

PY410 / 505  
Computational Physics 1

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# Probabilistic methods

Casino Monte Carlo



- A large number of processes in nature are random
- To some extent, we're all basically familiar with it
  - Radioactive decay
  - Flipping a coin
- In fact, quantum mechanics inherently is random

# Probabilistic methods

- What computational issues can come up with random numbers?
  - Computers are, by very nature, NOT random
  - So we need to make them LOOK random
  - The question is, how random is “random enough”?
- If you want something truly random, you’ll need to hook your computer up to a Geiger counter or something, and count decays (say, from atmospheric muons)

- Sounds silly, but it isn’t



# Probabilistic methods

- Chapter 7 in Numerical Recipes deals with generating random numbers
  - “Deviates”
- How to formalize random number generation (RNG)?
  - Given a set  $S$  of  $N$  uniformly distributed random numbers  $x_1 \dots x_N$ , then they must satisfy :
    - Given  $n$  generated numbers, the next number  $x_{n+1}$  must be independent and uncorrelated
    - $x_{n+1}$  should be equally likely to be a member of  $S$
  - For the example of an unbiased coin toss,  $S$  has 1 or 0 (heads or tails)
  - Each toss is independent of the previous and so a priori equally likely to be 1 or 0

# Probabilistic methods

- Why do computers have trouble here?
- They are inherently deterministic at the present time (that's why they're so great to use!)
- We generate “pseudo” random numbers
  - The statistical properties (i.e. equally likely in some region of interest) is “good enough”, but “good enough” depends on the situation
- For instance, if we use linear congruential algorithms of the type  $x_{n+1} = (ax_n + c) \bmod m$ ,
  - This generates a sequence of random integers in the set  $\{0, 1, \dots, m-1\}$

“multiplier”

“increment”

“modulus”

# Probabilistic methods

- Why does this help?
  - We typically choose  $a$ ,  $c$ , and  $m$  to be large(ish) relatively prime numbers
  - If we inappropriately choose  $a, c$ , or  $m$ , we can have repetition
  - If any  $x_n = x_0$ , then the process will repeat
  - We initialize this with a “seed”  $x_0$ , and away goes the sequence

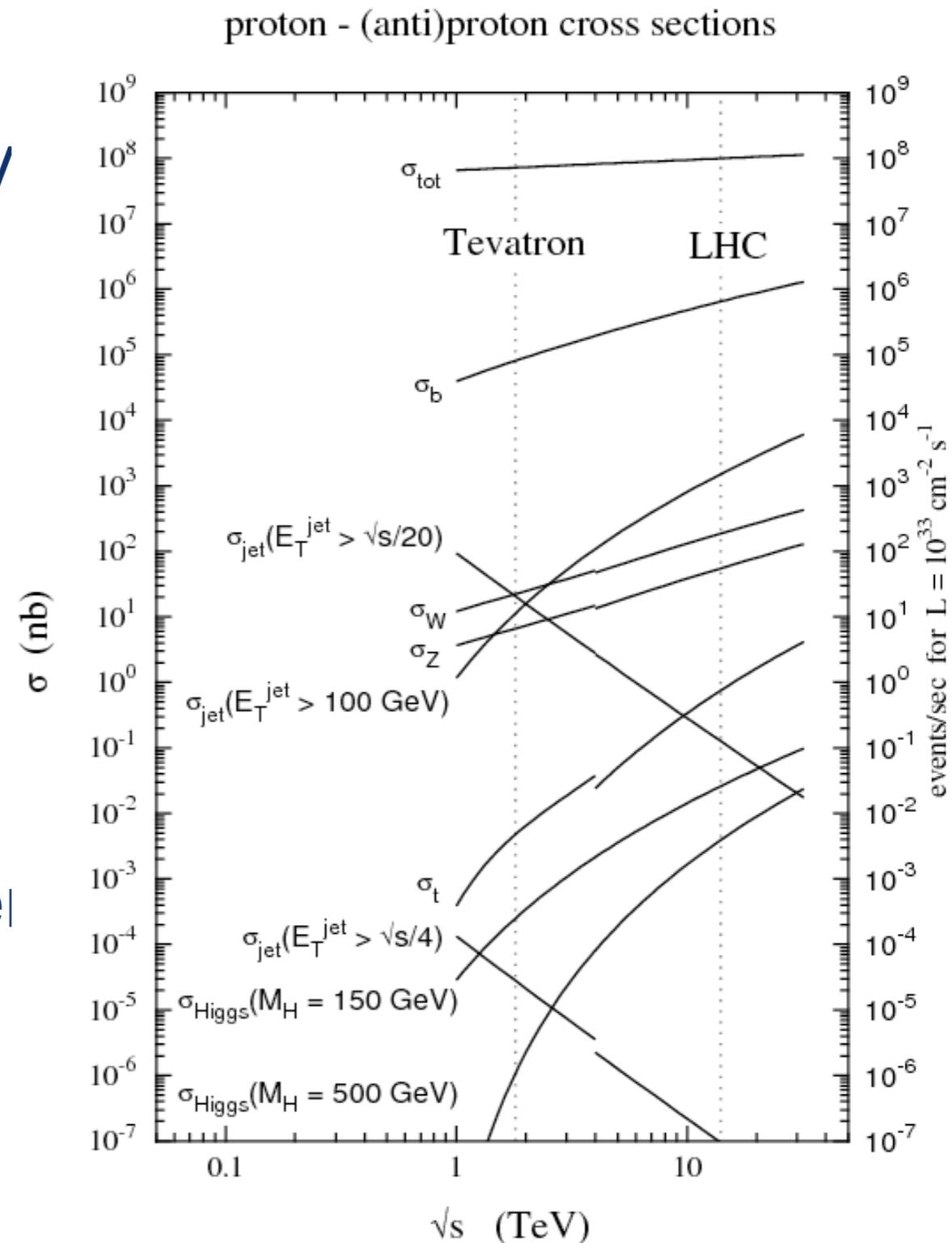
Source	$m$	(multiplier) $a$	(increment) $c$	output bits of seed in $rand() / Random(L)$
<i>Numerical Recipes</i>	$2^{32}$	1664525	1013904223	
Borland C/C++	$2^{32}$	22695477	1	bits 30..16 in $rand()$ , 30..0 in $lrand()$
glibc (used by GCC) <sup>[5]</sup>	$2^{31}$	1103515245	12345	bits 30..0
ANSI C: Watcom, Digital Mars, CodeWarrior, IBM VisualAge C/C++ <sup>[6]</sup>	$2^{31}$	1103515245	12345	bits 30..16
Borland Delphi, Virtual Pascal	$2^{32}$	134775813	1	bits 63..32 of $(seed * L)$
Microsoft Visual/Quick C/C++	$2^{32}$	214013 (343FD <sub>16</sub> )	2531011 (269EC3 <sub>16</sub> )	bits 30..16
Microsoft Visual Basic (6 and earlier) <sup>[7]</sup>	$2^{24}$	1140671485 (43FD43FD <sub>16</sub> )	12820163 (C39EC3 <sub>16</sub> )	

# Probabilistic methods

- There are several industrial-strength generators on the market
- But, if you need “really really” random numbers, use with extreme care
  - C++11, R, Python, Ruby, IDL, Maple, Matlab, GNU MPAL, BOOST, Glib, and NAG :
    - [http://en.wikipedia.org/wiki/Mersenne\\_twister](http://en.wikipedia.org/wiki/Mersenne_twister)
    - Long period of  $2^{19937} - 1$
    - Passes lots of randomness tests
    - NOT suitable for cryptography : observing a certain number of iterations will allow you to predict the rest of the sequence
  - Numerical recipes recommends
    - <http://en.wikipedia.org/wiki/Xorshift>
    - Period of  $2^{128} - 1$

# Probabilistic methods

- Example: the “sparky” trigger
- In particle physics, we throw away almost all of our data
- Only 1 in  $10^5$  is even “remotely interesting
- The really interesting stuff is only 1 in  $\sim 10^{16}$ !
- Given this, we have to often have “REALLY” random numbers to test
- So, we set up a spark chamber to actually generate random numbers so our triggers could perform adequate tests



# Probabilistic methods

- A few tests will be performed here :
  - The period of the generator should be much larger than the length of the generated sequence
  - A simple “eyeball test” (plotting  $(x_n, x_{n+1})$  as  $(x, y)$  pairs) should reveal no structure
  - The chi2 statistic should satisfy  $\frac{\chi^2}{\text{d.o.f.}} \simeq 1$  .

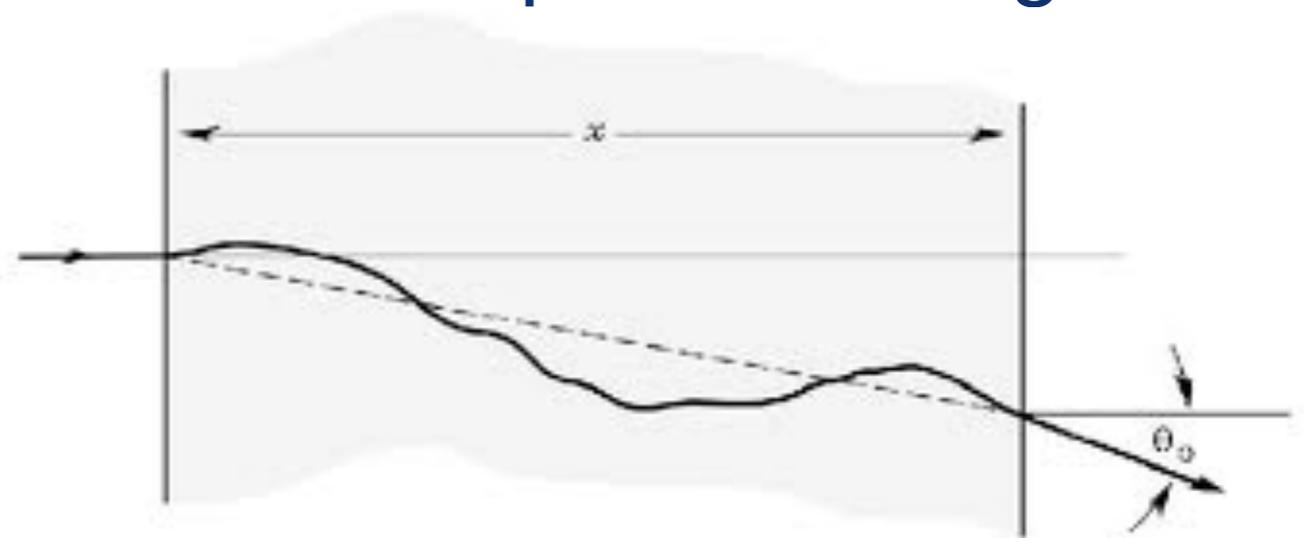
# Probabilistic methods

- Randomness is often used in simulations
- Lots of things are random in nature
- We often know their distribution, but cannot predict individual events :

– Standing a pencil on its side, what is the angle (wrt the desk) when it falls?

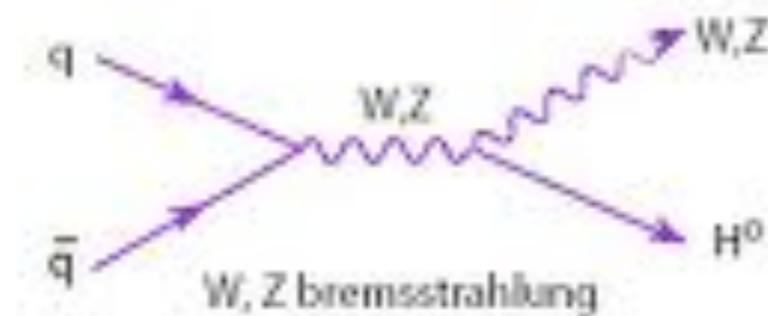
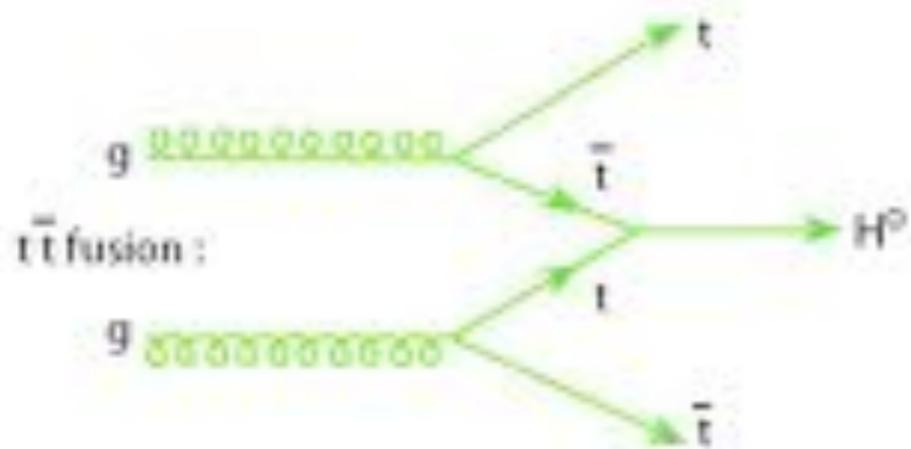
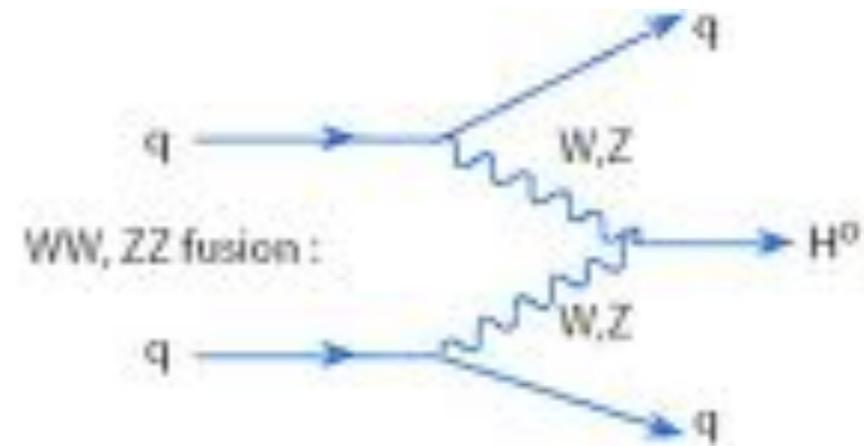
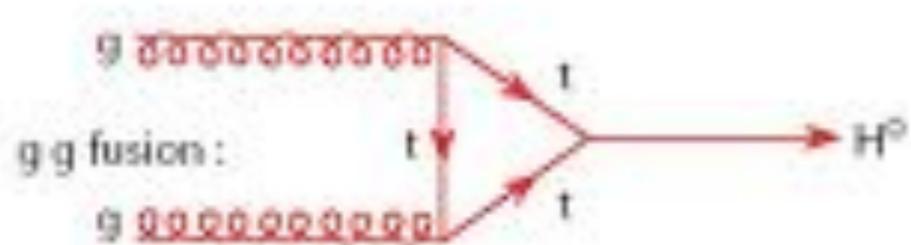


– What is the final angle between multiple scatterings of particles off of a lattice?



# Probabilistic methods

- So, to simulate these events, we need random (or pseudorandom) generators
- Example : Quantum mechanically, can produce Higgs bosons in several ways :



- Can't even ask (quantum mechanically) which will occur before it happens
- They all have a likelihood!
- So, to generate them, you have to randomly sample between these four (with appropriate weights)

# Probabilistic methods

- These methods are often called “Monte Carlo” after the Casino Monte Carlo in Monaco
- Much better name than “pseudo-random-number probabilistic event simulator”



# Probabilistic methods

- Why does this help?
- Suppose you take a system of 100 electrons in either “spin up” or “spin down” configurations
- Total number of states is  $= 2^{100} = 1.27 \times 10^{30}$
- Already intractable to list them (at one billion / second, would take  $4.02 \times 10^{13}$  years!)
- If you select one at random for any purpose, need to simply find a sufficiently good randomization algorithm!

# Probabilistic methods

- Markov-Chain MC (MCMC) :
  - Sequence of elements chosen from a fixed set using a probabilistic rule
  - Chain is constructed by adding the elements sequentially
  - Given the most recently added element, next element only depends on most recent addition
  - Formally, suppose  $x$  and  $y$  are members of a set  $S$
  - The transition probability function is

$$T(x \rightarrow y) , \quad \sum_y T(x \rightarrow y) = 1 .$$

- Example : Random walks

# Probabilistic methods

- Random walks are simple examples of MCMC's
- Suppose a walker can occupy any site on an infinitely-long 1-d lattice
- The walker tosses a coin and decides to go left or right (1 or 0)
- The transition probability is therefore

$$T(x \rightarrow y) = \begin{cases} \frac{1}{2} & \text{if } y = x - 1, \\ \frac{1}{2} & \text{if } y = x + 1, \\ 0 & \text{otherwise.} \end{cases}$$

- In equilibrium, suppose the lattice is periodic now, with  $L$  lattice points on a circle
  - Then we can take  $L \rightarrow \infty$  to examine the properties
- Probability is  $P(x) = \frac{1}{L}$  :  
(walker visits each site the same number of times)

# Probabilistic methods

- The walker's position after  $n$  steps depends on the sequence of tosses in the past, and cannot be predicted

$$x_n = \sum_{i=1}^n s_i, \quad s_i = \pm 1$$

- Over a large number of  $n$ -step walks the average is zero  
–(From symmetry : left and right are equally likely!)
- However, each  $x_n$  is not zero, and DOES increase:

$$\begin{aligned} \langle x_n^2 \rangle &= \left\langle \sum_{i=1}^n \sum_{j=1}^n s_i s_j \right\rangle \\ &= \left\langle \sum_{i=1}^n s_i^2 \right\rangle + \left\langle \sum_{i=1}^n \sum_{\substack{j=1 \\ j \neq i}}^n s_i s_j \right\rangle \\ &= n, \end{aligned}$$

Diffusion equation!

# Probabilistic methods

- RMS displacement is

$$\sqrt{\langle x_n^2 \rangle} = \sqrt{n} .$$

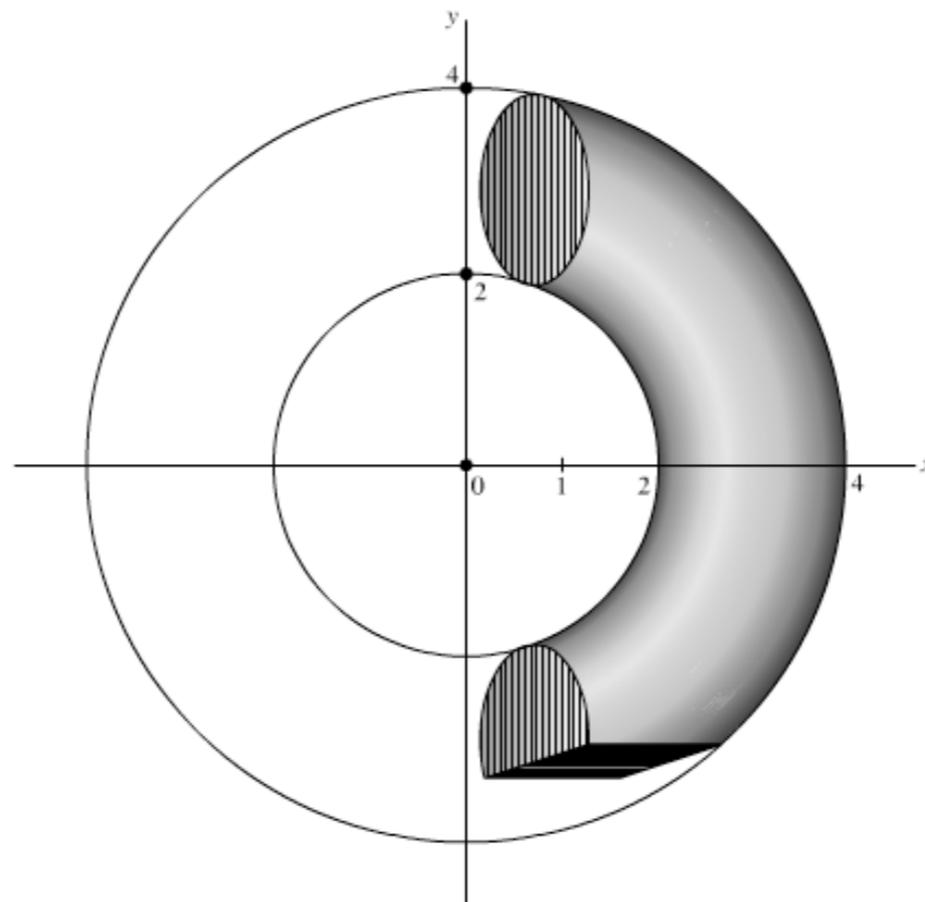
- Diffusion constant is

$$\langle x_n^2 \rangle = 2Dn .$$

- So, for a 1-d walk,  $D = 1/2$

# Probabilistic methods

- Another example of MC methods : integration of functions
- Sprinkle the area you're interested with a "dust" of points uniformly distributed
- The fraction "below the curve" (or in N-dim, within the surface) is the integral!



# Probabilistic methods

- Formally, the integral is :

$$I = \int_V d^d \mathbf{x} f(\mathbf{x})$$

- Now we choose  $N$  uniformly distributed points in  $V$ , and estimate

$$I \simeq \frac{V}{N} \sum_{n=0}^{N-1} f(\mathbf{x}_i)$$

- The error of repeating this  $M$  times is :

$$I_m = \frac{b-a}{N} \sum_{n=0}^{N-1} f(\mathbf{x}_{m,n}), \quad m = 0 \dots M-1$$

- Can just compute the mean and standard deviation:

$$\bar{f} = \frac{1}{MN} \sum_{m=0}^{M-1} \sum_{n=1}^N f(\mathbf{x}_{m,n})$$

$$\begin{aligned} \sigma_M &= \sqrt{\frac{1}{M} \sum_{m=0}^{M-1} I_m^2 - \left( \frac{1}{M} \sum_{m=0}^{M-1} I_m \right)^2} \\ &= \sqrt{\frac{1}{M} \sum_{m=0}^{M-1} \left( I_m - \frac{1}{M} \sum_{m'=0}^{M-1} I_{m'} \right)^2} \end{aligned}$$

# Probabilistic methods

- If the measurements are independent and randomly distributed about the mean then:

$$\sigma_M^2 = \frac{(b - a)^2}{N} \sigma_f^2$$

- where

$$\sigma_f^2 = \overline{f^2} - (\bar{f})^2 = \frac{1}{MN} \sum_{m=0}^{M-1} \sum_{n=1}^N f(\mathbf{x}_{m,n})^2 - \left( \frac{1}{MN} \sum_{m=0}^{M-1} \sum_{n=1}^N f(\mathbf{x}_{m,n}) \right)^2$$

- Thus for MC integration:

$$I = \int_V d^d \mathbf{x} f(\mathbf{x}) \simeq V \left[ \frac{1}{N} \sum_{n=1}^N f(\mathbf{x}_i) \pm \frac{\sigma_f}{\sqrt{N}} \right]$$

# Probabilistic methods

- Compare this with our previous integration methods
- For instance, Midpoint method

- We have 
$$\frac{\sigma_f}{\sqrt{N}} < \frac{\overline{f''}}{N^{2/d}}$$

- MC integration “wins” when  $d > 4$  (roughly)

# Probabilistic methods

- What if you want nonuniform samples?
  - Example : Gaussian distribution, exponential distribution, line segment
  - Still use the uniformly-distributed “deviates”
  - Then change variables ( $x \rightarrow y(x)$ ) to produce another distribution :

$$P(y) = P(x) \left| \frac{dx}{dy} \right| , \quad \int dy P(y) = \int dx P(x) = 1 .$$

# Probabilistic methods

- Uniform line segment :
  - Basically trivial, you just redefine  $(0,1)$  to  $(a,b)$  and you're done:

$$y(x) = a + (b - a)x, \quad \left| \frac{dx}{dy} \right| = \frac{1}{|b - a|}.$$

$$P(x) = \begin{cases} 1 & \text{for } 0 < x < 1 \\ 0 & \text{otherwise} \end{cases} \quad \Rightarrow \quad P(y) = \begin{cases} \frac{1}{|b-a|} & \text{for } a < y < b \\ 0 & \text{otherwise} \end{cases}.$$

# Probabilistic methods

- Exponential distribution :

$$P(y) = \frac{1}{\lambda} e^{-y/\lambda}, \quad 0 < y < \infty$$

- Then we have

$$y(x) = -\lambda \log(x), \quad \left| \frac{dx}{dy} \right| = \frac{x}{\lambda} = \frac{e^{-y/\lambda}}{\lambda} = P(y).$$

$$P(x) = \begin{cases} 1 & \text{for } 0 < x < 1 \\ 0 & \text{otherwise} \end{cases}$$

- so the distribution is

$$P(y) = \begin{cases} \frac{e^{-y/\lambda}}{\lambda} & \text{for } 0 < y < \infty \\ 0 & \text{otherwise} \end{cases}.$$

# Probabilistic methods

- Gaussian distribution (“normal” distribution)

$$P(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-(x-\mu)^2/(2\sigma^2)} ,$$

- Cannot convert with elementary functions
- However, consider the product of TWO Gaussian distributions in  $x$  and  $y$  :

$$\begin{aligned} \int dx \int dy P(x)P(y) &= \frac{1}{2\pi\sigma^2} \int dx \int dy e^{-[(x-\mu)^2 + (y-\mu)^2]/(2\sigma^2)} \\ &= \int_0^{2\pi} \frac{d\theta}{2\pi} \int_0^\infty \frac{dr^2}{2\sigma^2} e^{-r^2/(2\sigma^2)} , \end{aligned}$$

- We have switched to polar coordinates:

$$x - \mu = r \cos \theta , \quad y - \mu = r \sin \theta .$$

# Probabilistic methods

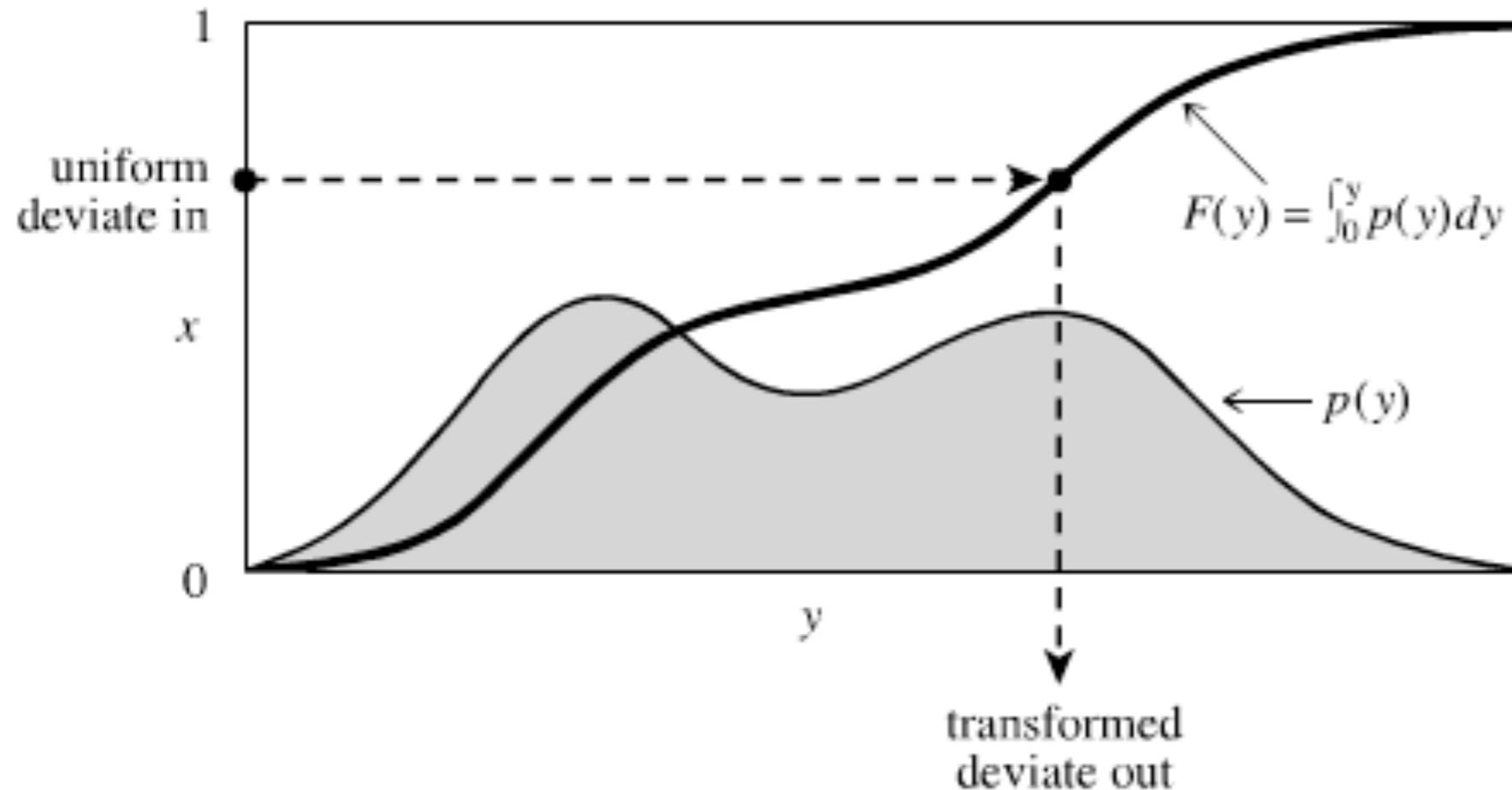
- This is the product of a uniform distribution on  $[0, 2\pi]$  with an exponential distribution in  $r^2$ !
- So we can generate theta and r, convert to x,y!

$$\theta = 2\pi r_1, \quad r = \sqrt{-2\sigma^2 \log r_2},$$

–“Box-Muller” algorithm

# Probabilistic methods

- General version of transformation method:



- Can be used even if you only know  $F(y)$  numerically

# Probabilistic methods

- To generate this kind of sequence, though, the Metropolis-Hastings algorithm works well
  - [http://en.wikipedia.org/wiki/Metropolis-Hastings\\_algorithm](http://en.wikipedia.org/wiki/Metropolis-Hastings_algorithm)
  - Type of Markov Chain Monte Carlo (MCMC)
  - One major advantage : it does not require the overall normalization of a distribution to be known to draw a random sequence from it!
    - Advantageous when using Bayesian statistics
- Original papers are from Metropolis et al and Hastings :
  - [J. Chem. Phys. 21, 1087 \(1953\)](#)
  - [Biometrika 57, 97 \(1970\)](#)

# Probabilistic methods

- This is a Markov Chain MC so the “next” step depends only on the “current” step
  - This is a type of random walk, similar to other MCMC methods
- Suppose we want to generate a sequence from a probability distribution  $P(\vec{x})$
- If we visualize this in d-dimensions, then  $P(x)$  is a mountainous terrain and the sequence is the steps of a hiker

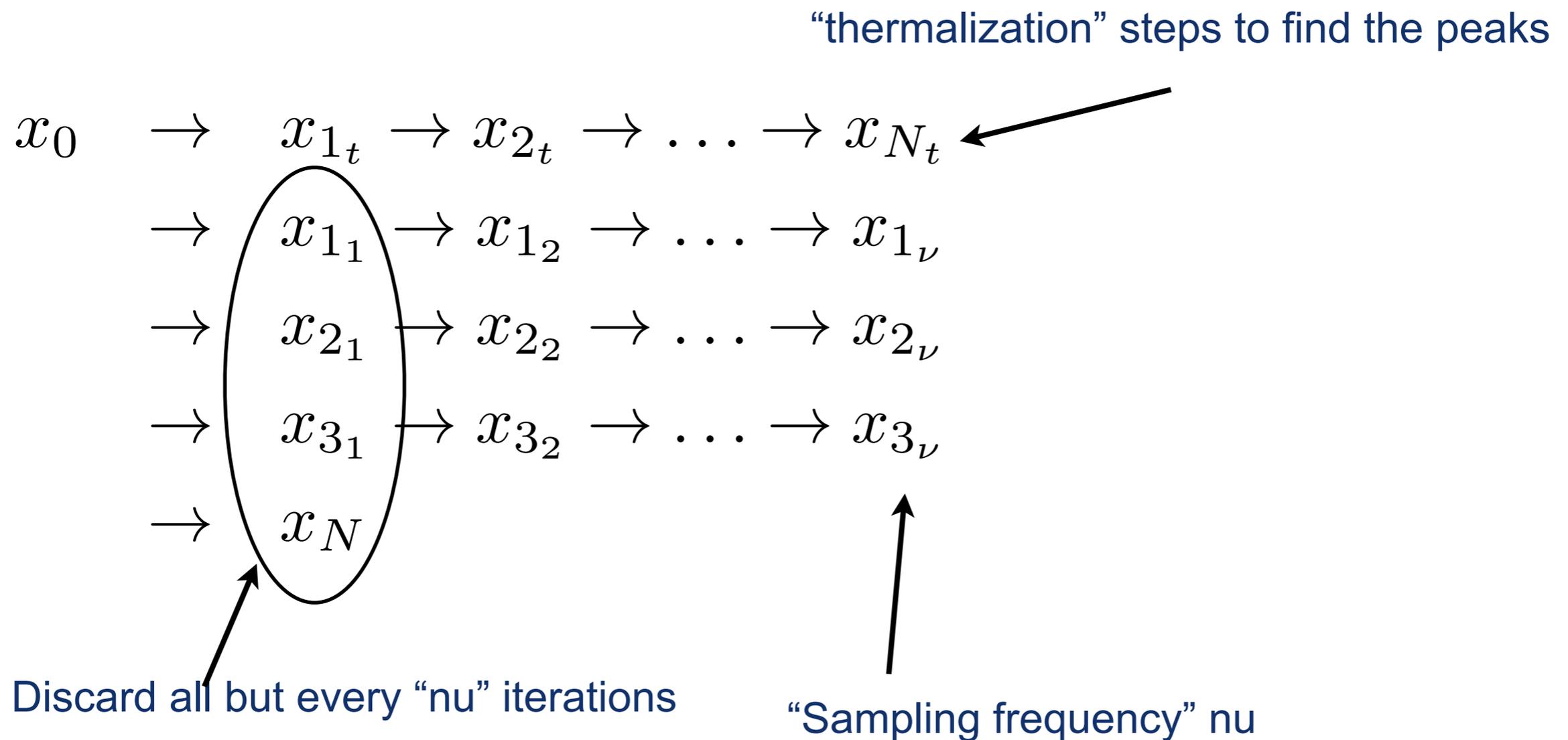


# Probabilistic methods

- Heuristically :
  - Explore the “terrain” a bit
    - Find peaks, valleys
  - Then generate the steps for the MC application
  - Record steps every so often
- As more and more sample values are produced, the distribution more closely approximates the desired distribution  $P(x)$

# Probabilistic methods

- More specifically, we have steps :



# Probabilistic methods

- Why sample?
  - Individual steps may be correlated
    - If walker's step size is much smaller than typical distances between "peaks" in the terrain
    - To emulate a "truly" random sequence, we don't want to get "stuck" between peaks

# Probabilistic methods

- Walking algorithm :
  - Suppose we have the walker at a point  $\vec{x}_{\text{current}}$
  - Then the next step  $\vec{x}_{\text{next}}$  is determined by
    - Choose a “trial” point randomly in the neighborhood of current  $x$  (  $\vec{x}_{\text{trial}}$  )
      - 1-d : maximum step size (delta) may be used, and choose trial  $x$  in a uniform interval  $[x_{\text{current}} - \delta, x_{\text{current}} + \delta]$
      - n-d : could try a fixed sphere of radius delta
    - Requirement is that the probability  $P_{\text{try}}(\vec{x}_{\text{current}} \rightarrow \vec{x}_{\text{trial}})$  should be **SYMMETRIC**:  $P_{\text{try}}(\vec{x} \rightarrow \vec{y}) = P_{\text{try}}(\vec{y} \rightarrow \vec{x})$ 
      - In other language, the conditional probabilities must be equal :
- $$P(x|y) = P(y|x)$$
- Calculate the next step by the ratio of probabilities (or “likelihoods”) :  $r = P(\vec{x}_{\text{trial}}) / P(\vec{x}_{\text{current}})$

# Probabilistic methods

- Given  $r = P(\vec{x}_{\text{trial}}) / P(\vec{x}_{\text{current}})$
- If  $r \geq 1$ , walker is moving “uphill” (toward higher probability)
  - Trial step is accepted, iterate
  - The “acceptance probability” is 1 by definition (it’s accepted,  $P_{\text{accept}}(\vec{x}_{\text{current}} \rightarrow \vec{x}_{\text{trial}}) = 1$  )
- If  $r < 1$ , walker is moving “downhill” (toward lower probability)
  - Accept only if displacement is not too large :
    - Generate uniform deviate  $n_{\text{random}}$
    - Accept if  $r > n_{\text{random}}$  , iterate
    - Reject otherwise, remain at current “step”
    - “Acceptance probability” in this case is  $r$

# Probabilistic methods

- What about those thermalization steps? Do they really thermalize?
- Imagine a large ensemble of walkers distributed with density  $\rho(\vec{x})$
- Want to show that the ensemble density becomes proportional to  $P(x)$  and then remain invariant
- According to the ergodic hypothesis, the average properties of an ensemble of systems should be the same as the average over time in a single system in thermal equilibrium  
– [http://en.wikipedia.org/wiki/Ergodic\\_hypothesis](http://en.wikipedia.org/wiki/Ergodic_hypothesis)
- So, we'll use the former to demonstrate the latter

# Probabilistic methods

- Recall the transition probability for a walker to move from  $x$  to  $y$  is determined by two consecutive and independent decisions:

$$T(\vec{x} \rightarrow \vec{y}) = P_{\text{try}}(\vec{x} \rightarrow \vec{y}) \times P_{\text{accept}}(\vec{x} \rightarrow \vec{y}) .$$

- If these  $T(x \rightarrow y)$  are symmetric, then:

$$\frac{T(\vec{x} \rightarrow \vec{y})}{T(\vec{y} \rightarrow \vec{x})} = \frac{P_{\text{accept}}(\vec{x} \rightarrow \vec{y})}{P_{\text{accept}}(\vec{y} \rightarrow \vec{x})} = \frac{P(\vec{y})}{P(\vec{x})} .$$

- Second equality can just be verified directly:

–if  $P(y) > P(x)$ ,  $P_{\text{accept}}(\vec{x} \rightarrow \vec{y}) = 1$  ,

$$P_{\text{accept}}(\vec{y} \rightarrow \vec{x}) = P(\vec{x})/P(\vec{y})$$

–if  $P(y) < P(x)$ ,  $P_{\text{accept}}(\vec{x} \rightarrow \vec{y}) = P(\vec{y})/P(\vec{x})$  ,

$$P_{\text{accept}}(\vec{y} \rightarrow \vec{x}) = 1_6$$

# Probabilistic methods

- The change in number density of walkers at  $\vec{x}$  when all the walkers in the ensemble take a step:

$$\begin{aligned}\Delta\rho(\vec{x}) &= \int d^d y [\rho(\vec{y})T(\vec{y} \rightarrow \vec{x}) - \rho(\vec{x})T(\vec{x} \rightarrow \vec{y})] \\ &= \rho(\vec{x}) \int d^d y T(\vec{y} \rightarrow \vec{x}) \left[ \frac{\rho(\vec{y})}{\rho(\vec{x})} - \frac{P(\vec{y})}{P(\vec{x})} \right] .\end{aligned}$$

- If  $\rho(\vec{y})/\rho(\vec{x})$  exceeds “equilibrium value”  $P(\vec{y})/P(\vec{x})$ :
  - will get a + correction
- If  $\rho(\vec{y})/\rho(\vec{x})$  is too small :
  - will get a - correction
- If  $\rho$  is proportional to  $P$ , we’ll have “detailed balance”:

$$\rho(\vec{y})T(\vec{y} \rightarrow \vec{x}) = \rho(\vec{x})T(\vec{x} \rightarrow \vec{y})$$

# Probabilistic methods

- Choices to make :
  - Step size
  - Thermalization steps
  - Sampling frequency
- All of these depend on the problem at hand
- You have to generally use trial and error to make sure there aren't large dependencies on the parameters

# Probabilistic methods

- An excellent example of using probabilistic methods is in statistical mechanics
- Makes sense : it's all about probability and statistics!
- A few definitions :
  - Microstates : configuration of a system
  - Probability of  $i$ th microstate is

$$p_i = \lim_{\mathcal{N} \rightarrow \infty} \frac{n_i}{\mathcal{N}},$$

Number of systems in the  $i$ th microstate



– Average value (e.g. for energy)  $\langle E \rangle = \sum_i p_i E_i$  .

– Variance is  $\langle (E - \langle E \rangle)^2 \rangle = \sum_i p_i E_i^2 - \left( \sum_i p_i E_i \right)^2$  .

# Probabilistic methods

- Examine the canonical and microcanonical ensembles:

## –Canonical :

- fixed number of constituents (N), fixed volume (V)
- exchange energy with thermal reservoir at temperature T
- Probability of a microstate with energy E is Boltzmann distribution:

$$p_i = \frac{e^{-E_i/(k_B T)}}{Z} ,$$

- Partition function is:

$$Z(N, V, T) = \sum_i e^{-E_i/(k_B T)} ,$$

## –Microcanonical :

- fixed number of constituents (N). fixed total energy (E). fixed volume (V)
- Probability of a microstate with energy E is :

$$p_i = \begin{cases} \frac{1}{\mathcal{N}} & \text{if } E = E_i \\ 0 & \text{otherwise} \end{cases} .$$

# Probabilistic methods

- Example of a system we can investigate is the hard-disk gas in two dimensions
  - Investigated by Metropolis et al with the MC methods above
  - J. Chem. Phys. 21, 1087 (1953)
  - “Equation of State Calculations by Fast Computing Machines”
    - “Fast” here meaning it could multiply two 40-bit ints in 1 millisecond (1 kHz)
    - For contrast, one core on your phone is 2.4 million times faster

# Probabilistic methods

- Look at the system as a Maxwell-Boltzmann gas at fixed volume and temperature

- Energy is 
$$E = \text{K.E.} + \text{P.E.} = \frac{m}{2} \sum_i \mathbf{v}_i^2 + \sum_{\text{pairs } ij} U(r_{ij}),$$

- Pairwise potential energy function :

$$U(r) = \begin{cases} 0 & \text{if } r > \sigma \\ \infty & \text{if } r \leq \sigma \end{cases},$$

- In an ensemble of systems, at temperature  $T$ , probability that the system has energy  $E$  is a Boltzmann distribution

$$\begin{array}{c} \nearrow \\ \text{Number of} \\ \text{microstates} \end{array} \sim W \exp \left[ -\frac{E}{k_B T} \right] = \exp \left[ -\frac{E - TS}{k_B T} \right] = \exp \left[ -\frac{F}{k_B T} \right], \begin{array}{c} \nwarrow \\ \text{Free} \\ \text{energy} \end{array}$$

Entropy

# Probabilistic methods

- Partition function is

$$Z = \sum_E W(E) \exp \left[ -\frac{E}{k_B T} \right] .$$

- The equation of state relates pressure  $p$ , volume  $V$  and temperature  $T$ :

$$pV = Nk_B T \left. \frac{\partial \log Z}{\partial \log V} \right|_{T,N} .$$

# Probabilistic methods

- MC simulation of hard disks from Metropolis et al:
  - Close-pack  $N=224$  disks of diameter  $d_0$  as follows

1090 METROPOLIS, ROSENBLUTH, ROSENBLUTH, TELLER, AND TELLER

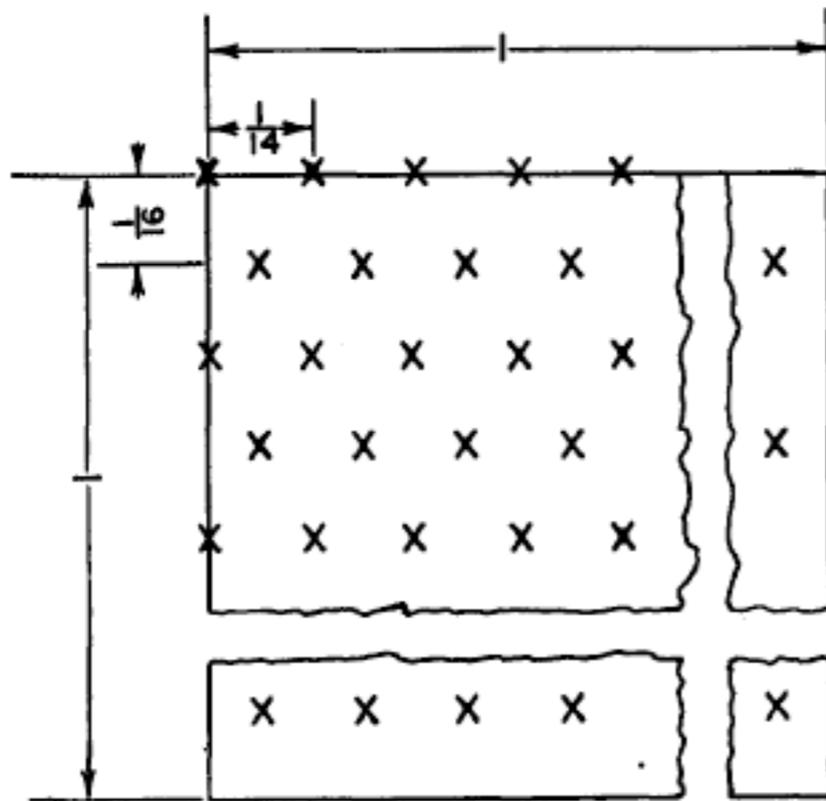


FIG. 2. Initial trigonal lattice.

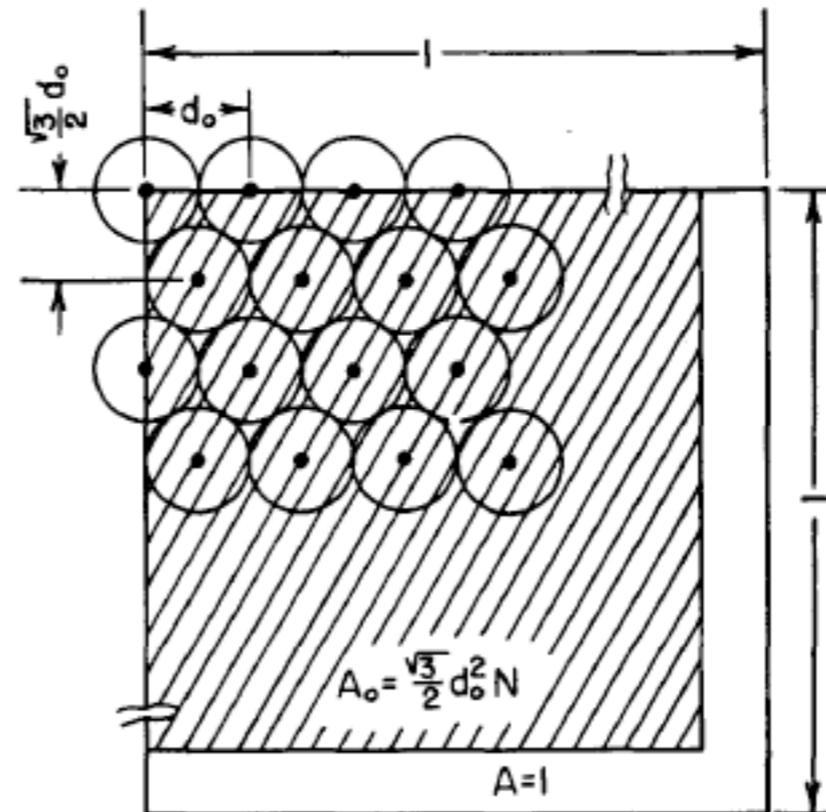
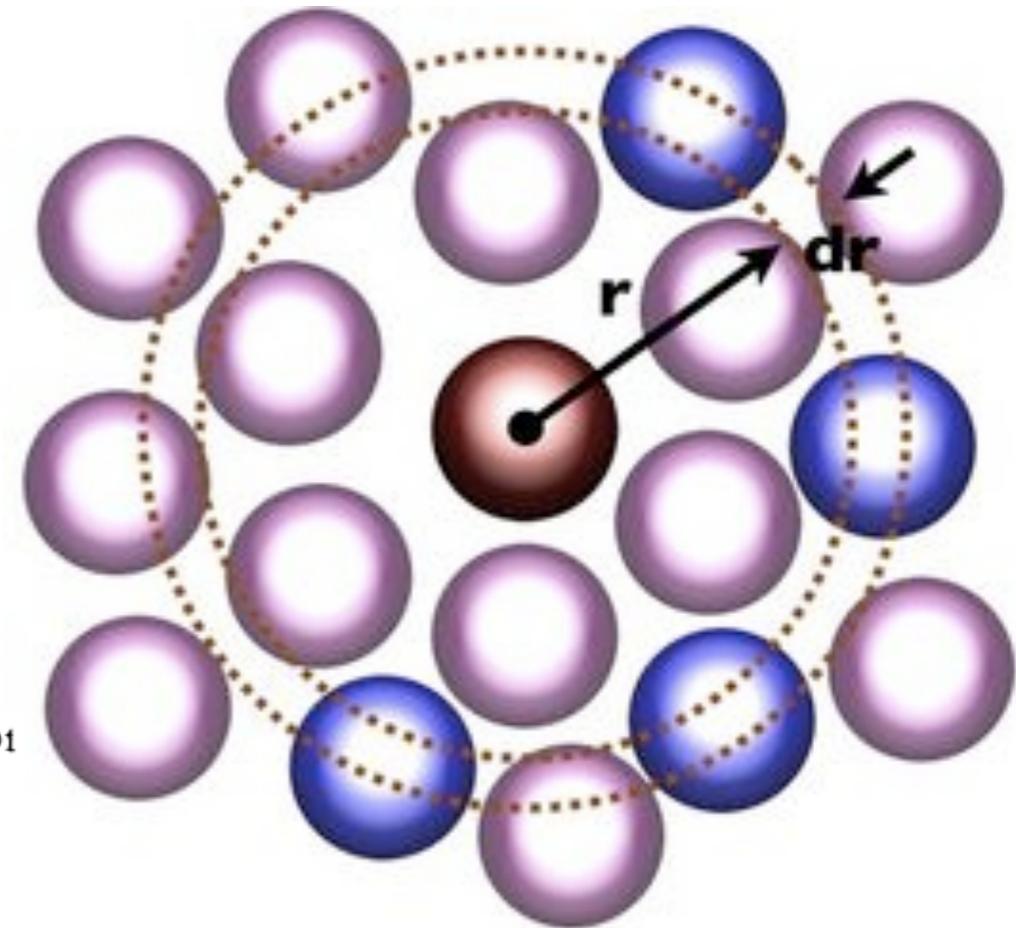


FIG. 3. The close-packed arrangement for determining  $A_0$ .

- Fix  $A_0 = \frac{\sqrt{3}d_0^2 N}{2} = 1$  and vary  $d_0$ , hence number is  $N$
- Apply periodic boundary conditions

# Probabilistic methods

- Radial distribution function measures correlations between particles separated at distance  $r$
- Can be used to distinguish solid, liquid, gases
  - See Gould-Tobochnik Chapter 8 Section 8.5
- Equation of state is deduced from the radial distribution function:



CALCULATION OF STATE BY FAST MACHINES

1091

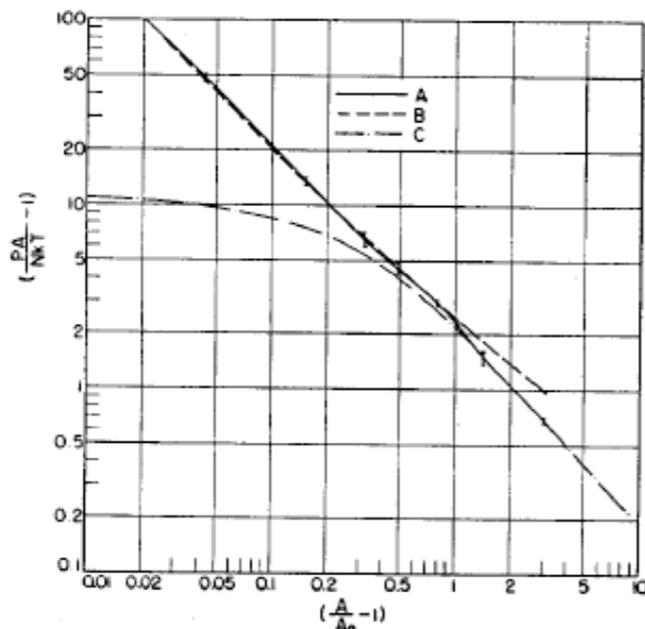


FIG. 4. A plot of  $(PA/NkT) - 1$  versus  $(A/A_0) - 1$ . Curve *A* (solid line) gives the results of this paper. Curves *B* and *C* (dashed and dot-dashed lines) give the results of the free volume theory and of the first four virial coefficients, respectively.

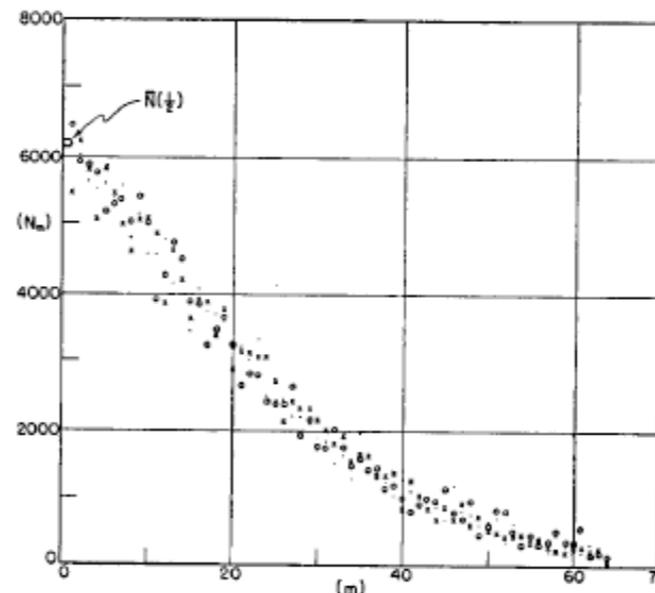


FIG. 5. The radial distribution function  $N_m$  for  $\nu=5$ ,  $(A/A_0) = 1.31966$ ,  $K=1.5$ . The average of the extrapolated values of  $N_m$  in  $\bar{N}_1=6301$ . The resultant value of  $(PA/NkT) - 1$  is  $64\bar{N}_1/N^2(K^2 - 1)$  or 6.43. Values after 16 cycles,  $\bullet$ ; after 32,  $\times$ ; and after 48,  $\circ$ .

# Probabilistic methods

- They showed

$$PA = Nk_B T \left( 1 + \frac{\pi d_0^2 \bar{n}}{2} \right), \quad \text{where } \bar{n} = n(d),$$

- Using the Virial Theorem :

– States that  $\text{K.E.} = N \times \frac{1}{2} m \bar{v}^2 = pA + \frac{1}{2} \left\langle \sum_i \mathbf{r}_i \cdot \mathbf{X}_i^{\text{int}} \right\rangle$ .

- Derivation is based on this figure, defines variables involved in a collision between two disks:

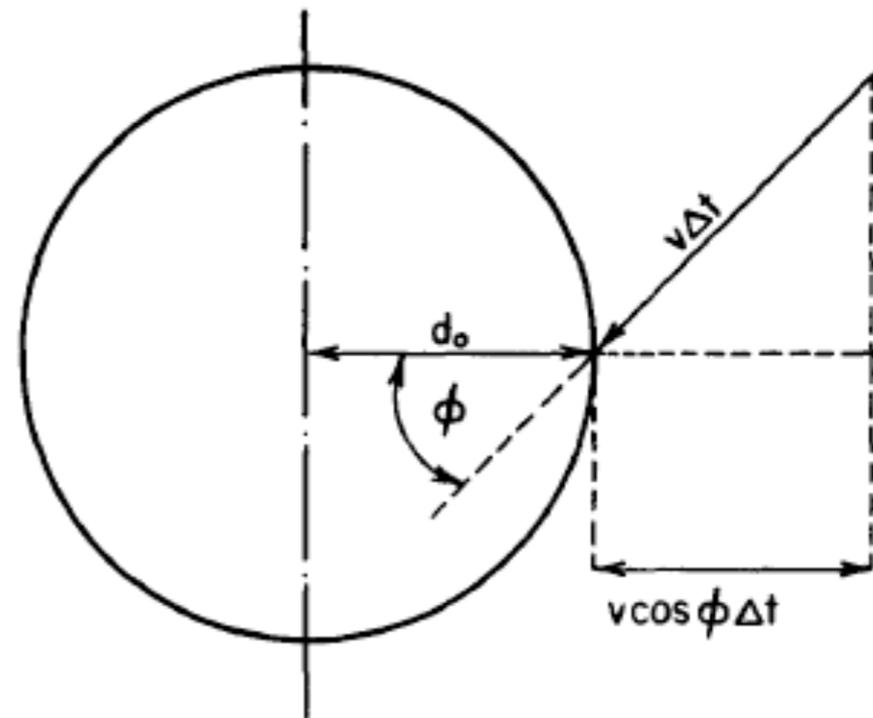


FIG. 1. Collisions of rigid spheres.

# Probabilistic methods

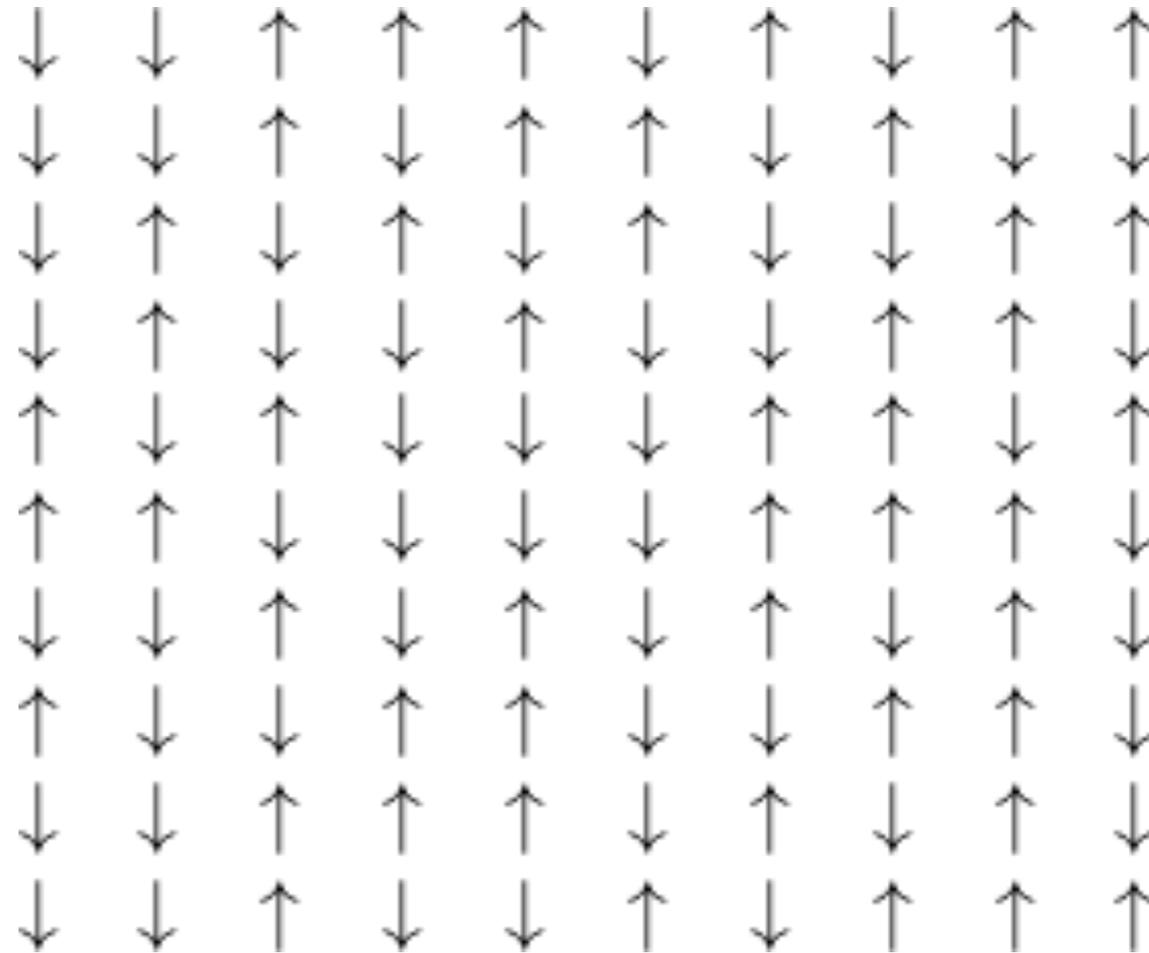
- They showed

$$\left\langle \sum_i \mathbf{r}_i \cdot \mathbf{X}_i^{\text{int}} \right\rangle = -\frac{1}{2} \sum_i \sum_{j \neq i} r_{ij} F_{ij} = N \times \frac{1}{2} m \bar{v}^2 \times \pi d_0^2 \bar{n} .$$

- MC simulation measured the radial distribution function as a histogram:
  - For each MC configuration
    - For each disk:
      - Divide region from  $r=d$  to  $r=r_{\text{max}}$  into 64 annular zones of equal area
      - Count disks in each zone, store in histogram
  - Average over configurations
  - Fit histogram to model function and extrapolate to  $r=d$

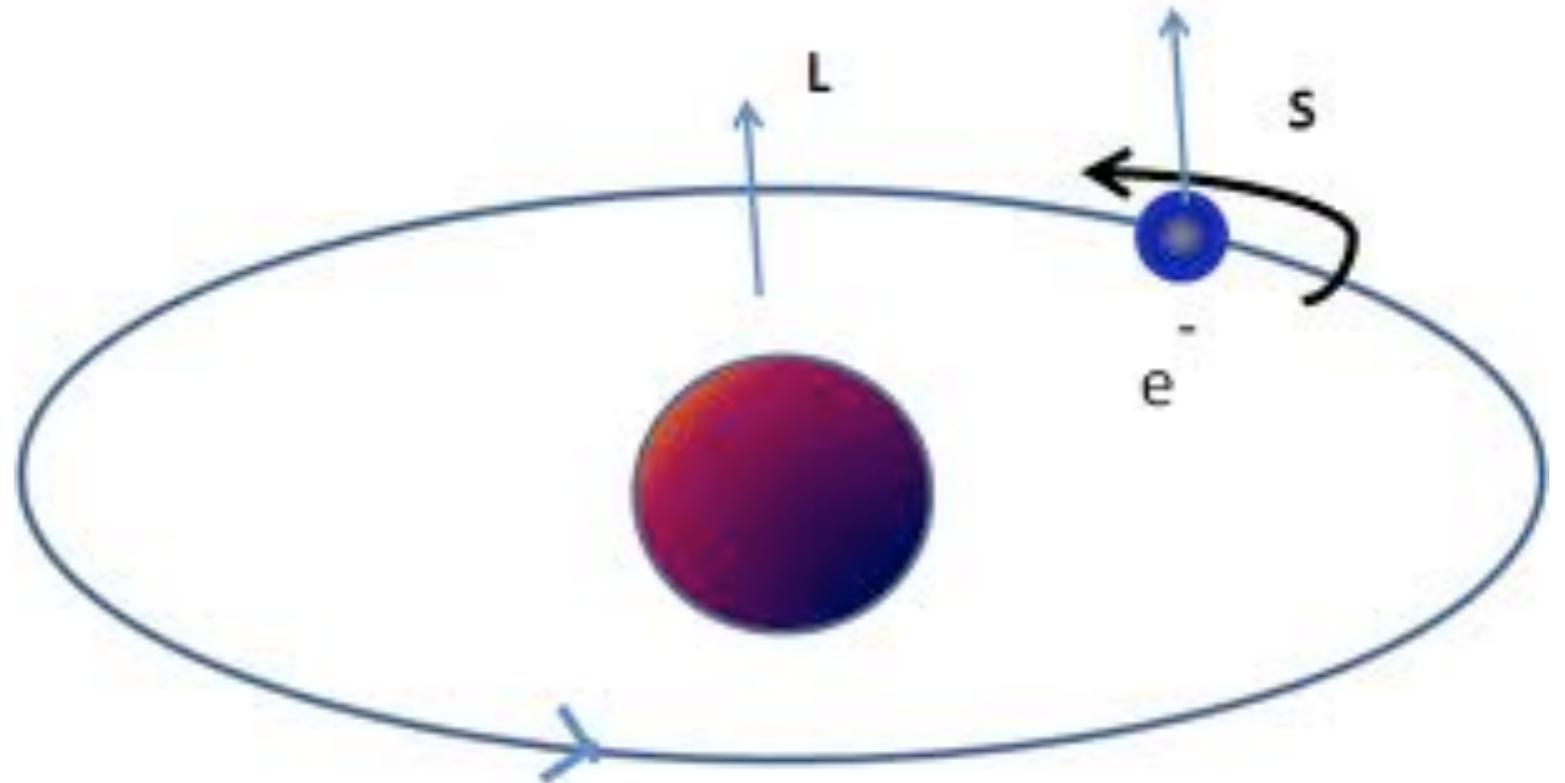
# Probabilistic methods

- Another very popular application of probabilistic methods is the simulation of idealized magnetic materials
- We use the Ising Model
  - [http://en.wikipedia.org/wiki/Ising\\_model](http://en.wikipedia.org/wiki/Ising_model)
- William Lenz assigned this to his student Ernst Ising as a PhD thesis problem
- This was solved by Ising exactly for 1-d
  - Didn't exhibit ferromagnetism
  - Details of 1-d Ising model : Gould-Tobochnik Chapter 5.
- Kramers, Wannier, Onsanger, et al worked on the 2-d model
  - Richer phenomenology :  
ferromagnetic phase at low temperatures,  
paramagnetic phase at high temperatures
  - Second order phase transition at the  
Curie temperature  $T_c$



# Probabilistic methods

- Recall : magnetism caused by charged particles “spinning” in closed orbits or about their axes
- For elementary particles, of course, we mean “spinning” in the quantum mechanical sense, not the rotational Newtonian sense
- So atoms will have both an “L” and and “S” contribution to the magnetic properties

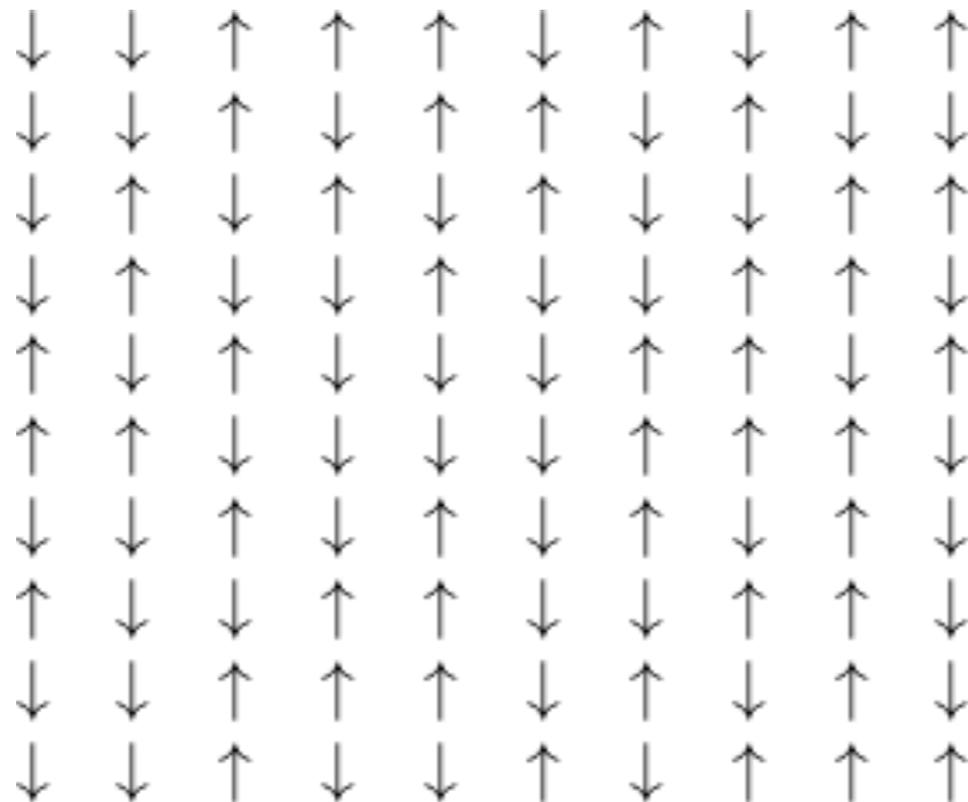


# Probabilistic methods

- We'll take a simple classical approximation called the "Ising spin" which has two values :

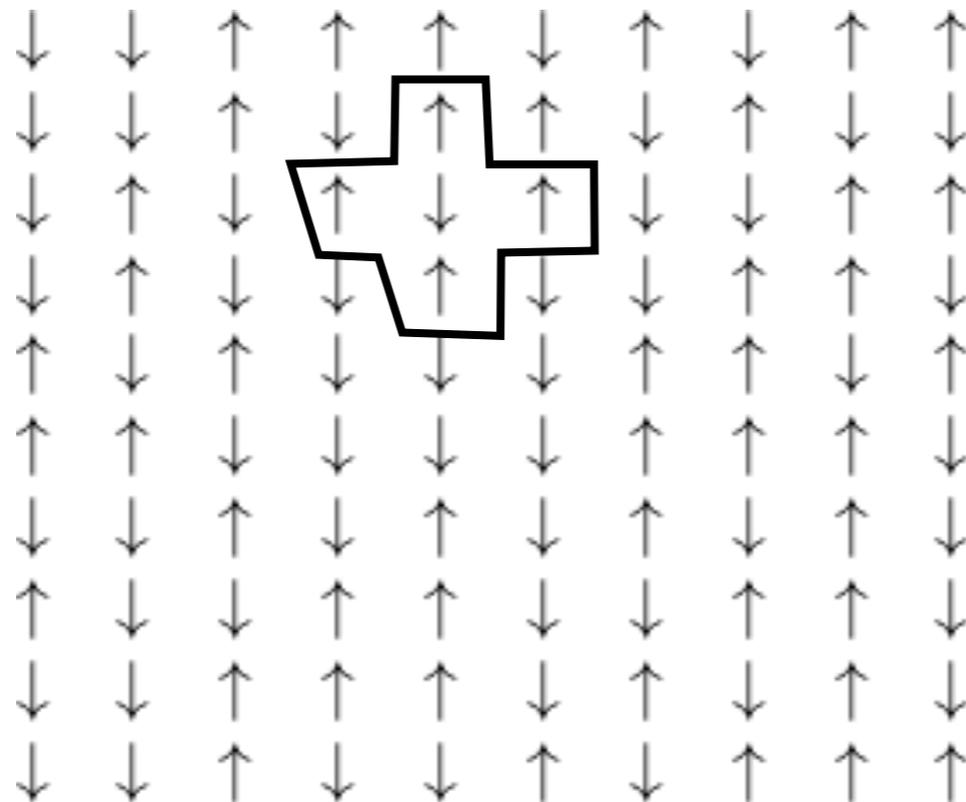
$$s_i = \begin{cases} +1, & \text{represents "spin up"} \uparrow \\ -1, & \text{represents "spin down"} \downarrow \end{cases}$$

- 2-d magnet is a set of  $N_s$  spins on a lattice



# Probabilistic methods

- The force between magnets falls of like  $r^{-3}$
- So, approximate that the spins only interact with their nearest neighbors :



# Probabilistic methods

- The interaction energy is proportional to the alignment of the spins :

$$E = -J \sum_{\langle ij \rangle} s_i s_j - H \sum_i s_i .$$

- If  $J > 0$  : ferromagnetic
  - Energy is minimized if the spins point in the same direction
- If  $J < 0$  : antiferromagnetic
  - Energy is minimized if spins locally point in the opposite direction
- “H” is an external field which couples to the total magnetization:  $M = \sum_i s_i$  ,
- Spins will align to this external magnetization

# Probabilistic methods

- For  $H=0$ , the system will be in one of two states :
  - Low temp (below Curie temperature) : magnetized
  - High temp (above Curie temperature) : sum of magnetization is zero
- Critical value is the second-order phase transition between ferromagnetic to paramagnetic phases

# Probabilistic methods

- We want to compute the observables (averages over the samples)
- Define a state as a specific set of spin values (up or down)

$$s_1 = +1, s_2 = -1, s_3 = +1, \dots, s_{N_s} = +1,$$

- Avg. of an observable calculated by weighting configurations by the Boltzmann factor
- Example : average magnetization :

$$\langle M \rangle = \frac{\sum_{\text{configs}} M e^{-E/k_B T}}{\sum_{\text{configs}} e^{-E/k_B T}}.$$

# Probabilistic methods

- Total number of configurations is very large
- If we have  $N$  spins, we have  $2^N$  configurations
- If  $L=20$ ,  $N=20*20 = 400$ , so  $2^{400}$  configurations (2.6e120 !!!)
- That's completely intractable to even list them, much less compute the total magnetization
  
- So instead, we use Monte Carlo methods
  - We generate a reasonable number of configurations at random
  - Use the Boltzmann factors to define the probability of each
  - Perfect for MC methods!

# Probabilistic methods

- The probability for each state is :

$$p(s_1, s_2, \dots, s_{N_s}) = \frac{e^{-E(s_1, s_2, \dots, s_{N_s})/k_B T}}{\sum_{\text{configs}} e^{-E/k_B T}} .$$

- Now, we need to generate  $N$  statistically independent configurations according to this probability
- The average magnetization and energy are :

$$\langle M \rangle = \frac{1}{N} \sum_{i=1}^N M(s_1^{(i)}, s_2^{(i)}, \dots, s_{N_s}^{(i)}) ,$$

$$\langle E \rangle = \frac{1}{N} \sum_{i=1}^N E(s_1^{(i)}, s_2^{(i)}, \dots, s_{N_s}^{(i)}) .$$

# Probabilistic methods

- Now use the Metropolis algorithm from last class to “map out” the configuration space
  - Choose initial configuration
  - For all spins :
    - Trial flip (+1 --> -1 or vice versa)
    - Compute change in energy
    - If  $w = e^{-\Delta E/k_B T} > r$ , flip the spin (where  $r$  is a uniform deviate)
- Make sure to pick out the thermalization and skip steps

# Probabilistic methods

- IT is expensive to compute all of the Boltzmann factors at each step
- Instead, we can realize that for a simulation at fixed T and H, there are only 10 distinct values
- Can precompute them and then just refer to that
- First, consider the sum of the 4 neighboring spins:

$$\sum_{\text{neighbors } j} s_j = s_{(i_x+1, i_y)} + s_{(i_x-1, i_y)} + s_{(i_x, i_y-1)} + s_{(i_x, i_y+1)} .$$

- The value is
  - $\sum_{\text{neighbors } j} s_j = +4$  if all four neighbors point up,
  - $\sum_{\text{neighbors } j} s_j = +2$  if three neighbors point up and one down,
  - $\sum_{\text{neighbors } j} s_j = 0$  if two neighbors point up and two down,
  - $\sum_{\text{neighbors } j} s_j = -2$  if one neighbor points up and three down,
  - $\sum_{\text{neighbors } j} s_j = -4$  if all four neighbors point down.

- Product can only have one of these five values!

$$s_i \sum_{\text{neighbors } j} s_j = +4, +2, 0, -2, -4 .$$

# Probabilistic methods

- Now, consider  $H \neq 0$
- Have terms like  $H^*s_i$
- For those we get the other 5 terms
- Can store this as a 2-d array of 5 rows and 2 columns

–First index :

$$2 + \frac{1}{2} s_i \sum_{\text{neighbors } j} s_j = 0, 1, 2, 3, 4 ,$$

–Second index :

$$\frac{1 + s_i}{2} = 0, 1 .$$

# Probabilistic methods

- Metropolis step :
  - Choose a spin at random
  - Take the metropolis step
  - If the spin is at the boundary, use periodic boundary conditions
- Because the MC step of a single spin-flip will be highly-correlated with “this” step, we need to make sure to choose sufficiently high sampling factors
  - Typically need AT LEAST  $N$  steps
  - Each spin has a chance to flip

# Probabilistic methods

- Below the Curie temperature, Kramers and Wannier showed that there is a duality argument to compute the exact value : Phys. Rev. 60, 252 (1941),

$$\frac{k_B T_c}{J} = \frac{2}{\log(1 + \sqrt{2})} = 2.269 \dots$$

- Onsager showed the same model in the thermodynamic limit  $N \rightarrow \infty$  with  $H=0$  (Phys. Rev. 65, 117 (1944) )

$$m = \lim_{N \rightarrow \infty} \frac{\langle \sum_i s_i \rangle}{N} = \begin{cases} \left[ 1 - \left\{ \sinh \left( \frac{2J}{k_B T} \right) \right\}^{-4} \right]^{1/8}, & \text{for } T \leq T_c \\ 0, & \text{for } T > T_c \end{cases}$$

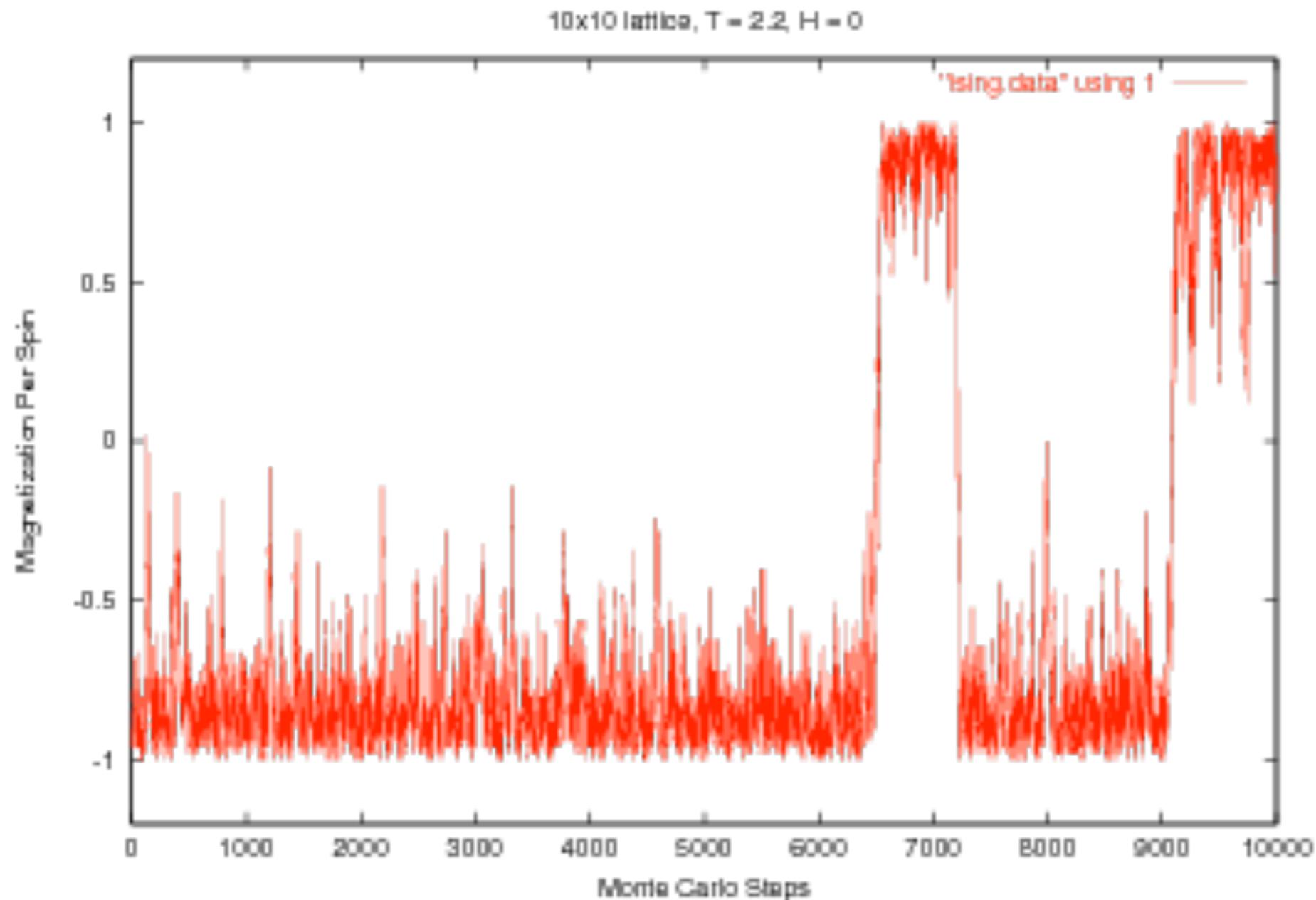
- Near the Curie temperature

$$m \sim (T_c - T)^\beta ,$$

- Beta = 1/8 for the 2-d Ising model

# Probabilistic methods

- Here are some numerical results for  $T = 2.2$ , below  $T_c$



- Note : for finite systems, spins can flip from time to time!

# Probabilistic methods

- And now for  $T=3.0$ , above  $T_c$ :

