



Licentiate Thesis

Examining Hausdorff dimension and Scaling behaviour in Worm algorithm

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Stockholm, Sweden 2018

Typeset in L^AT_EX

Akademisk avhandling för avläggande av teknologie licentiatexamen (TeknL) inom ämnesområdet teoretisk fysik.

Scientific thesis for the degree of Licentiate of Engineering (Lic Eng) in the subject area of Theoretical physics.

TRITA-FYS-2010:05

ISSN

ISRN KTH/FYS/--10:05--SE

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Printed in Sweden by Universitetsservice US AB, Stockholm May 2018

Abstract

2.3 The Metropolis Monte Carlo method The principle of importance sampling in statistical physics as suggested by Metropolis et al. [83] is as follows. In Fig. 2.1 we generated numbers x and from a uniform distribution. In the evaluation of something like (2.1), we could instead imagine generating the configurations $\mu \equiv \{\theta_i\}_{i=1}^N$ not uniformly, but in such a way that the number of hits and misses are roughly equal. The estimated expectation value after M generated configurations μ_n can be written

$$\langle O \rangle = \frac{\sum_{n=1}^M O_{\mu_n} e^{\beta H \mu_n}}{\sum_{n=1}^M e^{\beta H \mu_n}} \quad (1)$$

for states μ_n generated with uniform probability. If we instead were to draw configurations with probabilities according to their Boltzmann weights, the estimator is simply an unweighted arithmetic average

$$\langle O \rangle \approx \sum_{n=1}^M O_{\mu_n} \quad (2)$$

for states μ_n generated with probability $e^{-\beta \mu_n}$

The remaining question is then how one generates configurations with probability according to their Boltzmann weights. In practice, the importance sampling discussed above is obtained with a Markov chain through the Metropolis method [83]. Metropolis Monte Carlo works by proposing a new configuration j from an old configuration i with a proposal distribution $g(i \rightarrow j)$, which is accepted with a specified probability $A(i \rightarrow j) = \min(1, e^{-\beta \Delta E})$ where $\Delta E = E_j - E_i$. Consider the time evolution of the probability of being in state i ,

$$\frac{dp_i}{dt} = \sum_j [p_j P(j \rightarrow i) - p_i P(i \rightarrow j)] \quad (3)$$

where $P(i \rightarrow j)$ is the probability of transitioning from i to j . The first and second terms on the right-hand side correspond to the rate of transitioning into and out of the state i respectively. In equilibrium the time derivative is zero and (2.5) is fulfilled if (but not only if)

$$p_j P(j \rightarrow i) = p_i P(i \rightarrow j) \quad (4)$$

The condition of (2.6) is called detailed balance and is a sufficient but not necessary condition for Metropolis Monte Carlo given that states are generated ergodically, see [84, 88]. Ergodicity means that each state is reachable from every other state in a finite time, however a non-ergodic calculation may still yield useful information within an ergodic class [84]. In a simulation, the probability of transitioning from i to j is the product of the proposal and acceptance probabilities, that

is $P(i \rightarrow j) = g(i \rightarrow j)A(i \rightarrow j)$ which after insertion into the balance equation (2.6) gives

$$\frac{p_i}{p_j} \frac{g(j \rightarrow i) \min(1, e^{-\beta(E_i - E_j)})}{g(i \rightarrow j) \min(1, e^{-\beta(E_j - E_i)})} = \frac{e^{-\beta E_i}}{e^{-\beta E_j}} \quad (5)$$

assuming that the proposal distribution is symmetric. The Metropolis Monte Carlo method thus generates a chain of configurations where each configuration appears with a probability proportional to its Boltzmann weight. The estimation of thermal averages can thus be done via arithmetic averaging of the form (2.4) by starting with some initial configuration μ_0 and generating a chain μ_1, \dots, μ_M via the Metropolis scheme.

2.5 Phase transitions: definition and classification

The thermodynamical bulk free-energy density $f = \beta^{-1} \ln(Z)/V$ depends on the coupling parameters of the Hamiltonian, and thermodynamic functions are calculated via differentiation of the free energy. Such thermo-dynamic functions may exhibit discontinuities at certain sets of coupling parameters, for which the free energy is not analytic. Such sets can be taken as definitions of phase boundaries which separate phases (i.e. regions of analyticity). The crossing of a phase boundary is then understood as a phase transition. It should be noted that the definition in terms of analytic free-energy is not completely general, considering the BKT-transition. For further details of mathematical aspects of phase transitions, we refer to [94] and for a more physical discussion to [95].

By the Ehrenfest classification [96] phase transitions are classified by the order of the derivative of the free energy which has a discontinuity at the critical point. If a first order derivative (e.g. entropy, internal energy) is discontinuous, then the phase transition is of first order. If a second order derivative (e.g. heat capacity) is discontinuous, the phase transition is of second order, and so on.

The Ehrenfest classification is however discouraged [95, 98], for example there may be divergences rather than discontinuities in thermodynamic functions (as for the 2D Ising model without external field, where the heat capacity diverges logarithmically at T_c). The modern classification of phase transitions is binary where a phase transition is either ‘first-order’ (there is a latent heat for a temperature-driven transition) or ‘continuous’ (no latent heat) [98, 95].

5. Partition function A problem of statistical mechanics is to calculate the partition function, that is calculating the trace of $e^{-\beta H}$, a task most straightforwardly written as

$$Z = \sum \langle \alpha_n | e^{-\beta H} | \alpha_n \rangle \quad (6)$$

$$= \sum e^{-\beta E_i} \quad (7)$$

where the $|\alpha_n\rangle$ is a complete orthonormal set and E_i the energy eigenvalues of H . In this form, it is necessary to determine the eigenvalues E_i of the Hamiltonian operator, and then carry out the sum over all corresponding weights $e^{-\beta E_i}$, as

Feynman put it [132] a ‘hopelessly difficult’ calculation. Instead we reformulate the problem in terms of imaginary time path integrals [132, 69, 133].

Imaginary-time path integral reformulation of the partition function First, we outline the reformulation of (5.5) in terms of path integrals, with the Bose-Hubbard model (5.2) in mind. Following the derivation in [69], first denote the statistical operator It then follows that

$$\rho(\beta) = e^{-\beta H} \quad (8)$$

$$\partial \rho = -H \rho \quad (9)$$

0.1 Derivation for Ising 2D Worm

The Ising model energy

$$E = -J \sum_{\langle ij \rangle} S_i S_j \quad (10)$$

Let $K = \beta J$ where $\beta = 1/k_B T$.

$$\beta E = -K \sum_{\langle ij \rangle} S_i S_j \quad (11)$$

The partition function Z .

$$Z = \sum_{\text{all states}} e^{-\beta E} = \sum_{\text{all states}} e^{K \sum_{\langle ij \rangle} S_i S_j} = \sum_{\text{all states}} \prod_{\langle ij \rangle} e^{K S_i S_j} \quad (12)$$

Since $S_i S_j = \pm 1$ in the Ising model, Euler identities can be used to expand the exponential in (12).

$$\begin{aligned} e^{K S_i S_j} &= \frac{e^K + e^{-K}}{2} + S_i S_j \frac{e^K - e^{-K}}{2} \\ &= \cosh(K) + S_i S_j \sinh(K) \\ &= \{T = \tanh(K)\} \\ &= (1 + T S_i S_j) \cosh(K) \end{aligned}$$

For N spins there are $2N$ bonds, therefore the partition function is

$$\begin{aligned}
Z &= \sum_{\text{all states}} \Pi_{\langle ij \rangle} (1 + T S_i S_j) \cosh(K) \\
&= \cosh^{2N}(K) \cdot 2^N \left(2^{-N} \sum_{\text{all states}} \Pi_{\langle ij \rangle} (1 + T S_i S_j) \right) \\
&= \cosh^{2N}(K) \cdot 2^N Z'
\end{aligned}$$

And

$$\begin{aligned}
Z' &= 2^{-N} \sum_{\text{all states}} \Pi_{\langle ij \rangle} (1 + T S_i S_j) \\
&= 2^{-N} \sum_{S_1=\pm 1} \sum_{S_2=\pm 1} \dots \sum_{S_N=\pm 1} \left(1 + T \sum_{l=1} S_l S_{l+1} + T^2 \sum_{l=2} (S_l S_l)(S_{l-1} S_{l+1}) + \dots \right)
\end{aligned}$$

Where the sums $\sum_{l=L}$ should be interpreted as the sum over all sets where the link length is L . Link length is the coupling between S -terms as

$$\begin{aligned}
(S_1 S_2) &- \text{Open with length 1} \\
(S_1 S_2)(S_2 S_3) &- \text{Open with length 2} \\
(S_1 S_2)(S_2 S_3)(S_3 S_4)(S_4 S_1) &- \text{Closed with length 4}
\end{aligned}$$

Since $\sum_{S_i=\pm 1} S_i = 0$, only terms with an even number of S_i are contributing to Z' . We will call these terms closed, indicating that they represent a closed loop if we were to draw the link lengths between the sites. The sum over all contributing terms gives a factor of 2^N , canceling the 2^{-N} .

We can now rewrite Z' in terms of loop lengths.

$$Z' = \sum_L g(L) T^L \quad (13)$$

Where $g(L)$ is the number of loops with length L . Finally we can write the expression for the partition function.

$$Z = 2^N \cosh^{2N}(K) \sum_L g(L) T^L \quad (14)$$

0.1.1 Energy calculation

$$E = -\frac{1}{Z} \frac{\partial Z}{\partial \beta} = -\frac{J}{Z} \frac{\partial Z}{\partial K} \quad (15)$$

Therefore

$$\begin{aligned}
\frac{\partial Z}{\partial K} &= 2^N 2N \cosh^{2N-1}(K) \cdot \frac{\partial \cosh(K)}{\partial K} Z' + 2^N \cosh^{2N}(K) \cdot \frac{\partial Z'}{\partial K} \\
&= 2^N \cosh^{2N}(K) \left(2N \tanh(K) Z' + \frac{\partial Z'}{\partial K} \right) \\
&= 2^N \cosh^{2N}(K) \tanh(K) \left(2N Z' + \frac{1}{\tanh(K)} \frac{\partial Z'}{\partial K} \right)
\end{aligned}$$

Examine $\tanh^{-1}(K) \frac{\partial Z'}{\partial K}$.

$$\begin{aligned}
\tanh^{-1}(K) \frac{\partial Z'}{\partial K} &= \tanh^{-1}(K) \frac{\tanh(K)}{\partial K} \sum_L g(L) L \tanh^{L-1}(K) \\
&= \frac{1}{\tanh^2(K) \cosh^2(K)} \sum_L g(L) L \tanh^L(K) \\
&= \frac{Z'}{\sinh^2(K)} \frac{\sum_L g(L) L \tanh^L(K)}{\sum_L g(L) \tanh^L(K)} \\
&= \frac{Z'}{\sinh^2(K)} \langle L \rangle
\end{aligned}$$

And finally

$$E = -J \tanh(K) \left(2N + \frac{\langle L \rangle}{\sinh^2(K)} \right) \quad (16)$$

where

$$\langle L \rangle = \frac{\sum_L g(L) L \tanh^L(K)}{\sum_L g(L) \tanh^L(K)} \quad (17)$$

0.1.2 Heat capacity

$$C = \frac{\partial E}{\partial T} = -\beta^2 \frac{\partial E}{\partial \beta} = -K \beta \frac{\partial E}{\partial K} \quad (18)$$

Let $A = 2N + \frac{1}{\sinh^2(K)} \langle L \rangle$. Then

$$\frac{E}{J} = -\tanh(K) A \quad (19)$$

and

$$\frac{1}{J} \frac{\partial E}{\partial K} = -\frac{\partial \tanh(K)}{\partial K} A - \tanh(K) \frac{\partial A}{\partial K} \quad (20)$$

where

$$\begin{aligned}
\frac{\partial A}{\partial K} &= \langle L \rangle \frac{\partial \sinh^{-2}}{\partial K} + \tanh^{-1}(K) \sinh^{-2}(K) (\langle L^2 \rangle - \langle L \rangle^2) \frac{\partial \tanh(K)}{\partial K} \\
&= \frac{1}{\sinh^2(K) \tanh(K)} \left(-2\langle L \rangle + \frac{\langle L^2 \rangle - \langle L \rangle^2}{\cosh^2(K)} \right)
\end{aligned}$$

and finally

$$C = \frac{K^2}{\sinh^2(K)} \left(\frac{\langle L^2 \rangle - \langle L \rangle^2}{\cosh^2(K)} - E \tanh(K) - 2\langle L \rangle \right) \quad (21)$$

Preface

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Part I

Introduction and background material

Chapter 1

Introduction

Chapter 2

Worm Algorithm

Chapter 3

Graph Indexing

Chapter 4

Graph Division and Hausdorff Dimension

4.1 Fractals

Everyone agrees that the dimension of a point is zero, and that of a smooth line is one, but what about a set of points? A definition could be to say that the dimension is the minimum number of coordinates needed to describe every point in the set. Effectively, a point would describe itself, and a curve could be parametrized to the distance of some point on the same curve.

The situation is more complex when examining fractals. Take for example the Koch curve, it starts out as a line segment of length L_0 , and successively adds a ‘bump’, making the total length $L_1 = 4/3 \cdot L_0$. Continuing this trend gives for the n th iteration a line length of $L_n = (4/3)^n \cdot L_0$, and so for the final fractal the length is infinite. Now any two point on the curve has a distance of infinity between them, so parametrization is impossible, but the area is still finite, so the dimension should be somewhere between one and two.

A useful concept here is the similarity dimension, defined by the scaling of each iteration. If m is the number of similar elements after an iteration and r is the scaling factor, the dimension is defined as $m = r^d$, or equivalently

$$d = \frac{\ln m}{\ln r} \tag{4.1}$$

So for the Koch curve, each segment is divided into fourths with each having one third the length from the previous iteration, giving it a dimension of $\ln 4 / \ln 3 \approx 1.26$.

4.2 Box dimension

4.3 Graph Dividing Algorithm

In order to calculate the box dimension the lattice needs to be divided into boxes of decreasing size. A step by step instruction of a graph dividing algorithm is provided below, and an implementation in pseudocode is available in the Appendix at Section 7.1.

For brevity some abbreviations are introduced.

d = dimension	l_i = side length of the current box
l_0 = side length of the smallest box allowed	e_j^i = vector of length $l_i/2$ in the j 'th direction
$\text{perm}(v)$ = All permutations of v	s_i = starting site of the current box

1. If $l_i \geq l_0$, go to 2, else stop.
2. Save all sites in the current box, starting for s_i going l_i in d directions.
3. Find all starting points for new boxes.

- (i) Form the matrix $E^i = (e_0^i, e_1^i, \dots, e_d^i)^T$
- (ii) For all vectors v_k in $\text{perm}(0, 0, \dots, 0)$, $\text{perm}(1, 0, \dots, 0)$,
 \dots , $\text{perm}(1, 1, \dots, 1)$, create the new start s_k^i as

$$s_k^i = v_k E^i$$

4. For each start s_k^i :

- (i) $s_i = s_k^i$, $l_i = l_i/2$
- (ii) Go to 1.

Chapter 5

Connection between Hausdorff Dimension and Scaling Behaviour

Chapter 6

Summary and conclusions

Chapter 7

Appendix

7.1 Pseudo Code for Box Division Algorithm

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