

# **Nested Sampling for computational thermodynamics**

**(how to calculate phase diagrams without the fuss)**

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# Nested Sampling

To have an unbiased sampling technique, that can be automated:  
not specific to a give part of the PES, where no prior knowledge of structures are necessary.

- Nested sampling algorithm
- Calculate the phase diagram with nested sampling
  - metals, alloys
  - clusters
  - molecules
- Visualise the PES with nested sampling

# Nested Sampling

John Skilling, 2004, Bayesian statistics

“...to sample probability densities in **high-dimensional** spaces where the regions contributing most of the probability mass are **exponentially localised**.”

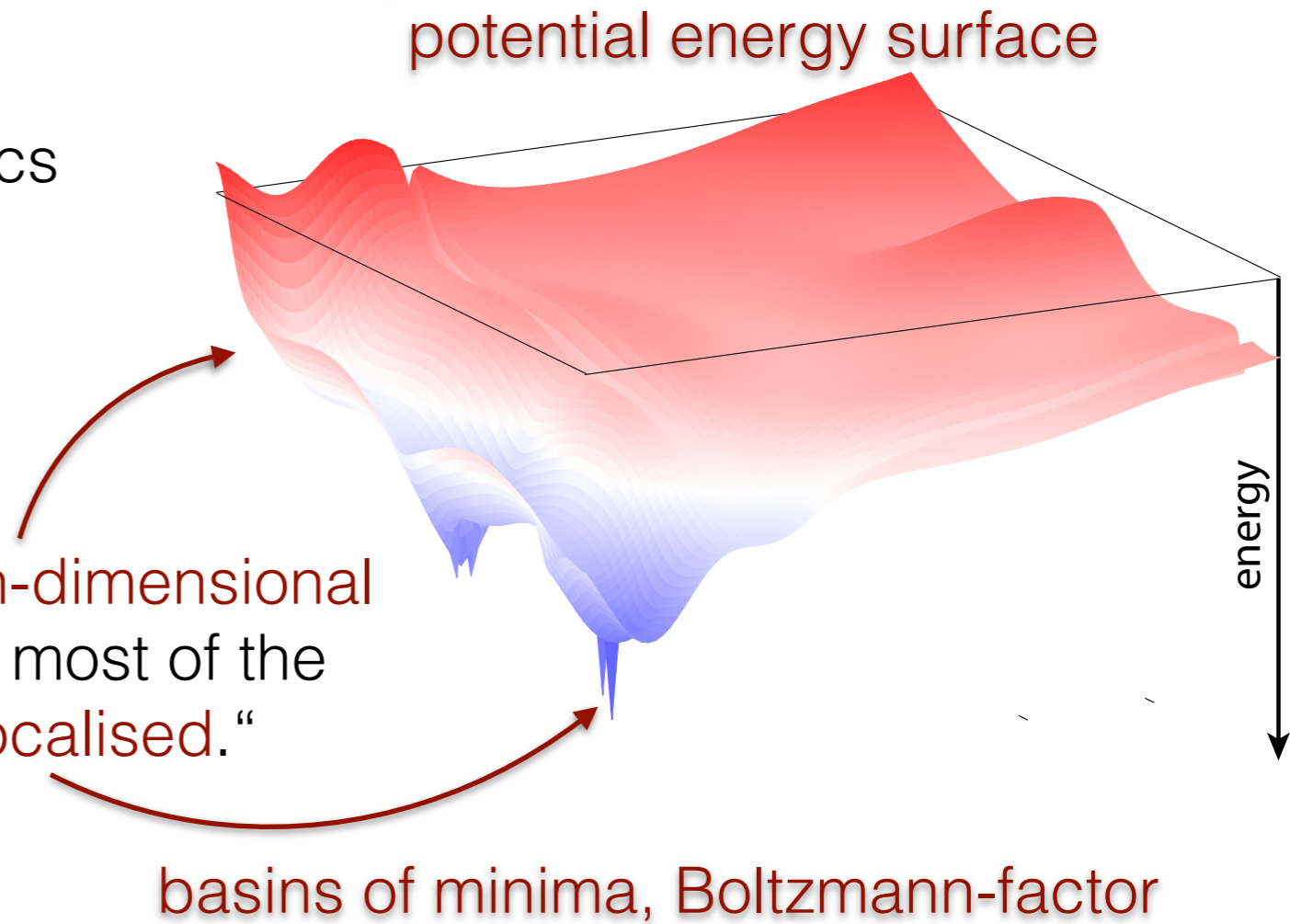
evidence ( $Z$ )      likelihood ( $L$ )      proportion of the prior distribution

$$Z = \int_0^1 L(X) dX$$

# Nested Sampling

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evidence ( $Z$ )

likelihood ( $L$ )

proportion of the prior distribution

$$Z = \int_0^1 L(X) dX$$

partition function

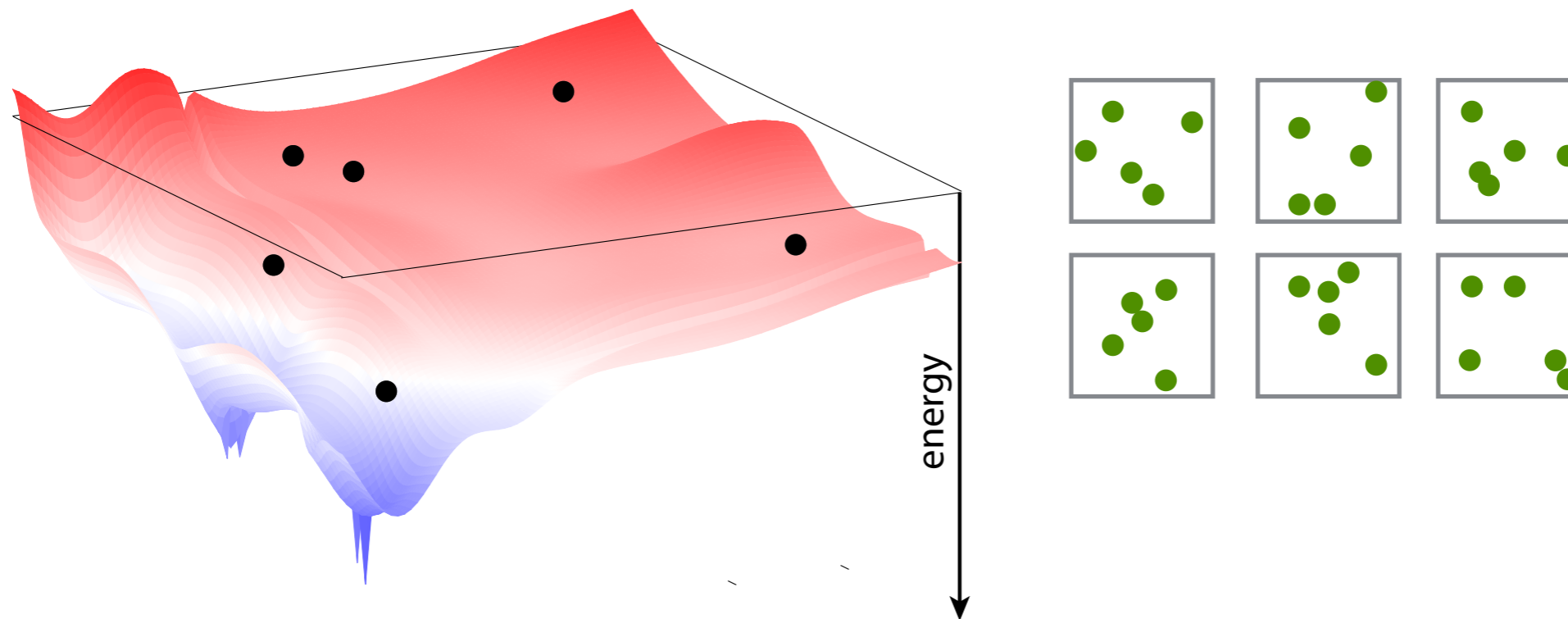
Boltzmann-factor  
(energy)

phase space volume  
(prior distribution is uniform)

# Nested Sampling Algorithm

Iterative algorithm, starting from the “top” (ideal gas) and going towards the “bottom” (global minimum), through a series of nested energy “contours”.

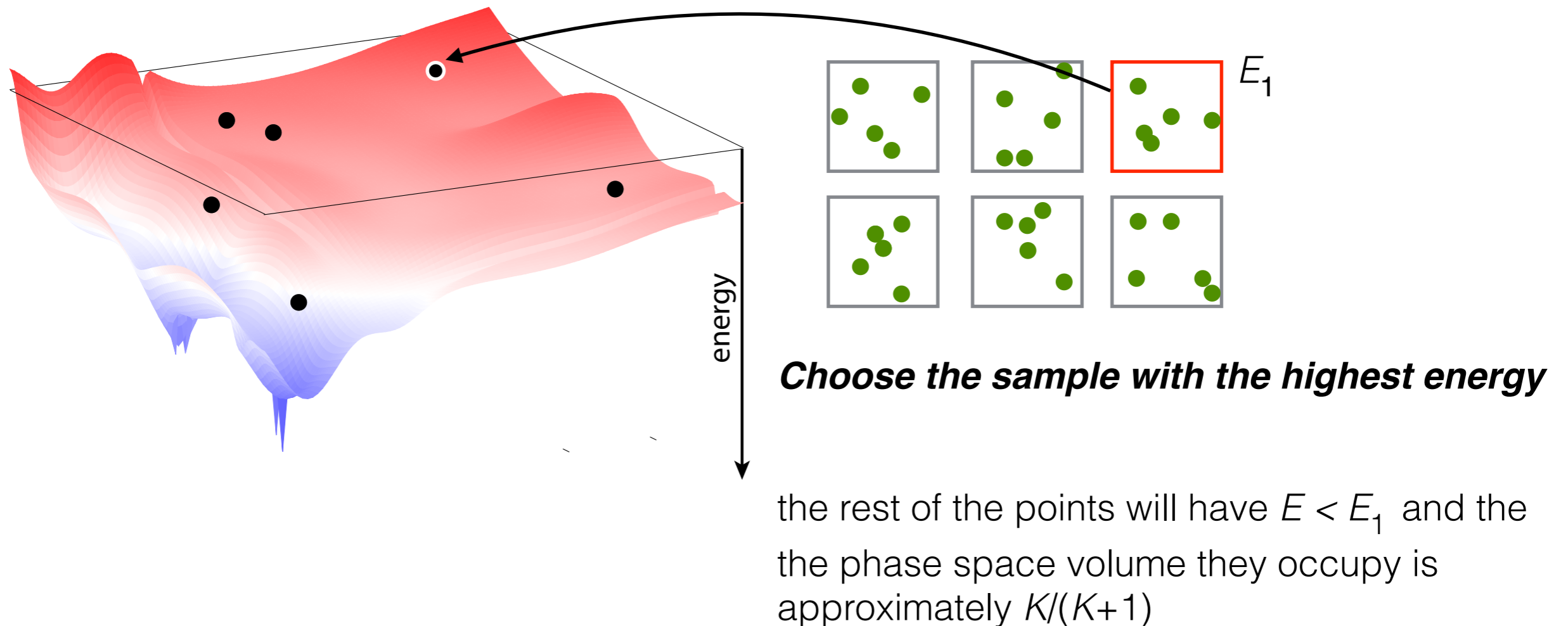
Generate  $K$  random samples uniformly in the total phase space volume



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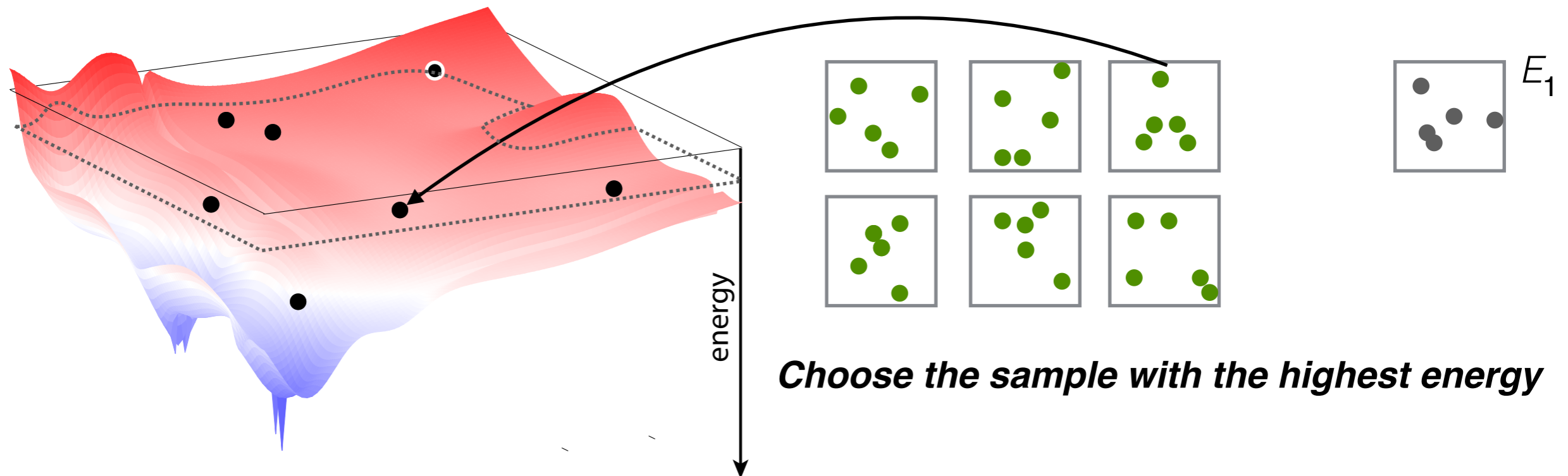
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**Choose the sample with the highest energy**

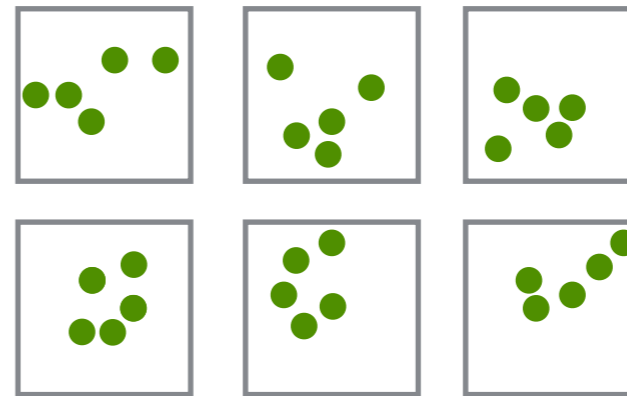
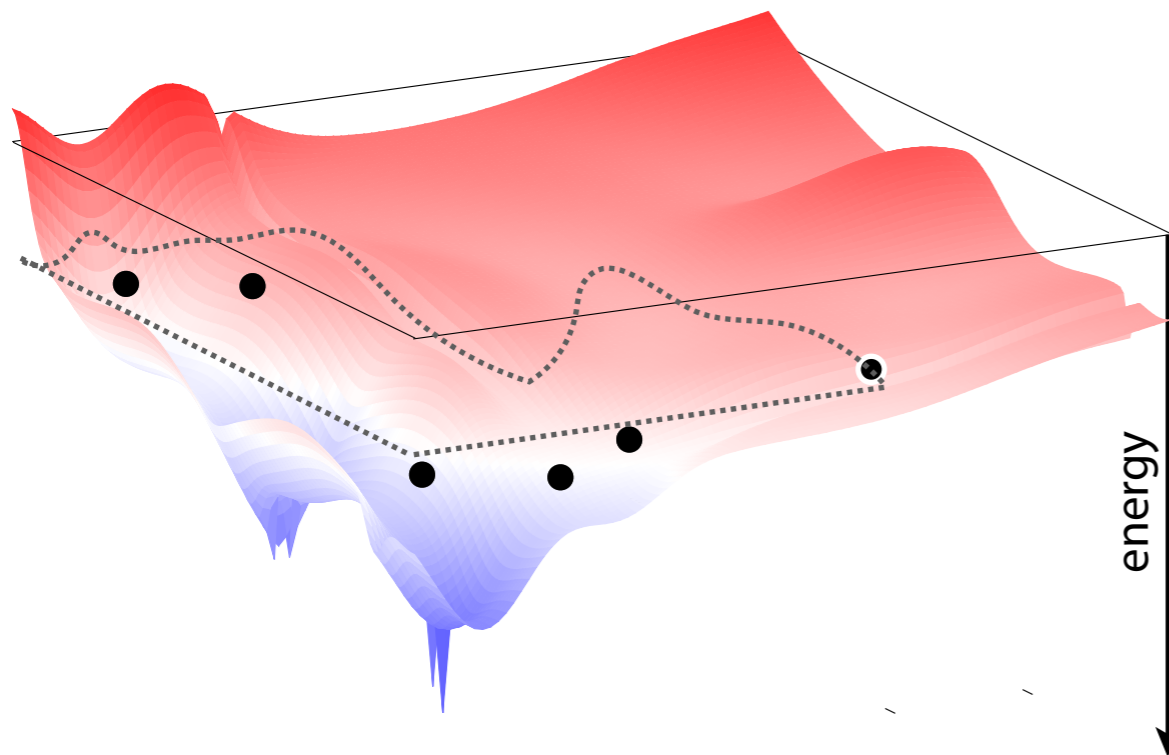
the rest of the points will have  $E < E_1$  and the phase space volume they occupy is approximately  $K/(K+1)$

**Generate a new sample uniformly with  $E < E_1$**

# Nested Sampling Algorithm

Iterative algorithm, starting from the “top” (ideal gas) and going towards the “bottom” (global minimum), through a series of nested energy “contours”.

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*Choose the sample with the highest energy*

*Generate a new sample uniformly with  $E < E_i$*

***Repeat this iteration many times...***

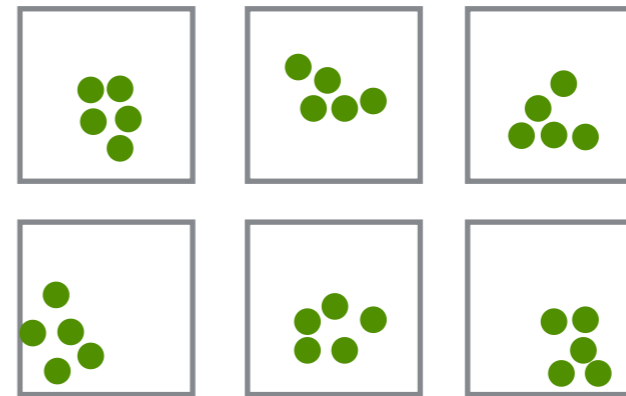
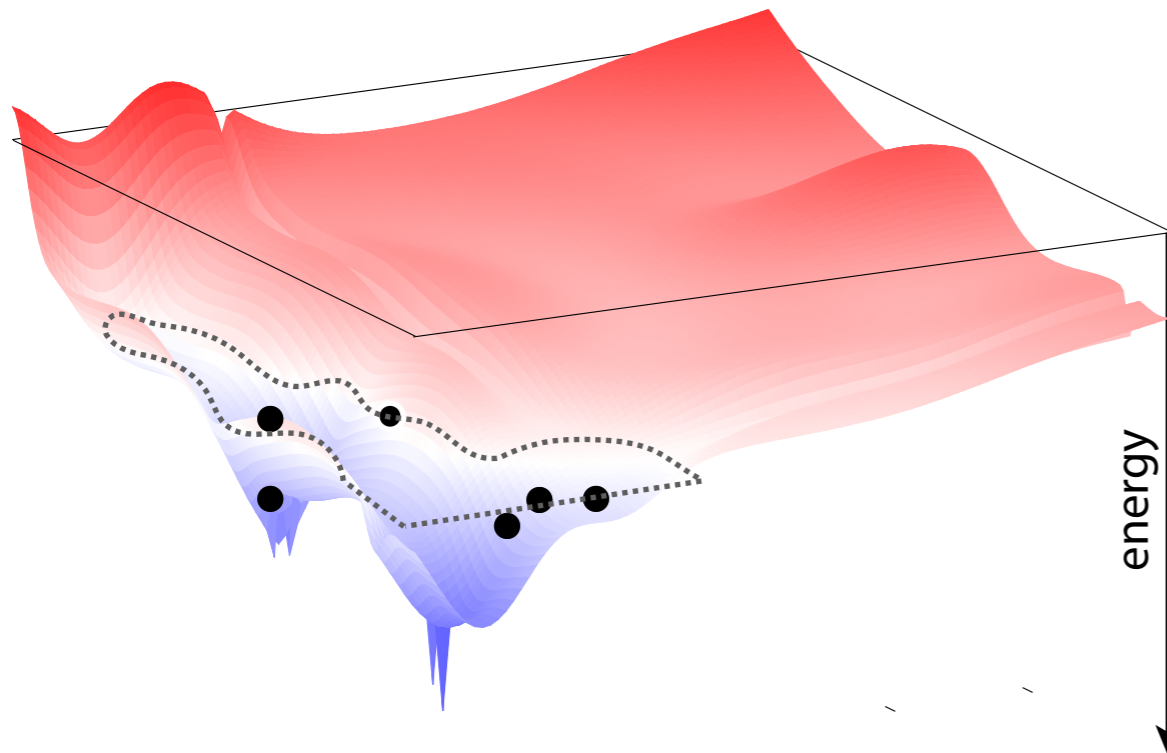
at the  $i^{\text{th}}$  iteration the samples will have

$E < E_i$  and phase space volume  $\sim [K/(K+1)]^i$

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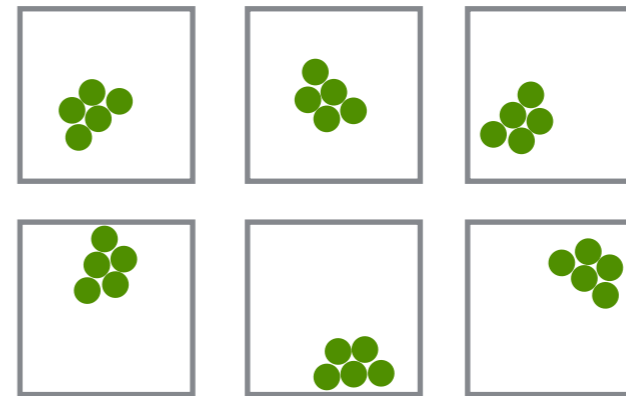
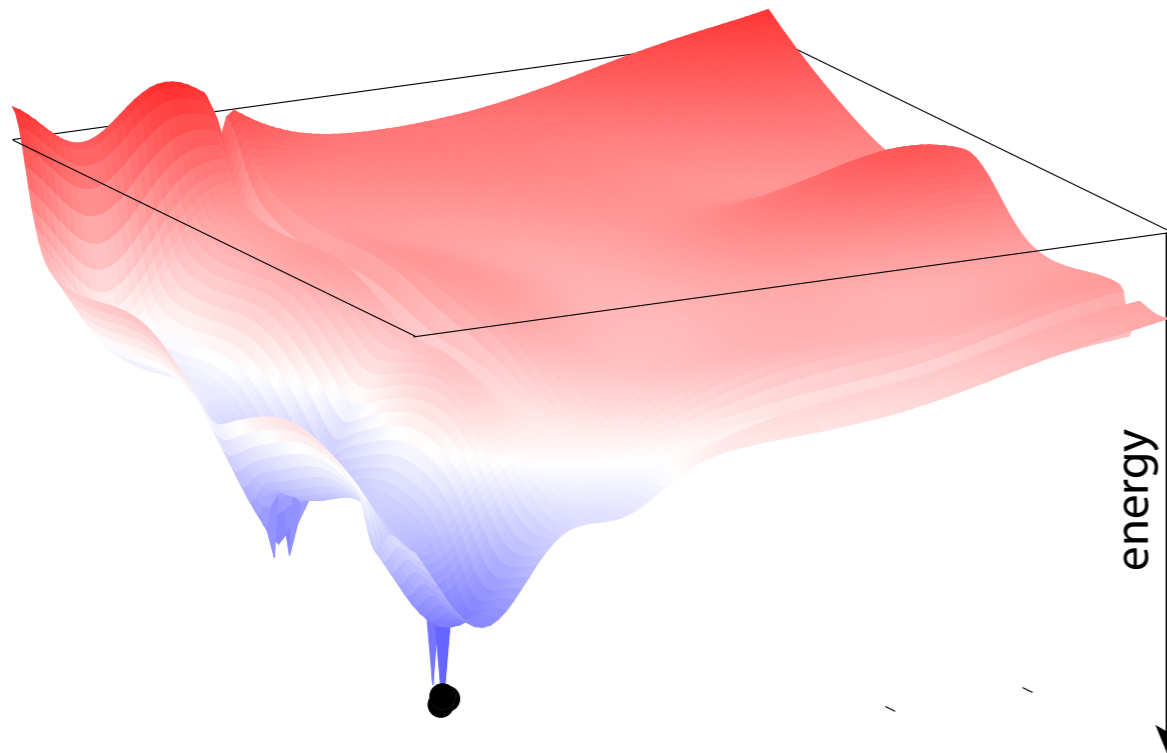
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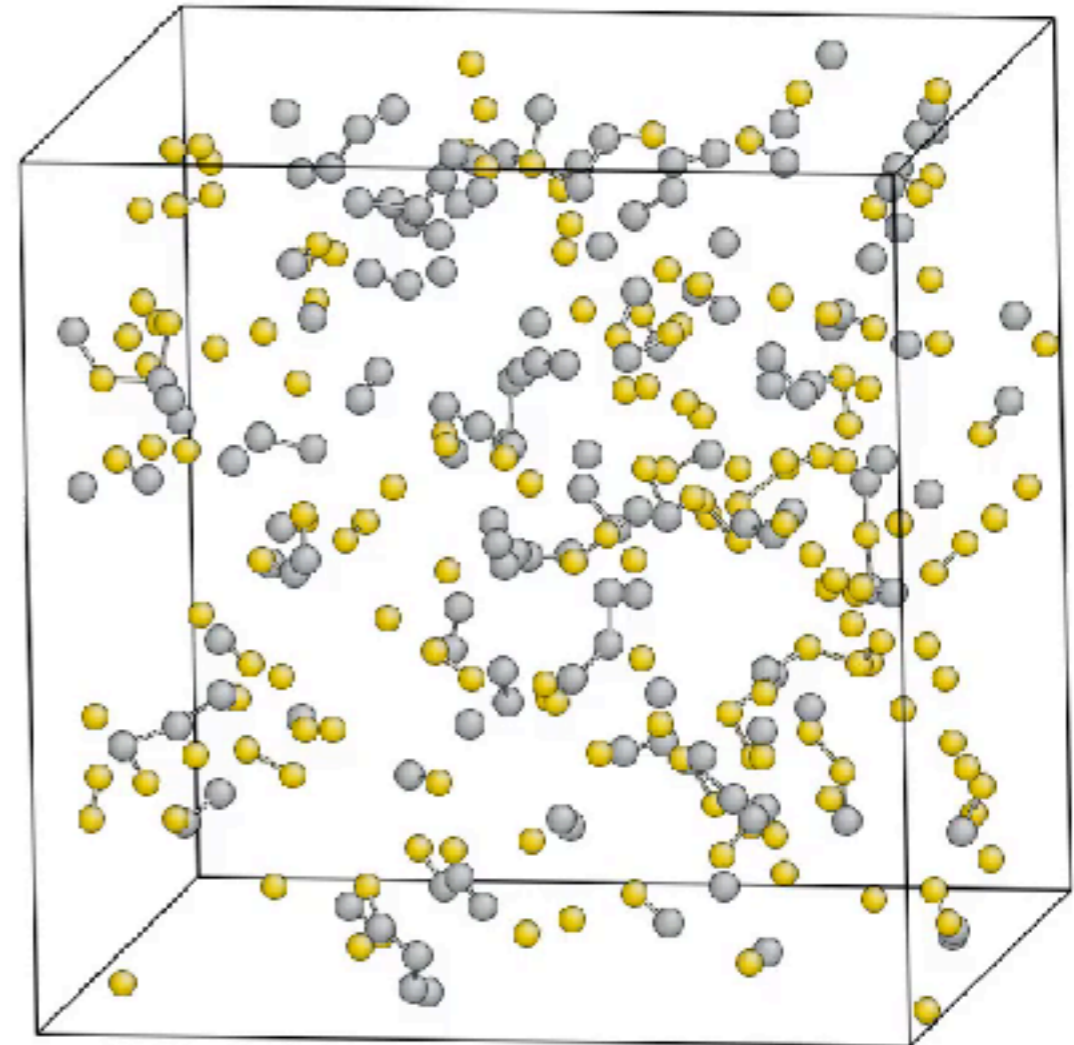
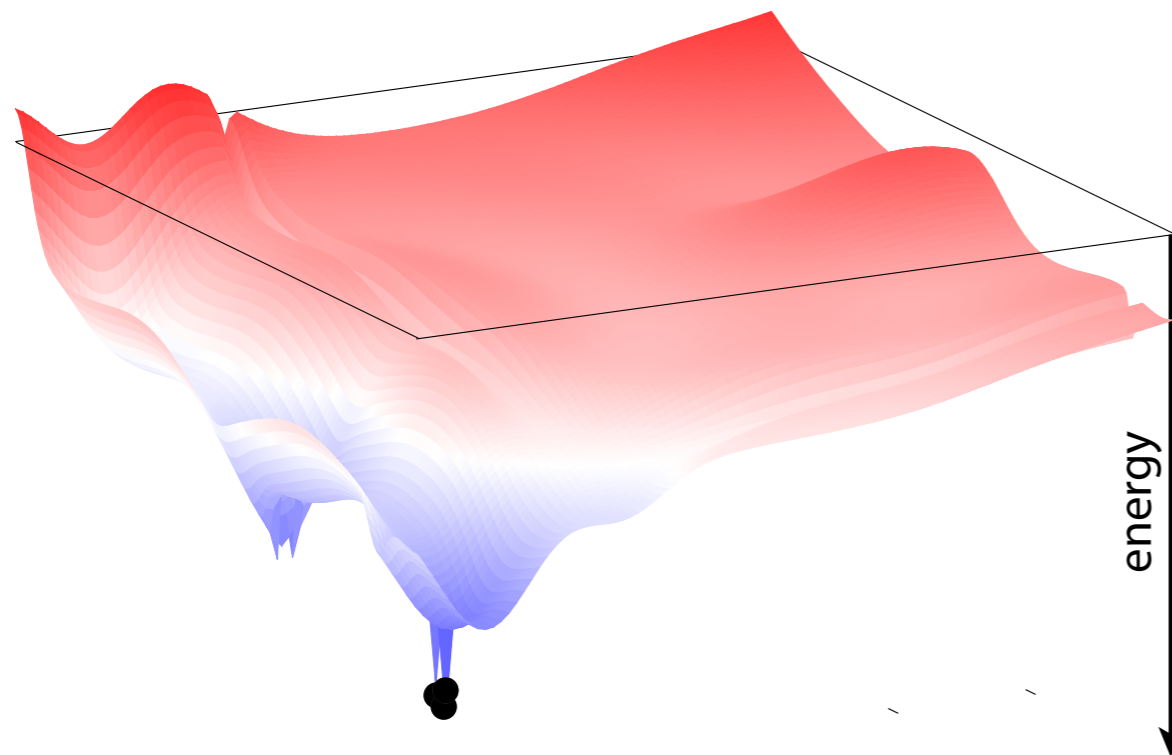
at the  $i^{\text{th}}$  iteration the samples will have  
 $E < E_i$  and phase space volume  $\sim [K/(K+1)]^i$

***Until the “bottom” is reached.***

# Nested Sampling Algorithm

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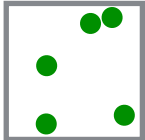
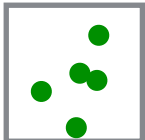



Generate  $K$  random samples uniformly in the total phase space volume



309 atoms (Cu and Pt) in fixed cell:  
cluster


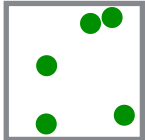
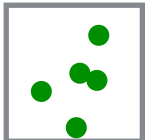



# Nested Sampling Algorithm

At the end we have a set of energies  $\{E_i\}$ , corresponding volumes  $\{[K/(K+1)]^i\}$  and configurations

$E_1$	$K/(K+1)$	
$E_2$	$K/(K+1) * K/(K+1)$	
$E_3$	$K/(K+1) * K/(K+1) * K/(K+1)$	
...		
$E_i$	$\{K/(K+1)\}^i$	
$E_{i+1}$	$\{K/(K+1)\}^{i+1}$	
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$E_n$	$\{K/(K+1)\}^n$	

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**Partition function:**

$$\begin{aligned}
 Z(\beta) &= Z_{\mathbf{p}} \sum_i w_i e^{-\beta E_i} \\
 &= Z_{\mathbf{p}}(\beta) \sum_i \left[ \left( \frac{K}{K+1} \right)^i - \left( \frac{K}{K+1} \right)^{i+1} \right] e^{-\beta E_i}
 \end{aligned}$$

**for any temperature**

# Nested Sampling Algorithm

At the end we have a set of  $\{E_i\}$  and corresponding volumes  $\{[K/(K+1)]^i\}$

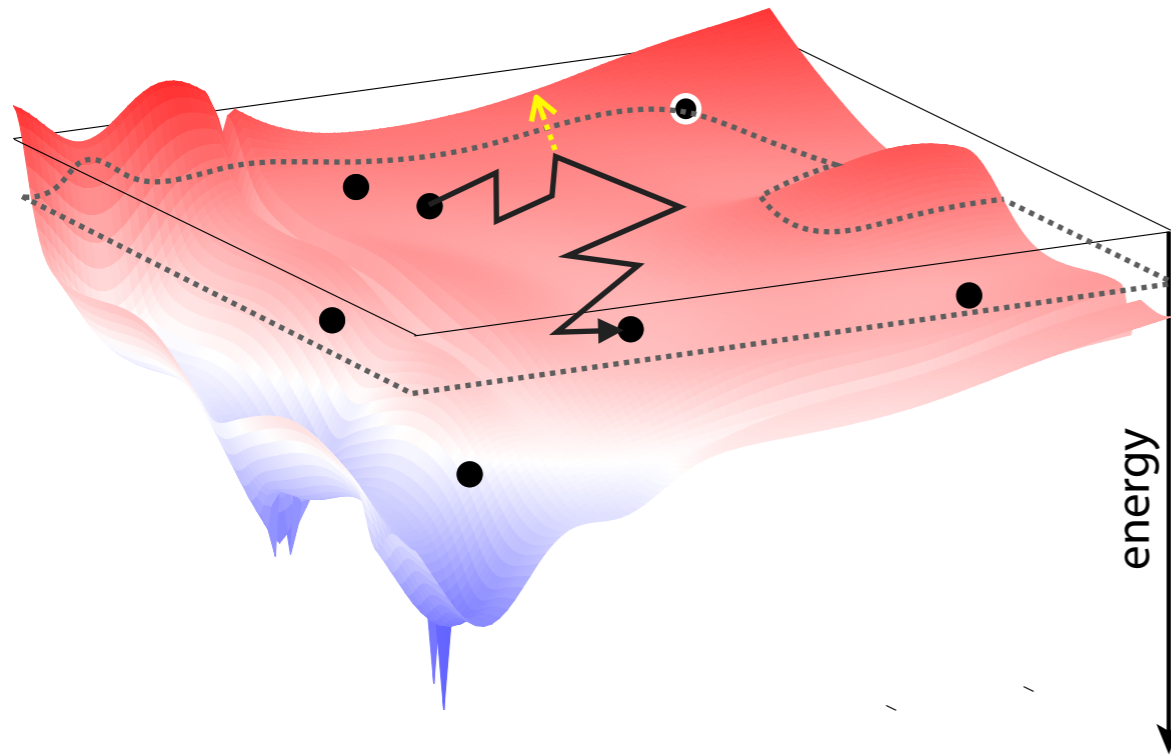
- Thermodynamic quantities as a simple post processing step

$$\begin{aligned} Z(\beta) &= Z_{\mathbf{p}} \sum_i w_i e^{-\beta E_i} \\ &= Z_{\mathbf{p}}(\beta) \sum_i [(K/(K+1))^i - (K/(K+1))^{i+1}] e^{-\beta E_i} \end{aligned}$$

$$C_V = \left( \frac{\partial U}{\partial T} \right)_V = - \left( \frac{\partial}{\partial T} \frac{\partial \ln Z}{\partial \beta} \right)_V$$

- Sampling itself independent from temperature
- No need for prior knowledge of the structures
- Can be done with both  $(N, V, T)$  and  $(N, p, T)$
- Easy control parameter of the sampling is  $K$ , called the “live set” or the “walkers” (is the resolution of the PES)

# Generating a new sample configuration



Clone a randomly selected sample and perform a random walk until it is independent from its parent configuration:

- atomic coordinates
- cell shape
- cell volume
- swap types

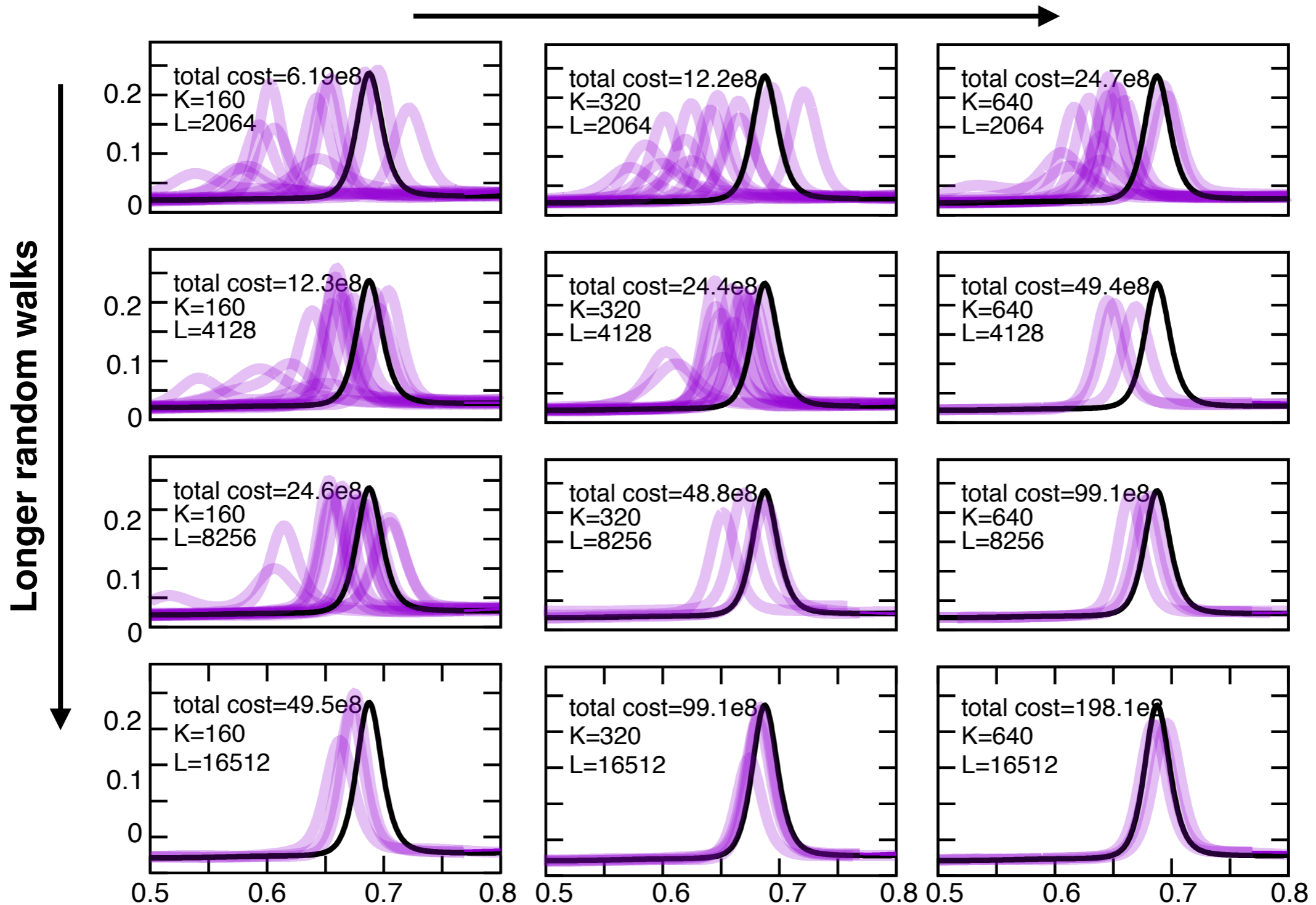
- **Markov Chain Monte Carlo (MCMC):**  
single particle and cell moves (volume, shear, stretch)
- **Total Enthalpy Hamiltonian Monte Carlo (TE-HMC)**  
short constant total energy MD trajectories
- **Galilean Monte Carlo (GMC)**  
all-atoms moves, along straight lines between elastic collisions (reflect the velocities to redirect the sample to allowed phase space region)

Length of the random walk,  $L$ , is the other control parameter.

# Convergence with $K$ and $L$

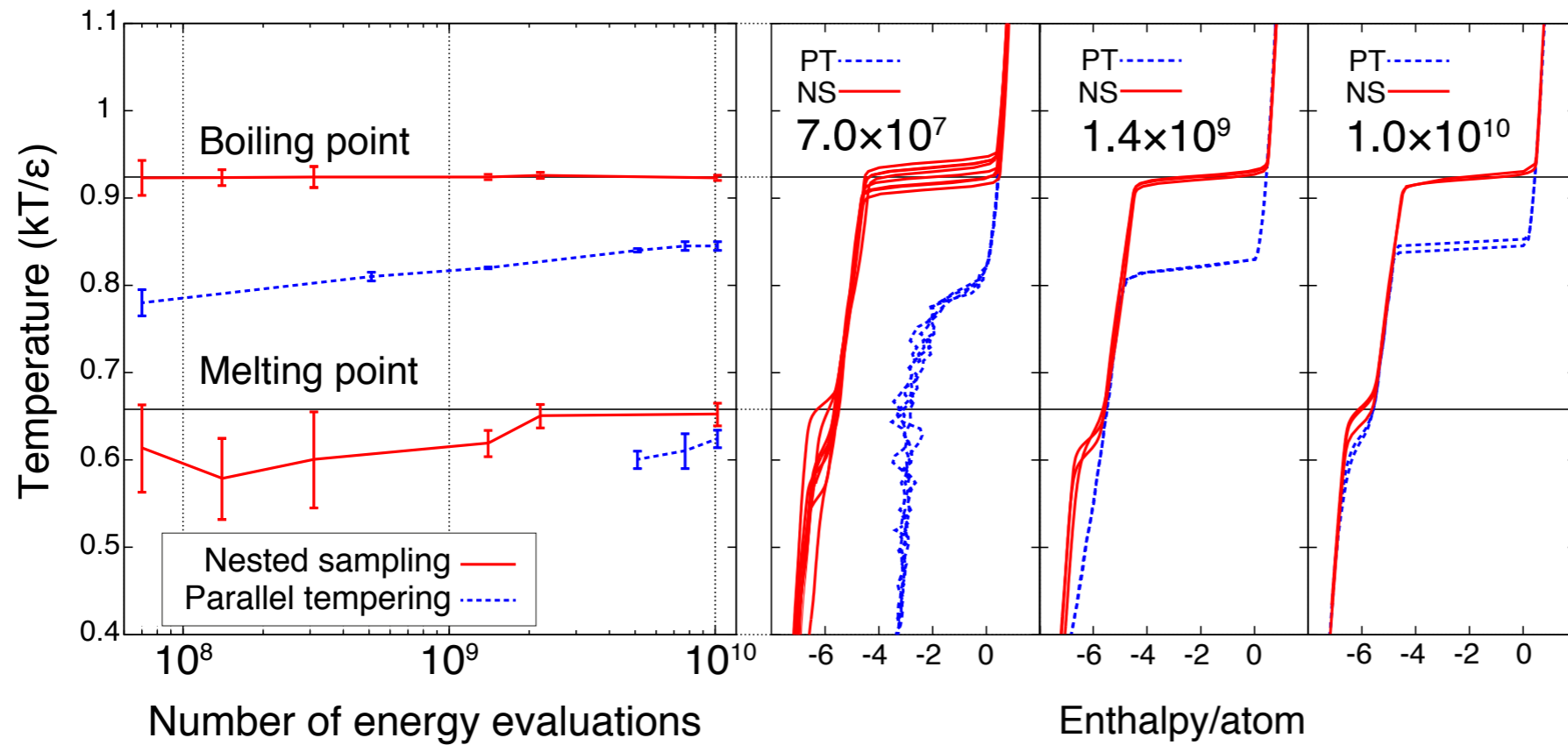
Melting peak on the heat capacity curve of 64 LJ atoms at constant pressure.

**Increased resolution = more walkers**



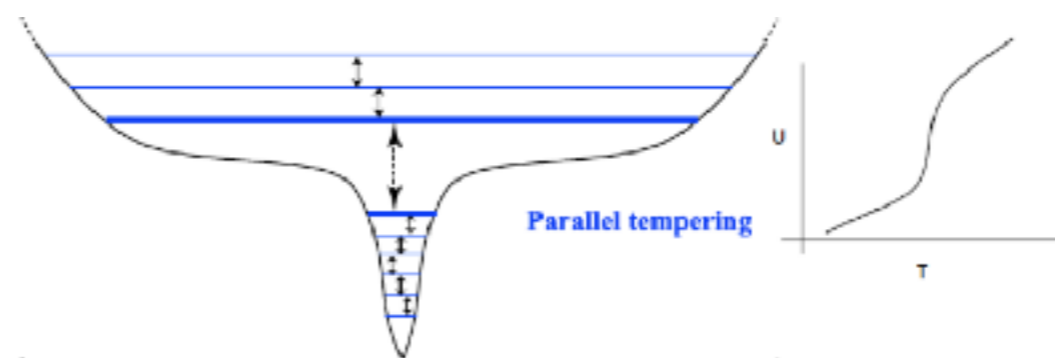
# Computational cost comparison

64 LJ atoms at constant pressure

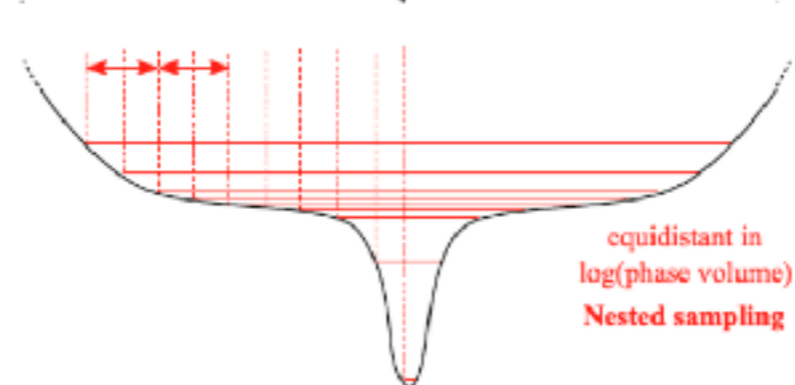


Nested Sampling vs. Parallel Tempering

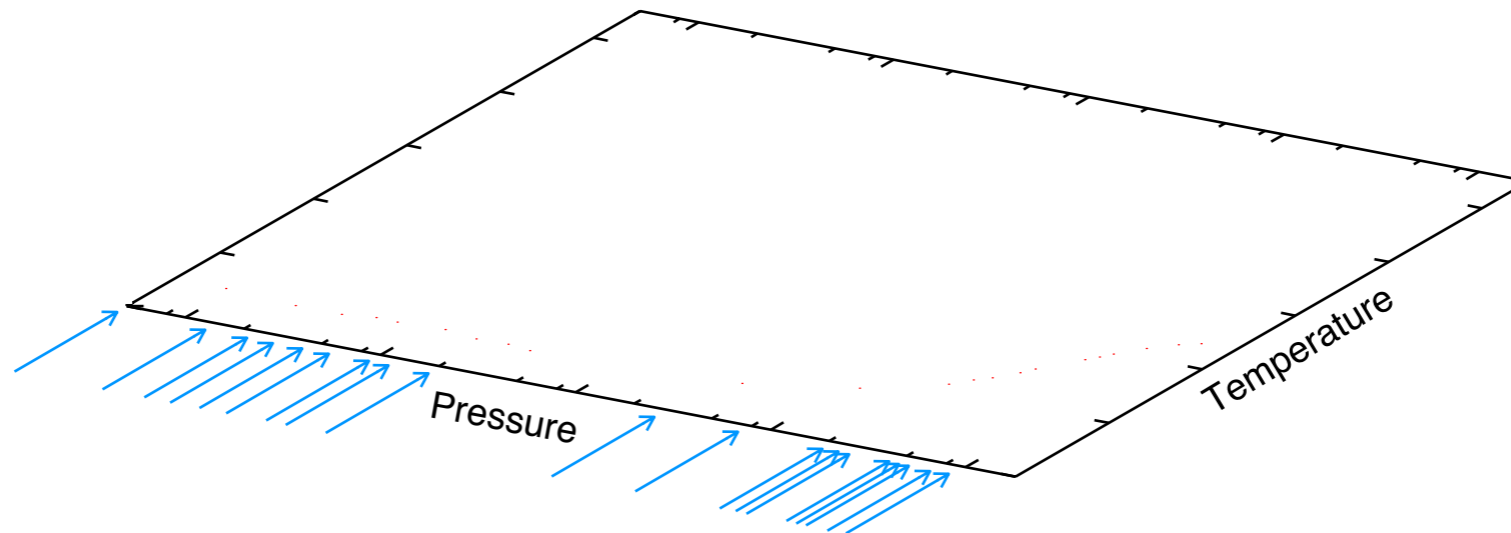
levels equidistant in temperature



levels equidistant in  $\log(\text{phase volume})$



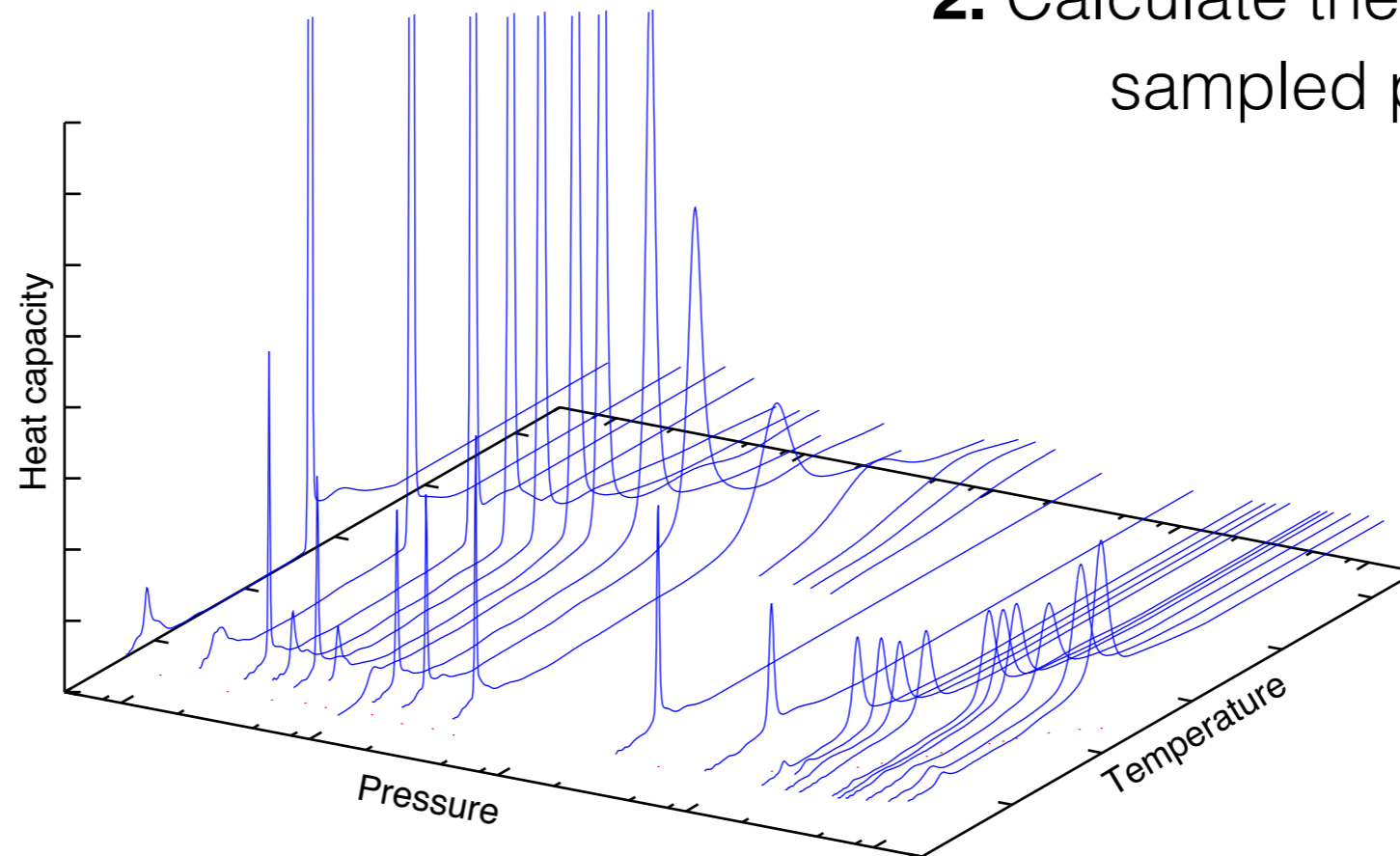
# $p$ - $T$ phase diagram “in three steps”



1. Choose a system (potential model) and a set of pressures. Perform a *nested sampling* calculation for each.

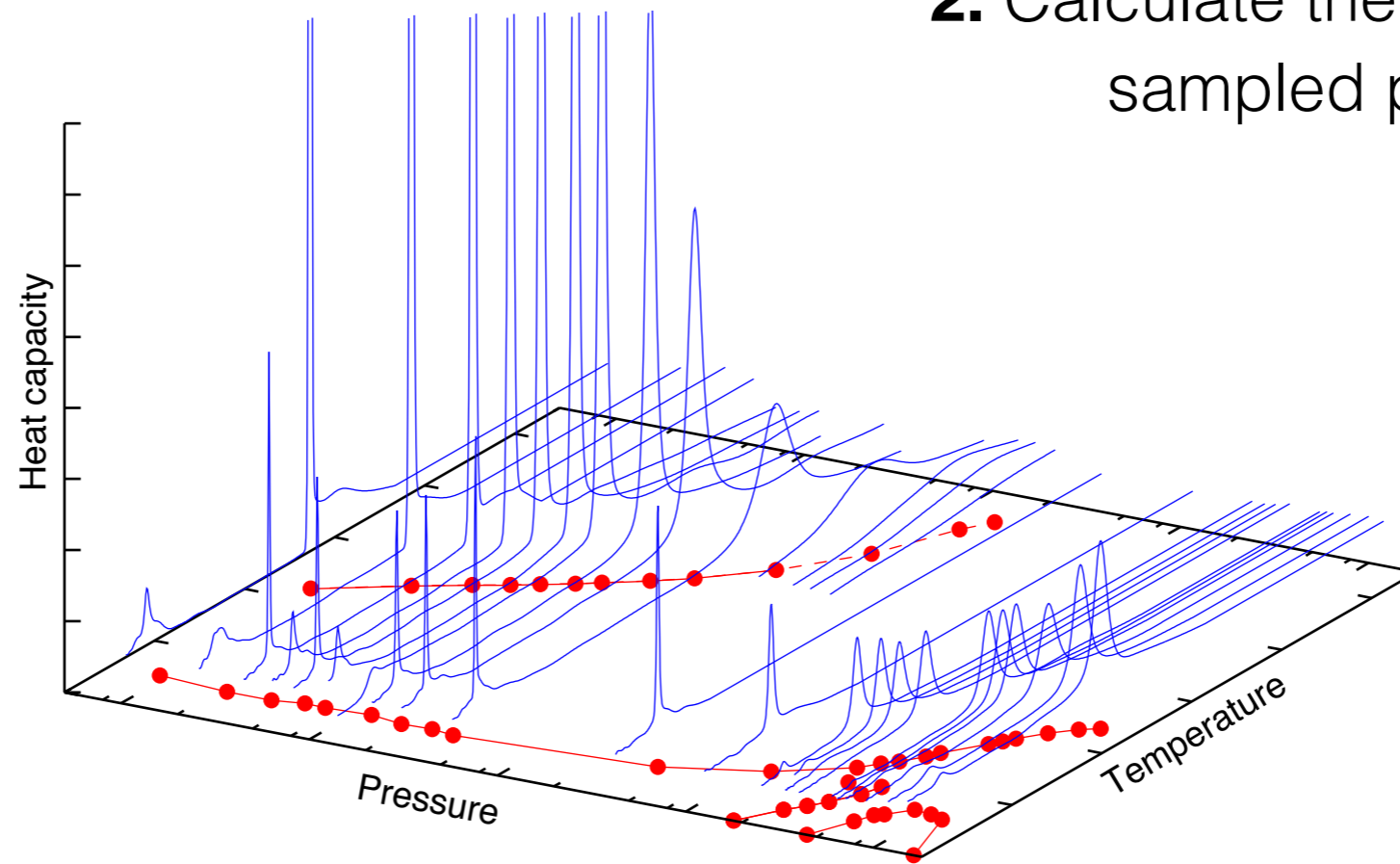
# $p$ - $T$ phase diagram “in three steps”

**2.** Calculate the  $c_p(T)$  for every sampled pressure.



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# $p$ - $T$ phase diagram “in three steps”



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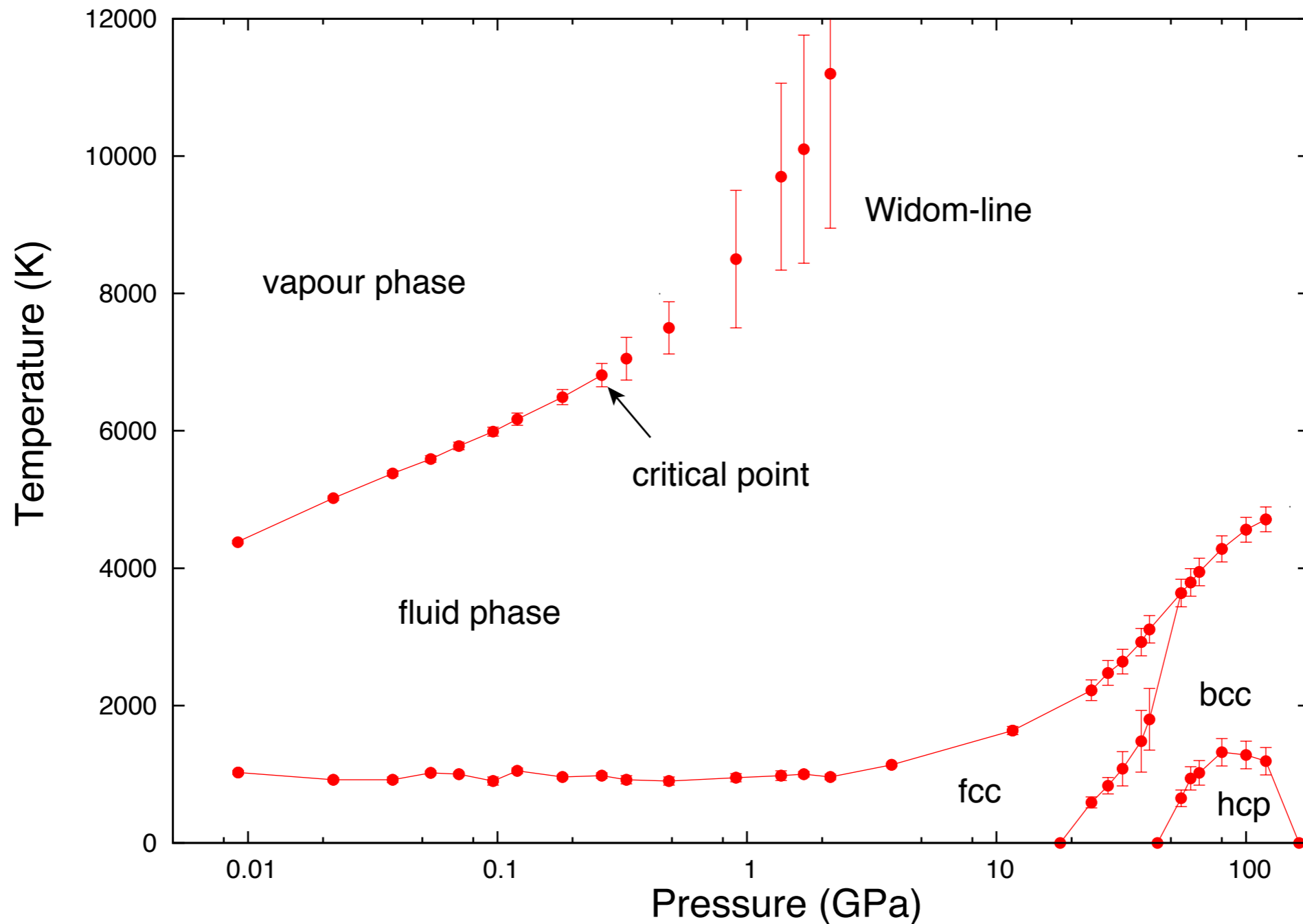
- 3.** Determine the location of maxima on the  $c_p(T)$  curve, showing the phase transitions.

$p$ - $T$  phase diagram!

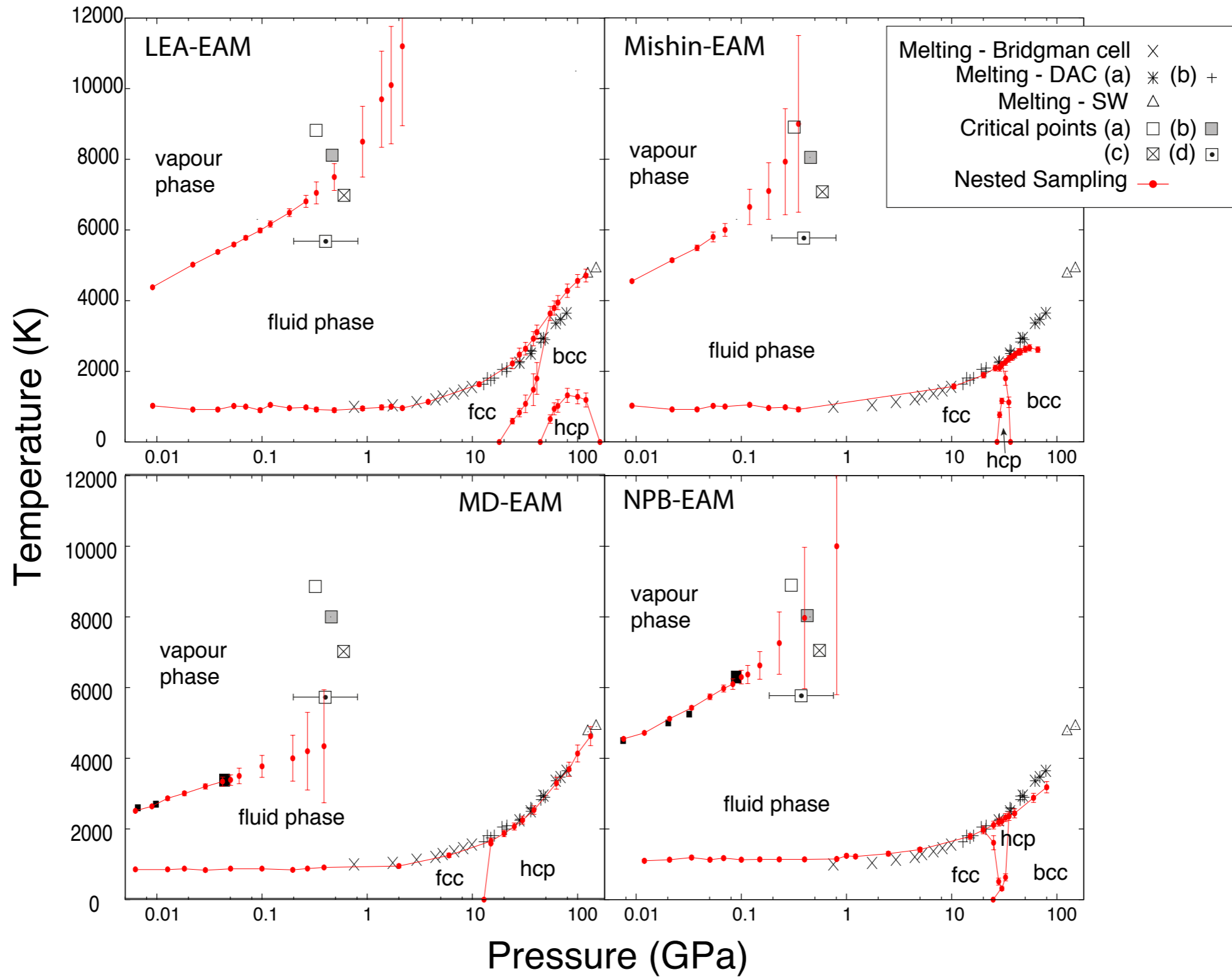
# Phase Diagram of Aluminium

Embedded atom model by Ercolessi and Adams

64 Al atoms in a simulation cell with variable size and shape

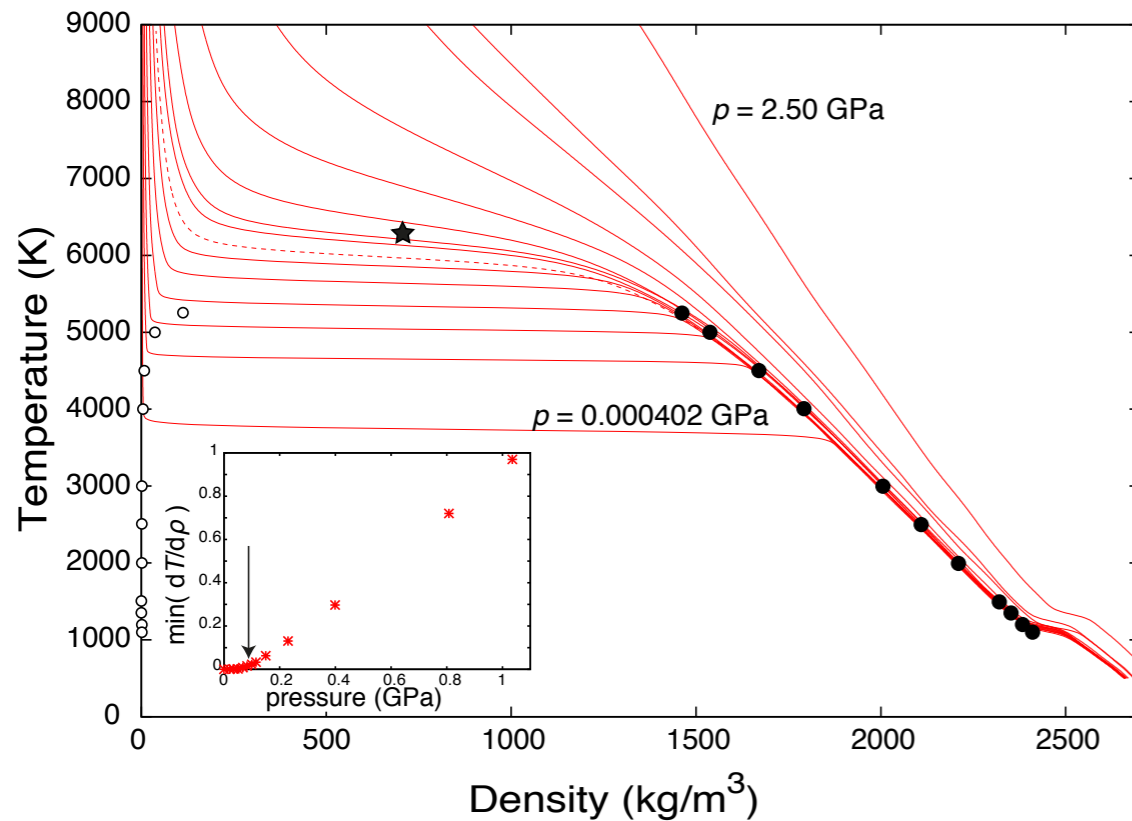


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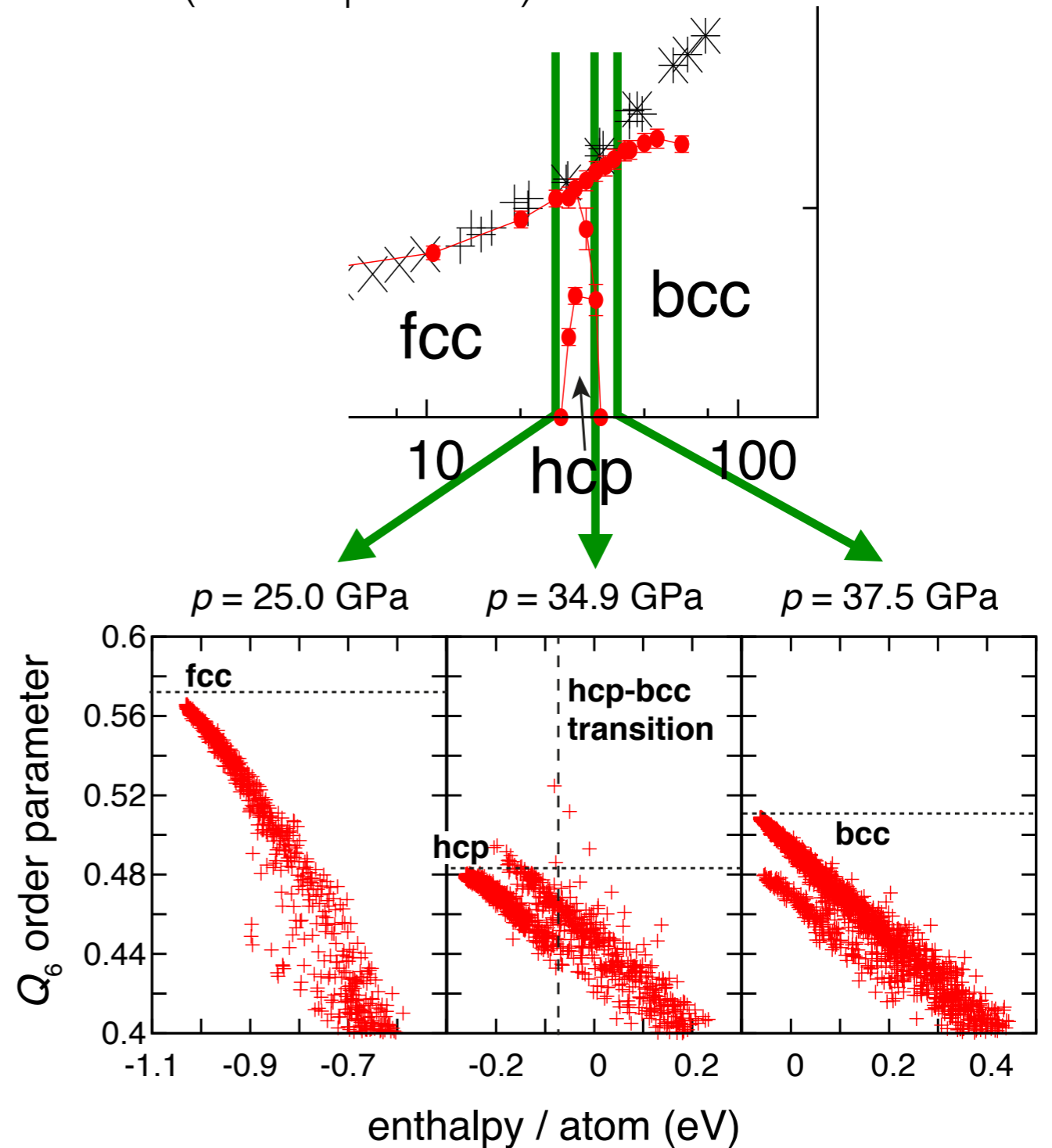


# Phase Diagram of Aluminium

Comparison to Gibbs ensemble MC:  
density-temperature phase diagram  
(NPB potential)



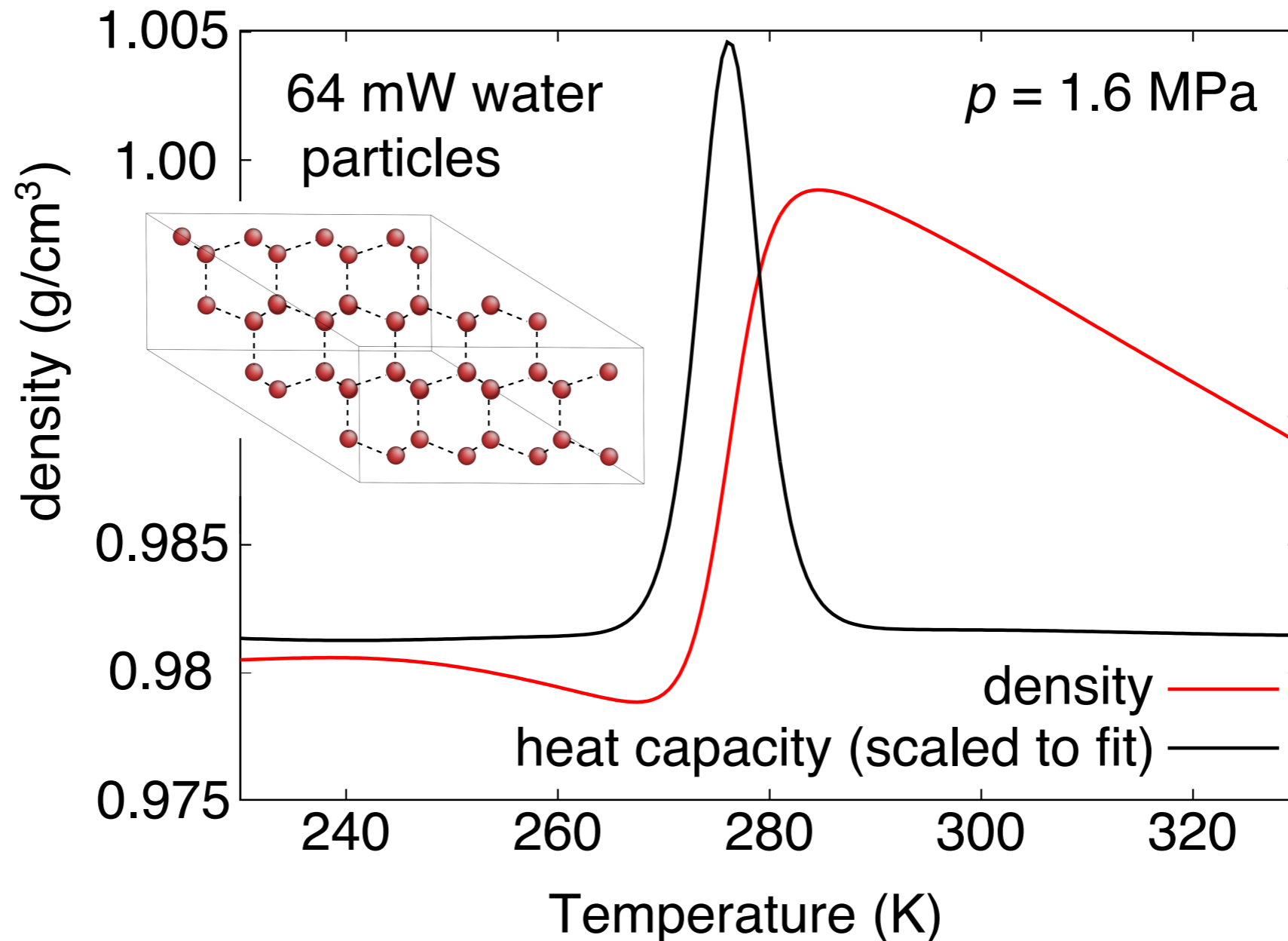
Solid-solid transition and  $Q_6$  bond  
order parameter  
(Mishin potential)





# Phase behaviour of water

coarse grain water model: mW (re-parametrised Stillinger-Weber Si)  
angular dependent term that encourages tetrahedral configurations



# Binary Lennard-Jones

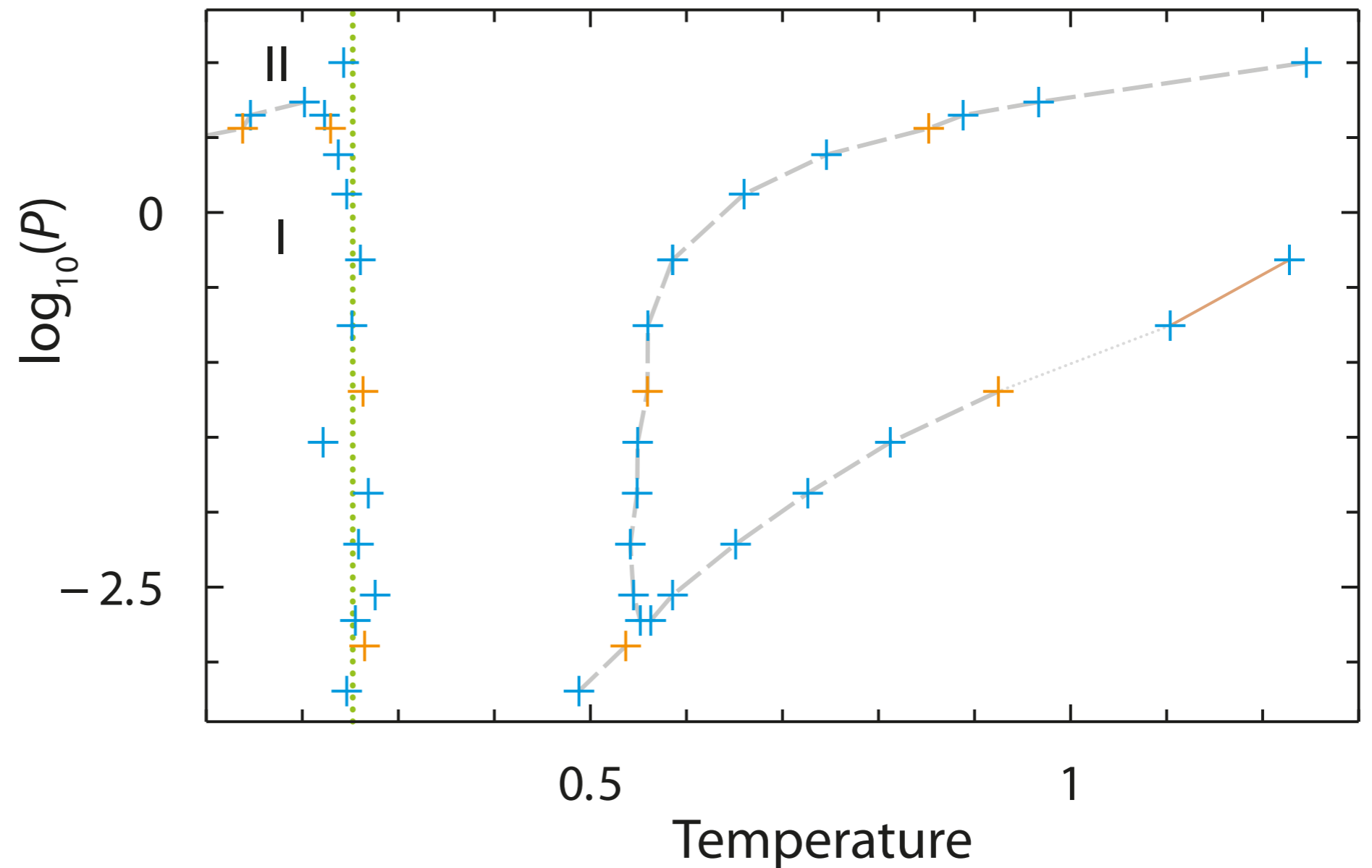
$$U_{\text{LJ}}(r) = 4\epsilon \left[ \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^6 \right]$$

$$\sigma_{\text{AA}} = \sigma_{\text{AB}} = \sigma_{\text{BB}}$$

same sizes

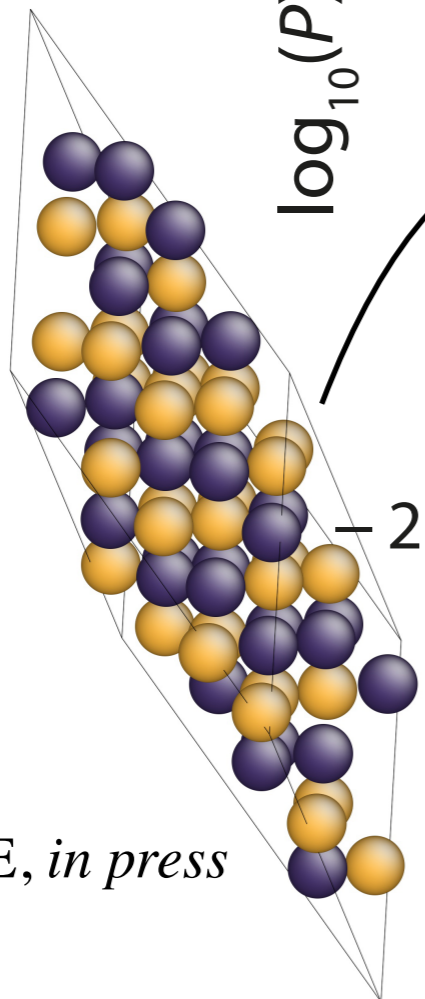
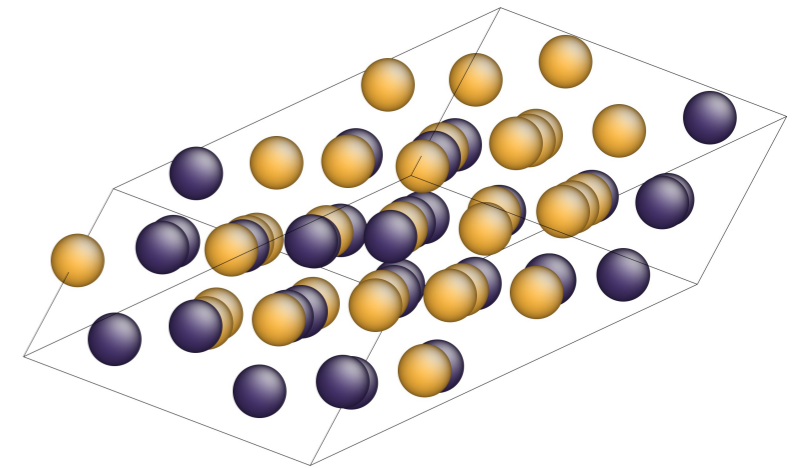
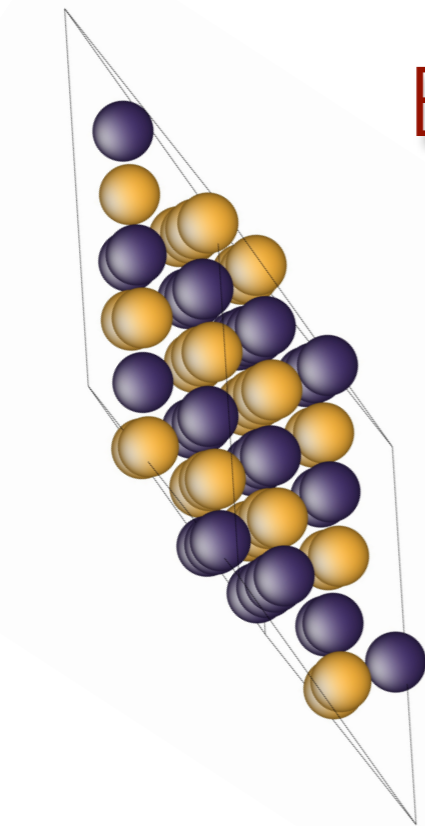
$$\epsilon_{\text{AA}} = \epsilon_{\text{BB}} = 0.5\epsilon_{\text{AB}}$$

different type preference



# Binary Lennard-Jones

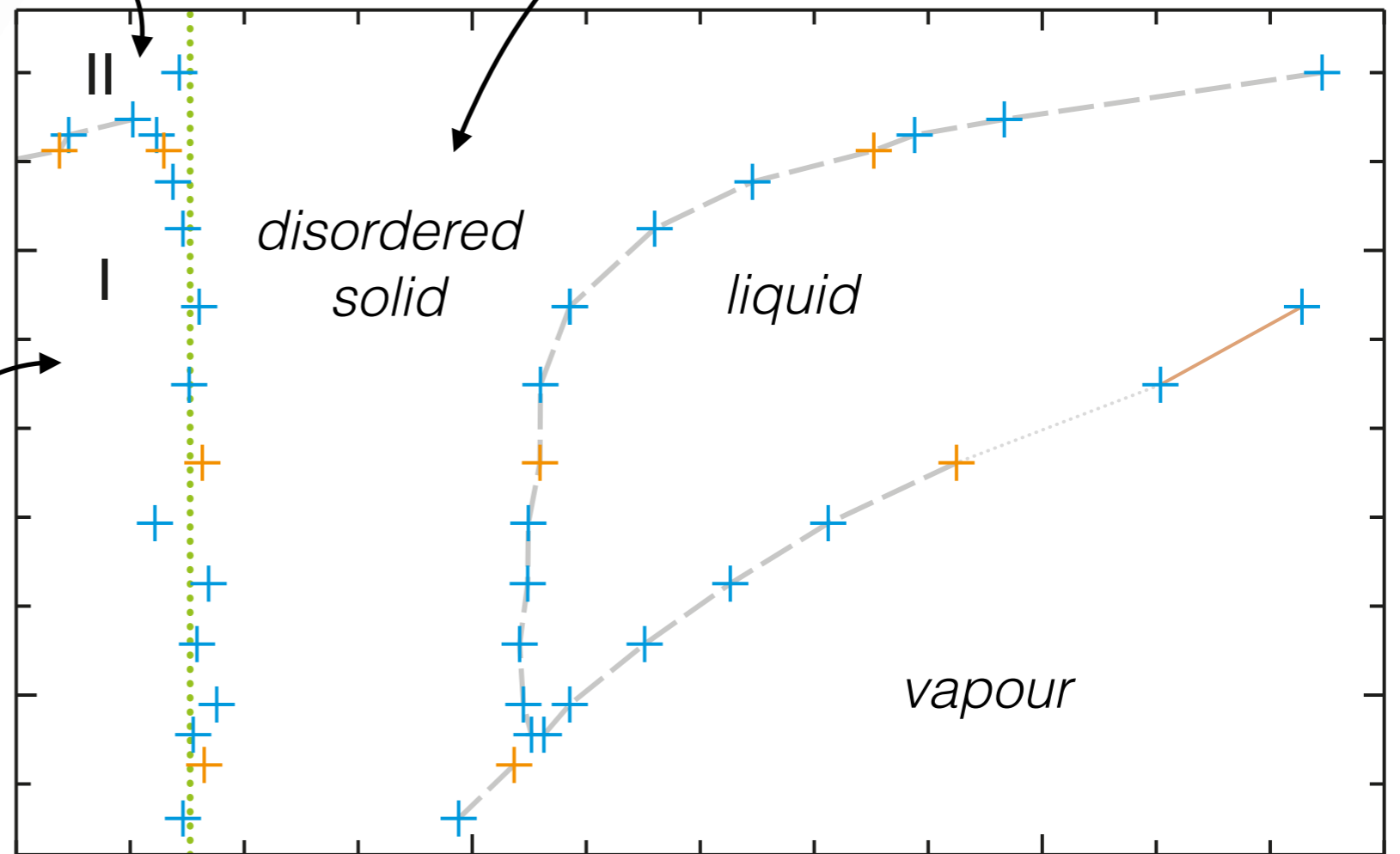
substitutional  
alloy, order-  
disorder  
transition



$\log_{10}(P)$

0

-2.5



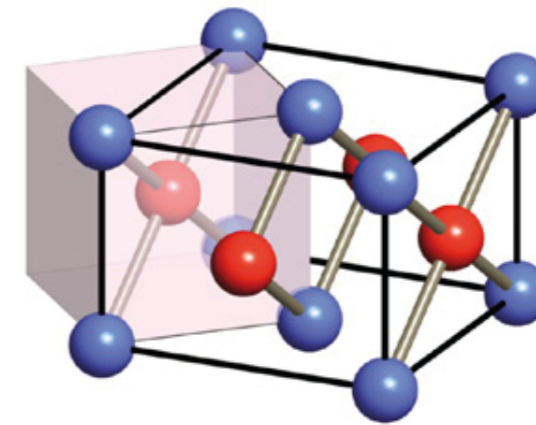
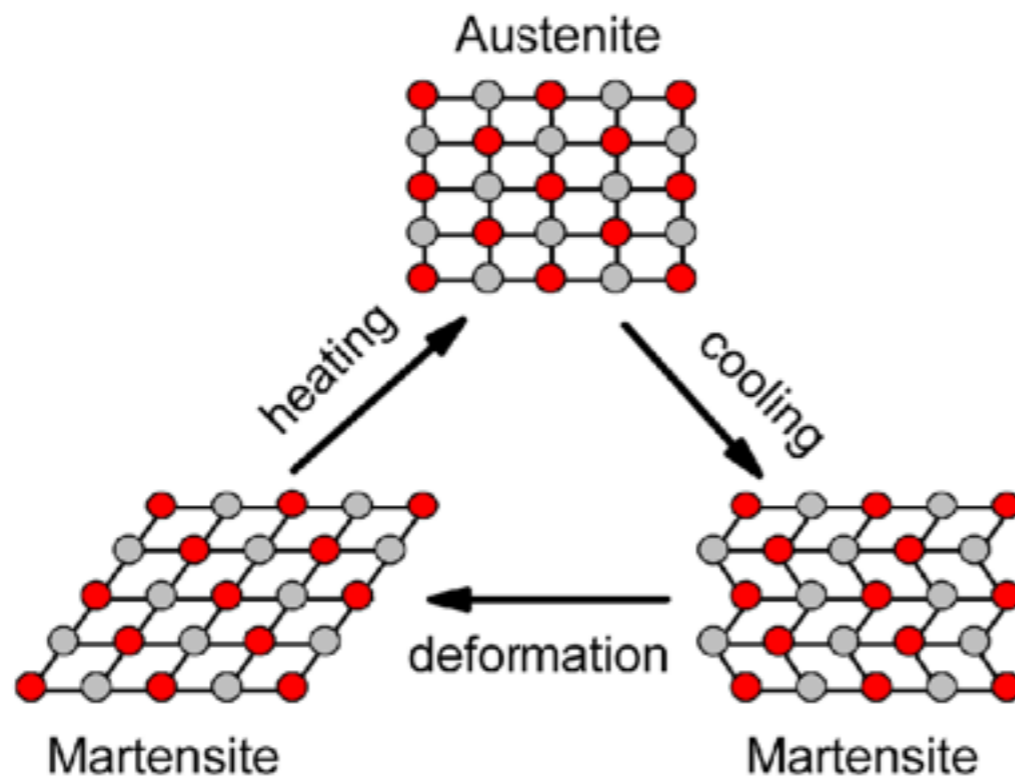
0.5

1

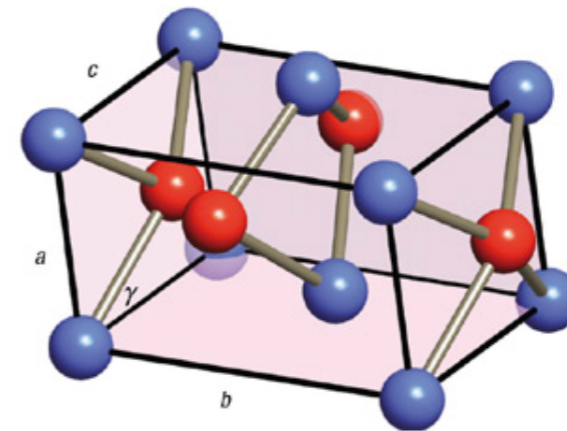
Temperature

# Martensitic Transition in NiTi alloys

shape memory alloy: “remembers” its original shape when deformed, and returns to it when heated



B2



B19'

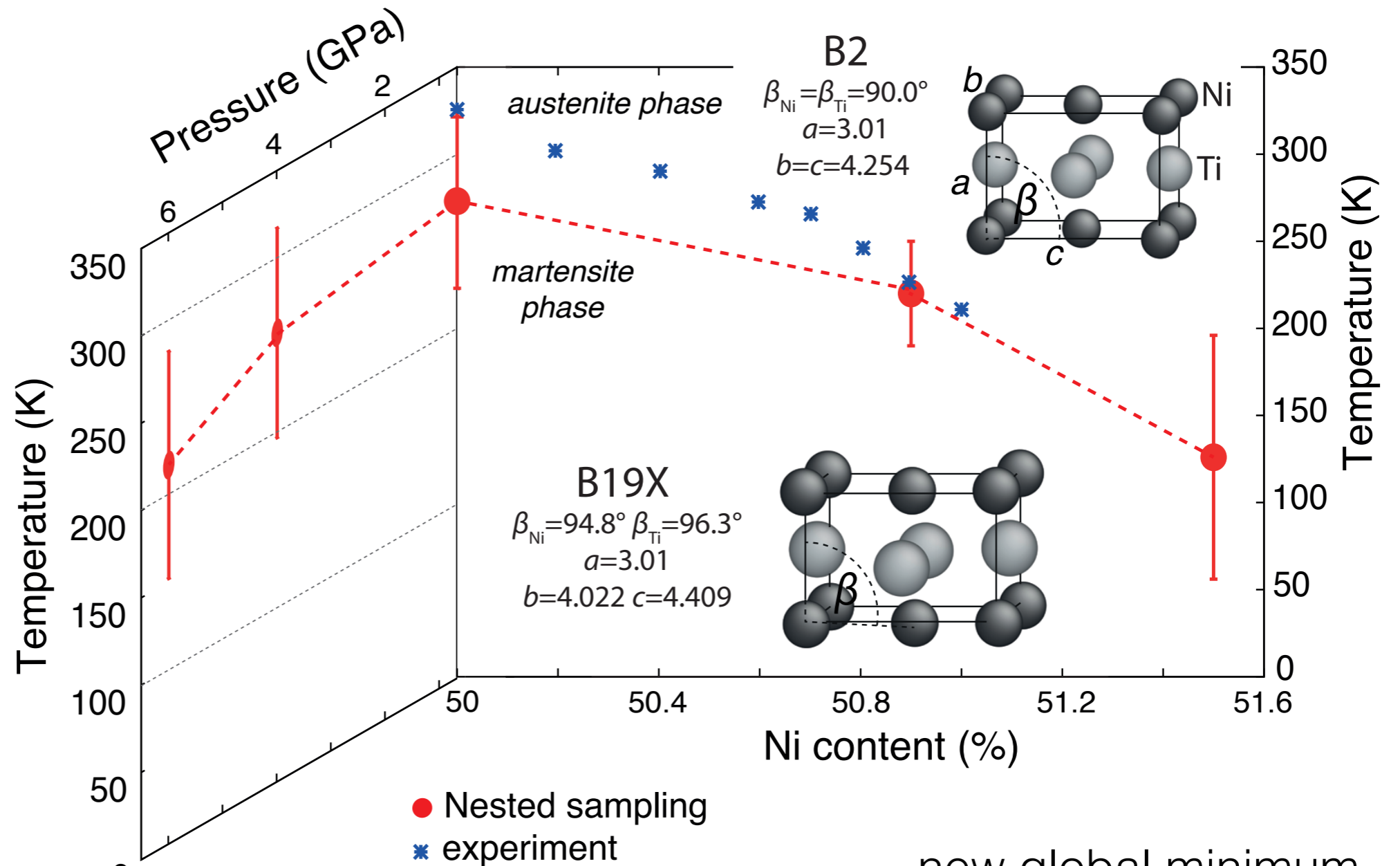
(from Huang et.al. Nat.Mat.2003)

# Martensitic Transition in NiTi alloys

pressure-temperature-composition phase diagram

Embedded atom model by Zhong et al.

64 and 108 atoms in a simulation cell with variable size and shape

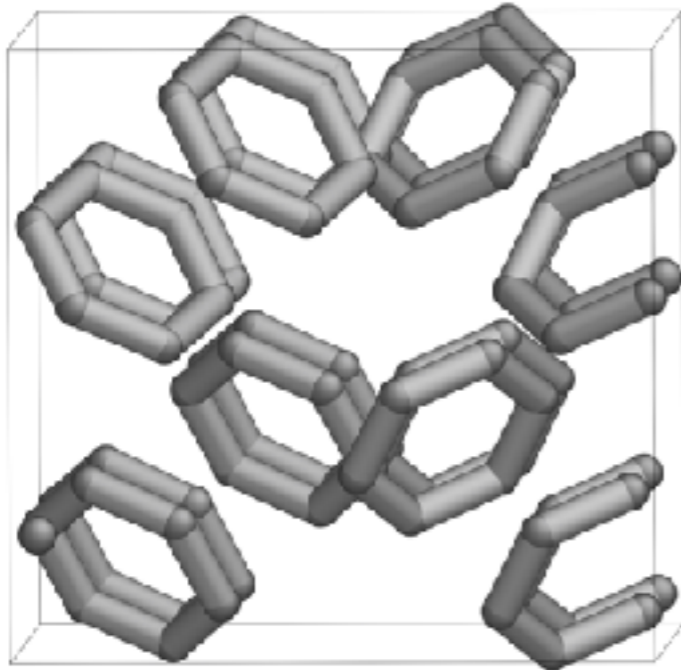


new global minimum structure found!

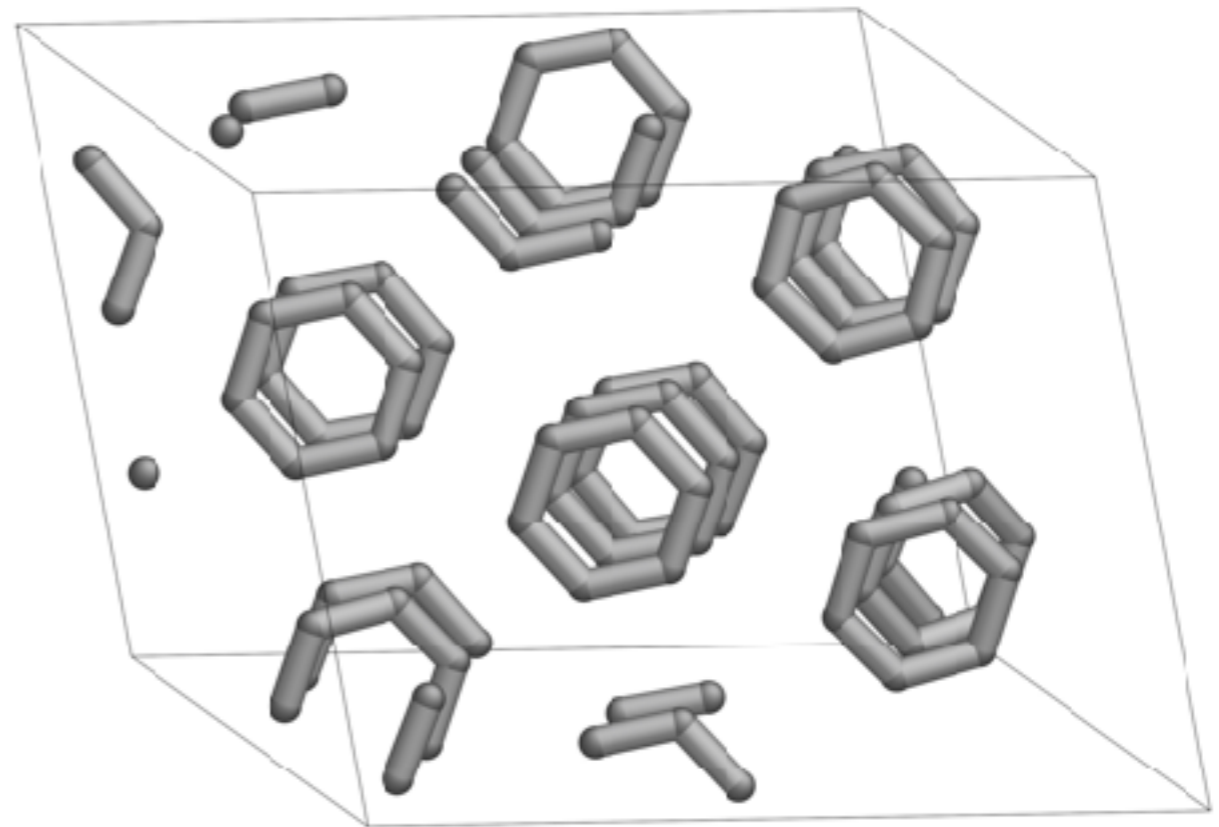
# NS with molecular systems

16 benzene molecules (united atom, LJ)

0.5 GPa



5.0 GPa

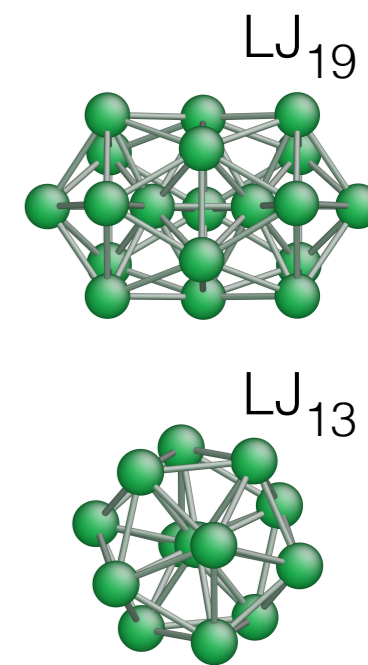
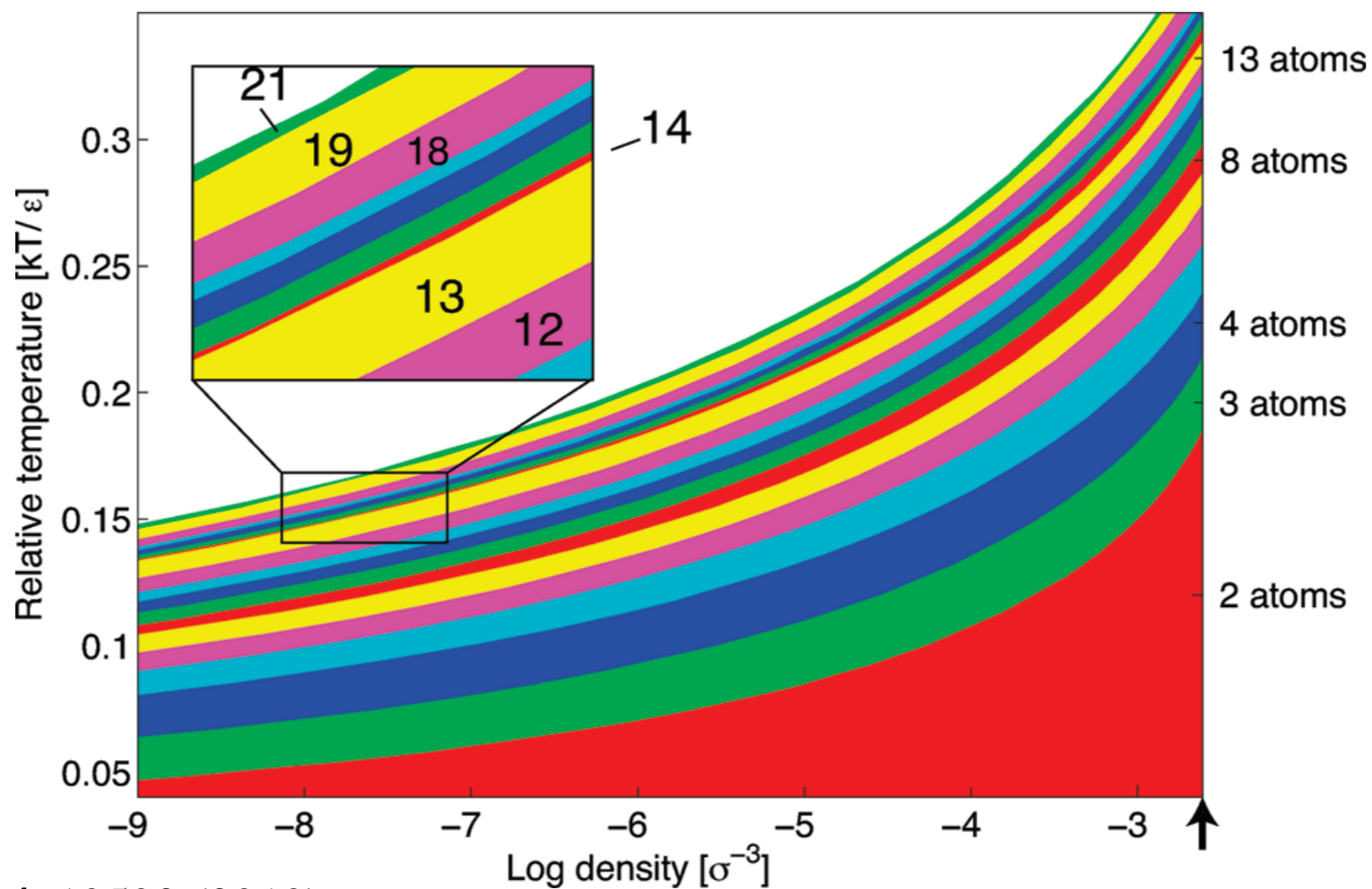


# Lennard-Jones clusters

$$U_{\text{LJ}}(r) = 4\epsilon \left[ \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^6 \right]$$

Studied clusters from size LJ<sub>2</sub> to LJ<sub>38</sub>

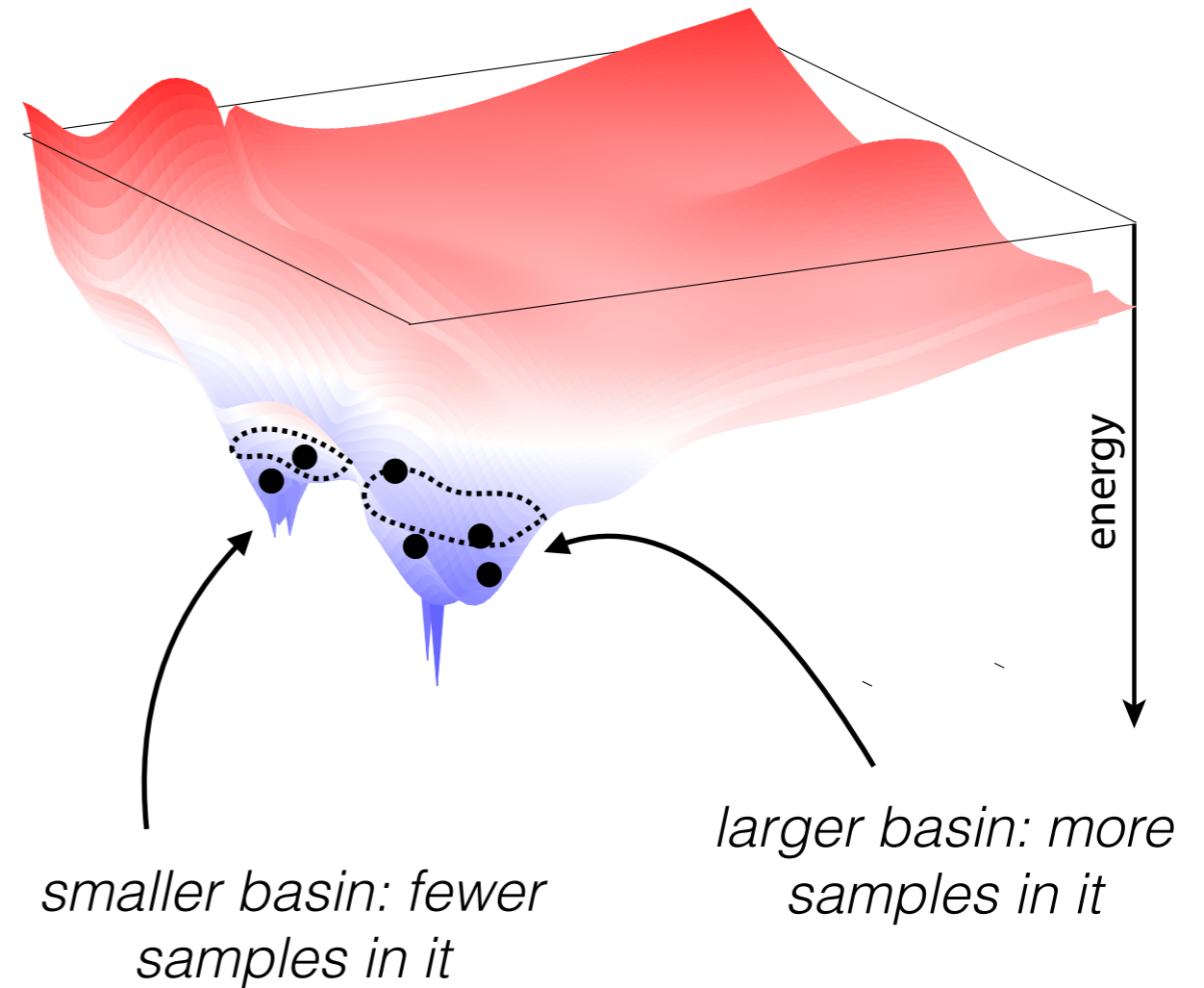
Stability of the Lennard-Jones clusters against the ideal gas



# Visualising the PES

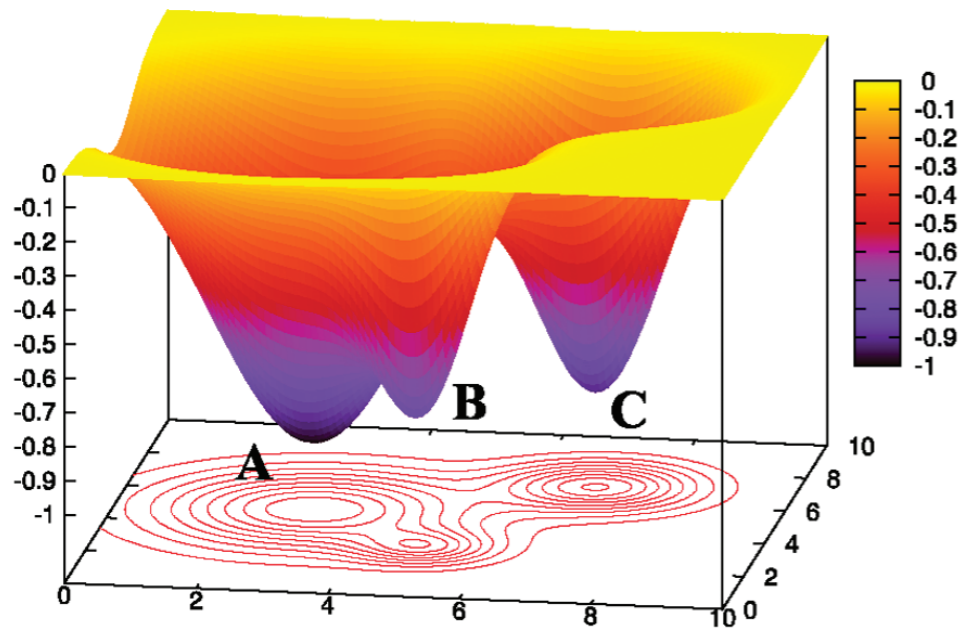
With a suitable metric  
categorise configurations  
and construct a graph

- identify different basins
- estimate saddle points
- sample distribution at a given energy level shows the relative phase space volume ratio of basins



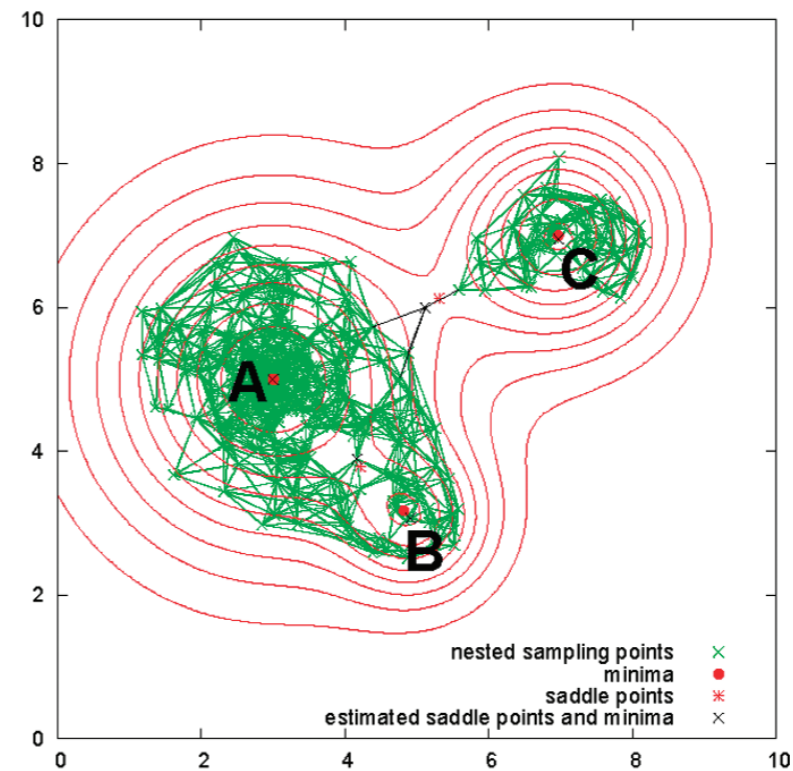
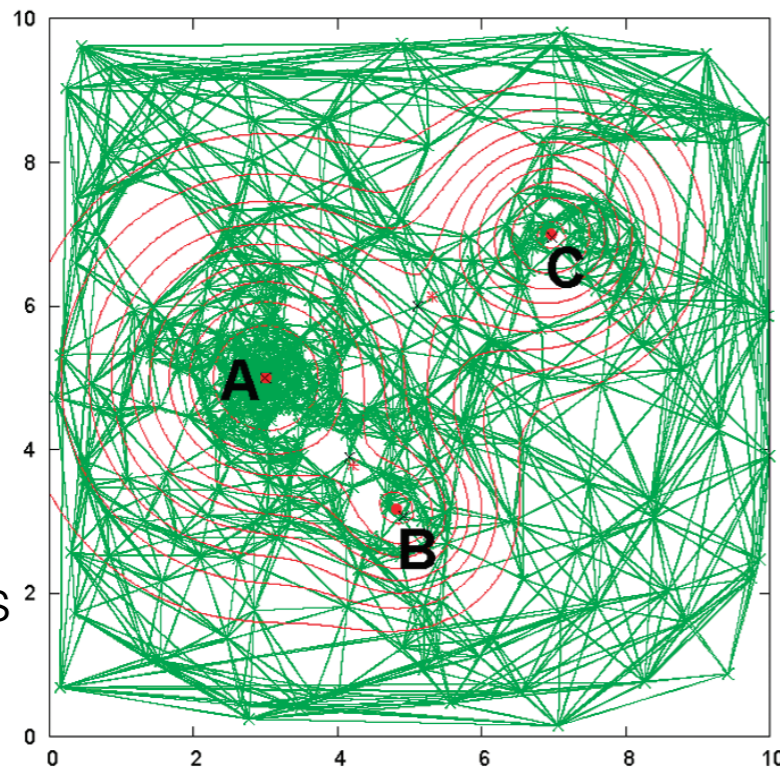
# Visualising the PES

Simple test landscape



After NS:

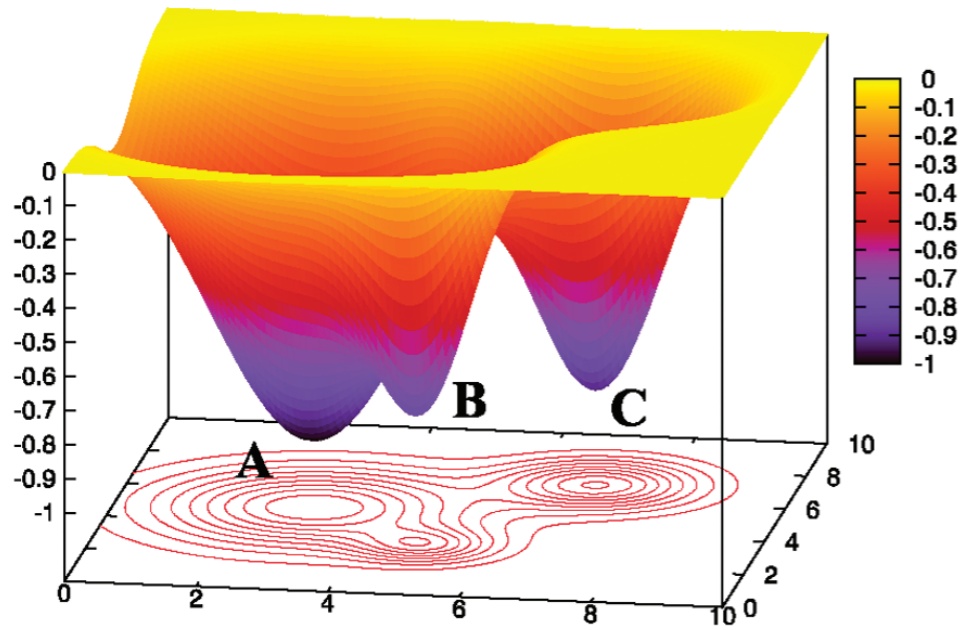
choose a *good* metric and produce a graph by connecting the  $k$  closest higher energy neighbours



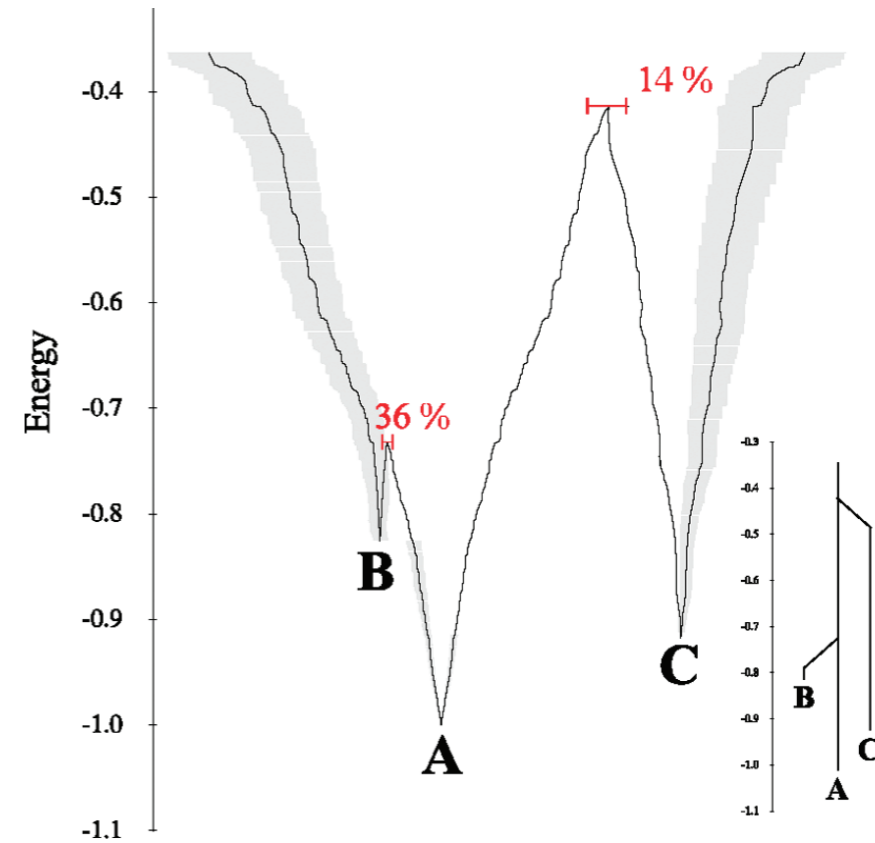
start deleting the points from the top (high energy) and see as the graph to split

# Visualising the PES

Simple test landscape

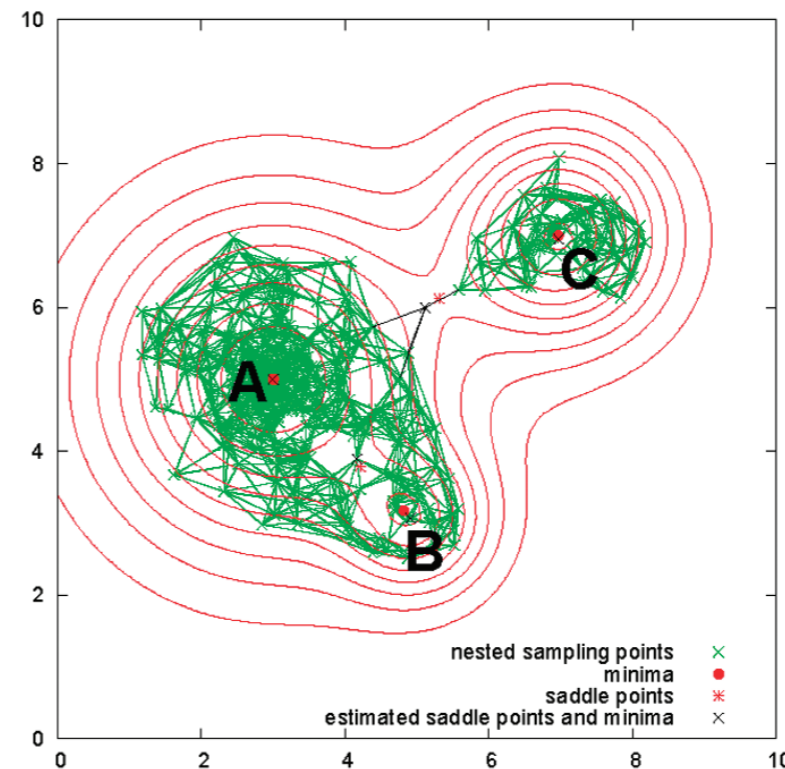
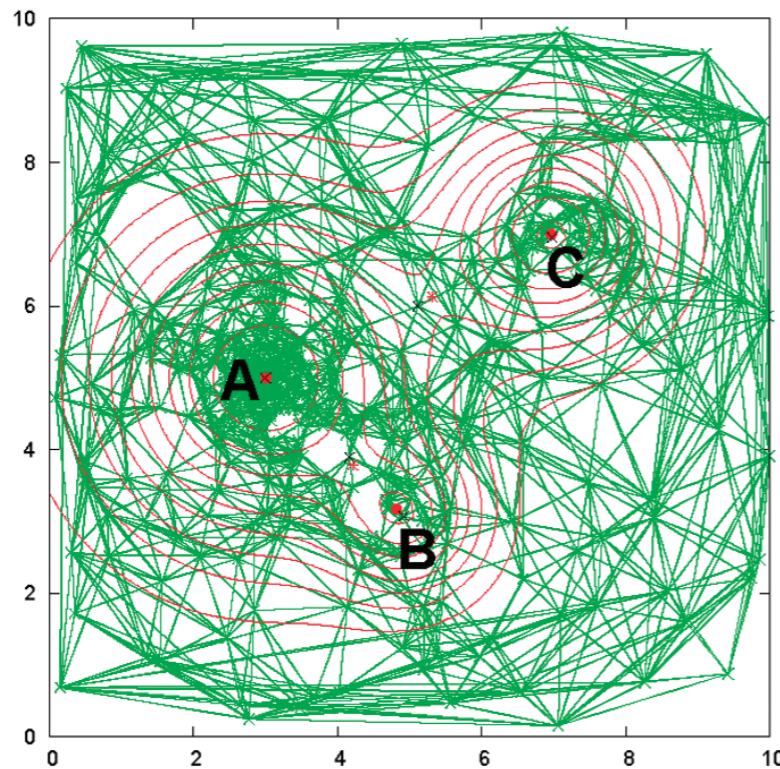


energy landscape chart



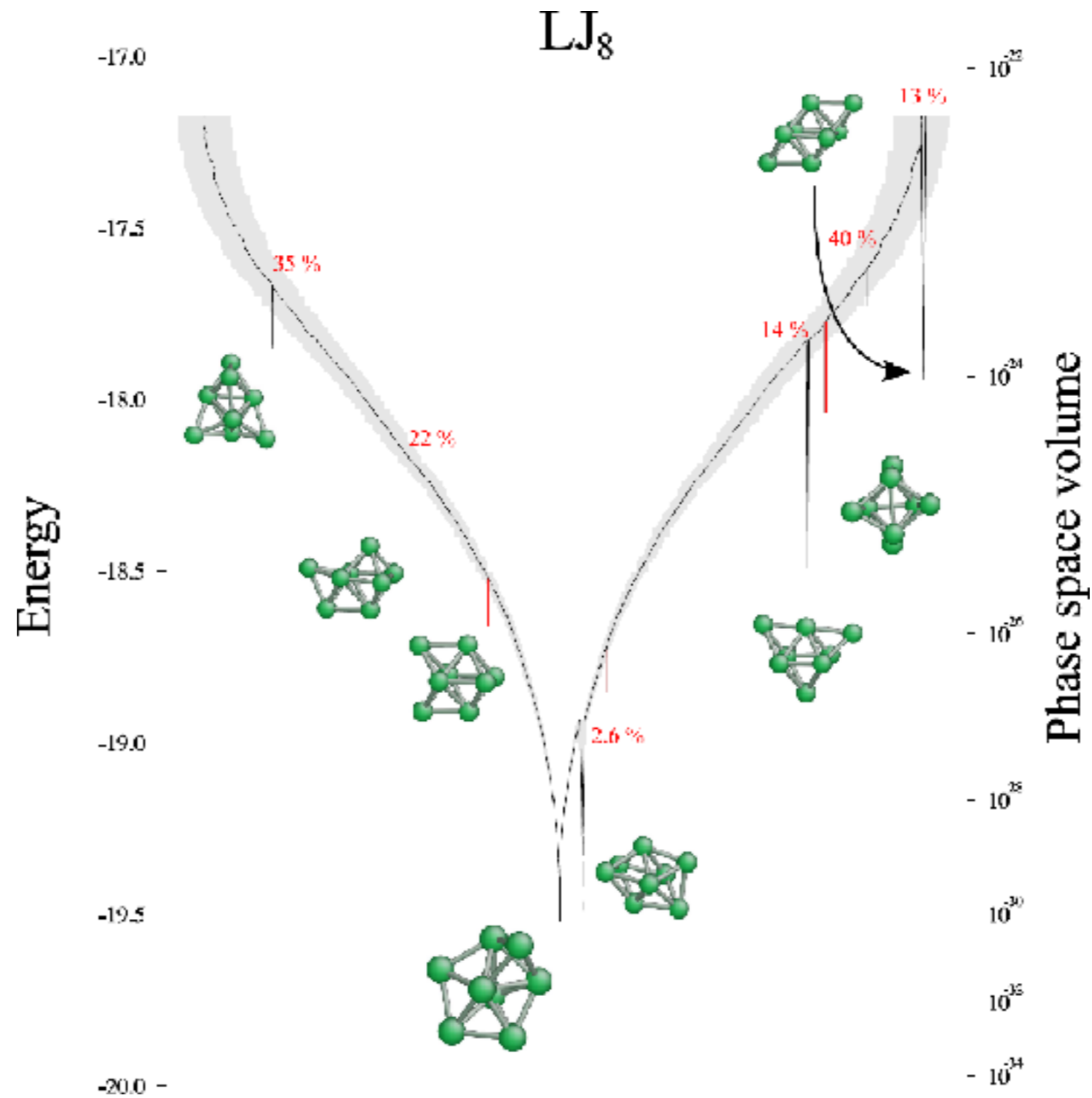
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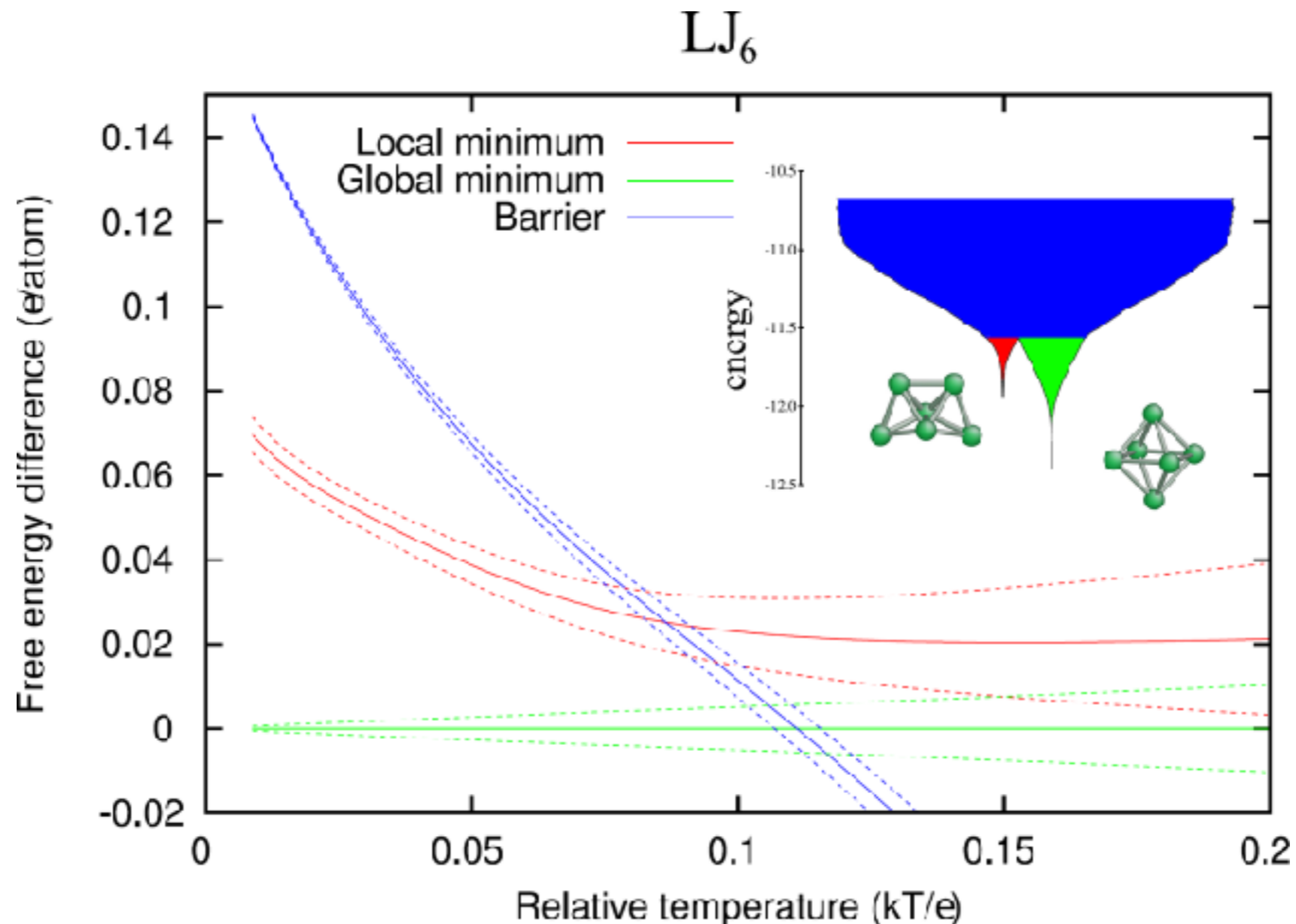
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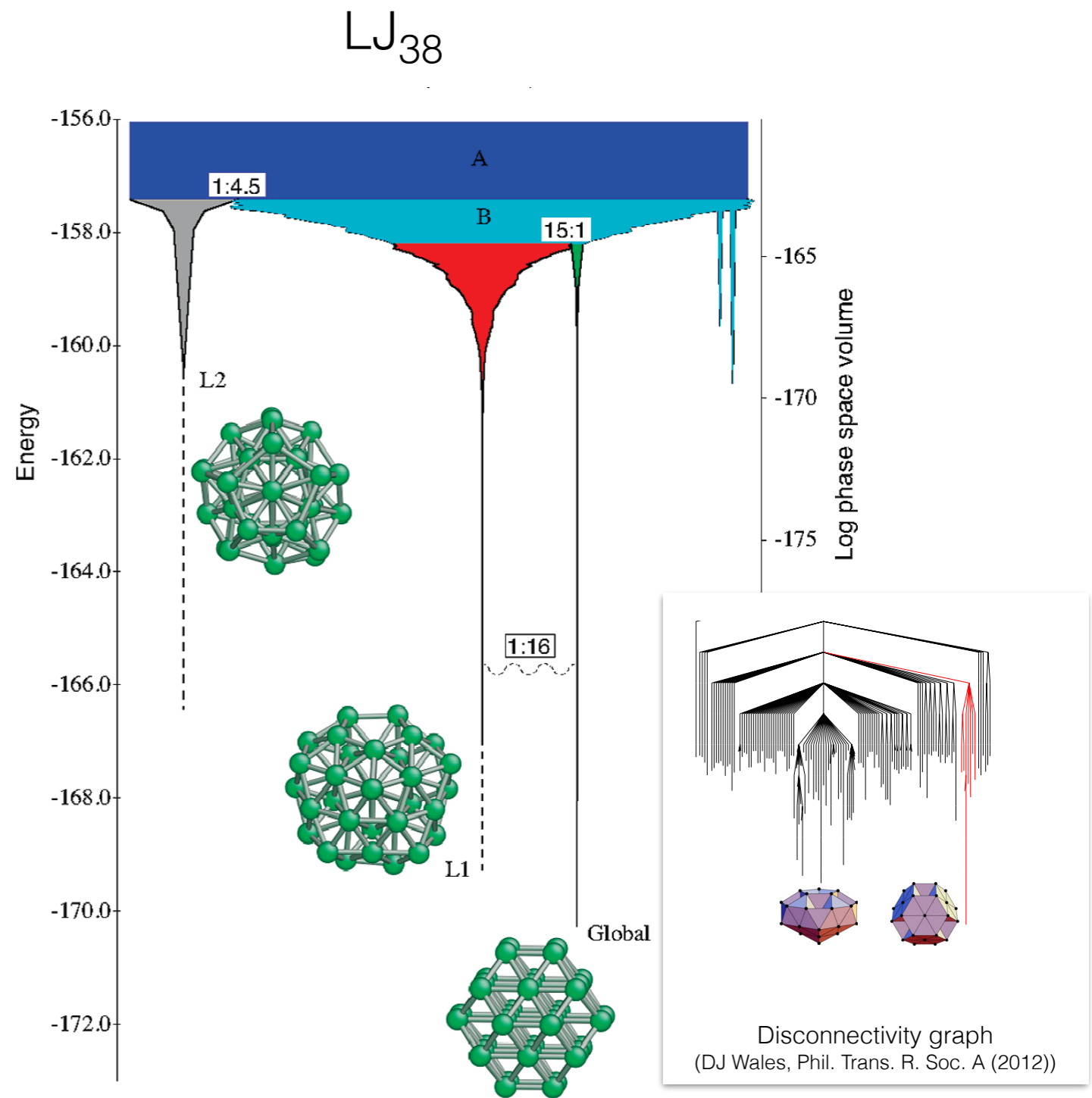
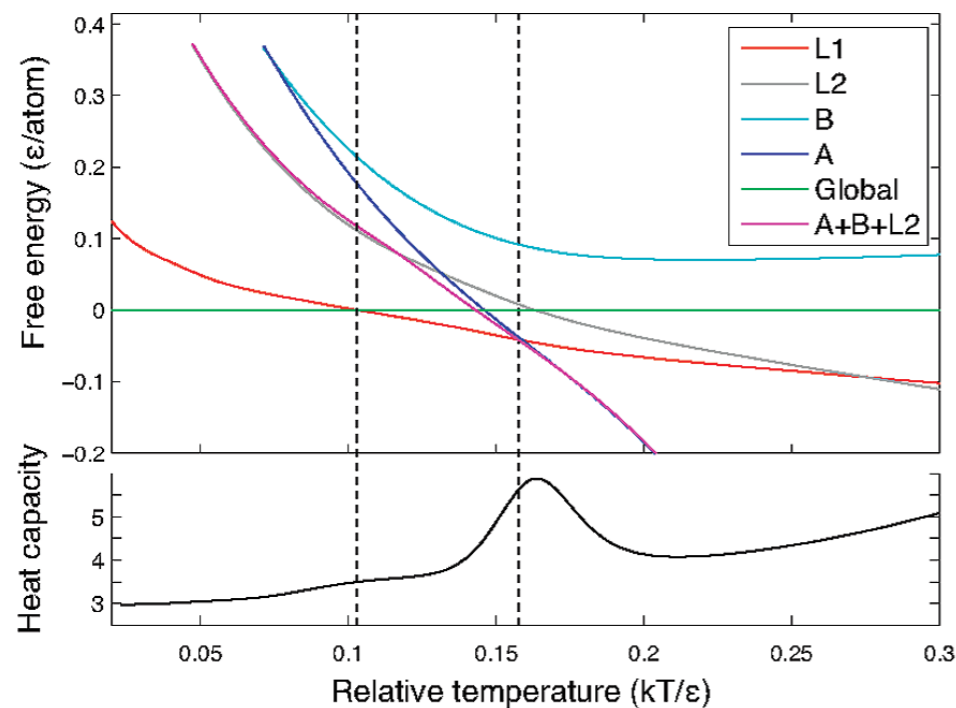
Using the landscape chart, calculate the free energy of the basins, determine the phase transitions without the need of externally defined order parameter.

$$A(\beta) = -\frac{1}{\beta} \ln Z(\beta)$$

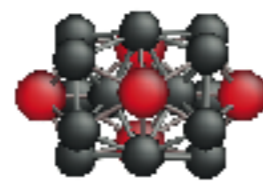
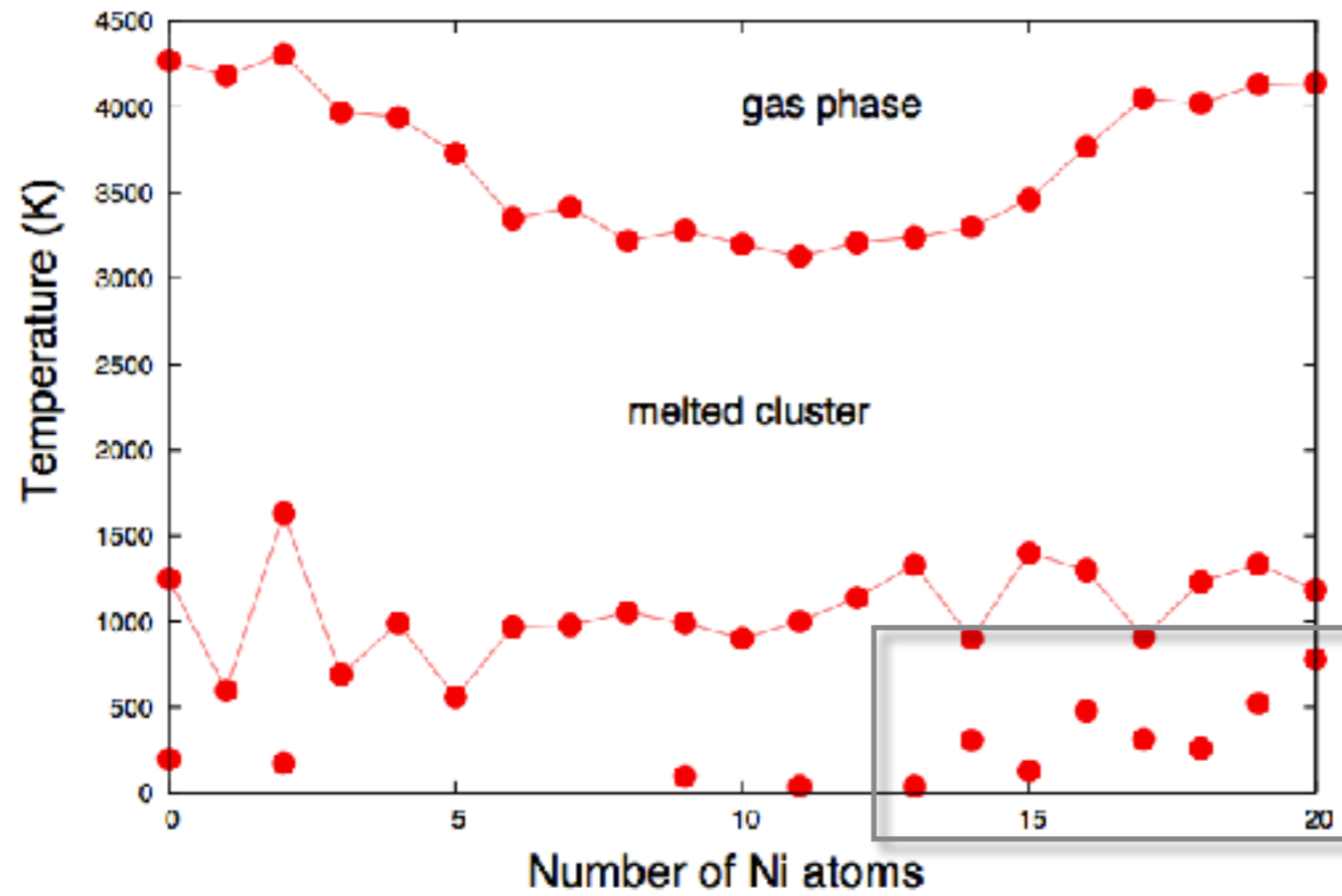


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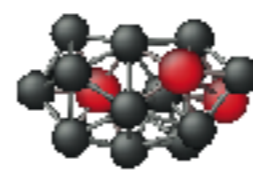
# Phase Diagram of Ni<sub>x</sub>Ti<sub>y</sub> cluster



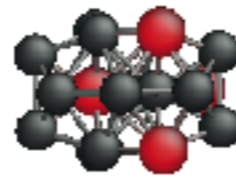
130K

Ni<sub>15</sub>Ti<sub>5</sub>

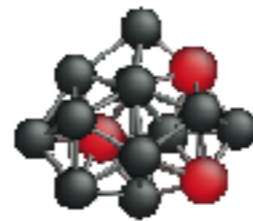
?



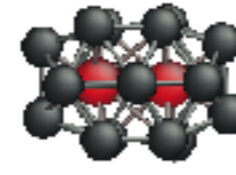
315K



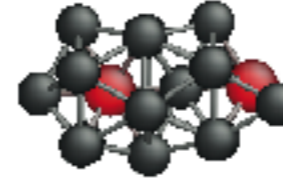
Ni<sub>16</sub>Ti<sub>4</sub>



Ni<sub>17</sub>Ti<sub>3</sub>



260K



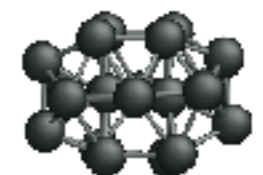
Ni<sub>18</sub>Ti<sub>2</sub>



525K



Ni<sub>19</sub>Ti<sub>1</sub>



780K



Ni<sub>20</sub>

# Conclusions

Nested sampling is an unbiased sampling technique,

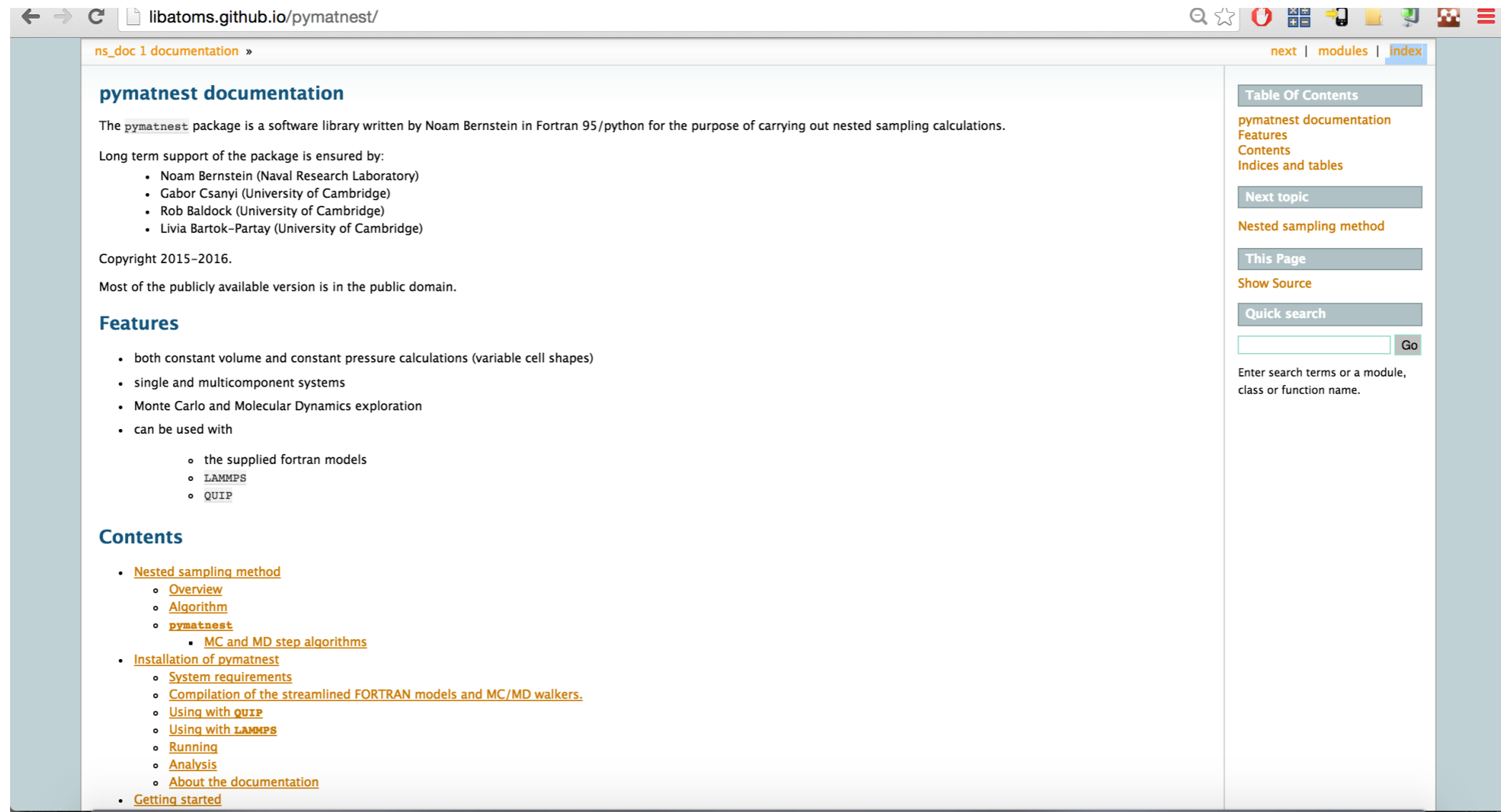
- sampling independent of temperature
- no prior knowledge of the phases necessary
- entire phase diagram can be calculated
- periodic systems, clusters, multicomponent systems
- next steps:
  - molecules
  - potential development

# Code release - pymatnest

Python code with an interface to QUIP and LAMMPS

<http://github.com/libAtoms/pymatnest>

<http://libatoms.github.io/pymatnest/>



The screenshot shows a web browser displaying the documentation for the pymatnest package. The browser's address bar shows the URL `libatoms.github.io/pymatnest/`. The page content includes a title "pymatnest documentation", a brief description of the package, a list of long-term supporters, copyright information, a "Features" section with a bulleted list of capabilities, and a "Contents" section with a hierarchical list of links to various parts of the documentation. A right-hand sidebar contains navigation links like "Table Of Contents", "Next topic", and a search box.

ns\_doc 1 documentation »

next | modules | index

## pymatnest documentation

The `pymatnest` package is a software library written by Noam Bernstein in Fortran 95/python for the purpose of carrying out nested sampling calculations.

Long term support of the package is ensured by:

- Noam Bernstein (Naval Research Laboratory)
- Gabor Csanyi (University of Cambridge)
- Rob Baldock (University of Cambridge)
- Livia Bartok-Partay (University of Cambridge)

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Most of the publicly available version is in the public domain.

## Features

- both constant volume and constant pressure calculations (variable cell shapes)
- single and multicomponent systems
- Monte Carlo and Molecular Dynamics exploration
- can be used with
  - the supplied fortran models
  - `LAMMPS`
  - `QUIP`

## Contents

- [Nested sampling method](#)
  - [Overview](#)
  - [Algorithm](#)
  - [pymatnest](#)
    - [MC and MD step algorithms](#)
- [Installation of pymatnest](#)
  - [System requirements](#)
  - [Compilation of the streamlined FORTRAN models and MC/MD walkers.](#)
  - [Using with QUIP](#)
  - [Using with LAMMPS](#)
  - [Running](#)
  - [Analysis](#)
  - [About the documentation](#)
- [Getting started](#)

Table Of Contents

[pymatnest documentation](#)

[Features](#)

[Contents](#)

[Indices and tables](#)

Next topic

[Nested sampling method](#)

This Page

[Show Source](#)

Quick search

Enter search terms or a module, class or function name.

# Acknowledgment

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