There are 7×10^{26018276} self-avoiding walks of 38 797 311 steps on \mathbb{Z}^3

Nathan Clisby MASCOS, The University of Melbourne

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Self-avoiding walks

- Enumeration
- Direct sampling and weighted sampling (PERM)
- Ingredients for efficiently estimating c_N
 - Global move (pivot)
 - Efficient data structure (SAW-tree)
 - Clever choice of observable
 - Minimizing statistical error
- Results and conclusion





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- A walk on a lattice, step to neighbouring site provided it has not already been visited.
- Models polymers in good solvent limit.
- Exactly captures universal properties such as critical exponents.
- N-step SAW on Z^d is a mapping ω : {0, 1, ..., N} → Z^d with |ω(i + 1) − ω(i)| = 1 for each i (|x| denotes the Euclidean norm of x), and with ω(i) ≠ ω(j) for all i ≠ j.
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Not a SAW





• The number of SAW of length N, c_N , tells us about how many conformations are available to SAW of a particular length:

 $c_N \sim A \ N^{\gamma-1} \mu^N \left[1 + ext{corrections}
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- For \mathbb{Z}^2 , $c_N = 1, 4, 12, 36, 100, 284, 780, 2172, \cdots$
- For \mathbb{Z}^3 , $c_N = 1, 6, 30, 150, 726, 3534, \cdots$
- γ is a *universal* exponent.
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- Rich and active research area (more than 1800 articles in Web of Science with SAW in title / abstract).
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- For 3d lattices: most powerful method "length-doubling algorithm", combines brute force enumeration with inclusion-exclusion. $O(\mu^n) \rightarrow O((\sqrt{2\mu})^n)^1$.
- I think there are strong prospects to apply length-doubling algorithm to other problems, and improve its efficiency.
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- Obvious approach: simple sampling. Generate a simple random walk of length N, calculate probability that RW is self-avoiding. Probability $= c_N/(2d)^N \approx 4.68^N/6^N$ for \mathbb{Z}^3 .
- Can improve slightly: forbid immediate reversals in the walk. Probability $= c_n/2d/(2d-1)^{N-1} \approx 4.68^N/6/5^{N-1}$ for \mathbb{Z}^3 .
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Counting SAW



• Rosenbluth sampling: only choose free edges.

- This introduces bias: compact walks which have few choices available are preferred.
- Correct bias by weighting walks.

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- Weights provide an estimator of c_N , $c_N = \langle W_N \rangle$.
- Two issues:
 - High variance (poor estimator of c_N
 - Attrition still occurs, since walks can become trapped. Can't sample truly long walks (ok up to N of the order of hundreds).





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SAW-tree C

- PERM: Pruned Enriched Rosenbluth Sampling, a variant of sequential importance sampling.
- Prune: low weight walks, either discard with *P* = 0.5 or double weight.
- Enrich: high weight walks, make copies, ensure total weight remains the same.
- PERM: sensible choices for enrichment ensure attrition is eliminated, variance reduced.
- Dramatically better than Rosenbluth sampling, arbitrarily large *N* achievable.
- Sophisticated choices for pruning and enrichment algorithms can reduce correlations and variance.



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SAW-tree

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- Variance of sample is reduced, but not eliminated. (In practice, variance can be essentially eliminated, at the expense of stronger correlation.)
- Intrinsic limit: CPU time O(N) to produce a single walk. (Prohibitive for truly large N).
- Will now describe a method that overcomes each of these deficiencies.





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- Utilise efficient data structures.
- Find a suitable observable, with low variance.
- Design computer experiment to minimise statistical error.
- Will see that working with *fixed length* walks confers dramatic advantage over growth algorithms.





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- Markov chain:
 - Select a pivot site uniformly at random.
 - Randomly choose a lattice symmetry g (rotation or reflection).
 - Apply this symmetry to one of the two sub-walks created by splitting the walk at the pivot site.
 - If walk is self-avoiding: accept the pivot and update the configuration.
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Example pivot move



• Pivots are rarely successful, $\Pr = O(N^{-p})$, $p \approx 0.11$ for \mathbb{Z}^3 .

- Every time a pivot attempt *is* successful there is a large change in global observables.
- Only need O(1) successful pivots before we have an *essentially new* configuration with respect to observables measuring size.

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An efficient data structure for SAW

• Represent SAW as a binary tree.

- Enables global moves like pivots to be performed in CPU time $T(N) = O(\log N)$.
- c.f. O(N^{1-p}) for hash table implementation².
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PERM

Pivot (SAW-tree)

Observable

Conclusion



SAW-tree representation of a walk.



How to calculate c_N ?

- Would like to apply pivot algorithm in canonical ensemble.
- Approach: measure probability that object from larger set is a SAW, $|S| = P(x \in S | x \in T) |T|$, with |T| known.
- Obvious choice: concatenating pairs of SAWs. Every M + N-step walk can be split into M and N step subwalks $\Rightarrow c_{M+N} \leq c_M c_N$ for all M, N.
- S_N set of walks of length N.
- $|S_{M+N}| = P(\omega_1 \circ \omega_2 \in S_{M+N} | (\omega_1, \omega_2) \in S_M \times S_N) | S_M | | S_N$
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A long N step walk can be successively subdivided into smaller pieces.





















$$\langle B_{36,36} \rangle = \frac{c_{72}}{c_{36}c_{36}}$$

Iterate to obtain estimates for c_N for longer walks.

$$c_{N} = \frac{c_{N}}{c_{N/2}^{2}} \cdot \frac{c_{N/2}^{2}}{c_{N/4}^{4}} \cdots \frac{c_{2k}^{N/2k}}{c_{k}^{N/k}} c_{k}^{N/k}$$
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where c_k is known.

- Telescoping, with length doubling at each iteration.
- C.f. sequential growth, N steps, product of N factors





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• Could choose m, n = 36 (longest known for \mathbb{Z}^3):

Pivot

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$$\log \mu_N = \frac{1}{k} \log c_k + \frac{1}{2k} \log \langle B_{k,k} \rangle + \frac{1}{4k} \log \langle B_{2k,2k} \rangle + \cdots$$
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Counting SAW 24 / 30

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• Need to calculate $\langle B_{k,k} \rangle$, $\langle B_{2k,2k} \rangle$, \cdots

• Use pivot algorithm / SAW-tree.

- How many pivots must be completed before two walks are "essentially new" configurations with respect to observable *B*?
- Shape of walks close to the joint clearly important.
- Uniform pivot sites: $\tilde{\tau}_{int} = \Omega(N)$.
- Choose distance from joint uniformly from all distance scales,
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- Counting SAW 26 / 30



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- Partition CPU time amongst different terms to minimize overall statistical error (short test run).
 - $\sigma^2 = \sum \frac{a_i^2}{t_i}$ Total time $t = \sum t_i$ $\Rightarrow t_i = \frac{a_i}{\sum a_i}t,$ $\sigma = \frac{\sum a_i}{\sqrt{t}}$
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- Dominated by low k contribution, appropriate partitioning of effort reduced error by $O(\sqrt{\log N})$. Relative error in c_N proportional to 1/k. Counting SAW 26 / 30





• Can make unbiased estimates of c_N , for N up to 10^9 or so.

- Can push calculation to sufficiently large N s.t. asymptotic corrections for μ completely eliminated.
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Results

- Calculated log c_N with relative error of approximately 4×10^{-9} up to N = 38797311 (about 60000 CPU hours).
- Concentrated on \mathbb{Z}^3 because asymptotic behaviour for \mathbb{Z}^2 well understood from series.
- $c_{9471} = 1.43323(8) \times 10^{6352}$
- $c_{38797311} = 7 \times 10^{26018276}$. Confidence interval of mantissa is (6.6, 8.2).
- For comparison, see³. Relative error from PERM and related algorithms of the order of 10⁻³ for short walks of 100 steps. Not a fair comparison:

Not much CPU time used, i.e. not serious computers

■ Estimates would degrade for large N. Best cases error increasing as O(√N).

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- Different ingredients fit together to produce extremely accurate estimates.
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- Efficient data structures help.
- Can you do better than incremental growth? (fusing objects and doubling size, or splitting in two)
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