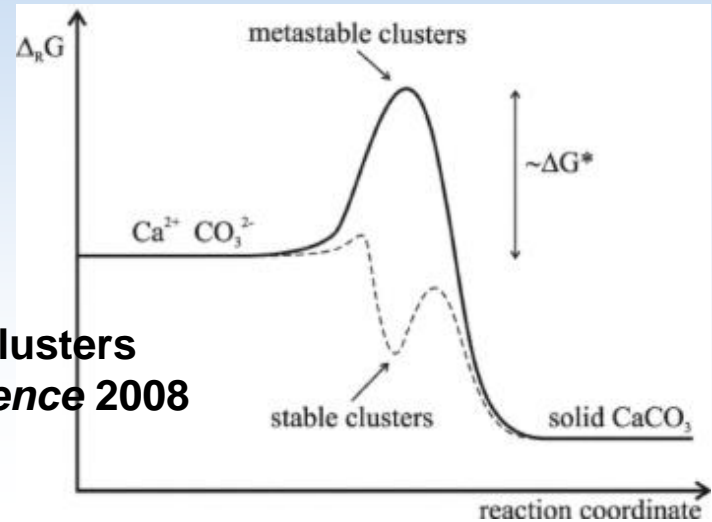
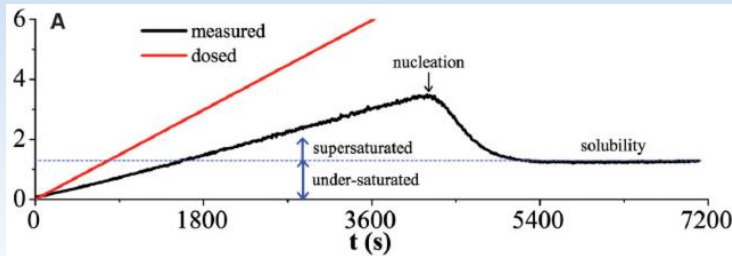
- Classical Nucleation Theory

$$\Delta G = \frac{4}{3} \pi \rho \Delta \mu r^3 + 4 \pi \gamma r^2$$

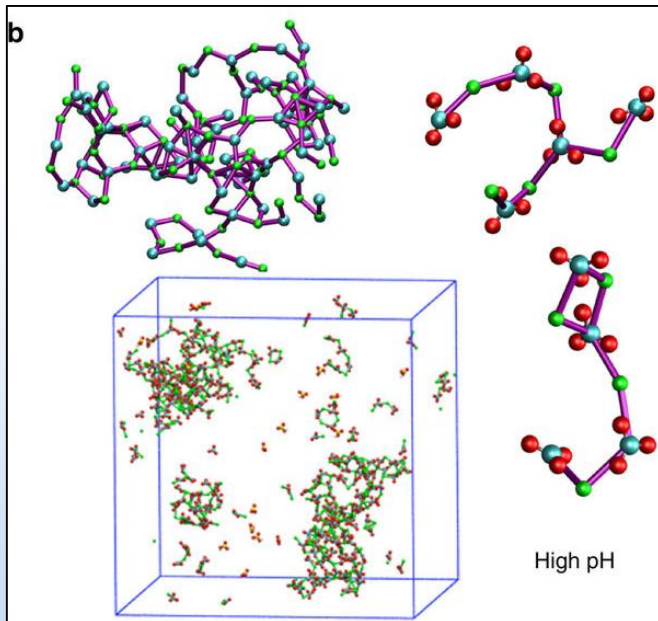
- Fundamentally Stochastic



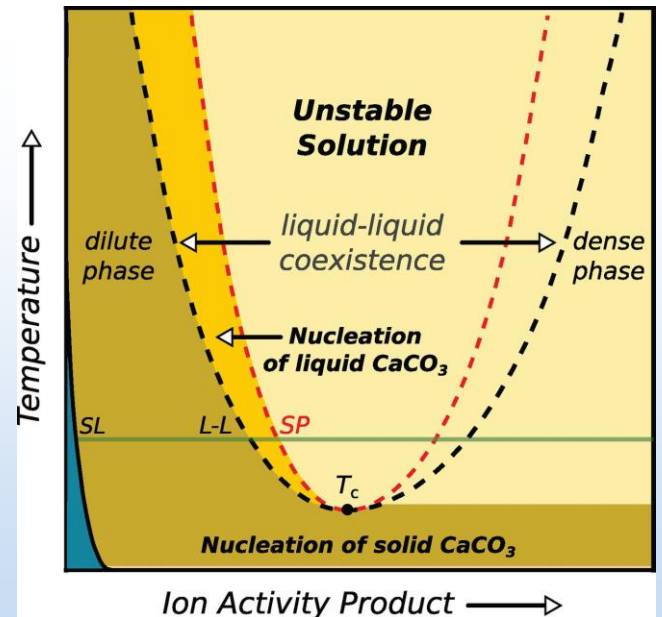
hACC nucleation: classical?



pre-nucleation clusters
Gebauer *et al.*, *Science* 2008



DOLLOP, Raiteri *et al.*,
Nature Comms, 2011

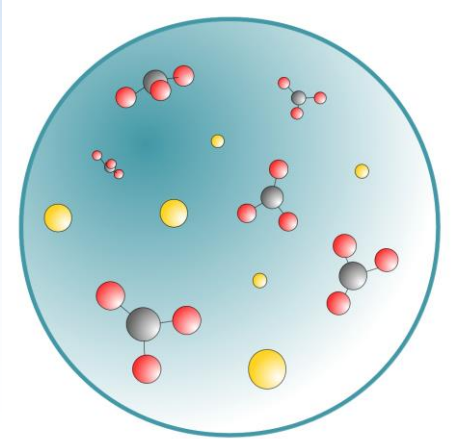


dense liquid phase
Wallace *et al.*, *Science*, 2014

Nature of the clusters?

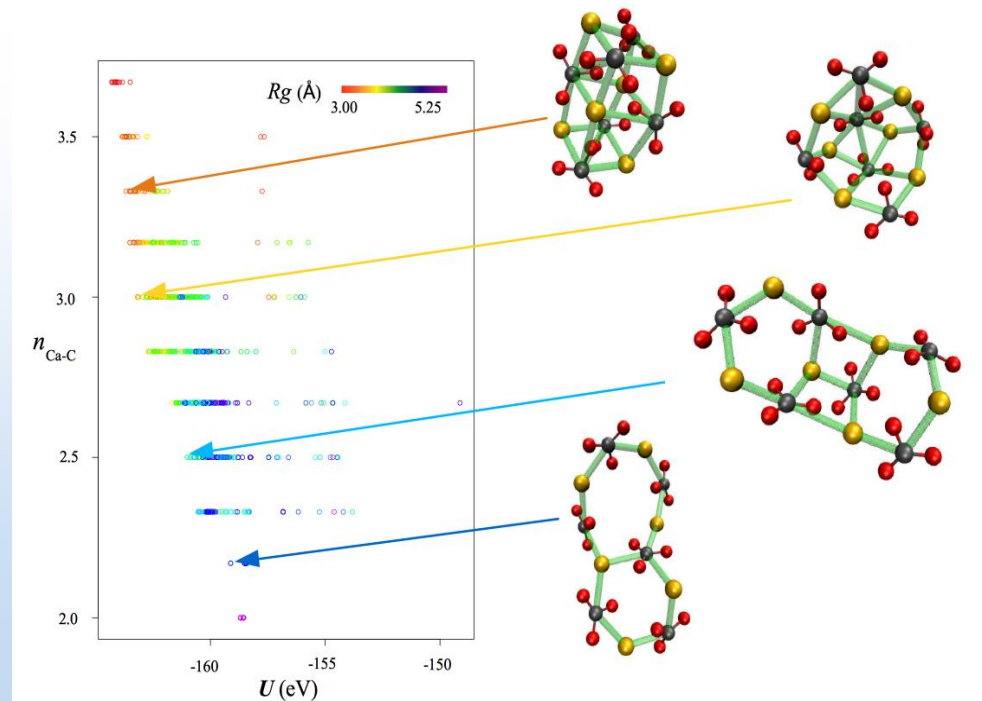
- spontaneous clusters dominate?
- Random Structure searches
 - random arrangement of ions; minimise; repeat
- Intelligent Water Drop algorithm
 - nature-inspired global optimisation
 - erosion of soil to define river valleys
 - applications to binary LJ mixtures and Janus particles
 - (not yet carbonates)

Random Structure Searches

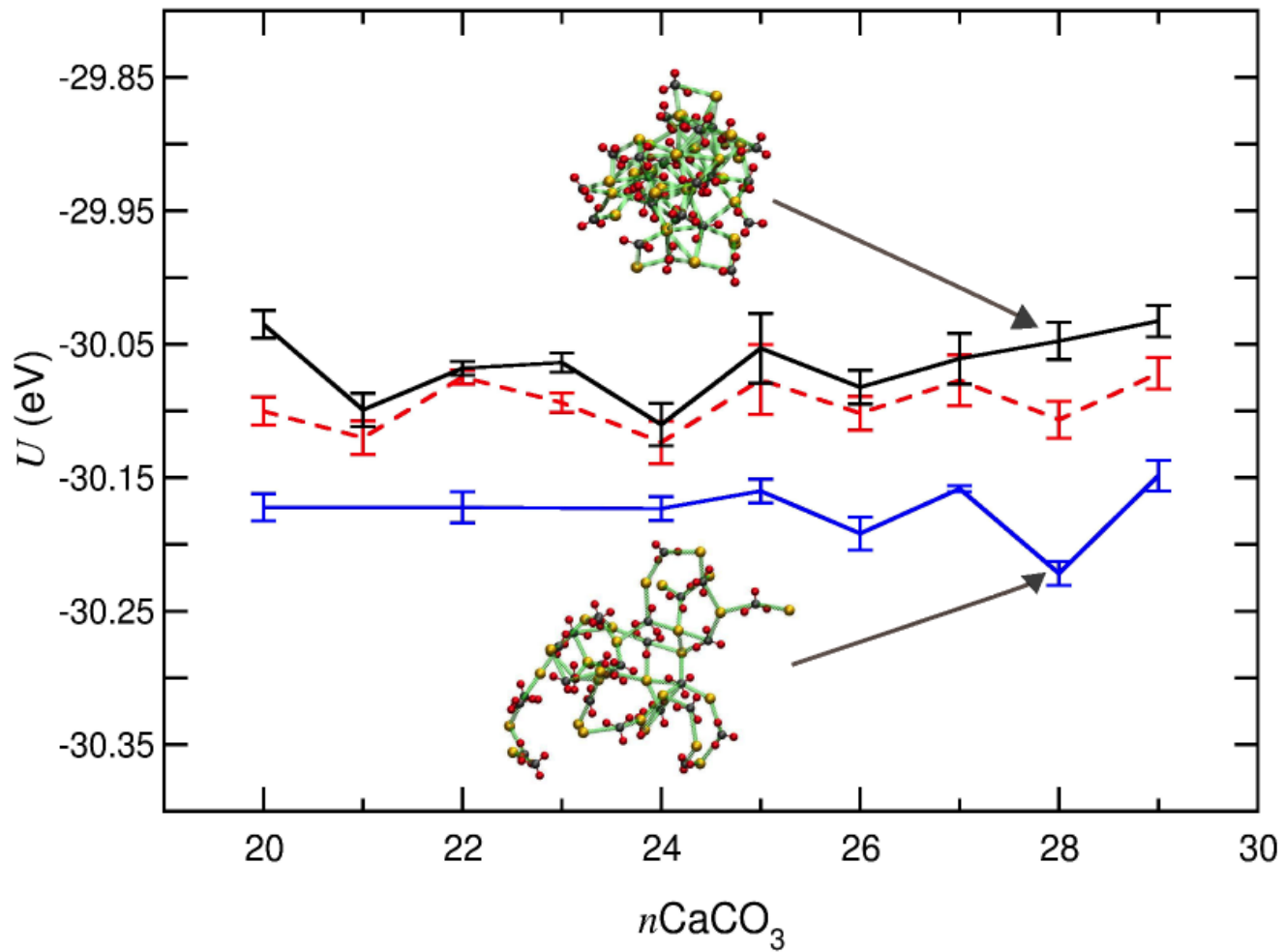


- random arrangement of ions within sphere
- Conjugate Gradient optimisation
- ~10,000 initial structures for each cluster size (1–40 formula units)

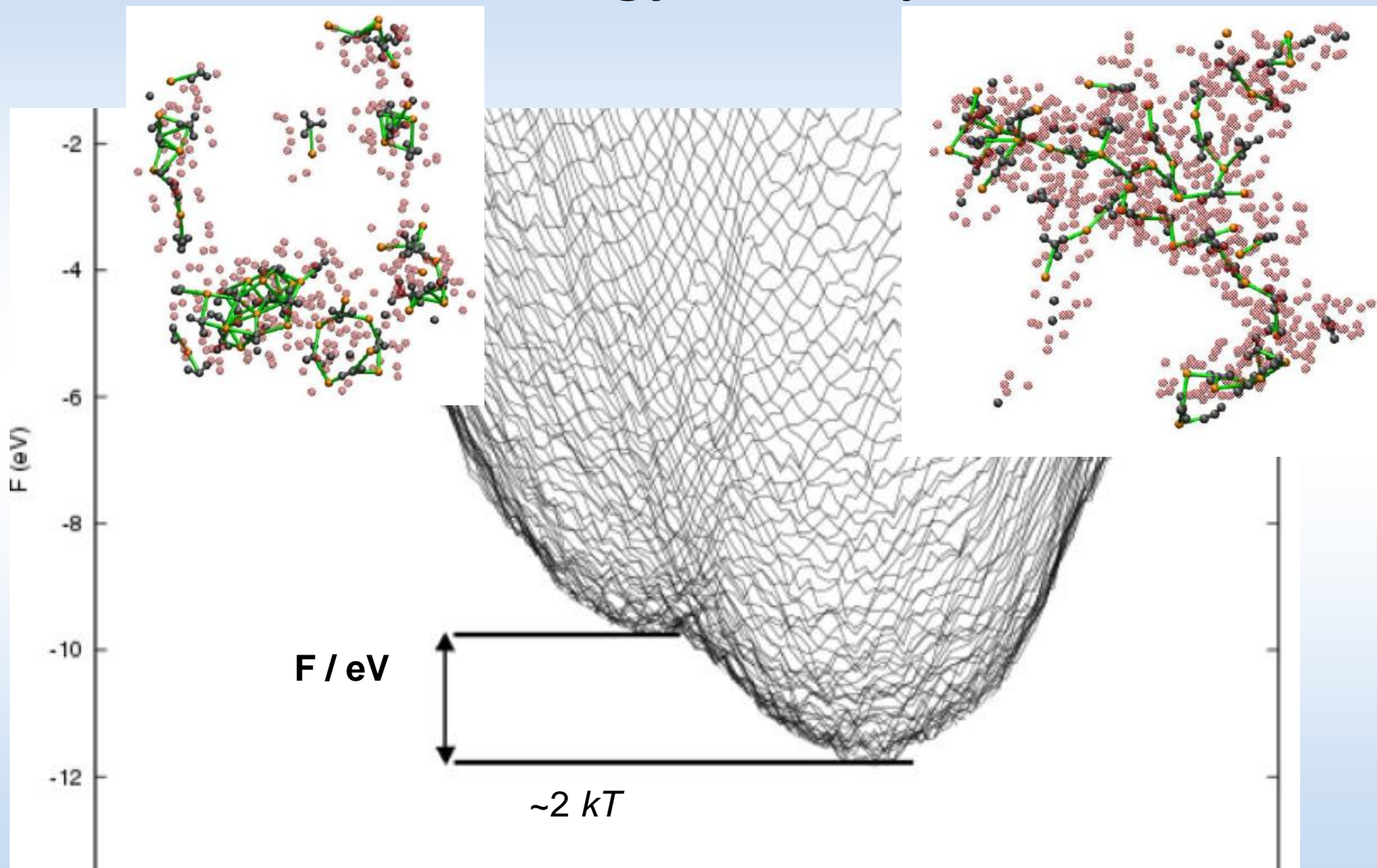
- variety of cluster properties
 - e.g. energy vs radius of gyration



Energetics of solvated clusters

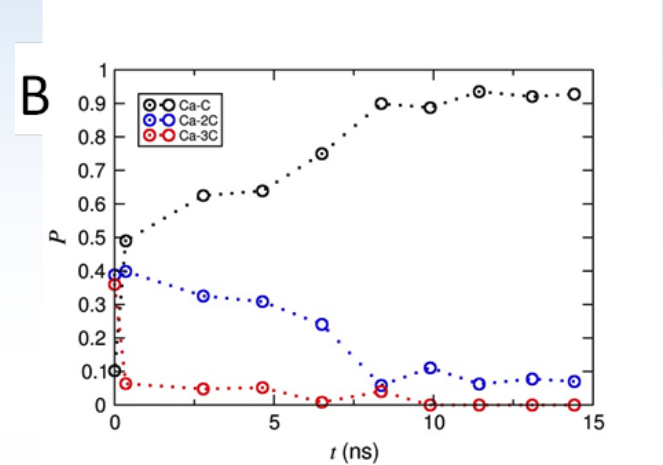
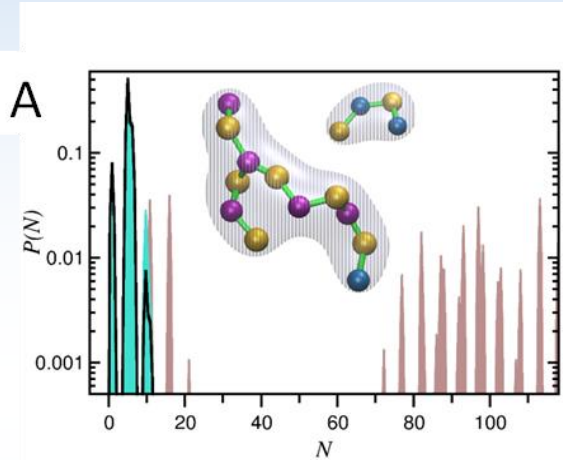


Free energy Landscape

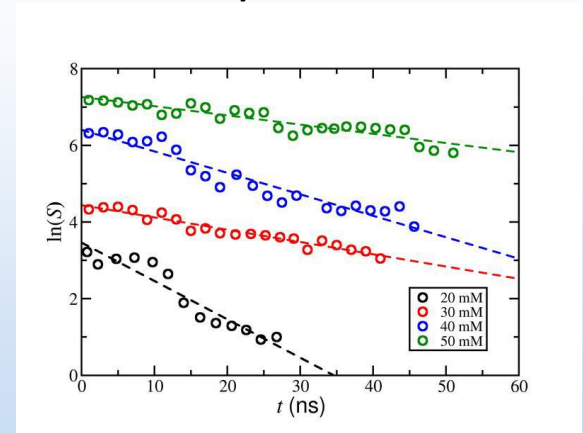
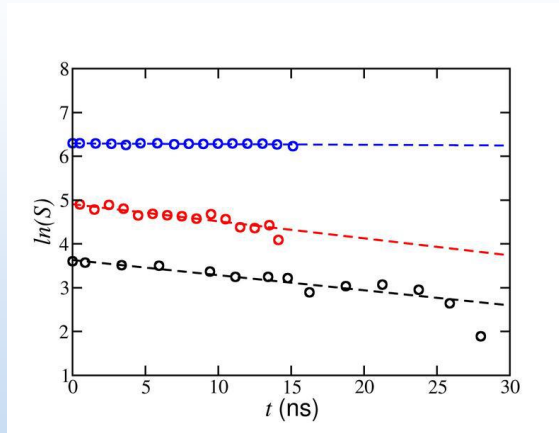


Long timescale MD: 20–50 mM

- “slow” dissolution: convergence from dispersion or initial clusters



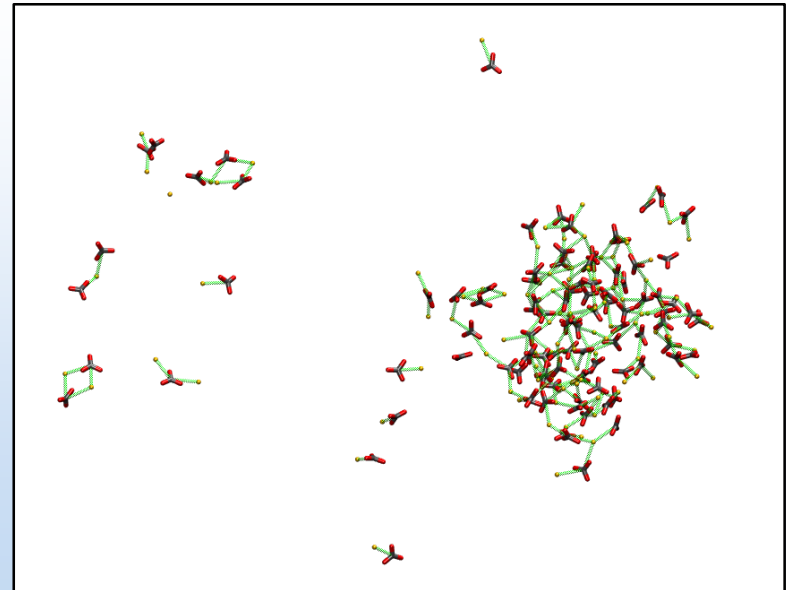
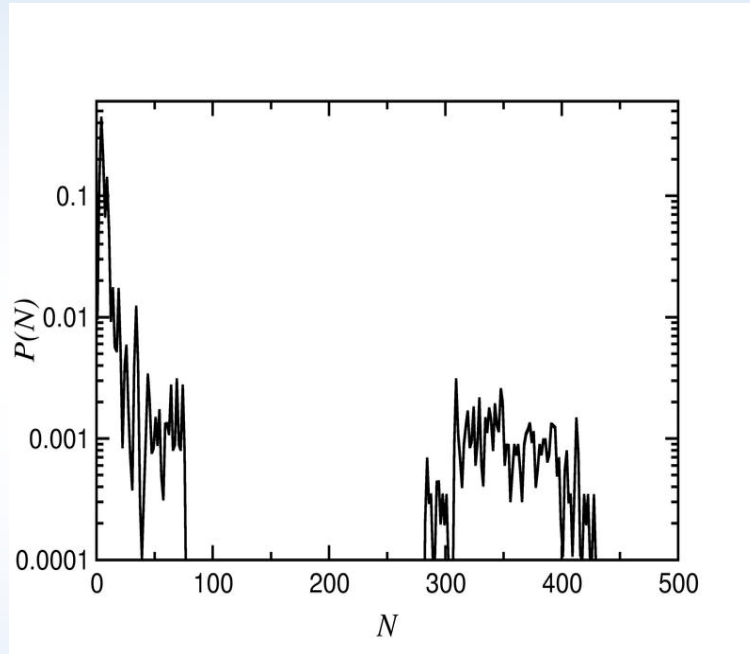
- dissolution slows with cluster size (concentration)



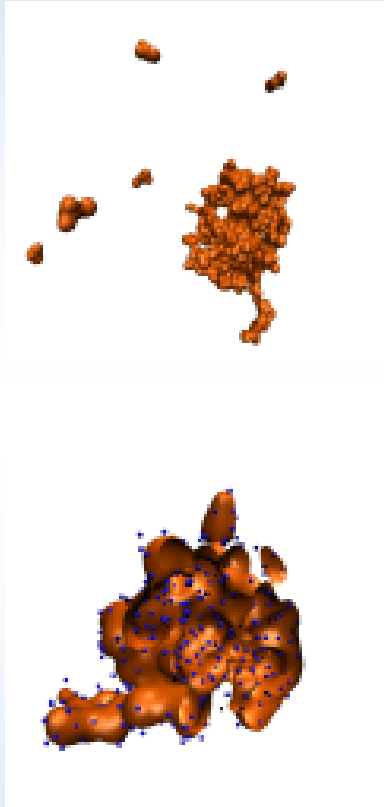
- (requires free energy barrier)

Long timescale MD: *ca.* 0.5 M

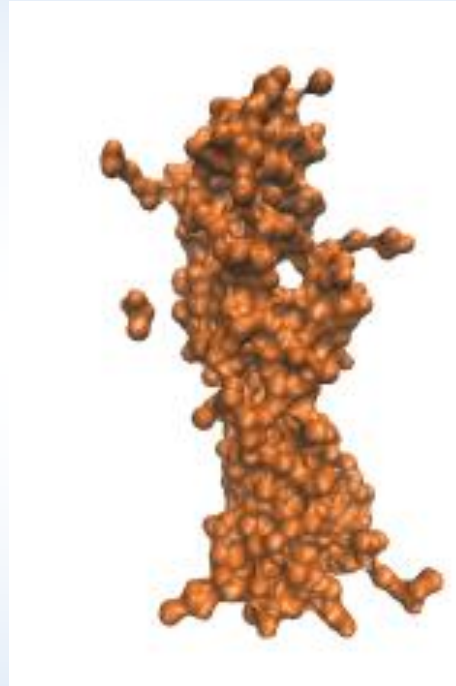
- stable “DOLLOP” behaviour; large clusters



And higher concentrations: 0.5–1.5 M



0.57 M
stable clusters
DOLLOP



1.1 M
infinite, liquid like
(phase transition?)

Conclusions

- Source of new mechanistic insight
 - kinetic inhibition; eggshell proteins
 - does your predictive model have the right features?
- Can capture some elements of intrinsic complexity
 - inhibitor polydispersity; nanoparticle structure
 - need for intelligent sampling
- Progress through data dimensionality reduction
 - excellent methods for a handful of dimensions
 - need to identify key dimensions
 - need to recognise and avoid pitfalls of too few (or wrong) dimensions