

# Mathematical Aspects of Computational Chemistry

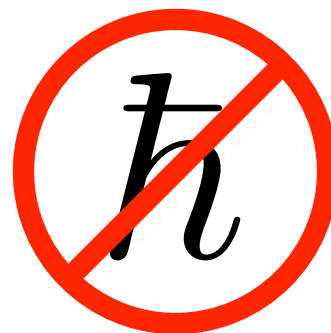
Mark Miller  
Durham Chemistry  
[m.a.miller@durham.ac.uk](mailto:m.a.miller@durham.ac.uk)

15 June 2017

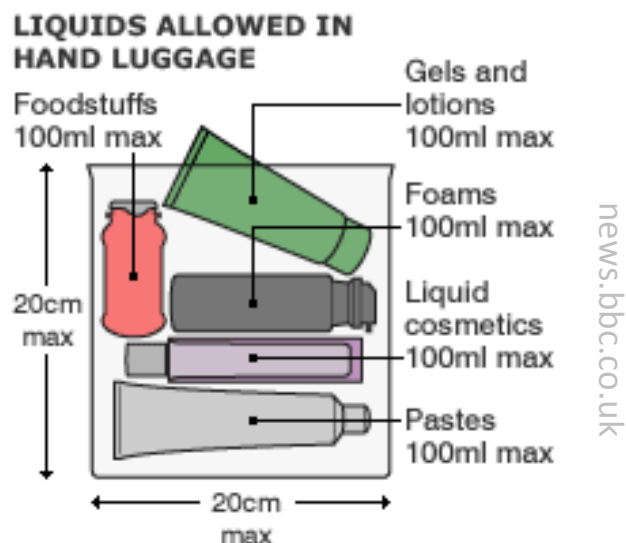


# Research themes

- Computational chemistry (classical versus quantum)



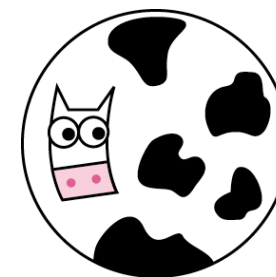
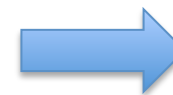
- Soft matter



- Coarse-grained modelling

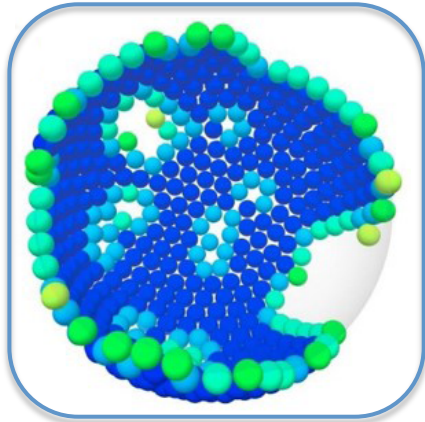


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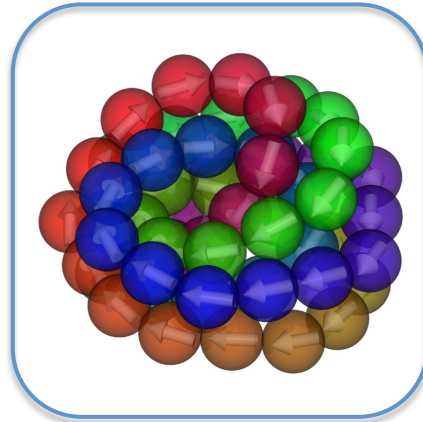


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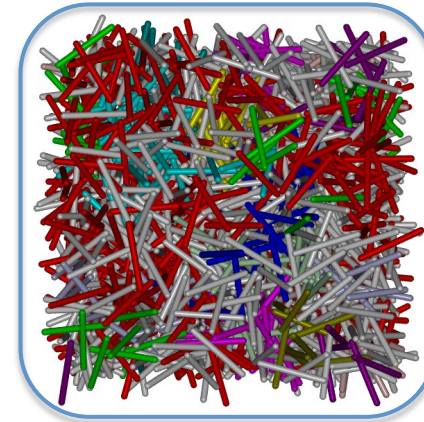
# Survey of projects



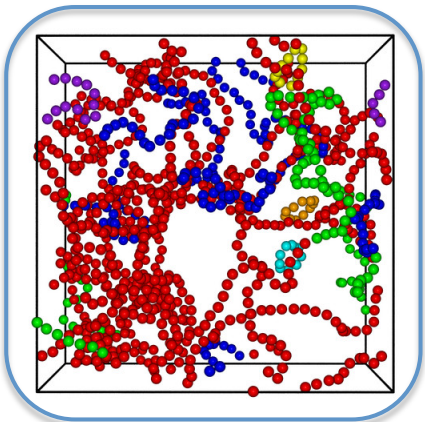
curved surfaces



knots



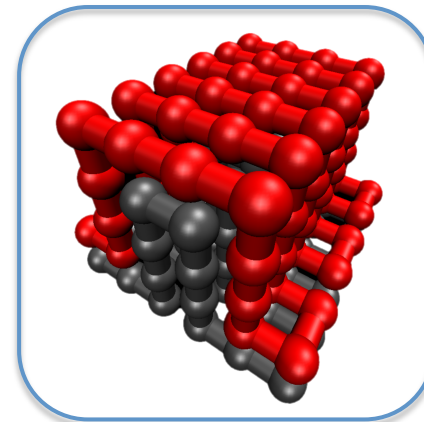
percolation



gels



self assembly



bio-inspired

# 1. Molecular Simulation

Energetic interactions

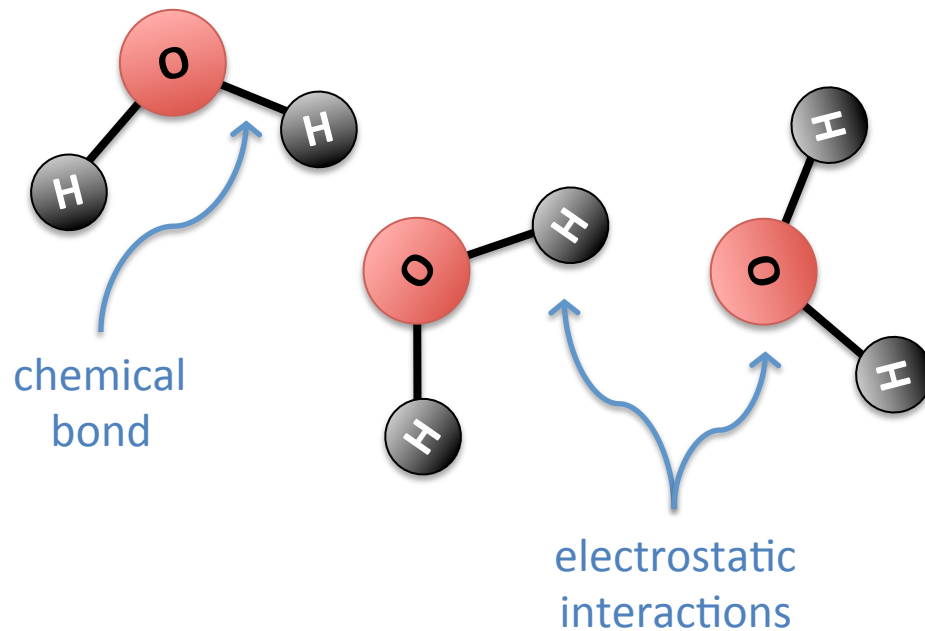
+

General laws of physics



Complex physical and chemical behaviour

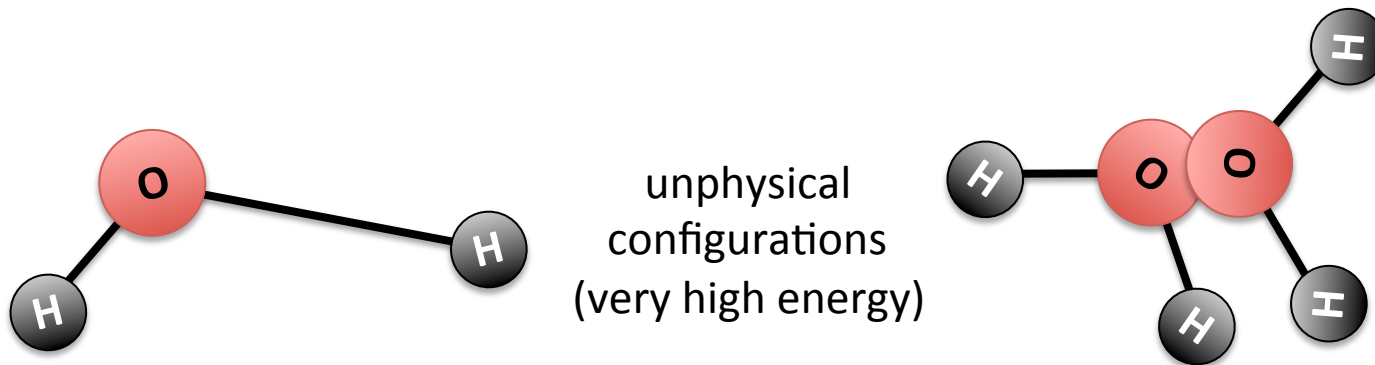
e.g. Newton's laws



- Freezing
- Boiling
- Viscosity
- Solvent properties
- + much more

# Molecular Simulations [cont.]

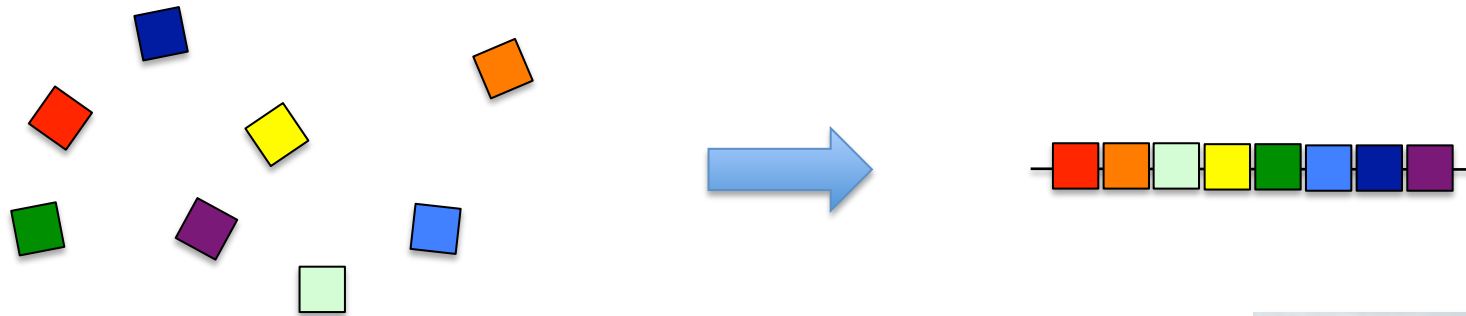
- Configuration  $\mathbf{X}$  of  $N$  atoms:  $\mathbb{R}^{3N}$
- Potential energy:  $3N$ -dimensional scalar field  $V(\mathbf{X})$
- Equilibrium statistical weight  $\rho^{\text{eqm}}(\mathbf{X}) \propto e^{-V(\mathbf{X})/kT}$



- Only a tiny fraction of configuration space is relevant
- Must navigate correctly:  $\hat{A}\rho^{\text{eqm}} = \rho^{\text{eqm}}$
- and efficiently:  $\lim_{n \rightarrow \infty} \hat{A}^n \rho = \rho^{\text{eqm}}$

## 2. A Constrained Optimisation

- Addressable self-assembly
  - Spontaneous formation of a target structure
  - All components of target are unique



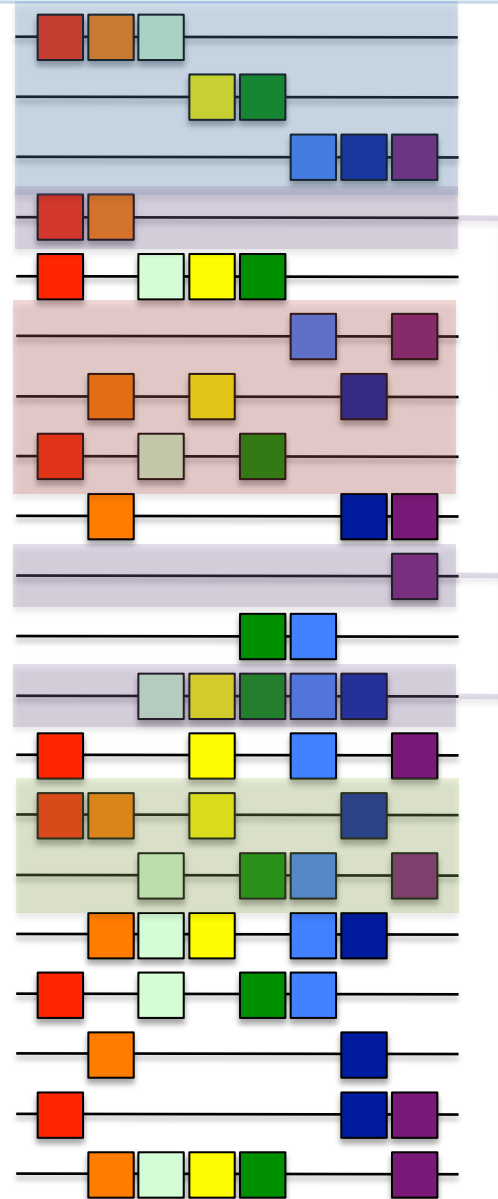
- Possible obstacles
  - Incorrect fragments (wrong partners bind)
  - Multiple correct fragments compete



Dr David Bourne

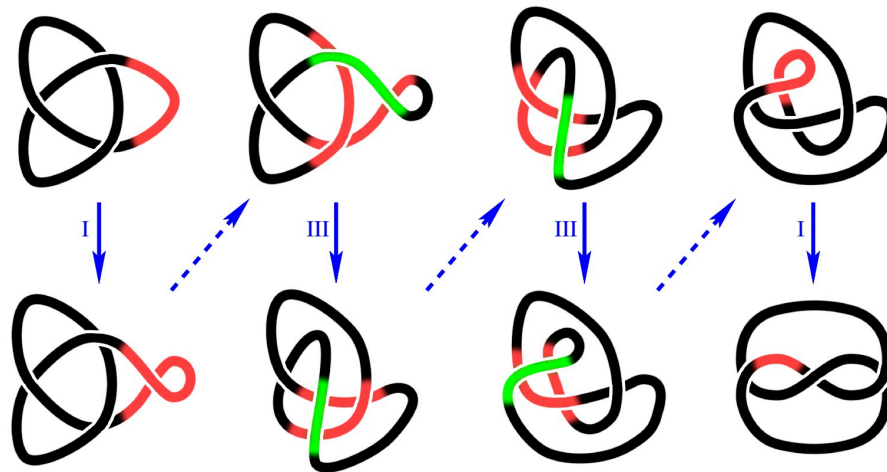
# A Constrained Optimisation [cont.]

- How frustrated are a given set of fragments?
- Combine into largest possible sets:
  - using each fragment no more than once
  - with no repeats of a given component in a set
- Constrained integer optimisation
- Quantify “completability”



# 3. Topological Invariants

- Knots and links
  - Identification by topological invariants
  - Polynomial invariants: Jones, HOMFLY...

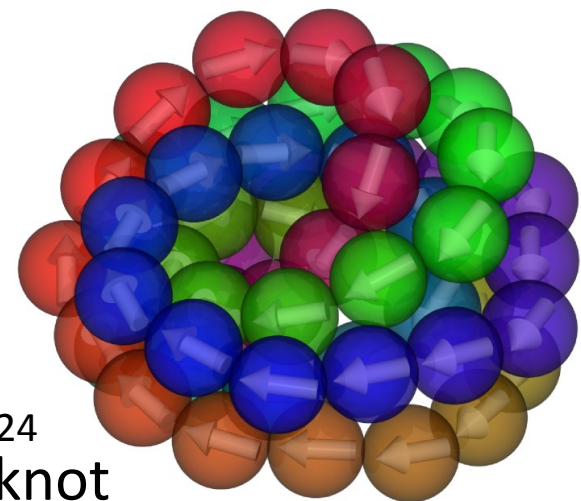


trefoil knot

$$V(t) = t + t^3 - t^4$$

- Knotted clusters
  - Energy optimisation
  - Rearrangement pathways

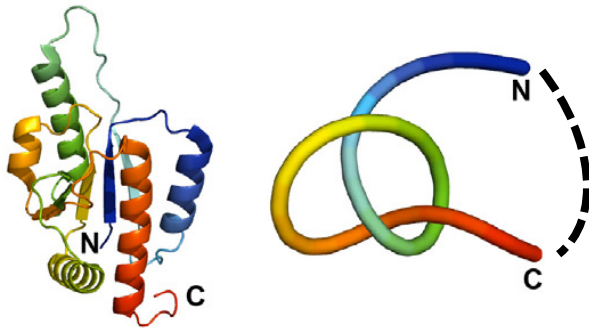
$10_{124}$   
torus knot





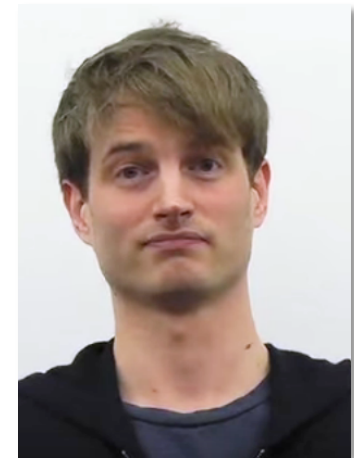
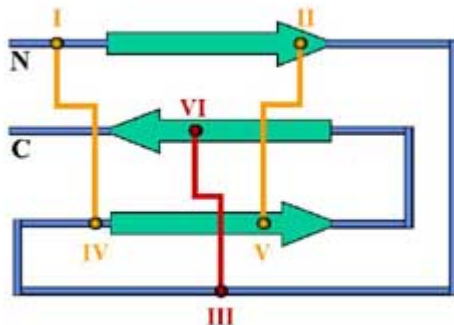
# Topological Invariants [cont.]

- Knotted proteins
  - Closure needed for open chain



Lim & Jackson  
J Phys Cond Matt **27** 354101 (2015)

- Bonding networks
  - Topological invariants of (branched) graphs
  - Yamada polynomial



Dr Andrew Lobb