

Simulating models of polymer collapse

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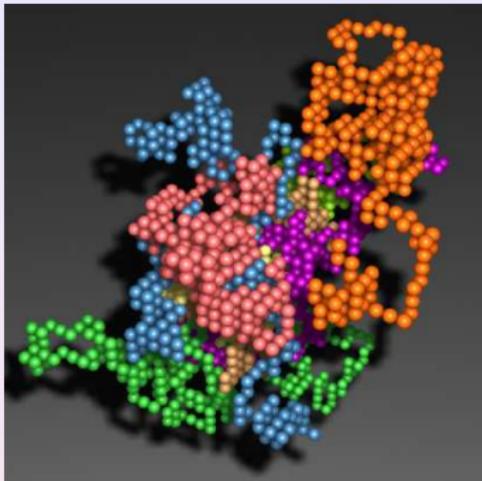
- Polymers in solution:
 - Equilibrium statistical mechanics, lattice model
- Algorithm:
 - Stochastic growth & flat histogram (PERM/flatPERM)
- Simulation of the canonical model:
 - Interacting self-avoiding walks (ISAW)

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 - Interacting self-avoiding walks (ISAW)
- Applications:
 - Protein groundstates (HP model)
 - Bulk vs surface phenomena:
 - confined polymers, force-induced desorption, interplay of collapse and adsorption
 - Comparison of alternative lattice models
 - Hydrogen-bond type interactions

Polymers in Solution

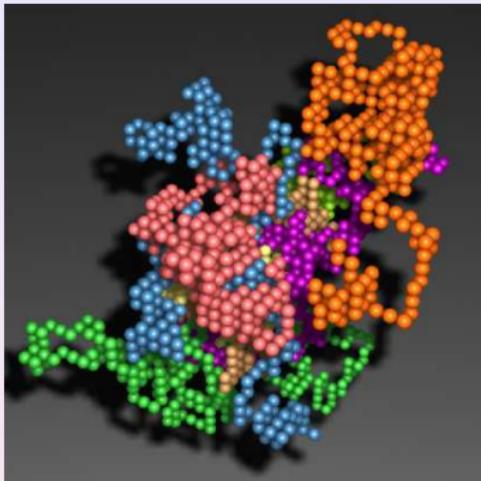
Modelling of Polymers in Solution

- Polymers:
long chains of monomers
- “Coarse-Graining”:
beads on a chain
- “Excluded Volume”:
minimal distance between beads
- Contact with solvent:
effective short-range interaction
- Good/bad solvent:
repelling/attracting interaction



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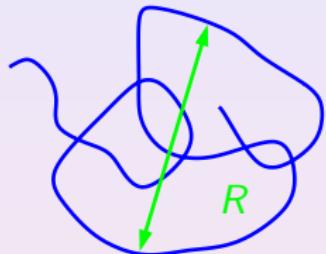


A Model of a Polymer in Solution

Random Walk + Excluded Volume + Short Range Attraction

Polymer Collapse, Coil-Globule Transition, Θ -Point

length N , spatial extension $R \sim N^\nu$



$T > T_c$: good solvent
swollen phase (coil)



$T = T_c$:
 Θ -polymer

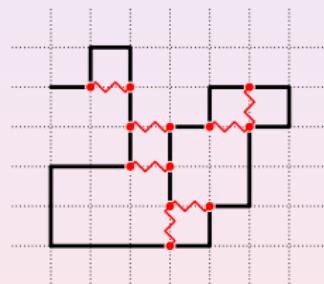
$T < T_c$: bad solvent
collapsed phase (globule)



The Canonical Lattice Model

Interacting Self-Avoiding Walk (ISAW)

- Physical space → simple cubic lattice \mathbb{Z}^3
- Polymer → self-avoiding random walk (SAW)
- Quality of solvent → short-range interaction ϵ



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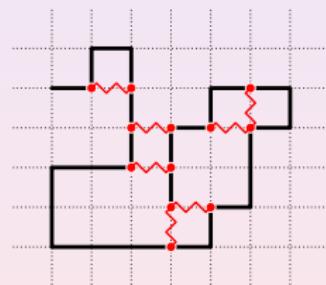
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$C_{N,m}$ is the number of SAWs
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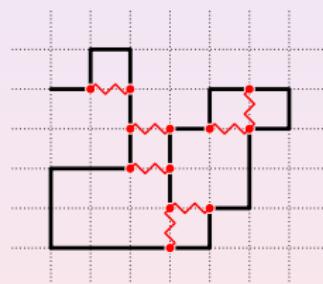
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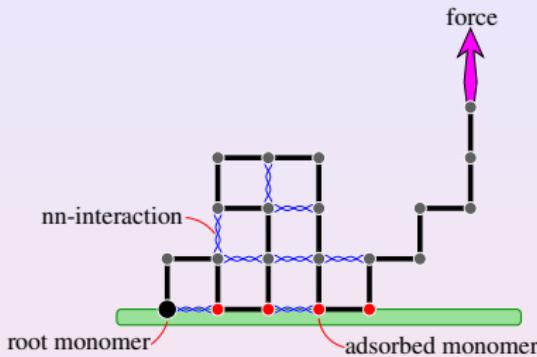


Thermodynamic Limit for a dilute solution:

$$V = \infty \quad \text{and} \quad N \rightarrow \infty$$

Extensions of the Model

- In addition to
 - solvent modelling
(bulk interaction)
- add
 - adsorption
(surface interaction)
 - micromechanical deformations
 - e.g. force on chain end
(optical tweezers)
- Complete description through three-dimensional density of states:
 - (a) bulk energy, (b) surface energy, (c) position of chain end



The Algorithm

PERM: “Go With The Winners”

PERM = Pruned and Enriched Rosenbluth Method

P Grassberger, Phys Rev E 56 (1997) 3682

- Rosenbluth Method: kinetic growth



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- Pruning: weight too small \rightarrow remove configuration occasionally

Current work: flatPERM = flat histogram PERM

T Prellberg and J Krawczyk, PRL 92 (2004) 120602

- flatPERM samples a generalised multicanonical ensemble
- Determines the whole density of states in *one* simulation!

Algorithm details

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- Add pruning/enrichment with respect to ratio
 $r = W_N^{(S+1)} / C_N^{\text{est}}$
 - Number of samples generated for each N is roughly constant
 - We have a flat histogram algorithm in system size

From PERM to flatPERM

- Consider athermal case
 - PERM: estimate number of configurations C_N
 - $C_N^{est} = \langle W \rangle_N$
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 - thermal PERM: estimate partition function $Z_N(\beta)$
 - $Z_N^{est}(\beta) = \langle W \exp(-\beta E) \rangle_N$
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 - $r = W_N^{(i)} \exp(-\beta E^{(i)}) / Z_N^{est}(\beta)$
- Consider parametrisation \vec{m} of configuration space
 - flatPERM: estimate density of states $C_{N,\vec{m}}$
 - $C_{N,\vec{m}}^{est} = \langle W \rangle_{N,\vec{m}}$
 - $r = W_{N,\vec{m}}^{(i)} / C_{N,\vec{m}}^{est}$

Why Simulations?

- Most interesting open questions for dense and geometrically restricted configurations

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There is little theory and this is notoriously difficult to simulate

Simulations and Results

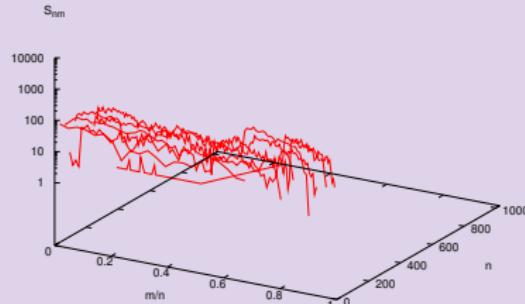
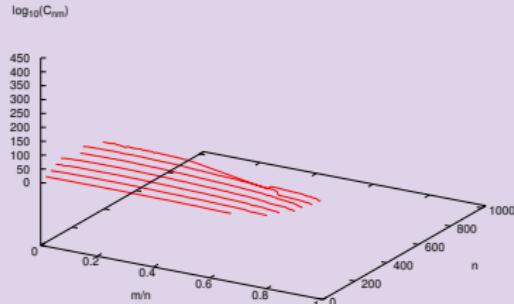
To stabilise algorithm (avoid initial overflow/underflow):

Delay growth of large configurations

Here: after t tours growth up to length $10t$

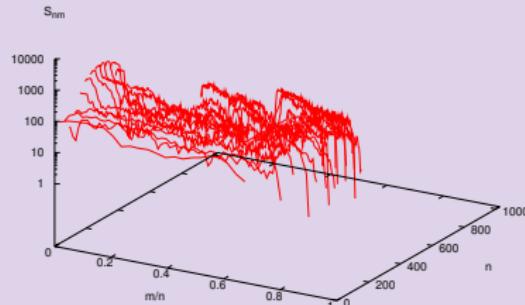
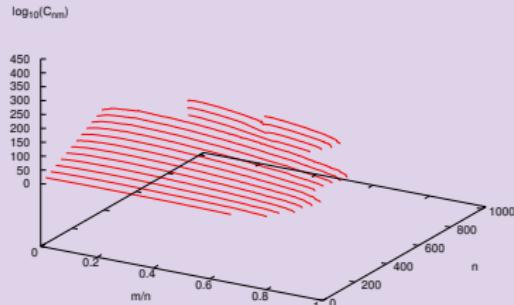
2d ISAW simulation up to $N = 1024$

Total sample size: 1,000,000



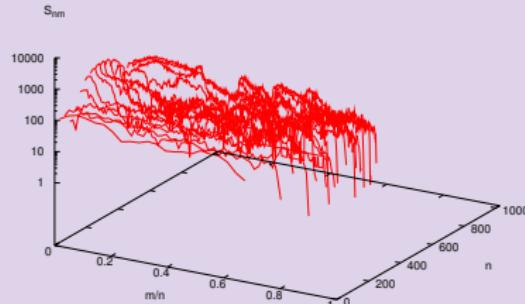
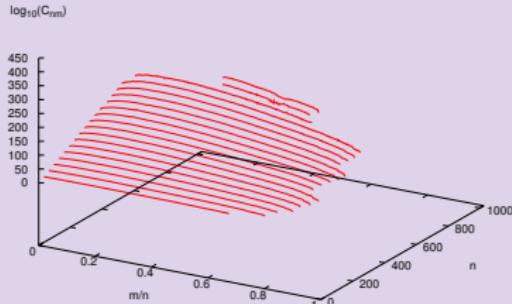
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Total sample size: 10,000,000



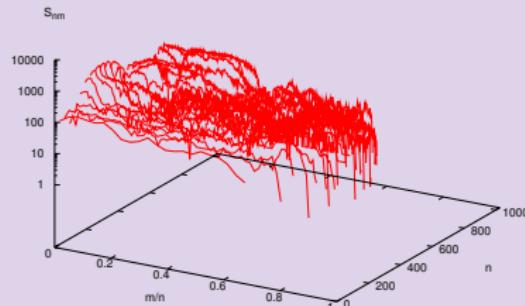
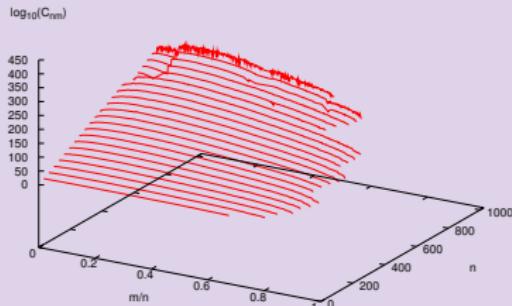
2d ISAW simulation up to $N = 1024$

Total sample size: 20,000,000



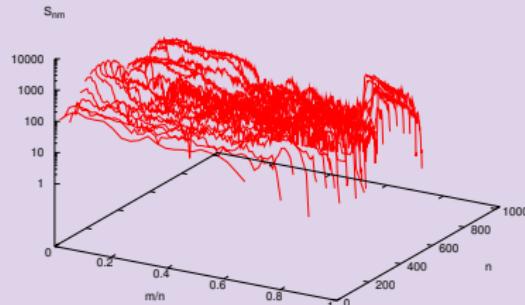
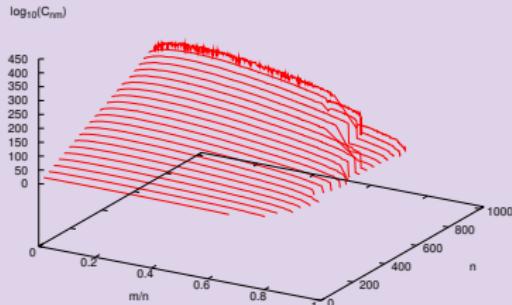
2d ISAW simulation up to $N = 1024$

Total sample size: 30,000,000



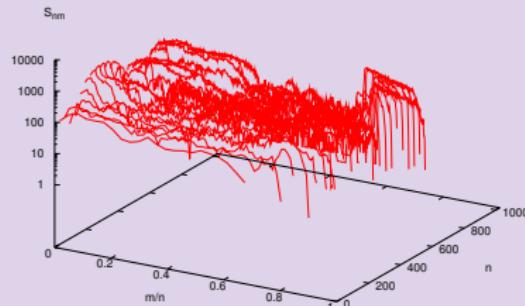
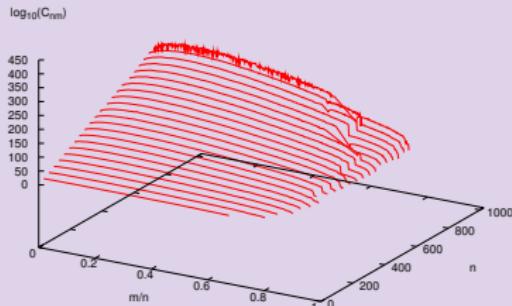
2d ISAW simulation up to $N = 1024$

Total sample size: 40,000,000



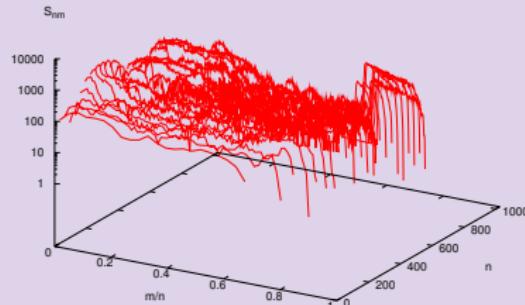
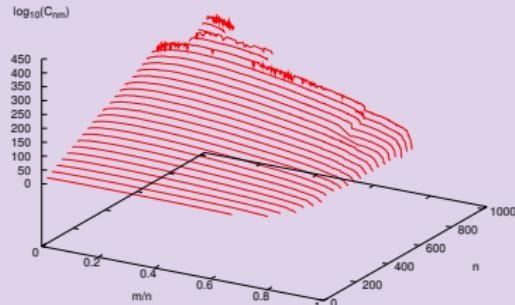
2d ISAW simulation up to $N = 1024$

Total sample size: 50,000,000



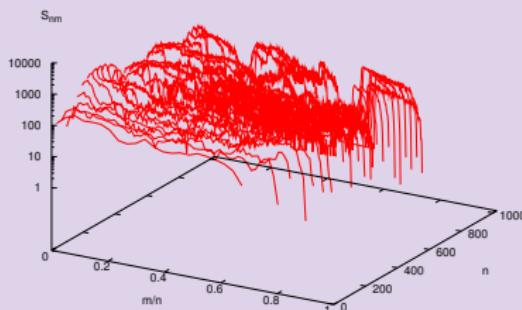
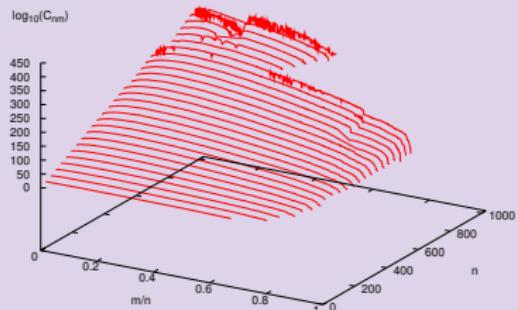
2d ISAW simulation up to $N = 1024$

Total sample size: 60,000,000



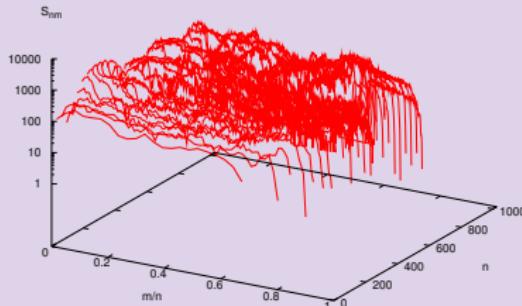
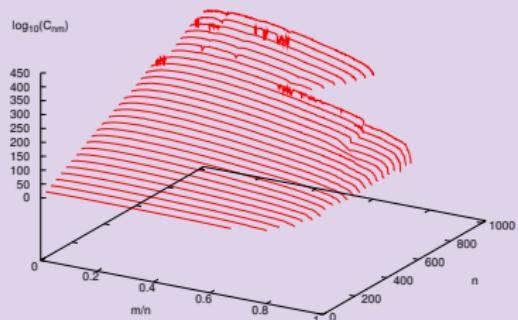
2d ISAW simulation up to $N = 1024$

Total sample size: 70,000,000



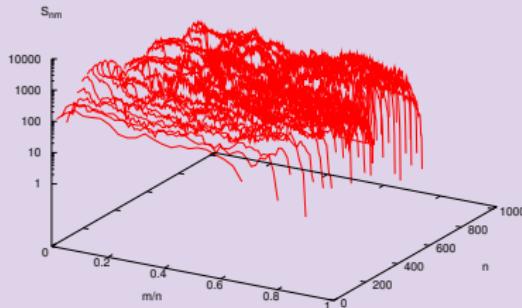
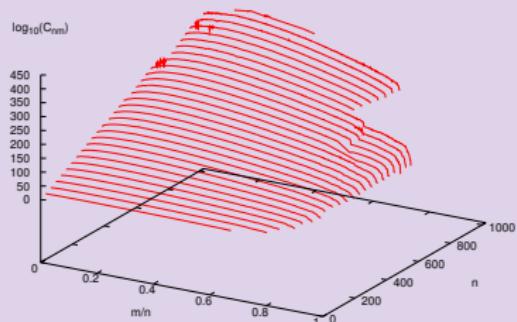
2d ISAW simulation up to $N = 1024$

Total sample size: 80,000,000



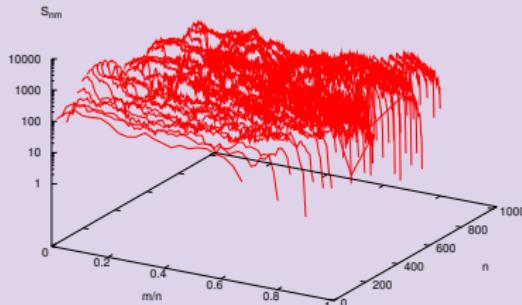
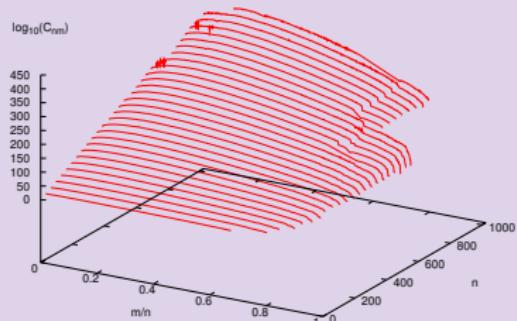
2d ISAW simulation up to $N = 1024$

Total sample size: 90,000,000



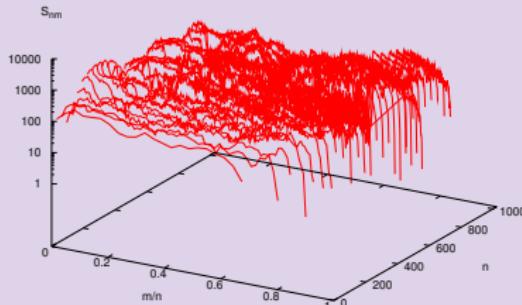
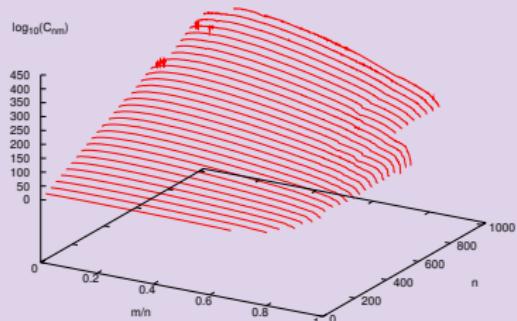
2d ISAW simulation up to $N = 1024$

Total sample size: 100,000,000



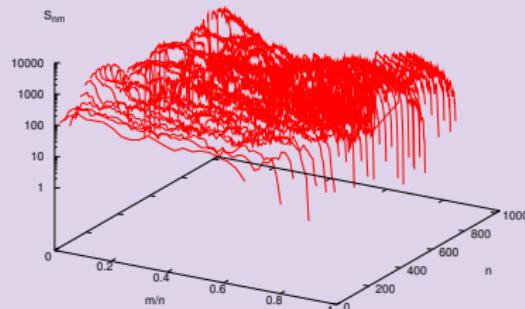
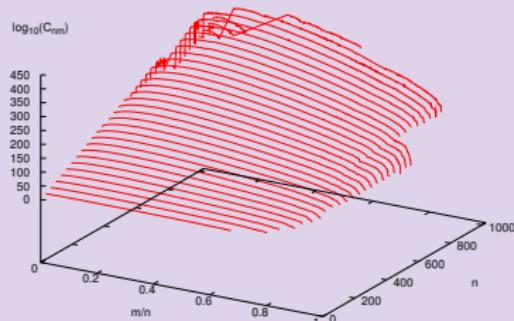
2d ISAW simulation up to $N = 1024$

Total sample size: 110,000,000



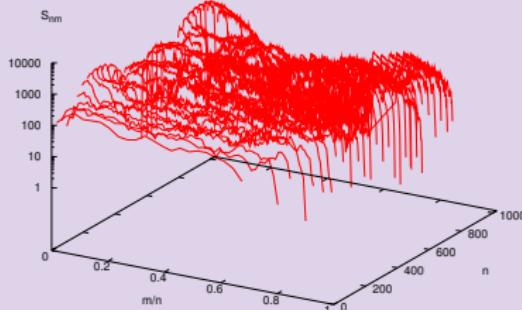
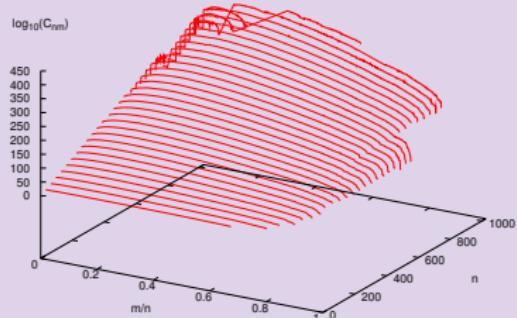
2d ISAW simulation up to $N = 1024$

Total sample size: 120,000,000



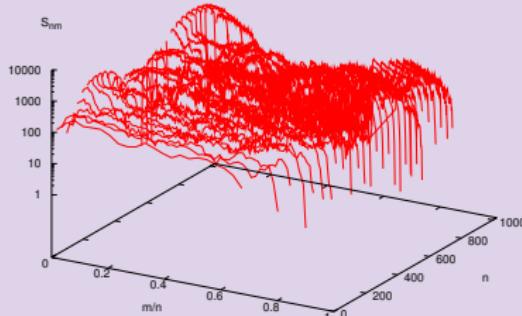
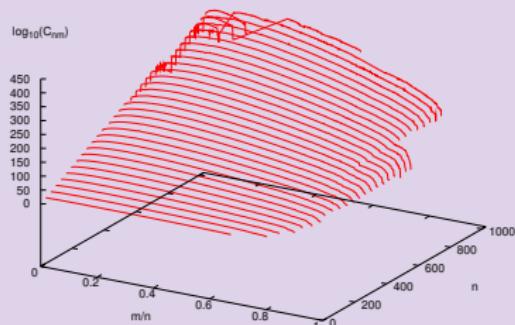
2d ISAW simulation up to $N = 1024$

Total sample size: 130,000,000



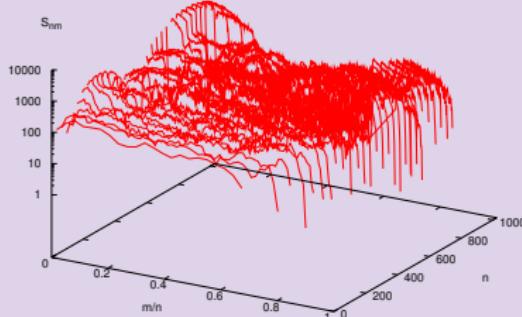
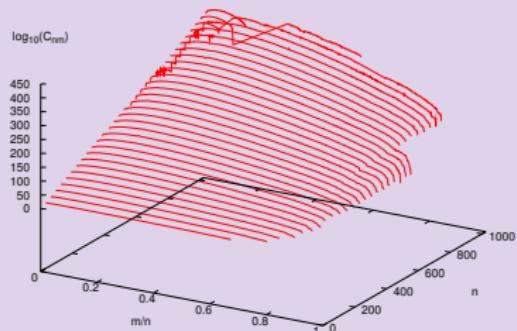
2d ISAW simulation up to $N = 1024$

Total sample size: 140,000,000



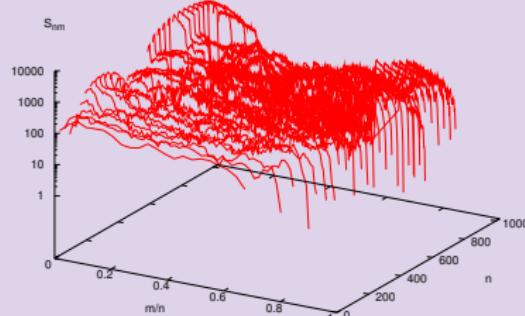
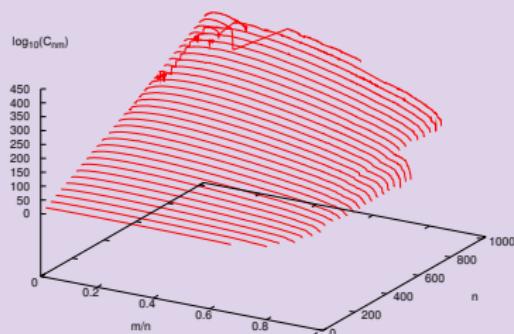
2d ISAW simulation up to $N = 1024$

Total sample size: 150,000,000



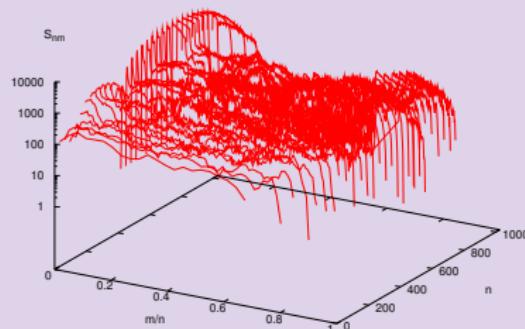
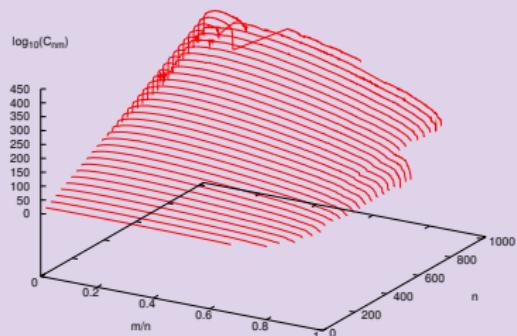
2d ISAW simulation up to $N = 1024$

Total sample size: 160,000,000



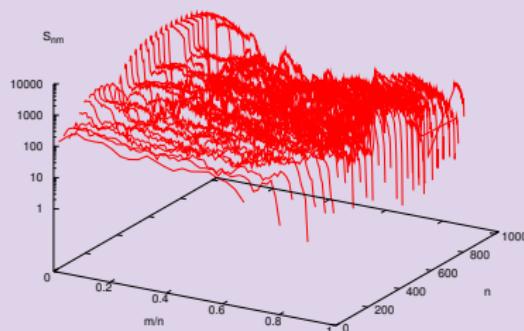
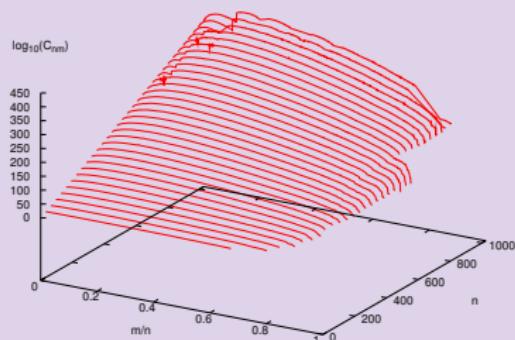
2d ISAW simulation up to $N = 1024$

Total sample size: 170,000,000



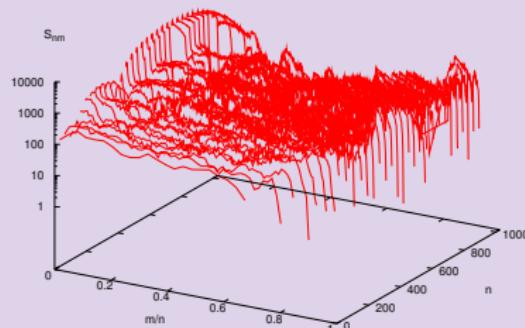
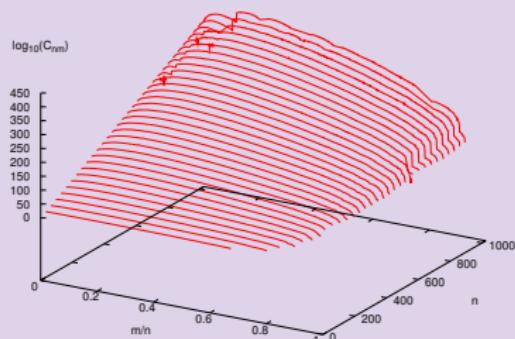
2d ISAW simulation up to $N = 1024$

Total sample size: 180,000,000



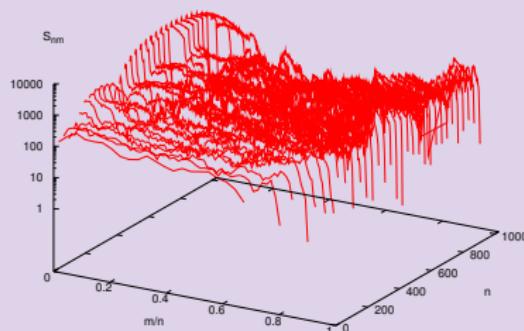
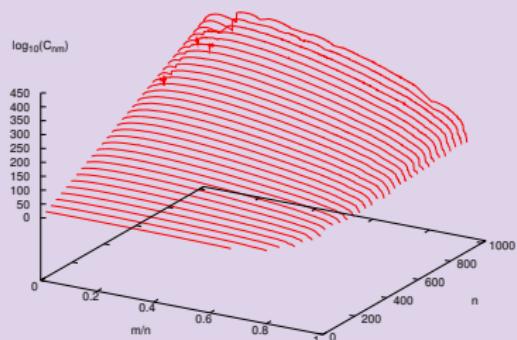
2d ISAW simulation up to $N = 1024$

Total sample size: 190,000,000



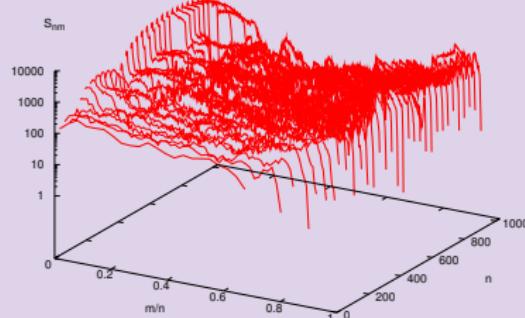
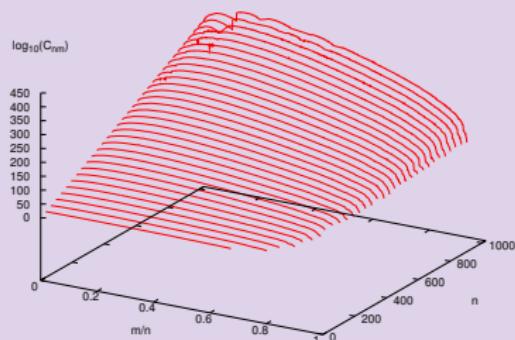
2d ISAW simulation up to $N = 1024$

Total sample size: 200,000,000



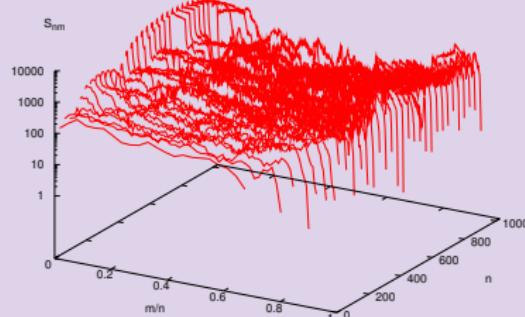
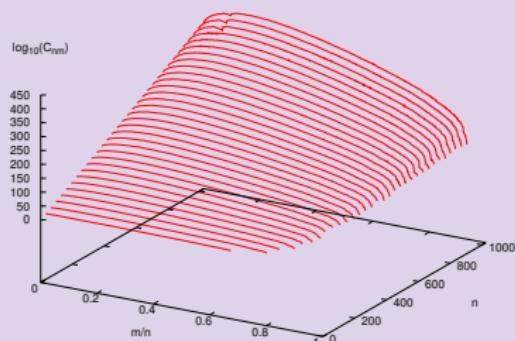
2d ISAW simulation up to $N = 1024$

Total sample size: 210,000,000



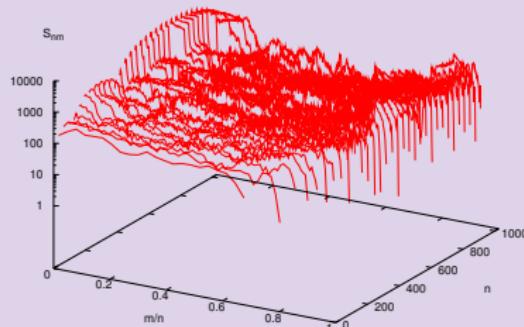
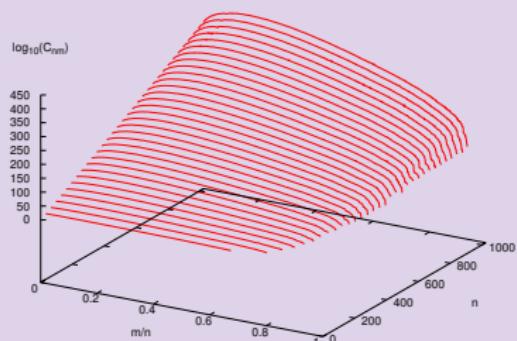
2d ISAW simulation up to $N = 1024$

Total sample size: 220,000,000



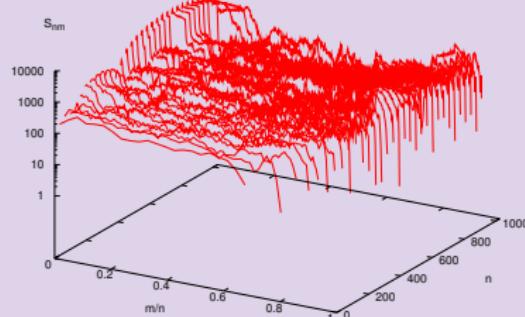
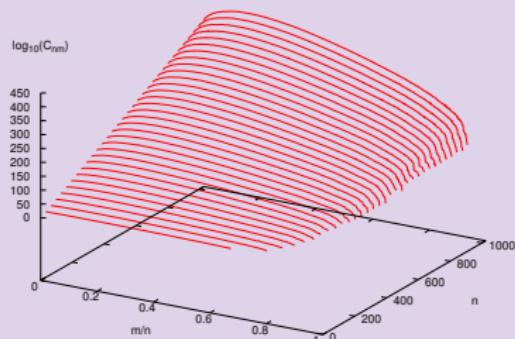
2d ISAW simulation up to $N = 1024$

Total sample size: 230,000,000



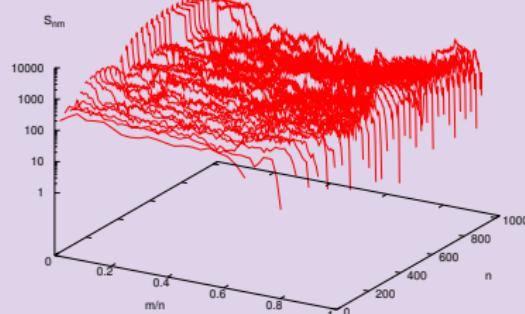
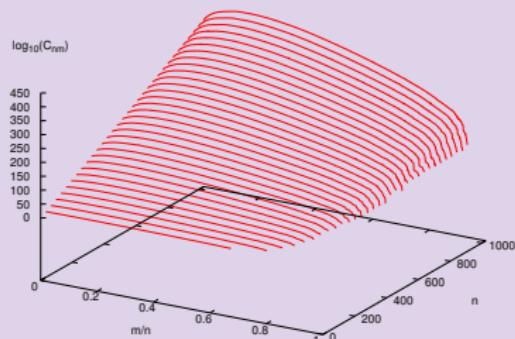
2d ISAW simulation up to $N = 1024$

Total sample size: 240,000,000



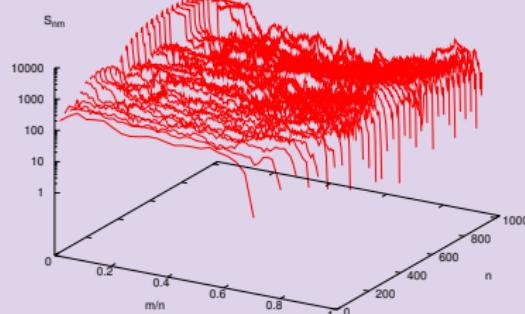
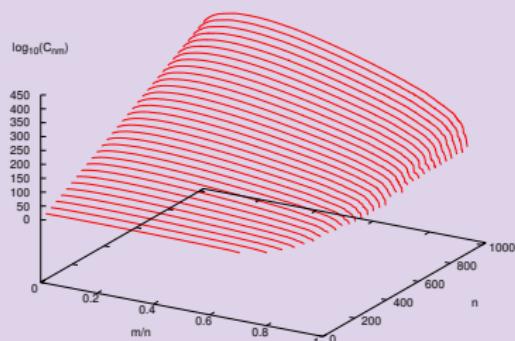
2d ISAW simulation up to $N = 1024$

Total sample size: 250,000,000



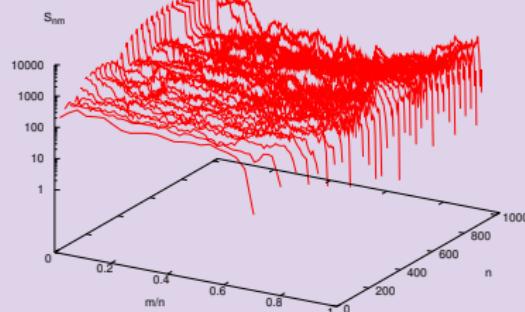
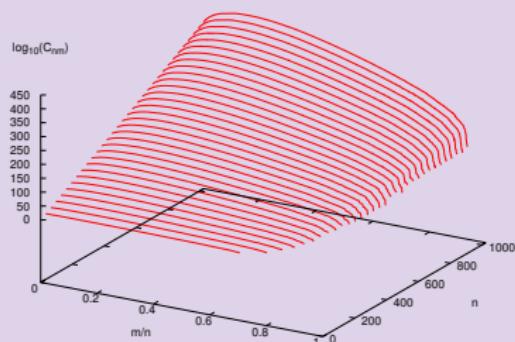
2d ISAW simulation up to $N = 1024$

Total sample size: 260,000,000



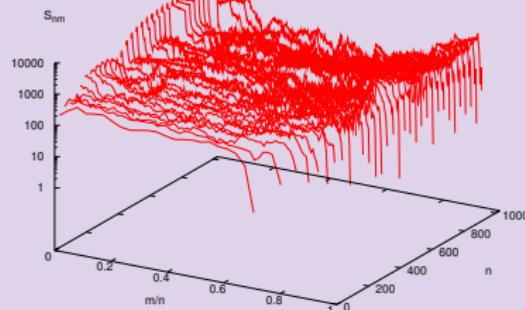
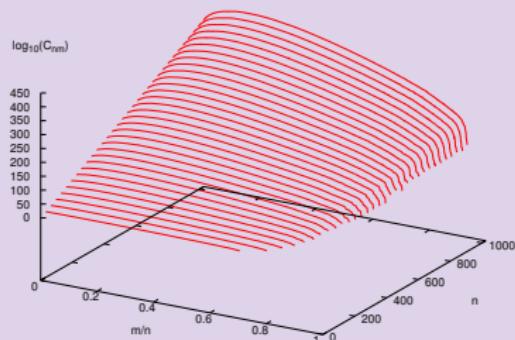
2d ISAW simulation up to $N = 1024$

Total sample size: 270,000,000



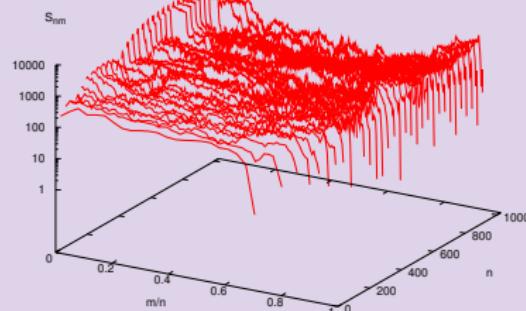
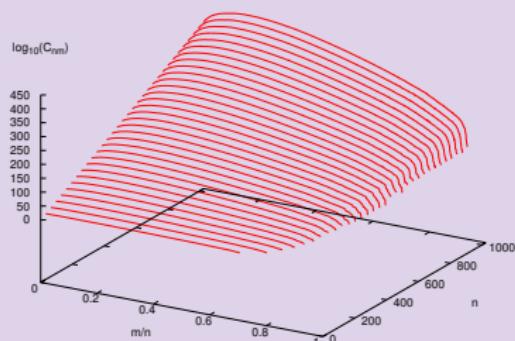
2d ISAW simulation up to $N = 1024$

Total sample size: 280,000,000



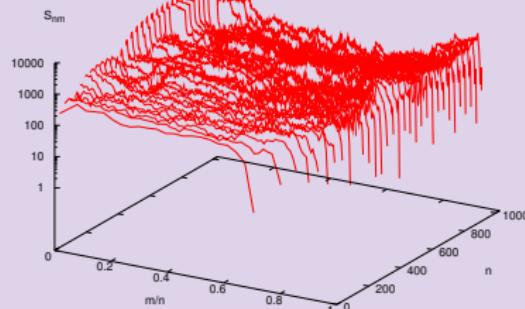
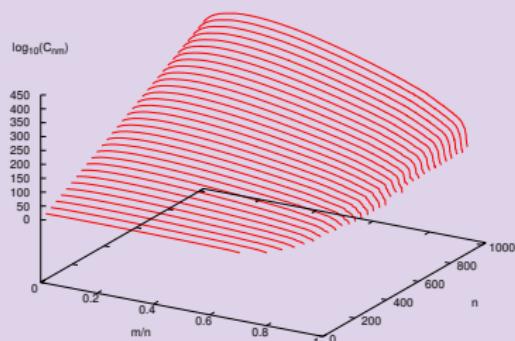
2d ISAW simulation up to $N = 1024$

Total sample size: 290,000,000



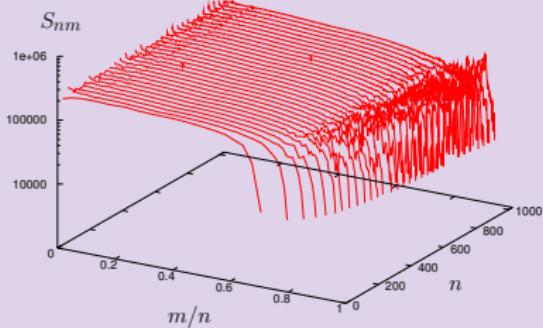
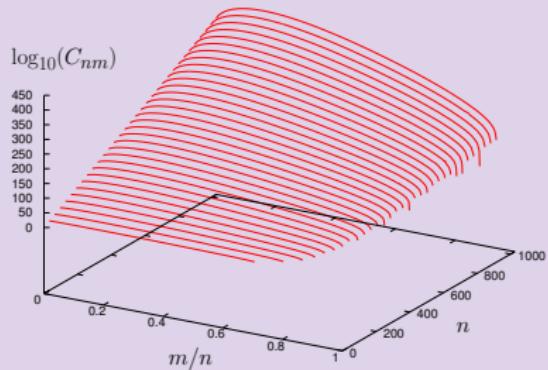
2d ISAW simulation up to $N = 1024$

Total sample size: 300,000,000

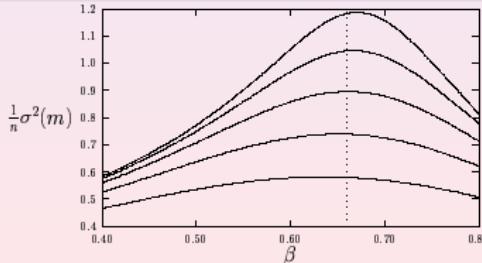


ISAW simulations

T Prellberg and J Krawczyk, PRL 92 (2004) 120602



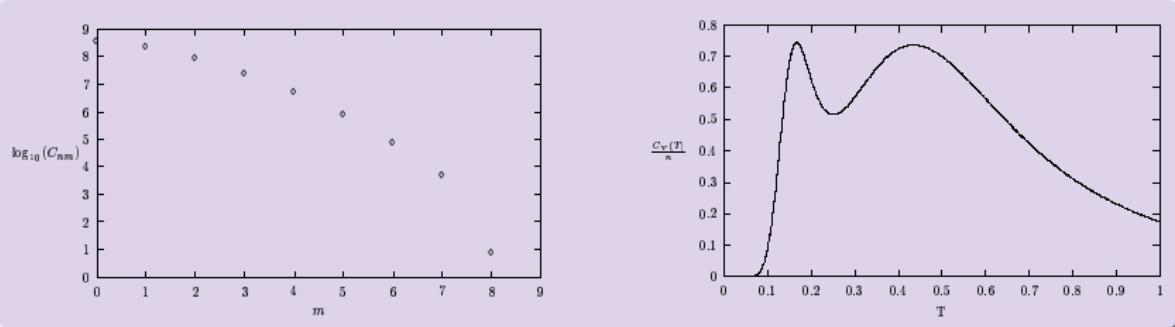
- 2d ISAW up to $n = 1024$
- One simulation suffices
- 400 orders of magnitude
(only 2d shown, 3d similar)



HP model simulations

T Prellberg et al, Computer Simulation Studies in Condensed Matter Physics XVII, pages 122-135, Springer Verlag, 2006

- Engineered sequence HPHPHHPHPHHPPH in $d = 3$:

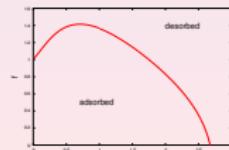
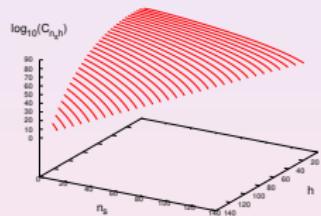
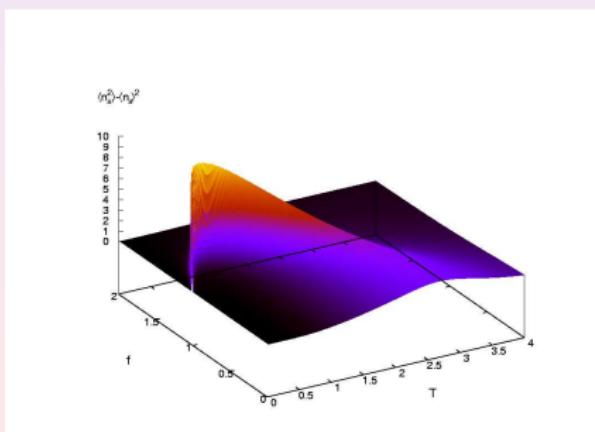


- Investigated other sequences up to $N \approx 100$ in $d = 2$ and $d = 3$
- Collapsed regime accessible
- Reproduced known ground state energies
- Obtained density of states $C_{n,m}$ over large range ($\approx 10^{30}$)

2-Dimensional Density of States

J Krawczyk et al, JSTAT (2004) P10004

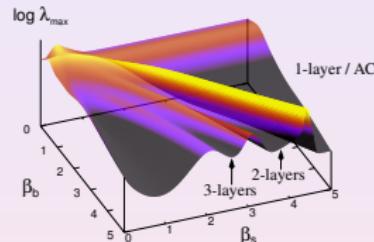
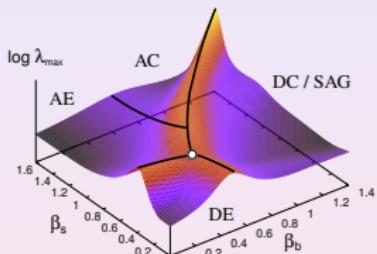
- Force-induced desorption of adsorbed polymers
 - Relevance: optical tweezers, AFM; related to DNA unzipping
- 3-dim polymer in a half space, one simulation, up to $n = 256$
 - Fluctuations of surface coverage



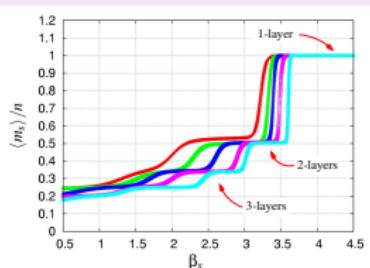
2-Dimensional Density of States

J Krawczyk et al, Europhys. Lett. 70 (2005) 726-732

- Layering transitions of adsorbed polymers in poor solvents



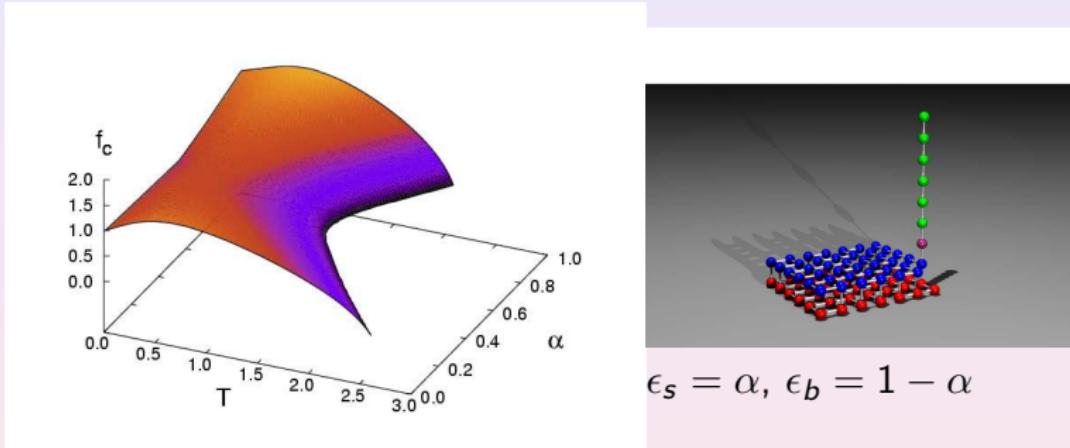
- whole phase diagram at once
- low temperatures accessible
- hierarchy of layering transitions
- resolved controversy over “surface attached globule”



3-Dimensional Density of States

J Krawczyk et al, JSTAT (2005) P05008

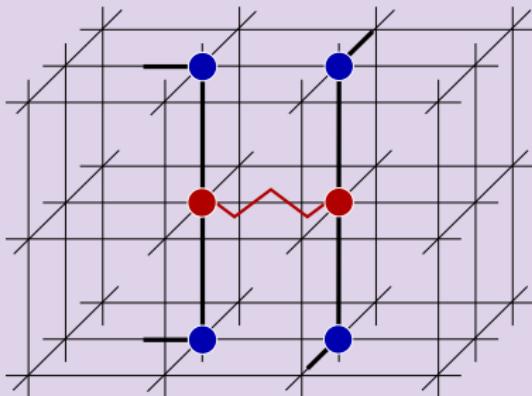
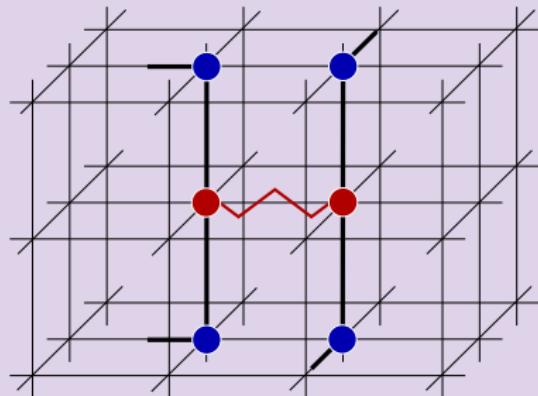
- Pulling adsorbing and collapsing polymers off a surface



- simulations up to $n = 91$ (4-dimensional histogram)
- interplay of (both force-induced and thermal) desorption ($\alpha = 1$) and stretching ($\alpha = 0$)

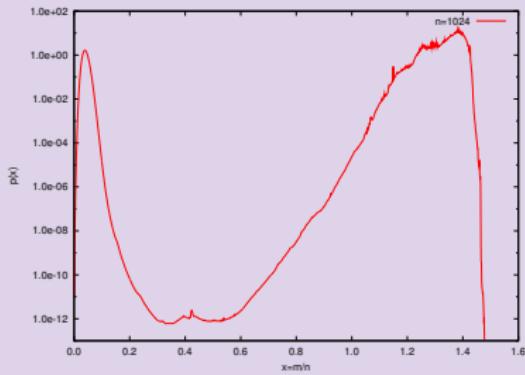
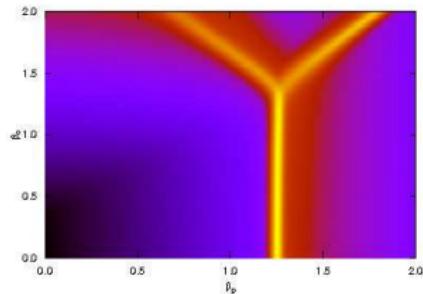
Hydrogen-bond type interactions

J Krawczyk et al, in preparation

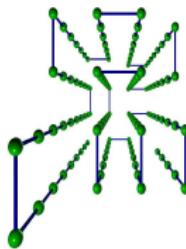
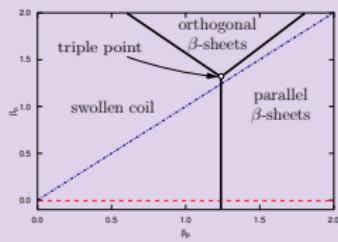
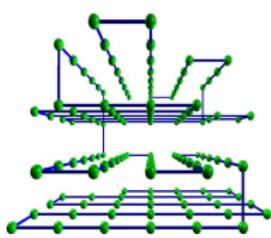
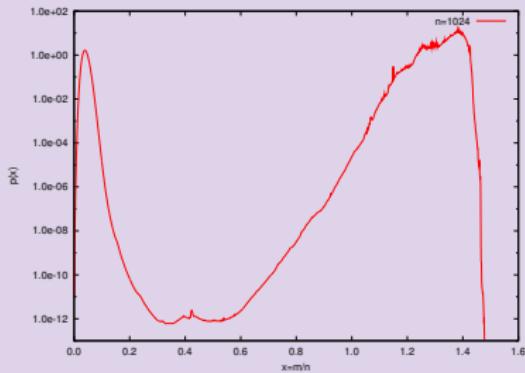
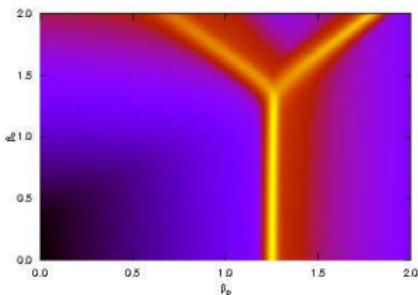


Introduce orientation-dependent (parallel/orthogonal) interactions between *straight* segments of the walk

Hydrogen-bond type interactions (ctd.)



Hydrogen-bond type interactions (ctd.)



Acknowledgements

Joined work with A.L. Owczarek, A. Rechnitzer, J. Krawczyk

- The algorithm:

- T. Prellberg and J. Krawczyk, "Flat histogram version of the pruned and enriched Rosenbluth method," Phys. Rev. Lett. 92 (2004) 120602; selected for Virt. J. Biol. Phys. Res. 7 (2004)
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- Alternative lattice models
 - A. L. Owczarek and T. Prellberg, "Collapse transition of self-avoiding trails on the square lattice," Physica A 373 (2007) 433-438
 - J. Krawczyk, T. Prellberg, A. L. Owczarek, and A. Rechnitzer, "On a type of self-avoiding random walk with multiple site weightings and restrictions," Phys. Rev. Lett. 96 (2006) 240603; selected for Virt. J. Biol. Phys. Res. 12 (2006)

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- Hydrogen-bond type interactions:
 - J. Krawczyk, A. L. Owczarek, T. Prellberg, and A. Rechnitzer, "Simulation of Lattice Polymers with Hydrogen-Like Bonding," preprint

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