# Physics 381/382/504 Lab Handout

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### 1 Introduction

The Ising model is one of the simplest physical models that exhibits a phase transition. It consists of a d-dimensional square lattice of "particles," with a "spin" (either up or down) and a magnetic moment that aligns with its spin. The particles (also known as lattice sites) interact with each other through a nearest-neighbor interaction of strength J and are coupled to an external magnetic field with strength B. For a lattice of sidelength N, we then have the Hamiltonian:

$$H = -J\sum_{\langle i,j\rangle} \sigma_i \sigma_j - B\sum_i \sigma_i \tag{1}$$

where  $\sigma_i = \pm 1$  denotes the *i*-th spin state and  $\langle i, j \rangle$  denotes a sum over nearest neighbors (in 2-dimensions there are 4 nearest neighbors). Analytic solutions to the Ising model are usually considered in the limit  $N \to \infty$ .

For J > 0, the spins tend to align, but at finite temperature, thermal fluctuations will prevent them from doing so. For  $d \ge 2$ , the system exhibits a phase transition at zero applied field and a critical temperature  $T_c$ , such that for  $T > T_c$ , the system is dominated by thermal fluctuations and no overall magnetization is observed, while for  $T < T_c$ , the spins tend to align and the system develops a spontaneous magnetization. You will simulate the Ising model using Markov Chain Monte Carlo (MCMC) techniques, and one of your first tasks will be to measure  $T_c$ .

A collection of exact results for the Ising model is presented at [1]. For a good introduction to Monte Carlo sampling techniques in general, see [2].

<sup>&</sup>lt;sup>1</sup>We will soon specialize to the case d=2, but the same questions you will investigate here can be asked for the Ising model in any number of dimensions. For d=1,2 and  $d\geq 4$ , it turns out the Ising model is exactly solvable, meaning all the quantities you will compute from simulation can be solved for analytically. The case d=3 is an area of current research.

## 2 Overview of Statistical Mechanics

This section is not intended to be a complete exposition of statistical mechanics, but merely to remind you of a few of the basic ideas and definitions. For a much more thorough reference, consult any textbook on thermodynamics or statistical mechanics, such as [3].

The most important prediction of statistical mechanics for us is the Gibbs distribution, which states that the probability of finding a macroscopic system in a microscopic state of energy E is given by:

$$P(\text{state}) \propto e^{-\frac{E}{k_B T}}$$
 (2)

where  $k_B$  is Boltzmanns constant and T is the temperature of the system. In principle this makes it easy to compute various macroscopic properties of the system. For example, if we index the possible microstates by i and their energies by  $E_i$  then the expected value of some observable property (like the overall energy of the system) is given by averaging over all the microstates with weights given by Equation 2. Denoting the thermodynamic average at a temperature T by  $\langle \cdot \rangle_T$  (we will occasionally drop the T but it should always be understood that the thermodynamic average is temperature-dependent) and defining the thermodynamic beta  $\beta = 1/(k_B T)$  we have:

$$\langle E \rangle_T = \frac{\sum_i E_i e^{-\beta E_i}}{\sum_i e^{-\beta E_i}} \tag{3}$$

where the denominator is a normalization factor to deal with the fact that Equation 2 is only a proportionality, not an equality.

Now there is a very slick way to repackage this in terms of the partition function  $\mathcal{Z}$ , defined as

$$\mathcal{Z} = \sum_{i} e^{\beta E_i} \tag{4}$$

It is clearly a function of temperature because of the explicit dependence on  $\beta$ , but it is also implicitly a function of other external variables, such as applied magnetic field in the 2-D Ising model because each factor of  $E_i$  depends on these variables. Now many thermodynamic quantities can be realized as simple derivatives of the partition function. For example:

$$\langle E_T \rangle = \frac{1}{\mathcal{Z}} \sum_i E_i e^{-\beta E_i} = \frac{1}{\mathcal{Z}} \left( -\frac{\partial}{\partial \beta} \sum_i e^{-\beta E_i} \right) = \frac{1}{\mathcal{Z}} \left( -\frac{\partial}{\partial \beta} \mathcal{Z} \right) = -\frac{\partial}{\partial \beta} \ln \mathcal{Z} \quad (5)$$

since we assume each  $E_i$  does not depend on temperature.

The partition function can also be used to derive various useful relations among thermodynamic quantities. In this experiment, you will be interested in computing specific heat  $c_V$  and magnetic susceptibility  $\chi$ , defined as:

$$c_V = \frac{\partial \langle E \rangle_T}{\partial T}, \quad \chi = \frac{\partial \langle M \rangle_T}{\partial B}$$
 (6)

where M is the magnetization (per site) of the system and B is an externally-applied magnetic field. Trying to compute these in this way would be difficult because of the large errors associated with derivatives of numeric quantities, but using the partition function we can find more useful forms for them. We begin with  $c_V$ :

$$c_V = \frac{\partial \langle E \rangle_T}{\partial T}, c_V = \frac{\partial \langle E \rangle_T}{\partial T} = \frac{\partial \langle E \rangle_T}{\partial \beta} \frac{\partial \beta}{\partial T} = -\frac{\beta}{T} \frac{\partial \langle E \rangle_T}{\partial \beta}$$
(7)

$$= \frac{\beta}{T} \frac{\partial}{\partial \beta} \left( \frac{1}{\mathcal{Z}} \frac{\partial}{\partial \beta} \mathcal{Z} \right) \tag{8}$$

$$= \frac{\beta}{T} \left( \frac{1}{\mathcal{Z}} \frac{\partial^2}{\partial \beta^2} \mathcal{Z} - \frac{1}{\mathcal{Z}^2} \left( \frac{\partial \mathcal{Z}}{\partial \beta} \right)^2 \right) \tag{9}$$

Note that the second term in the second factor in Equation 9 is simply  $\langle E \rangle_T^2$  while the first term is:

$$\frac{1}{\mathcal{Z}}\frac{\partial^2}{\partial\beta^2}\mathcal{Z} = \frac{1}{\mathcal{Z}}\frac{\partial^2}{\partial\beta^2}\sum_i e^{-\beta E_i} = \frac{1}{\mathcal{Z}}\sum_i E_i^2 e^{-\beta E_i} = \langle E^2 \rangle_T \tag{10}$$

so we obtain a more useful formula for  $c_V$ :

$$c_V = \frac{\beta}{T} \left( \langle E^2 \rangle_T - \langle E \rangle_T^2 \right) \tag{11}$$

A virtually identical derivation gives a similar formula for  $\chi$ :

$$\chi = \beta \left( \langle M^2 \rangle_T - \langle M \rangle_T^2 \right) \tag{12}$$

Throughout this experiment, you should use Equations 11 and 12 to compute  $c_V$  and  $\chi$  respectively.

The final thermodynamic quantity we will care about here is the correlation length  $\xi$ . In any system at a finite temperature there will be thermodynamic fluctuations, and often we are interested in the rough size of these fluctuations. To probe this, we look at spatial correlation functions, which measure how closely two different locations in our system are correlated. For the Ising model, we will be interested in the correlation function  $R(x) \equiv \langle \sigma(0)\sigma(x)\rangle$ , which measures how likely it is that two spins separated by a distance x (or, in our simulation number of lattice site) are aligned. Since the lattice is translation-invariant, we can always translate so one of the lattice sites are are considering is the origin). If  $\langle \sigma(0)\sigma(x)\rangle \sim 1$  then the two sites are highly correlated, while if  $\langle \sigma(0)\sigma(x)\rangle \sim 0$  then the two sites are uncorrelated. (If  $\langle \sigma(0)\sigma(x)\rangle \sim -1$ , the two sites are highly correlated but are likely to be anti-aligned rather than aligned.)

It is generally true that spatial correlation functions die off exponentially with some characteristic length scale  $\xi$ :

$$\langle \sigma(0)\sigma(x)\rangle \sim e^{-x/\xi}$$
 (13)

where  $\xi$  is known as the correlation length and will depend on temperature. For  $x \ll \xi$ , the correlation function will typically have some power-law behavior with some characteristic exponent typically written as:

$$\langle \sigma(0)\sigma(x)\rangle \sim \frac{1}{|x|^{d-2+\eta}}, \quad x \ll \xi$$
 (14)

where d is the dimension of the system. In this lab we will be interested in finding the correlation length of the system at various temperatures and also in finding the exponent near the phase transition.

### 3 Overview of Numerical Methods

This section is intended as an extremely brief overview of why numerical methods are necessary for this problem and how they work. I will skip most of the details since they are discussed quite well (and in the context of the Ising model) in [2].

In principle the rules of statistical mechanics are very straightforward. If we want to compute the thermodynamic average of some variable  $\mathcal{O}$ , we simply use the Gibbs distribution:

$$\langle \mathcal{O} \rangle_T = \frac{1}{\mathcal{Z}} \sum_i \mathcal{O}(i) e^{-i\beta E_i}$$
 (15)

In practice this is very difficult, since we need to sum over all the different microstates of the system, and any macroscopic system has an incredibly large number of possible microstates. Even a very simple system, such as a  $20 \times 20$  lattice of spins , has  $2^{20^2} \sim 10^{120}$  possible states, substantially more than the number of particles in the universe, so actually computing the thermodynamic average is impossible. Instead, we need to find a way to sample the Gibbs distribution in a way that gives a good approximation to the true thermodynamic average. This is where numerical methods come in.

In order to make our numerical approximations, we first note that we don't have access to the actual Gibbs probability distribution, given by

$$P(\text{state}) = \frac{1}{Ze^{-\beta E(\text{state})}} \tag{16}$$

since we do not know  $\mathcal{Z}$  We can, however, easily compute  $e^{-\beta E}$  for any given state, so we have access to a function which is proportional to the actual probability distribution we want to sample. This turns out to be enough. At this point there are several different numerical techniques we can use, such as *importance sampling*, rejection sampling, and the *Metropolis algorithm* (all of these are others are discussed in [2]). In this experiment you will use the last of these, and so that is all we will discuss here.

Before we continue however, it is worth understanding why the most naive sampling method, uniform sampling, does not work. Imagine trying to compute a thermodynamic average by selecting n random states, computing  $\mathcal{Z}$  based only on these states, and then

computing  $\mathcal{Z}^{-1}\Sigma_{j=1}^n \mathcal{O}(j)e^{-\beta E_j}$  as an approximation of  $\langle \mathcal{O} \rangle_T$ . The key problem with this is that although the number of states is very large, both  $\mathcal{Z}$  and the thermodynamic average are typically dominated by a much smaller set of states, known as the *typical set*. If our uniform sample doesnt happen to contain any of the states in the typical set, we will come up with a wildly inaccurate estimate for  $\langle \mathcal{O} \rangle_T$ . So just how much smaller does the typical set tend to be? In the case of the Ising model with  $N^2$  sites near the critical temperature, it turns out that the typical set is quite small. In order to hit it even once you need to draw on average  $2^{N^2/2}$  samples. For N=20, this is roughly  $10^{60}$  samples. Not quite the number of particles in the universe, but large enough that this should convince you that we need something more sophisticated than uniform sampling.

The basic idea of Metropolis sampling is to start somewhere in the state space and then explore the space in a way that tends to keep you near the areas of high probability density, since these are the areas that will dominate the thermodynamic average. It can be shown that over a long period of time this procedure gives an accurate sample of the underlying distribution, but we will not do so here. Instead, we will focus on how the algorithm works and what problems it can run into.

Metropolis sampling falls into a class of numerical techniques known as Markov Chain Monte Carlo (MCMC). This is because it explores the probability distribution by building up a Markov chain of samples. At each step, it only uses information about its current position to determine where it should move next. Over time, it will generate a set of samples  $\mathbf{x}(t)$  that will give an accurate value for thermodynamic averages. So how does it do this?

At each step t, a tentative step x' is generated (i.e. from the given state of the system, a spin is chosen to potentially be flipped). Then we compute

$$a = \frac{\exp(-\beta E(x'))}{\exp(-\beta E(x^{(t)}))}$$
(17)

where E(x') and  $E(x^{(t)})$  are the energies of the system in the trial state and current state respectively. If  $a \ge 1$  then the trial state is more probable than the one we are currently in (and thus closer to the typical set), and so we accept it and set  $x^{(t+1)} = x'$ . If a < 1 then the trial state is less probable than the one we are currently in, but that doesnt mean we should immediately rule it out. Instead, we want to explore the probability space, and so we accept the trial state with probability a. Importantly, if we reject the trial state we set  $x^{(t+1)} = x^{(t)}$ , so it is possible to have the same state for several steps in a row if that state is much more likely than the others close to it.

Clearly nearby samples are correlated, but over time the Metropolis algorithm will explore the state space and then a good approximation for  $\mathcal{Z}$  and the thermodynamic

<sup>&</sup>lt;sup>2</sup> Deriving these results uses some probability theory that we do not have here. [2] has slightly more information, but for good comprehension, you are recommended to use a book on information theory, such as [4]

average can be computed by:

$$\mathcal{Z} \approx \sum_{t} e^{-\beta E(x^{(t)})} \tag{18}$$

$$\mathcal{Z} \approx \sum_{t} e^{-\beta E(x^{(t)})}$$

$$\langle \mathcal{O}_{T} \rangle \approx \frac{1}{\mathcal{T}} \sum_{t} \mathcal{O}\left(x^{(t)}\right)$$

$$\tag{18}$$

where  $\mathcal{T}$  is the total number of samples generated by the run. In the second equation, note the absence of  $\mathcal{Z}$  and the Gibbs weight  $e^{-\beta E}$ . This is because the Metropolis algorithm automatically generates more samples in the regions of high probability density in precisely a way that eliminates the need for these factors.

Now that we have some idea of these techniques, it is worth pointing out a few things. First, although we have denoted our sample series above as  $\{x^{(t)}\}\$ , you should not interpret this as a time series. This is a walk through the probability space of the system, and although it is sometimes helpful to think about your experiment as simulating a spin lattice for some finite time period, you must remember that this isnt actually what you are doing. Second, the Metropolis algorithm must be given the chance to fully explore the probability space, otherwise the final average will be significantly biased by the initial state you set the system in. Third, although the algorithm will eventually find the typical set, it may take a while to do so. Once it finds the typical set the simulation is said to have *converged*, but before this happens, all the samples from the probability distribution are being drawn from low-probability areas dictated by your choice of initial conditions. For this reason, it is usually recommended that you allow the system to burn in for several thousand steps before you start collecting data.

For more information on how to implement the Metropolis algorithm in the case of the Ising model, see Section 5. For tips on how to better achieve convergence, see Appendix A.

#### Theory and Goals 4

From here on, it is easiest to choose a simple unit system. We first set J=1, which means all energies will be measured in units of J. We then set Boltzmanns constant,  $k_B = 1$ , and since  $k_B$  has units of energy over temperature, this amounts to measuring temperature in units of energy and thus in units of J. With these choices, all quantities in the problem become unitless, and we will assume this unit system throughout the rest of this handout. We also specialize to the case d=2.

The exact critical temperature of the 2D Ising model is given by:

$$T_c = \frac{2}{\ln\left(1 + \sqrt{2}\right)} \approx 2.269\tag{20}$$

Your first task is to simulate the Ising model and determine  $T_c$  numerically (with an uncertainty). A general characteristic of critical points is that many thermodynamic quantities in the system show nonanalytic behavior (discontinuities, divergences, etc.), and  $T_c$  can be determined by fitting theoretical functions to the behavior of these quantities in the regime near the critical point.

You should use at least three different system state variables (specific heat, magnetization, susceptibility, and correlation length) to determine  $T_c$ . Each experimental value for  $T_c$  should have an uncertainty of  $\lesssim 5\%$ , and your report should contain a full description and justification for your statistical analysis.

Your next task is to compare the expected theoretical forms for the heat capacity, magnetization, and magnetic susceptibility to a numerical calculation. Since the Ising model is solvable in two dimensions, there are known exact results for these that are valid at all temperatures. For example, in the  $N \to \infty$  limit, the energy per lattice site as a function of temperature, E(T) is given by:

$$E(T) = -2\tanh\left(\frac{1}{T}\right) - \frac{\sinh^2(2/T) - 1}{\sinh(2/T)\cosh 2/T} \left[\frac{2}{\pi}K_1(\kappa) - 1\right]$$
(21)

where,

$$\kappa = 2 \frac{\sinh(2/T)}{\cosh^2(2/T)} \tag{22}$$

$$K_1(\kappa) = \int_0^{\pi/2} \frac{\mathrm{d}\phi}{\sqrt{1 - \kappa^2 \sin^2 \phi}} \tag{23}$$

Here  $K_1(\kappa)$  is a complete elliptic integral of the first kind and can be numerically evaluated. The average magnetization per site has a much simpler form. For  $T > T_c$ , we have  $\langle M \rangle = 0$ , and for  $T < T_c$  we have:

$$|\langle M \rangle| = \left(1 - \left[\sinh(2/T)\right]^{-4}\right)^{1/8} \tag{24}$$

Knowing these formulas, it is possible to compute specific heat  $(c_V)$  and magnetic susceptibility  $(\chi)$  as:

$$c_V = \frac{\partial \langle E \rangle_T}{\partial T}, \quad \chi = \frac{\partial \langle M \rangle_T}{\partial B}$$
 (25)

Needless to say, these formulas are unwieldy. Luckily, your analysis will be made easy by restricting your attention to the vicinity of the critical temperature. It is a general result of the theory of critical phenomena that for temperatures T near  $T_c$ , virtually all the system state quantities of interest scale as some power law in the reduced temperature t:

$$t = \frac{T - T_c}{T_c} \tag{26}$$

The exponents in these power-law relations are known as *critical exponents*, and they are of principle interest, since they are are characteristics of the critical point itself, not

Definition	Theor. Values	
$c_V \propto  t ^{-\alpha}$	0	
$ M  \propto  t ^{\beta}$	1/8	
$\chi \propto  t ^{-\gamma}$	7/4	
$ M  \propto  B ^{1/\delta}$	15	
$\xi \propto  t ^{-\nu}$	1	
$\langle \sigma(0)\sigma(x)\rangle \propto  x ^{-d+1-\eta}$	1/4	
Name/definition		
specific heat		
magnetization		
magnetic susceptibility		
correlation length		
$(T-T_c)/T$		
	$c_V \propto  t ^{-\alpha}$ $ M  \propto  t ^{\beta}$ $\chi \propto  t ^{-\gamma}$ $ M  \propto  B ^{1/\delta}$ $\xi \propto  t ^{-\nu}$ $\langle \sigma(0)\sigma(x)\rangle \propto  x ^{-d+1-\eta}$ Name/definition  specific heat magnetization magnetic susceptibility correlation length	

Table 1: Theoretical values of some of the critical exponents in the d=2 Ising model as  $N\to\infty$ . In general,  $\alpha$ ,  $\gamma$ , and  $\nu$  are well-defined both above and below  $T_c$ , and the Renormalization Group (RG) approach to critical phenomena implies that in most cases (including the 2-D Ising model), they should be the same both above and below  $T_c$ . Note that  $\beta$  is only well-defined above  $T_c$ . The value of 0 given for  $\alpha$  means that  $c_V$  will not truly behave as a power law in |t| but rather  $c_V \propto \log(t)$  near  $T_c$ .

the Ising model in particular. In other words, there are several different systems you can write down with different degrees of freedom and different Lagrangians but which share some critical behavior and critical exponents near a phase transition.<sup>3</sup> The critical exponents you will be interested in are defined as follows, and Table 4 gives exact values for these exponents in the 2-D Ising model.

$$c_V \propto |t|^{-\alpha} \tag{27}$$

$$\chi \propto |t|^{-\gamma} \tag{28}$$

$$\xi \propto |t|^{-\nu} \tag{29}$$

$$|M| \propto |t|^{\beta} \tag{30}$$

where  $\xi$  is the correlation length and  $\beta$  is not to be confused with the thermodynamic  $\beta = 1/(k_BT)$ . In the above, it must be noted that the expression for |M| only makes sense below  $T_c$ . Further, near  $T_c$  (and within the correlation length), the spatial spin correlation function R(x) has the form:

$$R(x) = \langle \sigma(0)\sigma(x)\rangle \propto \frac{1}{|x|^{d-2+\eta}}$$
 (31)

where d=2 is the dimensionality of the system. Finally, for T slightly above  $T_c$ , the magnetization with an applied (static, homogeneous) magnetic field B has the form:

$$|M| \propto |B|^{1/\delta} \tag{32}$$

You should be able to use your simulated data to (1) verify that the above quntities truly do have power-law scaling behavior near  $T_c$  and (2) find the critical exponents discussed above. At minimum you should find  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\nu$ , and  $\eta$ , but if you have time, finding  $\delta$  is also fairly simple.

The general theory of critical phenomena also yields the following relations between the various critical exponents, where again d is the dimension of the system:

Rushbrooke's, Identity: 
$$\alpha + 2\beta + \gamma = 2$$
 (33)

Widom's Identity: 
$$\delta - 1 = \gamma/\beta$$
 (34)

Josephson's Identity 
$$2 - \alpha = d\nu$$
 (35)

Fisher's Identity: 
$$\gamma = (2 - \eta)\nu$$
 (36)

These imply that even though there are apparently six independent critical exponents listed above, there are only two independent exponents.<sup>4</sup> Once you have computed the

<sup>&</sup>lt;sup>3</sup> This is a highly nontrivial phenomenon, known as *critical universality*, and the reasons for it were only uncovered in the 1960s and 1970s, principally by Leo Kadanoff, Michael Fisher, and Ken Wilson. For their work they jointly received the Wolf prize in 1980, and for his work Wilson was awarded the Nobel Prize in Physics in 1982.

<sup>&</sup>lt;sup>4</sup> All this is essentially a consequence of the fact that you only need to tune two parameters (i.e. temperature to  $T_c$  and external applied magnetic field B to 0) to reach the critical point.

above critical exponents, verify that your system obeys these identities to within your experimental precision.

So far, everything in this section has assumed the infinite-size limit  $N \to \infty$ . Since this is a simulation-based experiment and you only have access to a finite-size computer and a finite amount of time, you will be simulating only a finite-size grid [5]. In this case, you will only have finitely-many degrees of freedom (namely the  $N^2$  lattice sites), and it is generally true that any system with only a finite number of degrees of freedom cannot have a true phase transition.<sup>5</sup> Luckily for us, such systems can come close to a phase transition, but there will still be some residual finite-size effects that will be most pronounced very near the critical temperature. For example, in a finite-size system, the specific heat, magnetic susceptibility, etc. cannot truly diverge, but instead will obtain some maximum value that will depend on the grid size N.

This behavior sets in when the system "realizes" it is finite, which happens roughly when the correlation length is on the order of the lattice size:  $\xi \sim N$ . Since  $\xi \propto |t|^{-\nu}$ , this gives an effective value  $t_{\rm eff}$  at which  $|t_{\rm eff}|^{-\nu} \sim N$ , and we expect that the various thermodynamic quantities will have their divergences cut off near this temperature. Thus we expect that the maximum value of  $\xi$  for example, instead of being infinite at  $T_c$ , will be given roughly by:

$$\xi(T = T_c) \sim (N^{-1/\nu}) \sim N^{\gamma/\nu} \tag{37}$$

This analytic method is known as *finite size scaling*, and provides another probe of the critical exponents. By running your experiment at several different grid sizes, you can extract / and compare it with your results from above. You can run a similar analysis for  $c_V$ . All of this finite-size analysis is optional, but if you have time it is highly recommended. It should help give you a sense for how what initially appears to be an obstacle (the fact that you have to limit yourself to a finite-size grid) can actually be used as a tool.

The above discussion has been fairly theoretical. We will now move into how to actually simulate the Ising model. If, after reading through that procedure, you still have questions about extracting any of the above quantities, check Appendix B, although be aware that you will likely need to experiment with things yourself. Getting a Monte Carlo simulation of the Ising model running and producing reliable results with good statistics will occupy the bulk of your time at first, and you will likely need to let your program run for an extended period of time in order to collect enough usable data to get reasonable precision on the various quantities you will measure.

<sup>&</sup>lt;sup>5</sup>This is fairly easy to see. A system with only finitely-many degrees of freedom only has finitely-many states it can possibly be in, and so the partition function must necessarily be analytic. An analytic partition function means that all macroscopic thermodynamic properties (various derivatives of the partition function) must be analytic, and thus a phase transition, characterized by non-analytic behavior in one or more thermodynamic quantities, is impossible.

# **Summary of Requirements:**

This simply restates the requirements for your experiment. All of the following should be included in your report (regardless of whether it is your formal or informal report).

- Compute  $T_c$  from at least three different system state variables (specific heat, magnetization, magnetic susceptibility, and correlation length). Each fit should have a precision of  $\lesssim 5\%$  and and you should fully explain and justify the statistical analysis that determines the uncertainty in your values.
- Verify power-law scaling behavior of  $c_V$ , |M|, (beneath  $T_c$ ),  $\chi$ , xi, and  $\langle \sigma(0)\sigma(x)\rangle$ . For the 2-D Ising model, note that  $c_V$  actually has logarithmic scaling rather than power-law scaling.
- Compute (with uncertainties) the critical exponents  $\alpha$ ,  $\beta$ ,  $\gamma$ , nu, and  $\eta$ . Compare them to the theoretical values in Table 1.
- Verify that to within your experimental precision the critical exponents that you find satisfy the known scaling relations given in Equations 33, 34, 35, and 36. If you did not find  $\delta$  you do not need to verify Equation 34.

If you have completed all of the above, consider experimenting with the some of the following:

- Experiment with varying the external applied magnetic field B and compute the critical exponent  $\delta$ .
- Investigate the finite-size scaling behavior of your system. Extract  $\gamma/\nu$  using this method as discussed above.

# 5 Code and Algorithm

Base code is available for you to get you started on this lab. The purpose of this code is to implement a 2D  $Markov\ Chain\ Monte\ Carlo\ (MCMC)$  algorithm and allow to you interface with the code to calculate the desired physical properties listed in the previous section.

The code generates a grid of size  $N_{\rm sites} = N \times N$  lattice locations/sites, where N is an input parameter. This "lattice" is selected with periodic boundary conditions – i.e. the bottom row "wraps around" to be next to the top row, as does the left column to be next to the right column. In effect, this places the lattice on a torus.

For any configuration of spins, function calls can calculate the following from the grid using the nearest neighbor interaction strength (J) and magnetic field strength (B).

<sup>&</sup>lt;sup>6</sup> Note that the code hard codes the value of J to 1. This is for the simplicity of coding, though in your lab write-up you will need to explain what this does to the units and values of B and T).

- 1. Average energy per site:  $H/N_{\text{sites}}$ , where H is calculated by Equation 1.
- 2. Average spin per site:  $\sum_{i} \sigma_i / N_{\text{sites}}$
- 3. The spacial spin correlation function:  $\langle \sigma(0)\sigma(k)\rangle_T = \text{avg}(R_i(k))$  where  $R_i$  is the auto-correlation for any given row/column i, calculated by

$$R_i(k) = \frac{1}{N} \sum_{i} (\sigma_{(i,j)} - \bar{\sigma}_i) (\sigma_{(i,j+k)} - \bar{\sigma}_i)$$

where  $\bar{\sigma}_i$  is the average of all spins in the row/column labelled by i and j + k is understood to be taken to be modulo N. Note that this is only calculated for j < N/2 because of the periodic boundary conditions.

The lattice may be evolved using MCMC steps, in which each step does the following:

- 1. Select  $n_{\rm s}$  lattice sites randomly, where  $n_{\rm s} \leq N_{\rm sites}$ .
- 2. For each selected site i:
  - (a) Determine the change in energy to the system if the site is flipped:  $\Delta E_i = 2(J\sigma_i(\Sigma_i\sigma_i) + B\sigma_i)$ , where j indexes the four adjacent sites.
  - (b) If  $E_i < 0$ , the new state is more probable, so add this site to the list of sites to flip.
  - (c) if  $E_i > 0$ , energy is required to flip the spin. The probability of having this energy is determined by the Boltzmann distribution. Therefore, add it to the list of sites to flip with the probability of  $\exp\left[-\Delta E(i)/(k_BT)\right]$ .
- 3. Flip all sites selected to be flipped.

For computational efficiency, a C++ library has been written to do the above. For the sake of convenience, a C wrapper has been written that allows Python to initialize and interact with the above library. Additionally, a Python program has also been written to drive the above library and submit multiple parallel runs of different ising MCMC lattice simulations and cataloging the final grid physical results in output text files.

<sup>&</sup>lt;sup>7</sup> Note that in the pure MCMC algorithm, only a single site are selected for every step with  $n_{\rm s}=1$ . However, for efficiency, it turns out that it's good enough to pick a small percentage of  $N_{\rm sites}$  and flip them. Be careful though. If you flip too few spins per step your simulation will take too long to get good results, and if you flip too many, the state will be too volatile and will not give a good walk throughout the probability distribution. The code default is  $n_{\rm s} \approx N_{\rm sites} \times 10\%$ .

<sup>&</sup>lt;sup>8</sup> In context of this literature, "chance to flip" means the chance of the probability distribution falling above a random number uniformly selected from the range of 0 to 1. Therefore, the program chooses "flip" if random number  $x \in (0,1)$  is less than  $\exp(-E_i/T)$ .

You are free to use the provided code at your convenience. Typically, your coding for this lab will consist of using Python to generate appropriate run conditions and sample outputs of the lattice at various steps. How you do this is largely up to you. In general, you are advised to start with smaller lattices, and smaller numbers of steps, and move to larger and longer runs when you have good code and are seeking more precision in your results (or perhaps are studying the effects of lattice size and run times on results).

The available code is posted on Github at https://git.yale.edu/phys382L/ising2D\_code. If you wish, you can download the code to your computer, compile the C++ library and use it locally. The Yale High Performance Computing Group (HPC) has also put the code onto the Grace cluster and will (or has already) generated a group folder for this class, so that you can ssh into Grace and work from there. This will allow you to use many nodes concurrently on the Grace cluster and run longer jobs than may be convenient on your own computer. You are referred to the GitHub link for more documentation. Note that there is also a pure python implementation of the Ising model, which you are also welcome to use. It is anticipated that this would be used primarily only when compiling the C++ library is troublesome.

### 5.1 Practical Introduction to ising.py

You are referred to the github readme file for a listing of the various code files. This subsection gives a little practical advice on how to get started using the code. A full code listing of ising.py is given in Appendix D.

#### Code Requirements

The ising.py requires python3 with the following modules:

- numpy: for various array and matrix functions
- matplotlib
- multiprocessing
- scipy

#### Running the code and input options

Issue the following command on the command line:

python3 ising.py [keyword1]:[value1] [keyword2]:[value2] ... [keywordN]:[valueN]

The keywords and values fill a dictionary which controls various options when running
the code. In ising.py this dictionary is named inp (for "inputs") and the defaults are
set in the function set\_input. The input options and defaults are listed in Table 2.

keyword	default value	use
N	10	ising lattice size = $\mathbb{N} \times \mathbb{N}$
n_steps	10000	number of steps in the MCMC algorithm
n_analyze	5000	num. steps to calc $\langle E \rangle$ , $\sigma_E$ , $\langle M \rangle$ , $\sigma_M$
flip_perc	0.10	ratio of lattice sites that may flip per step
plots	False	make plots at end of run
		note: cannot use with multiprocess
use_cpp	True	use the C++ library
multiprocess	False	use multiple cores concurrently
$skip\_prog\_print$	False	skip printing progress
$\mathtt{print}_{\mathtt{-}}\mathtt{inp}$	False	echo inp to command line
temperature array	options	
note: temperature	array is [t_min,t_max)	with step size of t_step
t_min	1.6	minimum temperature
t_max	3.6	maximum temperature
t_step	0.1	step size between t_min and t_max
		note: progress can only printed without multiprocess
options used in an	nealing	
$t_{-}top$	4.0	tempature from which to anneal
${\tt n\_burnin}$	2000	steps after annearling prior to n_analyze
В	0	constant mag. field
options used for re	esult printout	
$\mathtt{dir}\_\mathtt{out}$	data	name of output directory
$\mathtt{date\_output}$	False	prepend output dir with YYYMMDD-HHMMSS
file_prefix	empty string	prefix for output files
t_min	1.6	minimum temperature
	GRAMM	AR FOR INPUT VALUES
value	command-line input	notes
True	t or true	case insensitive
False	f or false	case insensitive
integer	integer	
float	decimal number	if there's no decimal point, defaults to integer
string	string	default value if none of above

Table 2: Default command line options for ising.py

### Functions in ising.py to Modify

#### • make\_B\_generator(inp, T\_final):

This function makes a python generator that returns a value for the magnetic field each time it is called. That generator's control flows from the top to the first yield. At that statement, it returns the yield value and pauses until it is called again with the next method. At that point it continues from the yield statement until the next yield statement, and so forth. It is called once per step of the Ising Lattice. Therefore, you should use this function to modify the evolution of the magnetic field through all inp['n\_steps'] of the simulation.

#### • make\_T\_generator(inp, T\_final):

This function is completely analogous to make\_B\_generator. Modify it to determine the temperature annealing.

#### • set\_input(cmd\_line\_args):

Modify this function if you wish to add new input parameters that can be modified from the command line. You may find it equally convenient to simply modify the code without new input parameters.

#### • run\_ising\_lattice(inp, T\_final, skip\_print=False):

This is where the actual body of the ising simulation is run. See in particular lines 194 to 222. It is likely that you may want to use some more sophisticated sampling than just averaging the final n\_steps values at the end of a run for determining some parameters/exponents.

#### • plot\_graphs(data):

You will most likely not modify this code. These are *very* simplistic plots just to get you started. You will probably make your plots from the output csv (comma separate variable) files. This has been successfully done with many different software packages, such as Matplotlib, MATLAB, and Excel, to name just a few.

### • print\_results(inp, data, corr):

You probably won't modify this code, except if you want to update the overly simplistic spin correlation output (the corr in the function) to be values averaged over many steps.

#### • remaining functions:

You are welcome, of course, to modify or copy any of the remaining code. This section is just to point out where you will likely spend your first efforts. It is anticipated that you can successfully ignore the remaining code. If this proves wrong, and you are feeling particularly helpful, you can also make pull requests

to the GitHub (https://git.yale.edu/phys382L/ising2D\_code) repository to improve the code.

### Where the code could improve

There are two major points in which the code could be improved. You are welcome to take a stab at either.

- 1. Implement a different algorithmic step for the Ising lattice that will be more resilient against critical slowing. As you will find out in this lab, the parameters you calculate are meaningful at the critical temperature T<sub>c</sub>. However, as you approach the T<sub>c</sub>, the evolution of the Ising Lattice slows down, which is known as "critical slowing down" [6]. There are some promising algorithms, such as the Swendsen-Wang and Wolff algorithms, which could improve results near the critical point. See [7].
- 2. Much simpler than the first point, make the code calculate the spin correlations for many steps and average them for the final values, instead of taking the value at the single final step alone.

# 6 Errata and Errata Reporting

The base files for this document are posted online at https://git.yale.edu/phys382L/ising2D\_handout. Feedback under the "Issues" and, from perhaps more proactive users, "Pull request" sections of the repository constitute the errata for this work. You are invited to provide constructive feedback to the same. For code errors and improvements, please submit likewise to the https://git.yale.edu/phys382L/ising2D\_code repository.

### References

- [1] http://www.scholarpedia.org/article/Ising\_model:\_exact\_results
- [2] D. J. C. MacKay, "Introduction to Monte Carlo Methods", Kluwer Academic Press (1998). Canvas link
- [3] R. K. Pathria and P. D. Beale, "Statistical Mechanics" 3rd ed., Elsevier (2011).
- [4] D. J. C. MacKay, "Information Theory, Inference, and Learning Algorithms", Cambridge Univ. Press (2003).
- [5] T. Paine (1776).

- [6] P.C. Hohenberg and B.I. Halperin, Rev. Mod. Phys. 49 (1977) 435.
- [7] R. H. Swendsen and J.-S. Wang, "Cluster Monte Carlo Algorithms", Physica A 167 (1990) 565-579. science direct link

# Appendix A: Tips for Convergence

As discussed in Section 3, it is important that the system converges at a given temperature before you start using that data to compute thermodynamic averages. Above  $T_c$ , convergence is easy to obtain, and a few thousand burn-in steps should be more than enough to ensure it. At low temperatures, however, the system has far less energy available to move around in the probability landscape, and so its easy for it to get stuck in a local minimum. This will often manifest itself as the development of large magnetic domains, macroscopic regions of your lattice that all have the same spin, but that do not have the same spin as their surroundings. Interestingly, normal magnetic materials have several different magnetic domains on a mesoscopic scale, but we want to avoid them in this experiment. What follows are a few different suggestions of ways to help improve convergence at low temperatures, ranging from the extremely brute force to the slightly-more-clever.

- 1. Its often very easy to check whether your system has converged or not by looking at its magnetization. At low temperatures we expect that all the spins should be roughly aligned, but if there are macroscopic domains present, then the average magnetization will be much closer to zero. A telltale sign for non-convergence is thus an absolute value for average magnetization that is much lower than the data at nearby temperatures would suggest. One very brute force option is thus to test whether your system has converged after each temperature run, and simply rerun it at that temperature if it hasnt. For a reasonably large range of parameters, this works alright because the chances of getting stuck in a local minimum of the Gibbs distribution and failing to converge isnt too high, so you may only need to rerun a few runs. Still, at very low temperatures this becomes extremely time-intensive.
- 2. Changing the percentage of accepted flips that you make each step can sometimes help achieve convergence. If the percentage is too low, the MCMC can only make very small steps in the Gibbs distribution, and so it takes a while for it to get out of a local minimum. If the percentage is too high, each step will be far too large and the system wont be able to park itself at the global minimum once it reaches it.
- 3. Most of the problems obtaining convergence stem from the fact that when you randomly initialize the spin matrix you can initialize it in a local minimum of the Gibbs distribution and at low temperatures the system doesnt have enough

energy available to get out of this initial area its stuck in. One way to combat this is to start the system at a much higher temperature than the one you are trying to measure it at. By starting it at a temperature above the critical point and then cooling it slowly down to the desired temperature you let it have more energy in the beginning to explore the whole probability landscape before slowly finding and settling into the global minimum. This technique is known as *simulated annealing*.

If you are implementing this, be careful not to bias the system! Once it has reached the desired temperature you should let it sit there for several thousand additional burn-in steps before collecting data so you can be sure it isnt still suffering from any of the effects of starting it at a higher temperature.

4. Since macroscopic magnetic domains are a significant problem when trying to obtain convergence, consider biasing your system from the start so one spin (either up or down) is favored, thus discouraging any domains of the other spin from forming. This can be done by subjecting your system to a small external magnetic field. Note though that applying a magnetic field changes the critical temperature and messes with other system state variables, so you should probably do something similar to the simulated annealing technique discussed above: Start your system with a small external magnetic field which you slowly cool to zero (or your actual desired magnetic field if youre doing a run at nonzero field). Then, let the system burn in for several thousand steps before starting to collect data.

This is by no means an exhaustive list, but should give you enough ideas to start attacking any convergence problems you might have.

# Appendix B: Computational Tricks

### Log Plots

Many of the relationships you expect to find throughout this experiment are of the form  $y = ax^b$  where a and b are unknown constants you wish to determine. Fitting to a function of this form is hard (and makes error analysis a pain), so convert it to a linear fit! Note that  $\ln y = \ln a + b \ln x$ , so plotting  $\ln y$  vs  $\ln x$  turns the annoying power-law fit into a trivial linear fit, and has the added benefit of making your error analysis much easier.

# Selecting the right range of data

The various critical exponent relationships summarized in Table 1 only hold near the critical temperature, so when fitting your simulated data to these relationships you should ignore data that is far from the critical temperature. Furthermore, data in the immediate vicinity of the critical temperature will suffer from incredibly large errors and potential finite size effects, so you should exclude a small window of data around the critical temperature when doing all data fits.

### Finding $\gamma$

Magnetic susceptibility typically suffers from very large errors near the critical temperature, making  $\gamma$  one of the harder exponents to get a good lock on. If youre suffering from this problem, try the following:

First recall the definition of magnetic susceptibility,  $\chi = \frac{\partial M}{\partial B}$ , so if we apply a very B small magnetic field (above the critical temperature where the average magnetization would otherwise be zero), we will in general have:

$$M = \chi B \propto |t|^{\gamma} B \propto |t|^{-\gamma} \tag{38}$$

This means that in order to compute  $\gamma$  we can apply a small magnetic field and sweep the temperature above the critical temperature. We can then measure magnetization and fit it to a power-law. Since errors in magnetization are typically smaller than errors in  $\chi$ , this can often help tame errors in  $\gamma$ .

If you are doing this, be sure that the magnetic field you apply is nt too large. Applying a magnetic field actually takes you away from the critical point, meaning even in the  $N\to\infty$  limit there is no divergence in any of the thermodynamic quantities. As long as the field is small this shouldn't give you any large errors in  $\gamma,$  but be careful not to stray into the large-B regime, and dont be too concerned if the critical temperature appears to be shifting around once you apply a magnetic field.

# Appendix C: Figures from Sample Student Data

What follows are a few graphs of sample student data so that you can get a sense for what your results may look like.

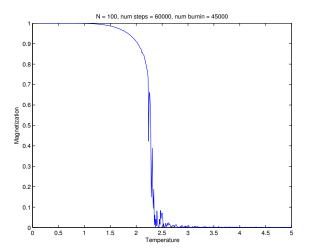


Figure 1: Average magnetization per site vs. temperature for N=100. Below  $T_c$ , the magnetization spontaneously achieves a nonzero value, while above  $T_c$  it is approximately zero (although fluctuations are much larger near  $T_c$  than they are far above it.

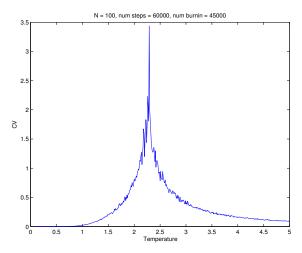


Figure 2: Specific heat vs. temperature for N=100. A divergence near the critical temperature is clearly visible.

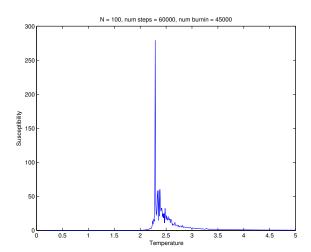


Figure 3: Magnetic susceptibility vs. temperature for N=100. A divergence near the critical temperature is clearly visible.

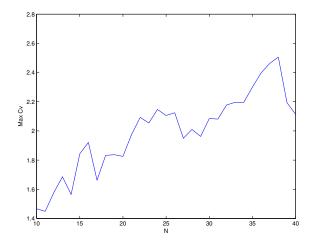


Figure 4: The maximum value of  $c_V$  obtained for a variety of grid sizes. Finite size effects are visible as the clear increase of maximum  $c_V$  with the size of the grid.

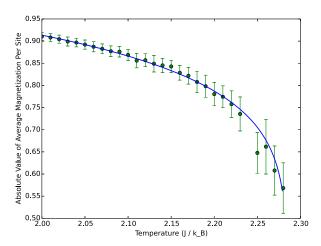


Figure 5: Magnetization vs. temperature for N=100 in a region near the critical temperature. Error bars and a power-law fit are shown; the latter can be used to determine  $\beta$ .

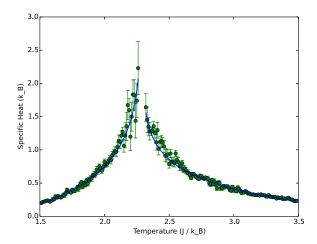


Figure 6: Specific heat vs. temperature for N=100. Error bars and two separate log fits are shown (one each for above and below the critical temperature) to confirm  $\alpha=0$ . Note that a window of data near the critical temperature has been excluded.

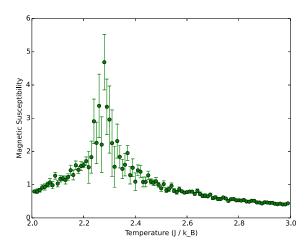


Figure 7: Magnetic susceptibility vs. temperature for N=100. Error bars are shown, but the relatively large errors near  $T_c$  make it difficult to obtain  $\gamma$  from this data.

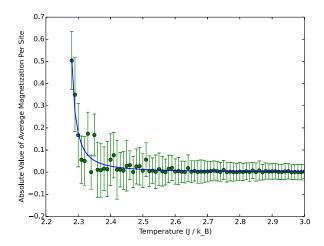


Figure 8: Magnetization vs. temperature for N=100 with an applied field of B=0.1 (in units of J). Error bars are shown, and the fit here was used to obtain  $\gamma$ .

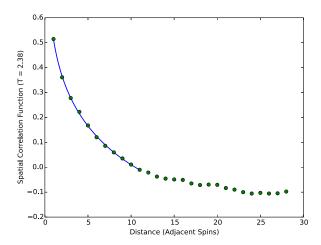


Figure 9: Spatial spin correlation function vs. distance (in units of the lattice spacing) at T=2.38. A fit to the first 11 data points gives a good estimate for  $\xi$  at this temperature, as well as an estimate for  $\nu$ .

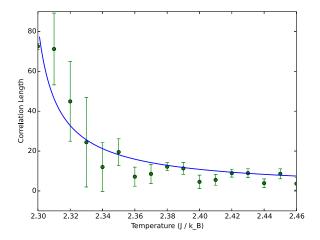


Figure 10: Correlation length  $\xi$  vs. temperature. Error bars are shown, and the fit here was used to obtain  $\nu$ .

# Appendix D: Code Listing of ising.py

```
import sys
2 import os
3 import time
  import csv
5 import logging
6 import numpy as np
7 import matplotlib.pyplot as plt
   import multiprocessing as mp
   from sys import exit, argv
   from IsingLattice_python import IsingLattice as IsingLattice_py
   # conditionally import python and C++ version of IsingLattice
   has\_cpp = False
   if os.path.isfile('ising_lattice_lib.so'):
14
       has\_cpp = True
       from IsingLattice_cpp
                                 import IsingLattice as IsingLattice_cpp
16
17
   ## main program
   def make_B_generator(inp, t_final=None):
       """Return a generator that makes values of B (magetic field in each
20
       note: you *should* play with this function
21
       default implementation: always return 0
23
       for val in range(inp['n_steps']):
           yield inp['B']
27
   def make_T_generator(inp, t_final):
28
       """Return a generator that makes values of T (temperature in each
29
       note: you *should* play with this function
30
       default implementation:
           start at T=t_top, and linearly decrease temperature to t_final
33
           hold at t_final for n_burnin
34
           hold at t_final for n_analyze
35
       n_slope = inp['n_steps'] - inp['n_burnin'] - inp['n_analyze']
       if n_slope < 0:
           print ('fatal_error:_n_steps_-_n_burnin_-_n_slope_<_0')
           print('terminating_program')
           exit(2)
41
42
       # get linearly decreasing values from t_top to t_final
43
44
       for val in np.linspace(start=inp|'t_top'|, stop=t_final, num=
          n_slope):
           yield val
```

```
46
       for val in range(inp['n_burnin']):
47
           yield t_final
48
49
       for val in range(inp['n_analyze']):
50
           yield t_final
   def set_input(cmd_line_args):
       """Parse command-line parameters into an input dictionary.
54
       provide default values
56
       note: feel free to add your own input parameters here,
57
             and them use them by calling them from the dictionary
58
             names 'inp' in the modules where they would prove useful
       input:
                 sys.argv
61
                 use syntax of keyword: value on command line
62
                 dict [key] = value
       return:
63
                 Any value which can be turned into a number will be a
       note:
           float
                 if it has a '.', otherwise it will be a float.
65
       22 22 22
67
68
       inp = dict()
       inp['t_min']
                          = 1.6
                                   # minimum temperature
70
       inp['t_max']
                          = 3.6
                                   # maximum temperature
       inp['t_step']
                          = 0.1
                                   # step size from min to max temperature
72
       inp['t_top']
                          = 4.0
                                   # start temperature (arbitrary; feel
           free to change)
       inp['N']
                                   \# sqrt(lattice size) (i.e. lattice = N^2
                          = 10
            points
       inp['n_steps']
                          = 10000
                                   # number of lattice steps in simulation
75
       inp['n_burnin']
                          = 2000
                                   # optional parameter, used as naive
           default
       inp['n_analyze'] = 5000
                                   # number of lattice steps at end of
           simulation calculated for averages and std.dev.
       # inp['J']
                            = 1.0
                                     # **great ** default value -- spin-spin
            interaction strength
       inp['B']
                          = 0.0
                                   # magnetic field strength
                          = 0.1
       inp['flip_perc']
                                   # ratio of sites examined to flip in
80
           each step
       inp['dir_out']
                          = 'data' # output directory for file output
       inp['plots']
                          = False
                                   # whether or not plots are generated
       inp['print_inp'] = False # temperature option
       inp['use_cpp']
                          = True
                                   # use 1 for True and 0 for False
85
86
       inp['date_output'] = False
87
       inp['file_prefix'] = '
       inp['multiprocess'] = False
89
```

```
inp['skip_prog_print'] = False
90
91
        for x in cmd_line_args[1:]:
92
            if ':' in x:
                try:
94
                     key, val = x.split(':')
                     try:
                         if '.' in val:
                              inp[key] = float(val)
98
                              print('%-20s'%('inp["%s"]'%key), 'set_to_float__
99
                                   , inp[key])
                          elif val.lower() = 'false' or val.lower() = 'f':
100
                              inp[key] = False
101
                          elif val.lower() == 'true' or val.lower() == 't':
                              inp[key] = True
                         else:
104
                              inp[key] = int(val)
                              print ('%-20s'%('inp["%s"]'%key), 'set_to_int____
106
                                  ', inp [key])
                     except:
107
                         inp[key] = val
108
                         print ('%-20s'%('inp["%s"]'%key), 'set_to_string_',
109
                             inp[key])
                 except:
                     print ('warning: _input_"%s" _not_added_to_arguments'%x)
            else:
112
                 print ('ignoring_command_line_input: _%s'%x)
113
114
        if inp['print_inp']:
            print('Printed_list_of_input_keys:')
            for key in sorted (inp.keys()):
117
                print('%-20s'%key,',',inp[key])
118
        return inp
119
120
    class check_progress(object):
121
        """A class that will print a simple status bar of percentage of
           work done"""
        def __init__(self , inp , T, skip_progress=False):
123
            self.skip_print = skip_progress
124
            if self.skip_print:
126
                 return
127
            self.start_time = time.time()
128
            self.n_steps = inp['n_steps']
            self.n_check = 1000
               'check_per_steps' in inp:
                 self.n_check = inp['check_per_steps']
            self.fmt_print = ('%ix%i_(T=%.2f)_steps:_%%7i/%7i,_%%5.1f%%%__
                     'run_time: \_\%8s_\_est.time-to-go: \_\%8s'\%
134
                     (inp['N'], inp['N'], T, self.n_steps))
135
```

```
self.n_called = -1 \# will progress untill n_called = n_steps
136
            self.check()
137
           # print(self.fmt_print)
138
139
             if False:
140
           #
                  print ('%ix%i IsingLattice. Finished %10i / %10i steps
141
               (\%4.1 \, \text{f}\%\%).
                       'Started '+str(time.strftime('%m-%d %H:%M:%S')))
                # dir_out += str(time.strftime("_%Y%n%d-%H%M%S"))
143
        def check(self, final=False):
144
            if self.skip_print:
145
                return
146
147
            self.n_called += 1
            if not self.n_called \% 1000 = 0 and not final:
150
                return
            ratio = float (self.n_called) / self.n_steps
            if ratio == 0:
                time_pass_str = '00:00:00'
154
                est_str = 'n/a'
            else:
                time_pass = time.time() - self.start_time
                time_pass_str = time.strftime('%H:%M:%S', time.gmtime(
158
                    time_pass));
                est\_time = (1-ratio)*time\_pass/ratio
159
                # print('est_time ', est_time)
                est_str = time.strftime('%H:%M:%S', time.gmtime(est_time))
161
            if final:
162
                print (self.fmt_print%(self.n_called, ratio*100.,
                    time_pass_str, 'done!'))
            else:
164
                print(self.fmt_print%(self.n_called, ratio*100.,
165
                    time_pass_str, est_str), end='\r')
166
   def run_ising_lattice(inp, T_final, skip_print=False):
167
        ''', 'Run a 2-D Ising model on a lattice.
168
        Return three objects (each a numpy.array, which you can treat
        as identical to a numpy.array)
170
            M_{avg}:
                         the average magnetization of each site for each
               n_analyze step
                          '..... ' energy
            E_{avg}:
                 correlation: an (N/2-1) array of the correlation function
               values at the final data frame
            Note that this will use the generators from functions:
175
            (1) make_T_generator
176
            (2) make_B_generator
177
178
179
```

```
time_start = time.time()
180
181
        lattice = None
182
        if inp['use_cpp'] and has_cpp:
183
             lattice = IsingLattice_cpp(inp['N'], inp['flip_perc'])
184
        elif inp['use_cpp'] and not has_cpp:
185
             print ('Warning: _although _use_cpp _is _set _to _1, _'
              _the_shared_library_IsingLattice.so_is_not_present.\n_'
             \verb|`_Therefore , \_the \_python \_implementation \_of \_Ising \_will \_be \_used. | |
             lattice = IsingLattice_py(inp['N'],inp['flip_perc'])
189
        else:
190
             lattice = IsingLattice_py(inp['N'],inp['flip_perc'])
191
        # Make the run loop
193
        try: # try loop that can be interrupted by the user
             #first loop through all steps up to n_analyze
195
             T_{generator} = make_{T_{generator}}(inp, T_{final})
196
             B_{generator} = make_{B_{generator}}(inp, T_{final})
197
             n_prior = inp['n_steps'] - inp['n_analyze']
198
199
             progress = check_progress(inp, T_final, skip_print)
200
             for T, B, step in zip(T_generator, B_generator, range(n_prior))
                 lattice.step(T,B)
203
                 progress.check()
204
            # loop through the analyze section of generators
206
             E_{avg} = []
207
             M_avg = []
             for T, B, step in zip (T_generator, B_generator, range (inp [ '
209
                n_analyze'])):
                 lattice.step(T,B)
210
                 E_avg.append(lattice.get_E())
211
                 M_avg.append(lattice.get_M())
212
                 progress.check()
213
             progress.check(True)
214
             spin_correlation = np.array(lattice.calc_auto_correlation())
216
             lattice.free_memory()
217
             return (
218
                 np. array (E_avg),
                 np. array (M_avg),
220
                 np.array(spin_correlation)
221
             )
        except KeyboardInterrupt:
224
             try:
225
                 lattice.free_memory()
226
             except:
227
                 pass
228
```

```
print ("\n\nProgram_terminated_by_keyboard._Good_Bye!")
229
             sys.exit()
230
231
    def plot_graphs(data): #T, E_mean, E_std, M_mean, M_std): #plot_graphs at
232
        dat = np.array(data)
233
        # print('data: ', dat)
        # print('x: ', dat[:,1])
        \# x = dat[:,1][0]
236
        # print('xlist: ', x)
237
238
        plt.figure(1)
239
        \# plt.ylim (0,1)
240
        plt.errorbar(dat[:,0], dat[:,1], yerr=dat[:,2], fmt='o')
        plt.xlabel('Temperature')
plt.ylabel('Average_Site_Energy')
243
        plt.figure(2)
244
        plt.errorbar(dat[:,0], np.absolute(dat[:,3]), yerr=dat[:,4], uplims
245
            =True, lolims=True, fmt='o')
        plt.xlabel('Temperature')
246
        plt.ylabel('Aveage_Site_Magnetization')
247
        plt.show()
    def get_filenames(inp): #make data folder if doesn't exist, then
250
       specify filename
         '''Generate the output file names for the EM (energy and megnetism)
251
             and SC (spin correlation) files '''
        try:
252
             dir_out = inp['dir_out']
             prefix = inp['file_prefix']
             if inp['date_output']:
                 dir_out += str(time.strftime("_%Y%m%d-%H%M%S"))
256
257
             if not os.path.isdir(dir_out):
258
                 os.makedirs(dir_out)
259
260
            # file name = [file_prefix]##_EM_v#.csv if only one temperature
261
                 (example: runA_4.20_EM_v0.csv)
                             [file_prefix]##T##_EM_v#.csv if there are two
262
                temperatures (example: runA_4.2T5.3_EM_v0.csv)
            # the other file name is identical, but with "SC" (for spin
263
                correlation)) instead of EM
             if inp['t_max'] <= inp['t_min']:
264
                 t_name = \%.2 f \% inp ['t_min']
             else:
                 t_name = '%.2fT%.2f'%(inp['t_min'],inp['t_max'])
268
            # print('%s%s_SC_v%i.csv'%(prefix,t_name,v))
269
             \mathbf{v} = 0
270
             while (os.path.isfile (os.path.join(dir_out, '%s%s_EM_v%i.csv'
271
                \%(\text{prefix}, t_{\text{name}}, v))) or
```

```
os.path.isfile(os.path.join(dir_out, '%s%s_SC_v%i.csv'
272
                       \%(\text{prefix}, t_{\text{-}}\text{name}, v)))):
                 v += 1
273
274
            return (os.path.join(dir_out, '%s%s_EM_v%i.csv'%(prefix,t_name
                ,v)) ,
                      os.path.join(dir_out, '%s%s_SC_v%i.csv'%(prefix,t_name
                          (v)
277
        except:
278
            print ('fatal: _Failed_to_make_output_file_names')
279
280
            sys.exit()
   def print_results(inp, data, corr):
        data_filename, corr_filename = get_filenames(inp)
        with open(data_filename, 'w') as f_out:
284
            writer = csv.writer(f_out, delimiter=',', lineterminator='\n')
285
            writer.writerow(['N', 'n_steps', 'n_analyze', 'flip_perc'])
286
            writer.writerow([inp['N'], inp['n_steps'], inp['n_analyze'],
                inp['flip_perc']]
            writer.writerow([])
            writer.writerow(['Temp', 'E_mean', 'E_std', 'M_mean', 'M_std'])
            for entry in data:
                 writer.writerow(entry)
291
            # for t, e_mean, e_std, m_mean, m_std in zip(T, E_mean, E_std,
292
                M_{-mean}, M_{-std}):
                   writer.writerow([t, e_mean, e_std, m_mean, m_std])
294
        with open(corr_filename, 'w') as f_out:
            writer = csv.writer(f_out, delimiter=',', lineterminator='\n')
            writer.writerow(['N', 'n_steps', 'n_analyze', 'flip_perc'])
297
            writer.writerow([inp['N'], inp['n_steps'], inp['n_analyze'],
298
                inp['flip_perc']])
            writer.writerow([])
299
            writer.writerow(['Temp']+['d=%i'%i for i in range(1,len(corr
300
                [0])+1)])
            for entry in corr:
301
                 writer.writerow(entry)
303
304
   def run_indexed_process( inp, T, data_listener):
305
   # def run_simulation(
306
   #
              temp, n, num_steps, num_burnin, num_analysis, flip_prop, j, b
307
        data_filename, corr_filename, data_listener, corr_listener):
        print ("Starting \_Temp\_{0}". format (round (T,3)))
        try:
            E, M, C = run_ising_lattice(inp, T, skip_print=True)
310
            data_listener.put(([T,E.mean(),E.std(), M.mean(), M.std()], [T
311
                ,]+[x[1] \text{ for } x \text{ in } C])
            # corr_listener.put([T,]+[x[1]] for x in C])
            print ("Finished \_Temp\_{0}". format (round (T,3)))
313
```

```
return True
314
315
         except KeyboardInterrupt:
316
              print ("\n\nProgram_Terminated._Good_Bye!")
317
              data_listener.put('kill')
318
             # corr_listener.put('kill')
             sys.exit()
         except:
322
              \log \operatorname{ging.error}(\operatorname{"Temp="+str}(\operatorname{round}(\operatorname{T},3)) + \operatorname{":\_Simulation\_Failed.\_No\_})
323
                 Data_Written")
              return False
324
325
    def listener (queue, inp, data):
326
         ''', listen for messages on the queue
         appends messages to data ','
328
        \# f = open(fn, 'a')
329
        # writer = csv.writer(f, delimiter=',', lineterminator='\n')
330
         while True:
331
             message = queue.get()
332
             # print('message: ', message)
333
             if\ message == \ 'kill':
                  data['data'].sort()
                  data['corr'].sort()
336
                  print_results(inp, data['data'], data['corr'])
337
                  print ('Closing_listener')
338
                  # print('killing')
                  break
340
              data ['data']. append (message [0])
341
              data['corr'].append(message[1])
             # print('--
                             ----\n ' , data )
344
    def make_T_array(inp):
345
         if inp['t_max'] \le inp['t_min']:
346
              return [inp['t_min'],]
347
         else:
348
              return np.arange(inp['t_min'], inp['t_max'], inp['t_step'])
349
350
351
    def run_multi_core(inp):
352
         print ("\n2D_Ising_Model_Simulation; _multi-core\n")
353
         T_{array} = make_{T_{array}}(inp)
354
355
        #must use Manager queue here, or will not work
356
         manager = mp. Manager()
         data_listener = manager.Queue()
        # corr_listener = manager.Queue()
359
         pool = mp. Pool (mp. cpu_count ())
360
361
        # arrays of results:
363
```

```
data = { 'data ':[], 'corr':[]}
364
        \# \operatorname{corr} = []
365
366
        #put listener to work first
367
        data_watcher = pool.apply_async(listener, args=(data_listener, inp,
368
             data,))
        # corr_watcher = pool.apply_async(listener, args=(corr_listener,
            inp, corr,))
        #fire off workers
371
        jobs = [pool.apply_async(run_indexed_process, args=(inp,T,
            data_listener)) for T in T_array]
373
        # collect results from the workers through the pool result queue
        [job.get() for job in jobs]
        data_listener.put('kill')
376
        pool.close()
377
378
    def run_single_core(inp):
379
        print ("\n2D_Ising_Model_Simulation; _single_core\n")
380
        # sequentially run through the desired temperatures and collect the
381
             output for each temperature
        data = []
        corr =
383
        for temp in make_T_array(inp):
384
            E, M, C = run_ising_lattice(inp, temp, skip_print=inp['
385
                skip_prog_print'])
            data.append( (temp, E.mean(), E.std(), M.mean(), M.std() )
386
            corr.append([temp,]+[x[1] for x in C])
        print_results(inp, data, corr)
390
        if inp['plots']:
391
            plot_graphs (data)
392
393
394
      __name__ = "__main__":
395
        """Main program: run Ising Lattice here"""
        inp = set_input(argv)
397
        if inp['multiprocess']:
398
            run_multi_core(inp)
399
        if not inp['multiprocess']:
            run_single_core(inp)
401
```