

# Monte Carlo Simulations with ProtoMS

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# Monte Carlo Simulations with ProtoMS

## Workshop Outline

<i>Time</i>	<i>Topic</i>
<i>10:00</i>	<i>Monte Carlo Simulation and the ProtoMS Code</i> <i>Dr. Chris Cave-Ayland, University of Southampton</i>
<i>10:15</i>	<i>Exercise 1</i> <i>Simulating a Protein-Ligand System</i>
<i>11:30</i>	<i>Introduction to GCMC</i> <i>Hannah Bruce-Macdonald, University of Southampton</i>
<i>11:45</i>	<i>Exercise 2</i> <i>GCMC Calculations for Water Binding</i>

# Monte Carlo Simulations with ProtoMS

## Development History



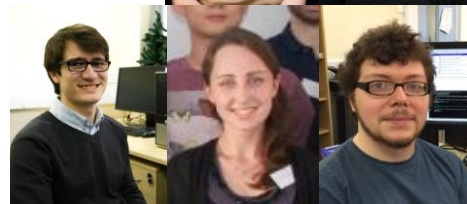
v1.0

2006



v2.4

2014



v3.3

v2.1

v3.0

2002



2013



2017  
(last week)



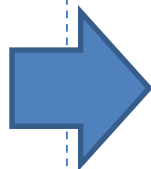
# Monte Carlo Simulations with ProtoMS

## Execution Outline

### Setup



- Parameterise small molecules
- Create input files
- Modular tools
- Master script
- Setup wizard



### Simulation

Fortran 77

- Metropolis Monte Carlo
- Alchemical Transformations
- **Grand Canonical Ensemble**



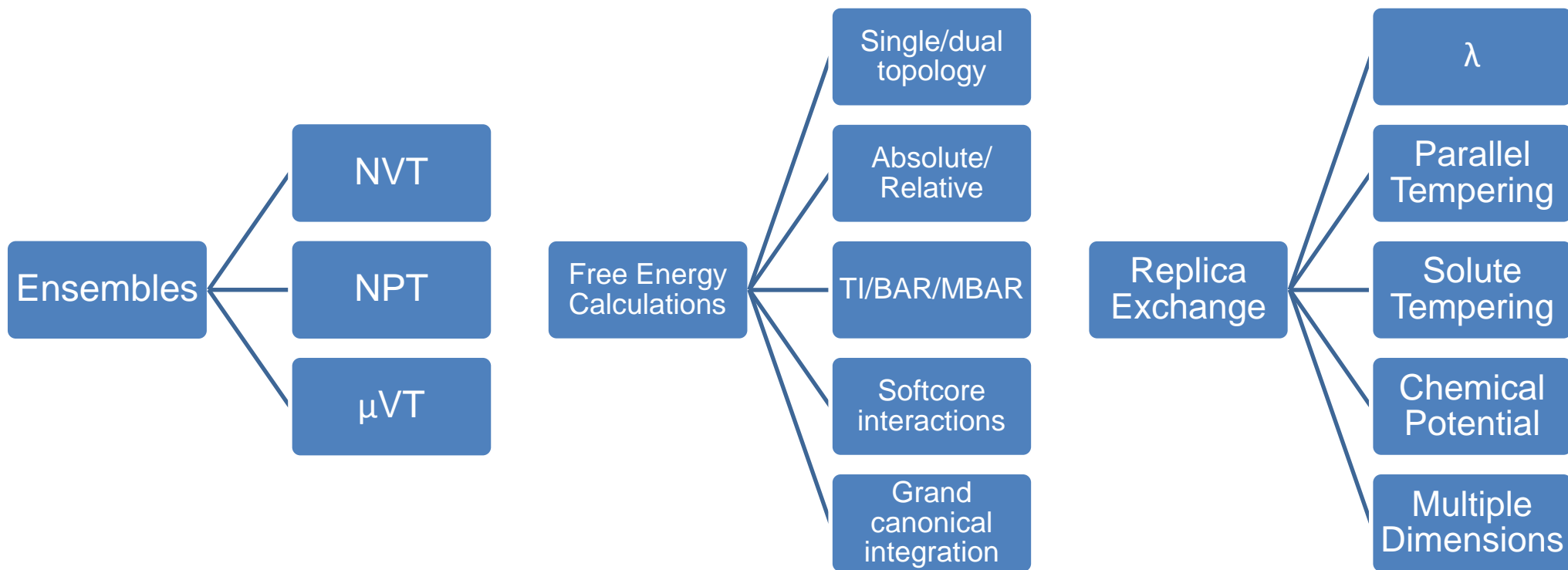
### Analysis



- Extract system properties
- Calculate free energies
- Clustering

# Monte Carlo Simulations with ProtoMS

## Code Capabilities



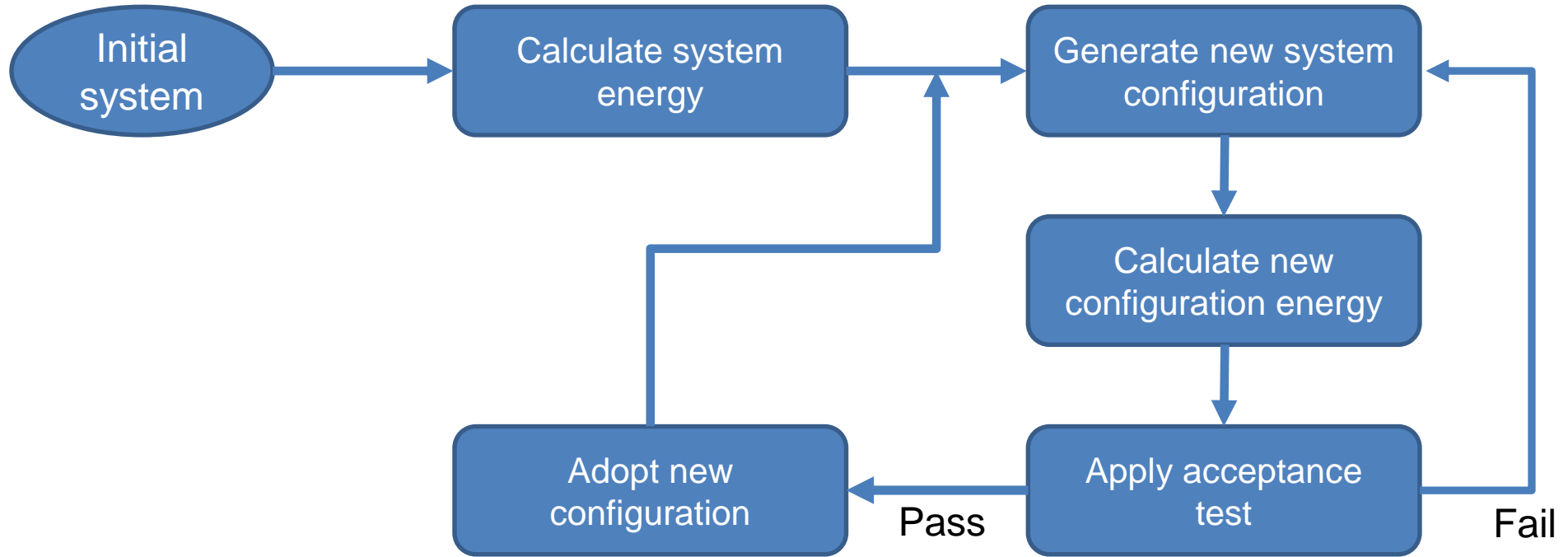
# Monte Carlo Simulations with ProtoMS

## Development Processes

- Developed with Mercurial, hosted on Bitbucket
- Bamboo continuous integration server
- SSI Collaboration
  - Test Suite
  - Usability review
- Licenced under GPL v2
- [www.protoms.org](http://www.protoms.org)
- ~10 publications – application note in preparation

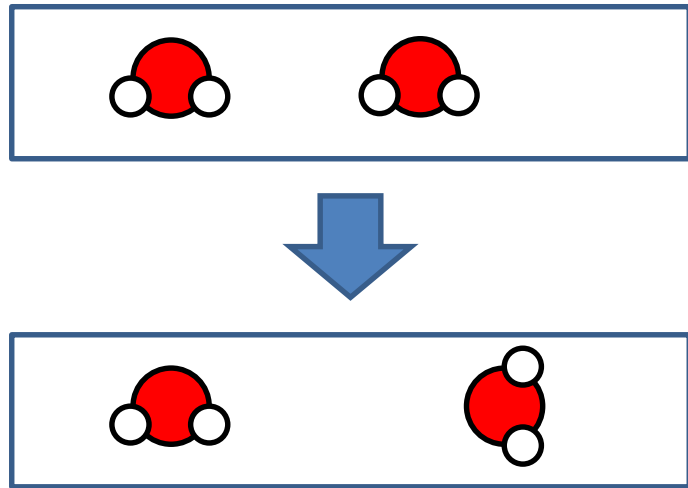
# Monte Carlo Simulations with ProtoMS

## Metropolis Monte Carlo Algorithm



# Monte Carlo Simulations with ProtoMS

## Move Types - Displacement



Test:

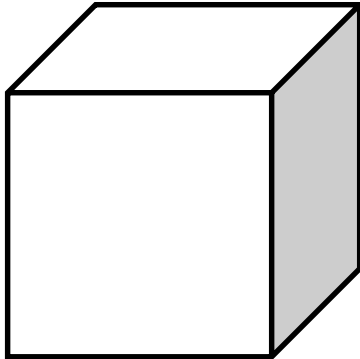
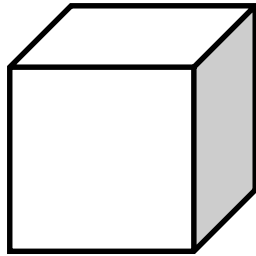
$$P_{accept} = \min(1, \exp[-\beta\Delta E])$$

- Single or multiple atoms
- Test assumes symmetric proposal distributions



# Monte Carlo Simulations with ProtoMS

## Move Types - Volume



Test:

$$P_{accept}$$

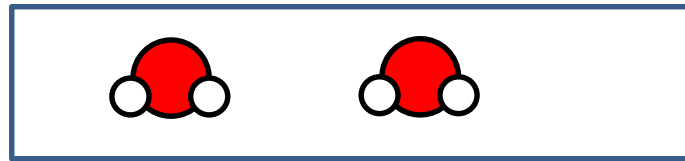
$$= \min(1, \exp[-\beta(\Delta E + p\Delta V) + N \ln \left(\frac{V_{new}}{V_{old}}\right)])$$

- Usually displace particles as well

# Monte Carlo Simulations with ProtoMS

## Move Types – Insertion/Deletion

Test:



Depends

- Variable particle number – GCMC

# Monte Carlo Simulations with ProtoMS

## MC Overview

### Advantages

- Only evaluate energy terms that change
- Free choice of move proposal distributions
- Can focus sampling on interesting parts
- Easy to impose constraints
- Vary particle number

### Disadvantages

- Parallelisation
- Need to define move proposal distributions
- Non-pairwise interactions

# Monte Carlo Simulations with ProtoMS

To get started:

<https://sites.google.com/site/protomsgcmcworkshop/>