Monte Carlo simulation of the effects of higher-order anisotropy on the spin reorientation transition in the two-dimensional Heisenberg model with long-range interactions

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The strength of perpendicular anisotropy is known to drive the spin reorientation in thin magnetic films. Here, we consider the effect different order anisotropies have on two phase transitions: the spin reorientation and the orientational order transitions. We find that the relative magnitude of different order anisotropies can significantly enhance or suppress the degree to which the system reorients. Specifically, Monte Carlo simulations reveal significant changes in the cone angle and planar magnetization. In order to facilitate rapid computation, we have developed a stream processing technique, suitable for use on graphics processing unit (GPU) systems, for computing the transition probabilities in two-dimensional systems with dipole interactions.

I. INTRODUCTION

In two dimensions, the effect of thermal fluctuations is enhanced. The number of possible symmetries is lower than in three-dimensional systems, and this reduced symmetry means that there are fewer degrees of freedom to absorb energy.[1–3] The discovery of a divergence in the susceptibility in the *XY* model,[4] caused by topological excitations,[5,6] meant that the existence and stability of spontaneous ordered states in two dimensions has been a rich and often contentious[3,4,7–15] area of theoretical interest. In particular, the dimensionality of the field,[15,16] finite size effects,[17] and anisotropies[18] can effect the phase diagram.

Two-dimensional systems can be realized experimentally as thin (typically <15 atomic layer) magnetic films. By varying the composition and thickness of thin films, a large variety of magnetic properties have been obtained.[19] It is possible to create films that strongly favor either in-plane or perpendicular orientation of the magnetization.[20] The functional dependence of energy on perpendicular magnetization varies and can be altered using ion beam irradiation.[21,22] In the presence of strong uniaxial anisotropy favoring perpendicular alignment, the competition between the entropy favored in-plane magnetization and the energetically favored out of plane state can lead to a temperature driven spin reorientation transition (SRT).[23–25] The nature of this transition is known to be dependent on the relative strengths of different orders of anisotropy.[26–28] In these systems, the ratio of long-range dipole coupling and short-range exchange coupling can lead to the formation of striped domains of alternating spin direction; either as the ground state[29,30] or as a mechanism of spin reversal.[31] Films have been produced with stripes observed running parallel with a common orientation,[30–32] as zigzags between regular defects[33] or forming complex patterns with no orientational order.[30,34] The width and mobility of stripes depend on temperature.[35] In particular, these systems can display strong thermal memory in which the domain configuration depends on the rate of heating or cooling.[36,37] Both the reorientation and stripe melting transitions have been studied analytically[38–41] and by using Monte Carlo simulation.[42,43]

Here, we propose a technique in which energy differences arising from long-range dipole coupling are approximated.

This approximation allows for computation to be parallelized significantly; reducing the computational time required. Having first examined the extent to which this approximation influences the results of simulation, the method is applied to a two-dimensional Heisenberg model where the effects of higher-order anisotropy are examined.

II. THEORY

A. Dipole coupling

Consider a thin ferromagnet, modeled as a two-dimensional square lattice of Heisenberg spins ($\vec{s} \in \mathcal{S}^2$). The spins experience a long-range dipole interaction,

$$H_D = \frac{C_D}{2} \sum_{n,m} \frac{1}{r_{nm}^3} (\vec{s}_n \cdot \vec{s}_m - 3\vec{s}_n \cdot \hat{r}_{nm} \vec{s}_m \cdot \hat{r}_{nm}), \quad (1)$$

where n and m represent vertices of the two-dimensional lattice, $\vec{r}_{nm}$ is the vector from n to m, and C_D is a constant, $C_D = (M^2 \mu_0)/(4\pi)$. In simulations, periodic boundary conditions are used to approximate an infinite system. The total system consists of a tiling of replicas. For simulation size $L \times L$, spins separated by a vector $\vec{G} = (aL, bL)$ with $a, b \in \mathcal{Z}$ being identical. To compute the infinite sum introduced by periodic boundary conditions, a new set of coordinates is introduced: $\vec{r}_{nm} = \vec{G} + \vec{\rho}_{nm}$. Here, $\vec{\rho}$ is restricted to $\vec{\rho} = (\rho_x, \rho_y)$ with $\rho_x, \rho_y \in [0, L]$. The dipole energy at a site n can then be written (taking the square lattice to be in the x-y plane):

$$H_n = \frac{1}{2} C_D \sum_{m \neq n} \sum_{\vec{G}} s_n^\alpha s_m^\beta \lim_{r \to 0} \partial_\alpha \partial_\beta \frac{1}{|\vec{\rho}_{nm} + \vec{G} - \vec{r}|}$$
$$+ C_D (s_n^\alpha)^2 \lim_{r \to 0} \partial_\alpha^2 \sum_{\vec{G} \neq 0} \frac{1}{|\vec{G} - \vec{r}|}, \quad (2)$$

where s_n^α represents $\vec{s}_n . \hat{\alpha}$ and expressions are summed over repeated greek indices. In order to achieve efficient computation, $\sum_{\vec{G}} \lim_{r \to 0} \partial_\alpha \partial_\beta \frac{1}{|\vec{\rho}_{nm} + \vec{G} - \vec{r}|}$ can be calculated in advance for all choices of n, m, α, and β. Since the sum is slow to converge, it can be split into a short-range real-space term and a long-range Fourier-space term according to the technique described by Harris[44] based on the analogous three-dimensional case

developed by Ewald.[45] Letting $f(\vec{r}) = \sum_{\vec{G}} \frac{1}{|-\vec{\rho}_{nm}+\vec{G}+\vec{r}|}$, one has $f = f_L + f_S$ with[46]

$$f_S(\vec{r}) = \sum_{\vec{G}} \frac{1}{|\vec{r} - \vec{\rho}_{nm} + \vec{G}|} \text{erfc}\left(\frac{|\vec{r} - \vec{\rho}_{nm} + \vec{G}|}{2\eta}\right) \quad (3)$$

and

$$f_L(\vec{r}) = 1/L^2 \sum_{\vec{k}} \tilde{h}^L(\vec{k},z) \exp(i2\pi\vec{k}\cdot\vec{r}), \quad (4)$$

where

$$\tilde{h}^L(\vec{k},z) = \frac{\pi}{k} e^{-i\vec{k}\cdot\vec{\rho}_{nm}} \left[e^{k|z-z_{nm}|} \text{erfc}\left(\frac{|z-z_{nm}|}{2\eta} + k\eta\right) \right.$$
$$\left. + e^{-k|z-z_{nm}|} \text{erfc}\left(\frac{-|z-z_{nm}|}{2\eta} + k\eta\right) \right]. \quad (5)$$

Despite the efficiency gained by precalculating the interactions and using this rapid summation technique, the calculation of the dipole interaction is still computationally intensive. In order to calculate the energy of a single spin in the system, one must calculate $N = L^2$ interactions, when calculating the energy of a state of the system C_2^N interactions are required. For a moderate system size $L = 64$, this equates to 4096 interactions for a single spin and over 8×10^6 for a single state.

B. Monte Carlo simulation

When considering a system at finite temperature, observable quantities O are calculated as expectation values of the Boltzmann distribution:

$$\langle O \rangle = \frac{\sum_i \hat{O}[\phi_i] \exp\left(\frac{-H_i}{k_B T}\right)}{Z}. \quad (6)$$

In order to approximate the properties of this distribution, a subset of possible states is selected using a Markov chain Monte Carlo method with transition function

$$P_T(\phi_j \to \phi_k) = \begin{cases} \exp\left[\frac{-(H_k-H_j)}{k_B T}\right] & \text{for} \quad (H_k - H_j) > 0, \\ 1 & \text{otherwise.} \end{cases} \quad (7)$$

Known as the Metropolis algorithm,[47] Eq. (7) does not define a method for selecting the prospective new state. There are numerous methods for constructing new states and the decision is based largely on the system being analyzed. In the case of magnetic systems, the simplest and most common choice is single spin flips. For a system of size N, one Monte Carlo step (MC step) requires N spin flips. Herein lies the computational difficulty, in order to complete one MC step in a two-dimensional system, one must calculate the energy of a single spin $N = L^2$ times. If there is dipole coupling present, each energy calculation requires L^2 interactions to be computed. In order to compute MC steps more efficiently, we present a stream processing algorithm in Sec. III that reduces this computational load.

III. GPU PARALLEL PROGRAMMING AND SIMULTANEOUS FLIPPING

In order to perform Monte Carlo simulations at an acceptable speed, we make use of a graphics processing unit (GPU). Most modern computing systems implement a single instruction single data execution model. In contrast, some systems implement a single instruction multiple data execution model, sometimes referred to as stream processing. In stream processing, a single function (kernel) is executed simultaneously on a large number of different inputs (the stream), the execution of each input (thread) is independent and there is no communication between threads.[48] Here, we present a stream processing algorithm to perform rapid Monte Carlo simulations exploiting the parallelism available in GPU cards. While our implementation makes use of GPUs, it is not entirely accurate to say that GPU chips implement a steam processing architecture. On modern GPUs, each multiprocessor has a small level one cache (typically tens of kilobytes) and a larger coherent level two cache (typically several hundred kilobytes). It is possible to pause execution of identical threads until all threads have reached a designated execution point, allowing one to avoid the race errors normally associated with different parallel threads reading and writing from the same memory location. In this sense, the GPU chips are capable of limited communication between threads.[49] For spin lattice models where interactions are limited to nearest or next-nearest neighbor, the problem of implementing a parallel GPU algorithm has been examined previously and several algorithms exist to distribute computation over single[50–52] or multiple GPUs.[53] In these cases, multiple single spin flips can be performed simultaneously (provided potential update sites are not nearest neighbors). In recent work, Campos *et al.*[54] have approached the problem of long-range coupling by parallelizing the long-range sum in three dimensions. Here, we go beyond the work of Campos *et al.* and focus on performing multiple simultaneous MC steps.

A. Algorithm

Here, we describe a method for parallelizing MC simulations in the presence of dipole coupling. A pseudocode implementation of the algorithm is given in Appendix B. The algorithm depends on the size of the system L and two parameters that will be defined below: l and P. For clarity of exposition, figures in this section will use a fixed small system size $L = 8$ and the parameters $l = 4$ and $P = 4$ (l and P need not be equal in general). When describing the algorithm we take the "host" to indicate any computation not performed on the GPU. Calculations run on host are implemented in the normal serial fashion. We will refer to the GPU card as the "device" and calculations run on device are implemented as parallel operations and have access to device memory (VRAM).

Initially, the current state of the system is held in the device memory, either from initialization or from the previous iteration of the algorithm. On the host, a site is selected at random, in Fig. 1, this site is denoted by a blue circle numbered 1. Additional sites are then selected at fixed multiples of l according to $i = (al,bl) + i_1$ for $a,b \in [0,L/l]$. These $n = (L/l)^2$ values of i are the update sites and are represented in Fig. 1 as circles numbered 2, 3, and 4. Next for each site, a new spin value is selected at random as the potential new spin values. In Figs. 1, 3, and 4, these potential new spin values are represented as diamonds. The location of the selected sites and the potential new spin values are copied to the device memory.
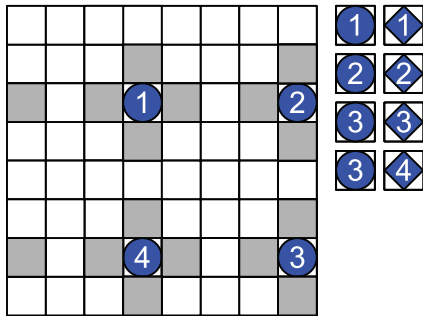
FIG. 1. (Color online) An 8×8 sample where in each square represents a spin site. Site 1 is chosen at random. Update sites 2–4 are selected at fixed distances from 1. In addition, four alternative spin values are also generated (indicated here as diamonds) giving a total of $(2L^2)/l^2 = 8$ energies that need to be calculated. Gray squares indicate the nearest-neighbor sites used in the calculation of short-range interactions.

The device then launches $2n$ threads to calculate nearest-neighbor exchange coupling.[55] The nearest-neighbor spins accessed by these threads are indicated as gray boxes in Fig. 1. Simultaneously, the system launches $2Pn$ threads to calculate the dipole interactions. Each thread calculates the interaction between a spin at an update site and spins in a vertical sub-section of the total system with width L/P. Figure 2 presents a subset of these threads that calculate the interactions associated with a single update site. When the above threads are complete, all interactions have been computed. For each of the n potential update sites, the results of both the current
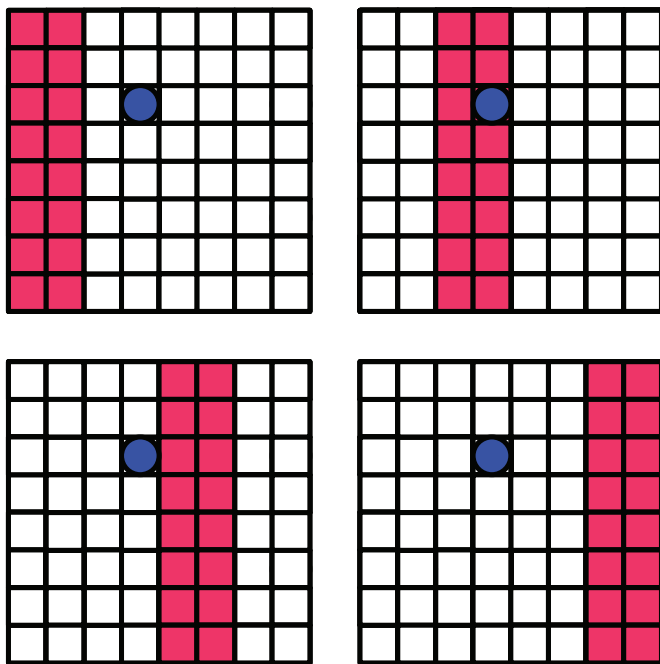

FIG. 2. (Color online) The dipole interactions between update site 1 (blue circle) and the rest of the system are parallelized into four threads, which compute a subset of the possible interactions shown here as pink shaded squares. Simultaneously, threads will be executing the calculation of dipole coupling for the other seven spin values.
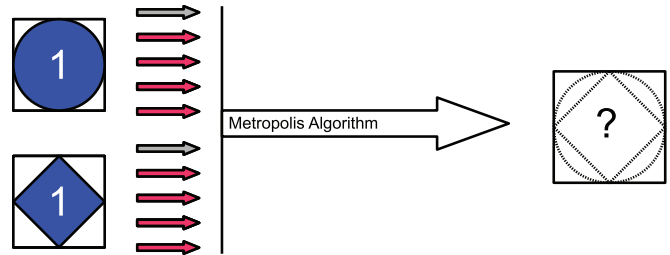

FIG. 3. (Color online) One of the n simultaneous spin flipping threads. The original spin value and the alternative spin value for site 1 each require five threads to compute all interactions. One thread to calculate the short-range interactions with gray squares in Fig. 1 (represented by a gray arrow) and $P = 4$ threads to calculate the dipole interactions with the subsystems shown as pink squares in Fig. 2 (represented by pink arrows). The results of the ten threads are then fed into a single thread that calculates the flipping probability for update site 1.

and the potential new spins are passed into a single thread. Each of the n new threads calculates any single-site energies (anisotropies and applied fields), then applies the metropolis algorithm and updates the state accordingly (see Fig. 3). In Fig. 4, all the threads executed in one iteration of the algorithm for the hypothetical small system are displayed.

B. Approximation

The algorithm presented here reduces the computation required by simultaneously executing $P \times (L/l)$ partial sums of size L/P. However, in doing so, an approximation has been made. In Fig. 5, a single thread will compute the dipole interaction between a potential update site (blue circles in the figure) and a subset of the sample of width L/P (pink). Some subsets will contain another potential update site (the interactions of which are being processed simultaneously another thread). If two spin updates were computed without parallelization, there are four possibilities for the new state ϕ_i given in Fig. 6: neither spin is flipped (ϕ_0), the first spin
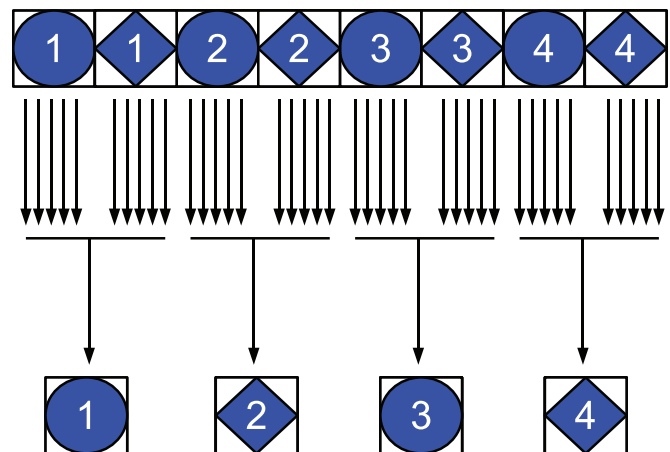

FIG. 4. (Color online) For each of the four current spins and each potential new spin, the five threads shown in Fig. 3 are computed and the results are fed into $n = 4$ threads that calculate the flipping probability. One of the possible 2^n possible resulting spin updates is shown.
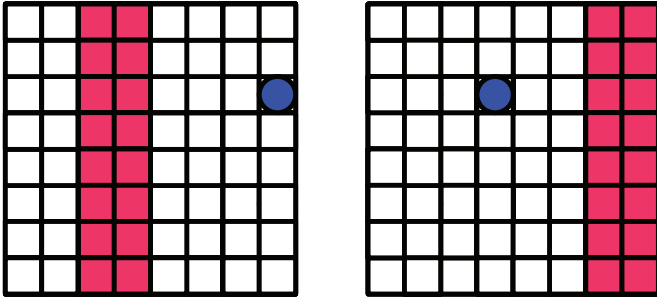
FIG. 5. (Color online) The interactions calculated by two of simultaneously executed threads. On the left, site 2 interacts with a section of the system that contains site 1. On the right, site 1 interacts with a section of the system that contains site 2.

flips (ϕ_1), the second spin flips (ϕ_2), or both spins flip (ϕ_3). Denote the probability that the first spin flips as P_1, the probability that the second spin flips given the first spin has flipped as P_{12}, and that the second flips given the first spin is not flipped as $P_{\bar{1}2}$. Then the probability that the system finishes in state ϕ_0, is $\mathcal{P}_0 = \neg P_1 \neg P_{\bar{1}2}$, where we have used $\neg P_1$ to denote the probability that P_1 does not occur $[\neg P_1 = (1 - P_1)]$. Similarly, $\mathcal{P}_1 = P_1 \neg P_{12}$, $\mathcal{P}_2 = \neg P_1 P_{\bar{1}2}$, and $\mathcal{P}_3 = \neg P_1 P_{12}$. If the two spins do not interact, then $P_{12} = P_{\bar{1}2}$ and depends only on the energies H_0 and H_2. In this case, simultaneously updating sites gives the same statistics as the conventional sequential flipping. If the spins do interact, as is the case with long-range coupling, then the above algorithm makes the approximation $P_{12} = P_T(\phi_1 \rightarrow \phi_3) \approx P_T(\phi_0 \rightarrow \phi_2)$ or $H_3 - H_1 \approx H_2 - H_0$.

In order to approximate, the size of this error first denote original spin values as $\vec{s}_1$ and $\vec{s}_2$ and the potential new values as $\vec{S}_1$ and $\vec{S}_2$. Then $\phi_0 = (\vec{s}_1, \vec{s}_2)$, $\phi_1 = (\vec{S}_1, \vec{s}_2)$, $\phi_2 = (\vec{s}_1, \vec{S}_2)$, and $\phi_3 = (\vec{S}_1, \vec{S}_2)$. Assuming that spins are not nearest-neighbors, then the error $\epsilon_{12} = |H_3 - H_1 - H_2 + H_0|$ depends only on the dipole energy. The error is

$$\epsilon_{12} = \frac{C_D}{r_{12}^3} |\delta_1 \cdot \delta_2 - 3(\delta_1 \cdot \hat{r}_{12})(\delta_2 \cdot \hat{r}_{12})| = \frac{C_D}{r_{12}^3} \mathcal{D}, \quad (8)$$

where $\delta_1 = \vec{s}_1 - \vec{S}_1$, $\delta_2 = \vec{s}_2 - \vec{S}_2$, and $\mathcal{D} = |\delta_1 \cdot \delta_2 - 3(\delta_1 \cdot \hat{r}_{12})(\delta_2 \cdot \hat{r}_{12})|$. We wish to find the maximal values of $\mathcal{D}$. There
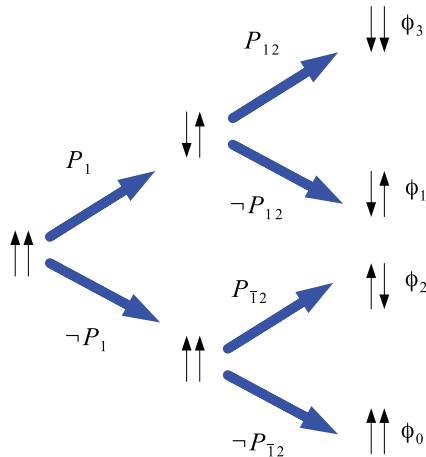

FIG. 6. (Color online) The four possible states that result from two potential spin flips and the associated probabilities.

are two cases in which local maxima occur. The first case, which we shall refer to as the perpendicular case, is $\delta_1 = \pm \delta_2 = \pm(0,0,2)$ corresponding to $\mathcal{D} = 2$. The second case, which we shall refer to as the planar case, is $\delta_1 = \pm \delta_2 = \pm 2\hat{r}_{12}$ corresponding to $\mathcal{D} = 4$.

Here, we have calculated the maximum error introduced by flipping two spins simultaneously. However, due to the periodic boundaries conditions described in Sec. II A, flipping a spin also flips its image in each replica that makes up the infinite system. For the perpendicular case, each replica introduces an additional error $\epsilon_R = (2C_D)/(|\vec{G} + \vec{r}_{12}|^3)$. For the in-plane case, the maximum $\mathcal{D} = 4$ requires δ_1 and δ_2 to be parallel to the vector connecting the updated spins, $\vec{r}_{12} + \vec{G}$, which will not be possible for all $\vec{G}$. To estimate the error for the in-plane case, let

$$\frac{\vec{r}_{12} + \vec{G}}{|\vec{r}_{12} + \vec{G}|} = (\cos(\gamma), \sin(\gamma), 0) \quad (9)$$

then fixing the $\delta_1 \parallel \delta_i$ gives $\mathcal{D} = 1 - 3\cos(\gamma)^2$. Since simultaneously flipped spins and their replicas will exist in all possible directions, we take the root-mean-square average with respect to γ giving $\mathcal{D} \approx 2$. Based on this argument, including all simulated spins, we estimate the maximum error in $H_3 - H_1$ as

$$\epsilon_{\text{Total}} = \frac{L^2}{l^2} \sum_{i=-n}^{n} \sum_{j=-n}^{n} \sum_{G} \frac{C_D}{[(il + G^x)^2 + (jl + G^y)^2]^{\frac{3}{2}}}. \quad (10)$$

The flipping probability will now be $\exp[\frac{-(H_k - H_j)}{k_B T} \pm \frac{\epsilon_{\text{Total}}}{k_B T}]$. This error places an upper bound on the error for a iteration of the algorithm spin flip, however, it does not preclude the possibility of large errors accumulated over the multiple spin flips that will be performed in simulation, typical simulations may involve tens of millions of single spin flips. Notably, the error depends on the value of l and selection of an appropriate value should try to minimize the error as much as possible, however, large values of l reduce the parallelism. In the work below, we consider the system size $L = 64$, restricting the choice of l to 2, 4, 8, 16, or 32.[56] In order to keep the error in flipping probability below 1% over the range of temperatures considered here, we select $l = 32$ for $C_D^{-1} k_B T = 0.1$ (the small value of T considered), the error in the flipping probability is approximately 0.21%. In contrast, the next smallest choice $l = 16$ corresponds to 10% error at $C_D^{-1} k_B T = 0.1$. In Appendix C, we present the results of simulating the two-dimensional Ising model with strong dipole interactions in order to attempt to place an upper bound on the accumulated error. We find that, when compared with conventional techniques, the algorithm produces a maximum error of around 8% on the obtained critical values.

With any parallel algorithm, one must ensure that the throughput is sufficient to overcome any memory and instructional latencies introduced by the parallelization. For the production runs below, $64 \times 4 = 256$ threads are executed in parallel and the GPU is far from being full loaded. This raises the reasonable question as to whether the GPU is experiencing sufficient throughput. In order to obtain some measure of the improvement, we compared the GPU simulation (Intel E5620 2.4 GHz, Tesla C2050 with blocksize = 128) with the

computation of the dipole sum implemented on the central processing unit (CPU). We found that the GPU simulation performed around 12 times faster. While not experiencing the huge improvement some algorithms have achieved, this performance enhancement is still significant.

IV. FOURTH-ORDER ANISOTROPIES IN THE HEISENBERG MODEL

In two-dimensional magnetic systems, the phase diagram depends on two ratios. The first is the ratio of perpendicular anisotropy to dipole coupling. In two dimensions, the dipole energy favors in-plane ordering of spins at $T = 0$,[42,43] however, a sufficiently strong perpendicular anisotropy can create a ground state with perpendicular spins. At finite temperature, the free energy $F = E - TS$ is minimized and the higher entropy in-plane state can become favored and the spin reorientation transition occurs. The other ratio determining the possible phases of the system is exchange to dipole coupling. For sufficiently strong dipole coupling, the system forms stripes in the ground state, with stronger dipole coupling favoring thinner stripes. The total energy for such a system is given by

$$H = \frac{J}{2} \sum_{\langle i,j \rangle} \vec{s}_i \cdot \vec{s}_j + \sum_i H_{A\,i}$$
$$+ \frac{C_D}{2} \sum_{i,j} \frac{1}{r_{ij}^3} (\vec{s}_i \cdot \vec{s}_j - 3\vec{s}_i \cdot \hat{r}_{ij} \vec{s}_j \cdot \hat{r}_{ij}), \quad (11)$$

where $H_{A\,i}$ is the single site magnetic anisotropy. In the absence of stripes, the spin reorientation transition is known to depend on the higher-order anisotropy terms.[23–25] We are interested on the effect of varying the ratio of second-to fourth-order anisotropy while keeping the anisotropic energy difference between the in-plane and out of plane spins constant. The anisotropy is defined as

$$H_{A\,i} = K[(1-a)(\vec{s}_i \cdot \hat{z})^2 + a(\vec{s}_i \cdot \hat{z})^4] \quad (12)$$

with $K < 0$ and $a > -1$. Here, K represents the strength of the anisotropy and a determines the ratio of fourth-order to second-order anisotropies. Theoretically, the dependence of spin orientation on higher-order anisotropy has been studied as a function of thickness[57] and temperature.[58,59] In these cases, the reorientation of spins is modeled as a competition between competing anisotropy terms which depend on temperature. In our simulations, the anisotropy is considered constant.

Previously, the case of $a = 0$ and varying K has been examined by Whitehead *et al.*[43] The broad thermal phase evolution, ordered stripes at low temperature followed by in-plane magnetization followed by the paramagnetic transition, is reasonably well understood.[41,43] The behavior near the SRT is not well understood, experiments performed on Pt/Co(0.5 nm)/Pt films by Bergeard *et al.*[60] have indicated long-time scale dynamics consisting of regions fluctuating between perpendicular stripes and in plane magnetic order. The authors note that near the SRT, quadratic coupling alone is not sufficient to account for this mixed behavior. Using ac susceptibility studies of striped phases in Fe/Ni films by Abu-Libdeh *et al.*,[36,37,61] have also indicated the presence of long-time scale dynamics. By varying the order of anisotropy,

we wish to investigate the nature of the phase transition between the striped and in-plane phases.

Here, increasing the value of a suppresses states with canting. We consider three choices of parameter a; $a = -1$ corresponding to a system that favors canting (F), $a = 2$ corresponding to a state with suppressed canting (S) and $a = 0$ corresponding to an intermediate propensity for canting (I).

We understand this as follows. Consider the restoring force due to anisotropy experienced by a spin slightly canted away from perpendicular alignment. For the intermediate case, the restoring force is given by the derivative of energy with respect to zenith angle $-\partial_{\theta_i} H_{A_i}|_{\theta_i=0} = -2K$ (with the same results for $\theta_i = \pi$), so for a small amount of canting away from the perpendicular alignment the change in energy is $\Delta H_{A_i}(\theta_i) = 2K\theta_i$. For the case of canting suppression, we have $-\partial_{\theta_i} H_{A_i}|_{\theta_i=0} = 0$ and so there is no force experienced for small spin canting. For the case of strong suppression, $-\partial_{\theta_i} H_{A_i}|_{\theta_i=0} = -6K$ and the restoring force is three times stronger than the equivalent quadratic anisotropy.

A. Results

Normalizing against the strength of the dipole coupling C_D to give dimensionless parameters, we define $\mathcal{T} = (k_B T)/C_D$, $\mathcal{J} = J/C_D$, and $\mathcal{K} = K/C_D$. The exchange coupling was fixed at $\mathcal{J} = 8.9$ giving a ground state with stripe width $w = 8$. The anisotropy strength was fixed at $\mathcal{K} = 15$.

The system is initialized in a perpendicular striped state, and then an ensemble is generated using the above parallel Monte Carlo algorithm with parameters $L = 64$, $P = 64$, and $l = 32$. In order to examine the convergence properties, we examine the results for varied equilibrium times at temperatures where we expect the slowest relaxation: at low temperatures and near the transition points. In almost all cases, energy converges rapidly, taking several hundred Monte Carlo steps. The morphological properties take longer. Slow relaxation of stripe patterns has been observed in studies of Ising systems.[62] However, in our simulations, we find that the in-plane magnetization is the slowest converging measure, often taking several tens of thousands of steps to equilibrate. There are, however, two regions where equilibrium is significantly slower. The first is near the sharp transition for the $a = 2$ case near $\mathcal{T} = 4$. At $\mathcal{T} = 3.5$ (just below the transition), all parameters converge rapidly. At $\mathcal{T} = 4$ (just above the transitions), the energy is slow to converge, taking approximately 5×10^4 steps. In this case, it takes 7.5×10^4 steps for the magnetization to reach convergence. The other region where some care is needed is the $a = -1$ case at $\mathcal{T} = 0.5$. Recalling $-\partial_{\theta_i} H_{A_i}|_{\theta_i=0} = 0$, spins have the ability to cant even at low temperatures, and the system carries a net magnetization. Since the system is initialized in a perpendicular state, the relaxation time required to form stripes is large, in this case, we find that the system requires approximately 1.25×10^5 steps to equilibrate. In Appendix A, we give the results of varied equilibrium times for these slowly converging temperatures. Here, we have taken a fairly course temperature spacing between simulations in order to explore the phase space of the system. If one were to attempt simulations closer to the transition points, it is likely that the required equilibrium
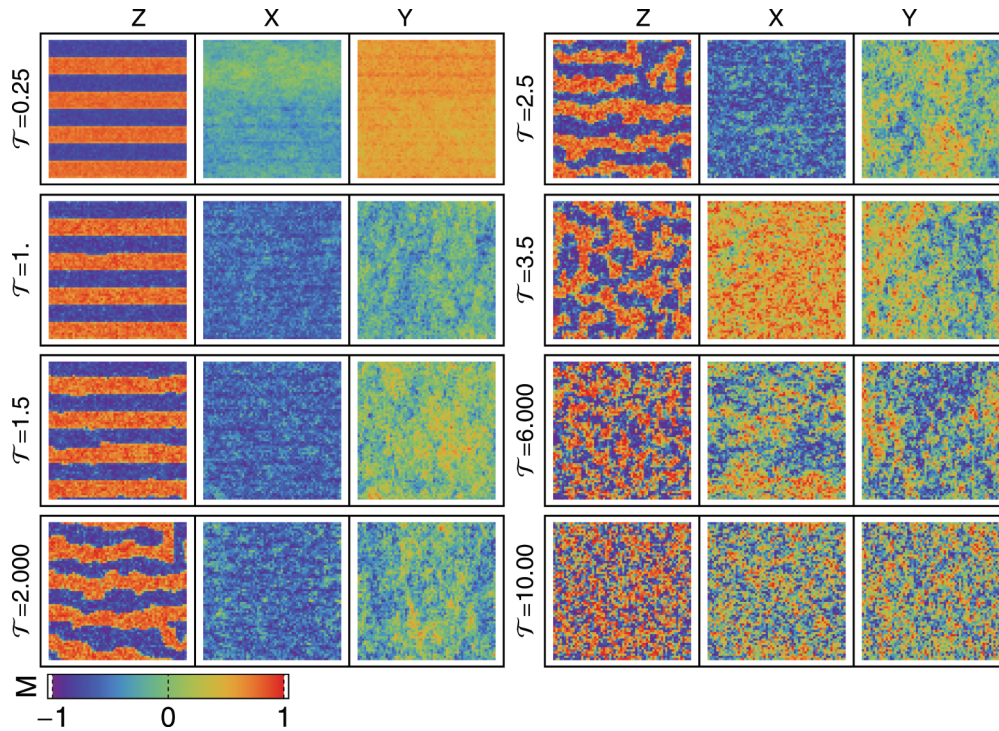
FIG. 7. (Color online) Example spin configurations for a canting favored system ($a = -1$). Each state is represented by three images, from left to right, showing the z, x, and y components of spins, respectively. Left column from top: $\mathcal{T} = 0.25, 1.0, 1.5, 2.0$. Right column from top: $\mathcal{T} = 2.5, 3.5, 6, 10$.

time will increase, particularly for the $a = 2$ case where we have already observed a large increase in the required equilibrium time. Once at equilibrium, an additional 5×10^4 steps are simulated with the state recorded every 50 steps to form an ensemble. This ensures that the correlation between the same spin in subsequent states of the ensemble is limited. Waiting 50 steps corresponds to an average value of the time correlation $\langle \vec{s}_i(t = t_o) \cdot \vec{s}_i(t = t_o + 50) \rangle \approx 0.5$ at $\mathcal{T} = 4$. The ensemble size $n = 1000$ ensures that the standard error in the thermal averages of order parameter O,

$$ \mathrm{SE}_{\langle O \rangle} = \left(\frac{\langle O^2 \rangle - \langle O \rangle^2}{n} \right)^{1/2}, \qquad (13) $$

remains smaller than the errors due to the algorithm discussed in Sec. III. We also calculate the errors using nonparametric jack-knife resampling,[63] in cases where the error is larger than the plot points, it is indicated on plots as error bars. One should still be careful when interpreting the results. Resampling only gives the deviation from the calculated values due to finite ensemble size, it can give no information about the cumulative error due to the algorithm discussed in Sec. III B.

Example states for each value of a are given in Figs. 7–9. For the case of favored spin canting, $a = -1$ (see Fig. 7), the system has stable stripes in the ground state, but the perpendicular magnetization is not saturated. The in-plane components display long-range ordering. As temperature is increased, the stripes display roughening, and then bridging is leading to the eventual loss of orientational order. As temperature is further increased, the in-plane order breaks into domains, before the system enters the high-temperature paramagnetic phase.

For the intermediate case, $a = 0$ (see Fig. 8), we observe perpendicular stripes at low $\mathcal{T}$. The boundaries of these stripes undergo roughening and eventual bridging, analogous to the canting favored case, before orientational order is destroyed with increasing temperature. Unlike the canting favored case, we note the presence of increased canting at the domain boundaries. Above this temperature, we observe a mixed phase in which perpendicular domains are interspersed with regions of in-plane magnetic order. As temperature is further increased, these domains become increasingly granular until the system reaches the paramagnetic limit.

When canting is suppressed, $a = 2$ (see Fig. 9), the system forms perpendicular stripes in the ground state. As the temperature is increased, the walls undergo roughening but not the bridging and gradual loss of orientational order displayed in the $a = 0$ and -1 simulations. Instead, the system undergoes a sudden transition into a state with only small regions of perpendicular alignment remaining and strong in-plane order. As temperature is increased, we observe an increasing number of perpendicular regions as the in-plane order breaks into domains. At high temperature, in-plane order is destroyed as the system becomes paramagnetic.

In order to locate domain walls, we define horizontal and vertical order parameters based on those described by Whitehead *et al.*:[43,64]

$$ n_v^z = \sum_i 1 - \mathrm{sgn}\left(s_i^z s_{i+\hat{y}}^z \right), \quad n_h^z = \sum_i 1 - \mathrm{sgn}\left(s_i^z s_{i+\hat{x}}^z \right), $$

$$ (14) $$

with analogous definitions for the x and y components of the spins. At low temperatures when the systems are dominated by
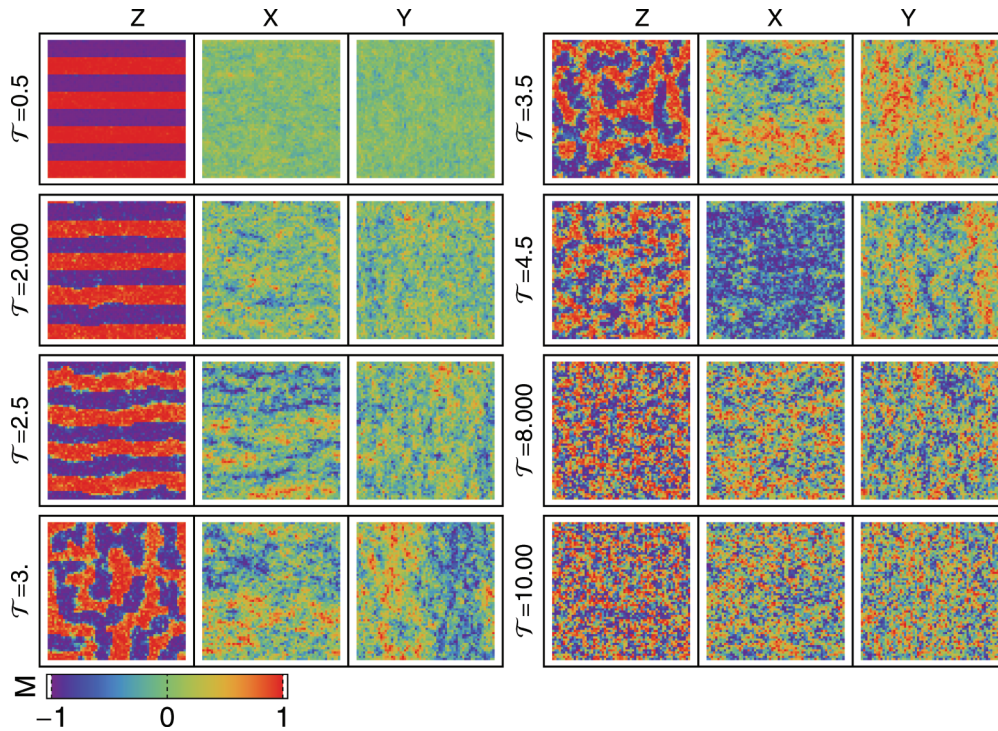
FIG. 8. (Color online) Example spin configurations for the intermediate case ($a = 0$). Left column from top: $\mathcal{T} = 0.5, 2, 2.5, 3$. Right column from top: $\mathcal{T} = 3.5, 4.5, 8, 10$.

perpendicular alignment, the density of perpendicular domain boundaries is given by $(n_v^z + n_h^z)/(4N)$. In Fig. 10, we see that for the intermediate and canting favored states, the wall density increases gradually with temperature. This represents

roughening and then the bridging before perpendicular order is lost and the wall density tends to the $T \to \infty$ value of $1/2$. In contrast, the canting suppressed state undergoes far less roughening and the wall density remains small until
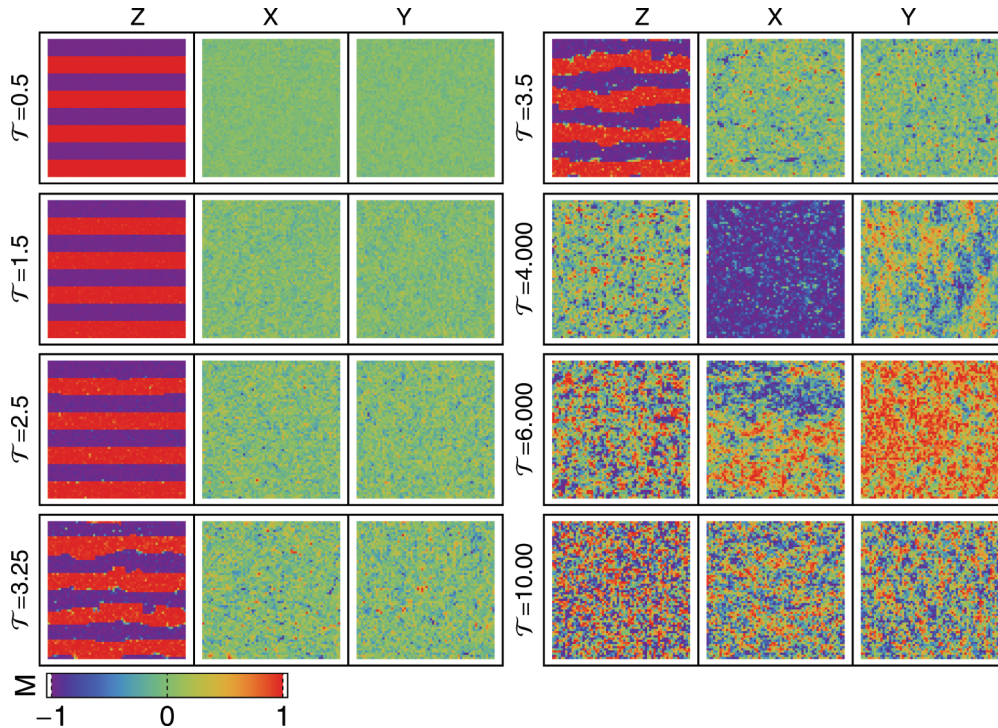

FIG. 9. (Color online) Example spin configurations for the canting suppressed case ($a = 2$). Left column from top: $\mathcal{T} = 0.5, 1.5, 2.5, 3.25$. Right column from top: $\mathcal{T} = 3.5, 4, 6, 10$.
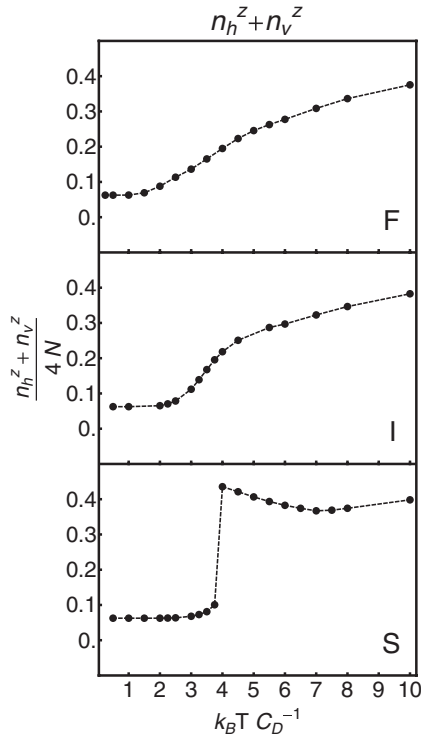
FIG. 10. Total wall length for the canting favored $a = -1$ (F), intermediate $a = 0$ (I), and canting suppressed $a = 2$ (S) cases.

the system undergoes a sudden change into the high-T state.

1. Orientational order transition

The orientational order is given by considering the ratio of horizontal and vertical domain boundaries,

$$\mathcal{O}_\alpha = \left\langle \frac{|n_h^\alpha - n_v^\alpha|}{n_h^\alpha + n_v^\alpha} \right\rangle \quad (15)$$

for $\alpha \in \{x, y, z\}$. In Eq. (14), we have defined separate order parameters for each component rather than defining the combined parameters $n_v = \sum_i 1 - \vec{s}_i \vec{s}_{i+\hat{y}}$ and $n_h = \sum_i 1 - \vec{s}_i \vec{s}_{i+\hat{x}}$. This definition is robust against canting, meaning reduction in $\mathcal{O}$ due to changing stripe morphology (such as coarsening or bending) can be distinguished from changes in the cone angle. Consider the low $\mathcal{T}$ states shown in Figs. 7 and 9, both states have stripes with no coarsening or bending and the definition given in Eq. (14) assigns the same order parameters to both states. In Fig. 11, this $\mathcal{O}_z$ is plotted for the three values of a. $\mathcal{O}_x$ and $\mathcal{O}_y$ were also calculated and were zero in all cases. For higher a, the stripes are stabilized at higher $\mathcal{T}$. For perpendicular stripes, high a values suppress small fluctuations at the stripe boundaries preventing the roughening and eventual bridging that leads to a loss of orientational order.

2. Spin reorientation transition

When the spins are not entirely perpendicular to the plane, it is possible for the system to acquire in-plane ferromagnetic order. If we define $M_x = 1/N \sum_i s_i^x$ and $M_y = 1/N \sum_i s_i^y$,
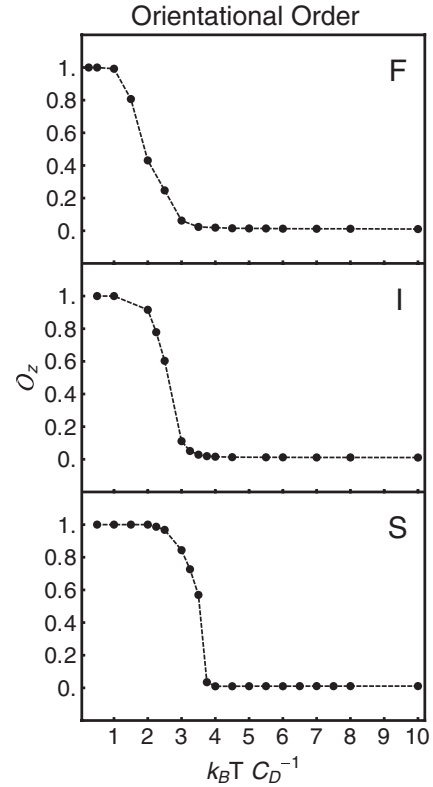

FIG. 11. The orientational order for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S).

then the parallel magnetization is given by

$$M_\parallel = \left\langle \left(M_x^2 + M_y^2 \right)^{\frac{1}{2}} \right\rangle. \quad (16)$$

The appearance of nonzero ferromagnetic order is not inherently indicative of global spin reorientation. Canted spin states and finite thickness domain walls can account for significant magnetic order.[43,65] To measure the degree to which the spins reorient as a function of temperature, we introduce the cone angle η:

$$\eta = \frac{1}{N} \sum_i \sqrt{(2/\pi)^2 \langle (\theta_i - \pi/2)^2 \rangle}, \quad (17)$$

where θ_i is the zenith angle of spin $\vec{s}_i$, $\theta_i = \arccos(s_i^z)$, and we have normalized η so that $0 \leqslant \eta \leqslant 1$. In Fig. 12, η is shown as a function of $\mathcal{T}$, along with the $T \to \infty$ value $\eta = \sqrt{(2/\pi)^2 [1/4(\pi^2 - 8)]}$. Despite the energy difference between parallel and perpendicular alignment remaining constants, the behavior varies dramatically with a.

For the canting favored case, $a = -1$, we see that by reducing the energy cost of canting spins, the sample does not experience a spin reorientation transition, instead it remains canted for all temperatures. In Fig. 13, it is seen that this canting allows the sample to have nonzero magnetic order at $\mathcal{T} = 0$. The appearance of maximum magnetic ordering at $\mathcal{T} = 0$ is characteristic of weaker values of $\mathcal{K} < 13$ and $a = 0$.[43] However, in these cases, the low $\mathcal{T}$ parallel magnetization does not occur simultaneously with orientational order. As $\mathcal{T}$ is increased, the degree of canting remains practically
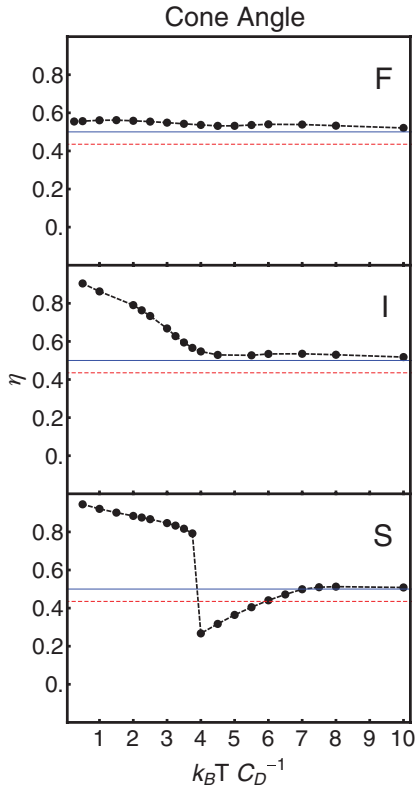
FIG. 12. (Color online) η as a function of $\mathcal{T}$ for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S). $\eta = 1$ indicates all spins point perpendicular to the plane ($\vec{s}_i \cdot \hat{z} = \pm 1$), $\eta = 0$ indicates all spins lie in plane ($\vec{s}_i \cdot \hat{z} = 0$). The red dashed line represents the $T \to \infty$ value. The solid blue line represents ($\vec{s}_i \cdot \hat{z} = 1/\sqrt{2}$).

unchanged with the orientational order decreasing over the range of $1 < \mathcal{T} < 2.5$.

For the intermediate case, $a = 0$, we observe results consistent with those of Whitehead *et al.*:[43] a loss of orientational order over a small region, $1 < \mathcal{T} < 3$, in which stripes also become slightly canted leading to a small in-plane magnetization. Above this temperature, the in-plane magnetization increases as η decreases, resulting in a peak in-plane magnetization of $M_\parallel \approx .46$ at $\mathcal{T} = 4.5$. For $\mathcal{T} > 4.5$, the $a = 0$ system is identical to the $a = -1$ system, η slowly decreasing and magnetic order gradually reduced to zero at $\mathcal{T} = 8$.

For the spin suppressed state, $a = 2$, we observe the same perpendicular ground state as in the $a = 0$ case. Unlike the previous cases, the orientational order is stabilized up to $\mathcal{T} = 4$, at which point the system undergoes a sharp transition to a parallel ferromagnetic state. In Fig. 12, we see that η simultaneously undergoes a sharp transition, representing the spin reorientation transition. Above $\mathcal{T} = 4$, we observe a gradual reduction in magnetic order until the system enters the paramagnetic state above $\mathcal{T} = 8$.

V. FLUCTUATIONS

In order to examine fluctuations near critical points, we calculate the autocorrelation,[66]

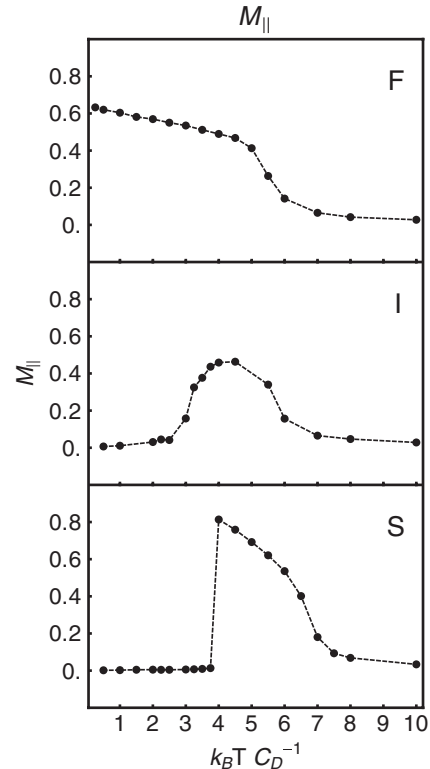$$\sigma^2(X) = \langle (X - \langle X \rangle)^2 \rangle, \qquad (18)$$

FIG. 13. $M_\parallel$ as a function of $\mathcal{T}$ for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S).

of the three order parameters $M_\parallel$, $\mathcal{O}^z$, and η, which we denote $\sigma_\parallel^2$, $\sigma_{\mathcal{O}}^2$, and σ_η^2, respectively. The small numerical values of these variances means that they are affected to a greater extent by errors introduced by the simultaneous flipping.

In Fig. 14, we observe peaks in $\sigma_\parallel^2$ associated with the loss of in-plane magnetization for each choice of a. In addition, we observe a smaller peak at around $\mathcal{T} = 3$ for the intermediate, $a = 0$, case corresponding to the formation of in-plane magnetic order. This low-temperature peak is absent in the canting suppressed case due to the first-order nature of the phase transition. The sharp in-plane transition does not correspond to a significant change in ferromagnetic ordering. In both the perpendicular striped phase and in-plane ferromagnetic state, the exchange energy is minimized for the majority of spins. In the canting favored state, the low-$\mathcal{T}$ transition is absent since maximum magnetic order occurs at the lowest temperature simulated $\mathcal{T} = 0.2$.

In Fig. 15, σ_η^2 is plotted for the three choices of a. For the canted favored case the fluctuations are small, culminating in shallow peak at $\mathcal{T} = 5$. This is consistent with the nearly homogeneous cone angle. For the intermediate case, the fluctuations displays a broad peak centered just below the minimum cone angle at $\mathcal{T} = 4$. For the canting suppressed case, there is a small peak corresponding to the initial reorientation of the spins followed by a broad peak as the cone angle starts to approach the high-temperature average.

In Fig. 16, the fluctuations of $\mathcal{O}_z$ are plotted as a function of temperature. In each case, the variance forms a peak
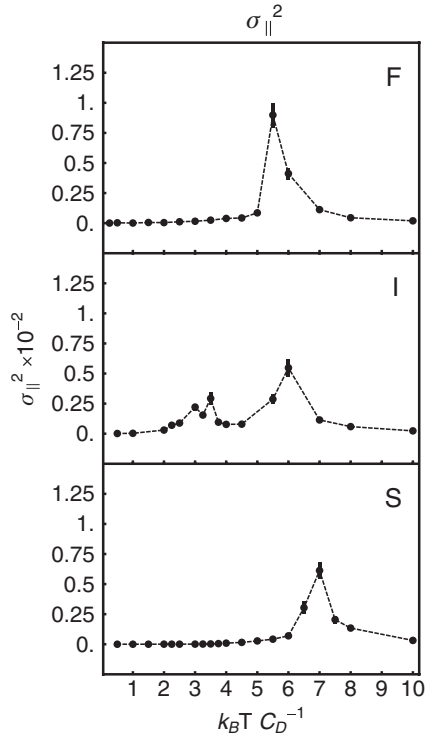
FIG. 14. $\sigma_{\parallel}$ as a function of $\mathcal{T}$ for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S).

corresponding to the loss of orientational order. Increased canting suppression corresponds to thinner peaks, as the transition occurs over a smaller temperature range. We note
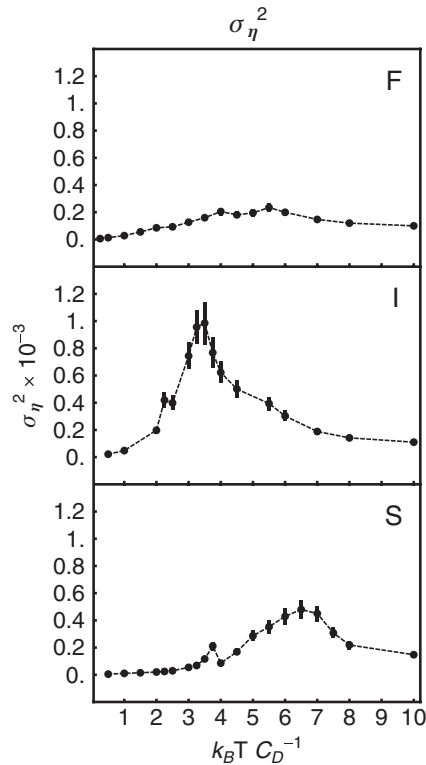

FIG. 15. σ_{η} as a function of $\mathcal{T}$ for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S).
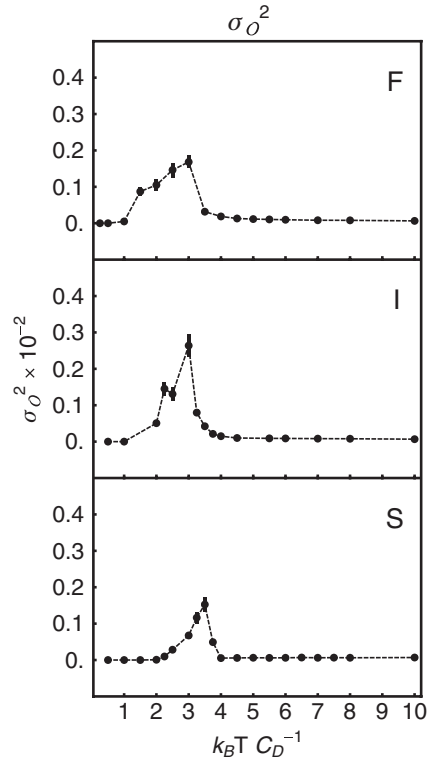

FIG. 16. $\sigma_{\mathcal{O}}$ as a function of $\mathcal{T}$ for favored canting $a = -1$ (F), intermediate canting $a = 0$ (I), and suppressed canting $a = 2$ (S).

also that the strength of the peak is smaller for the canting suppressed case, due to the fact that only roughening occurs but not bridging.

VI. CONCLUSIONS AND COMMENTS

Here, we have examined the nature of stripe formation and spin reorientation in the presence of strong perpendicular anisotropy. In order to do so, we have proposed an approximation that allows for a parallel algorithm for performing Monte Carlo simulations in cases where there is long-range coupling. We have argued that, since the dipole coupling contributes with a r^{-3} dependence, for appropriate choice of algorithm parameters the approximation is acceptable. This algorithm reduces significantly the computation time associated with increased system size. The algorithm has been applied to the case of Ising spins in the presence of dipole coupling and shown to be consistent with results obtained by conventional methods.

It is possible that further increases in computational efficiency might be gained through more efficient parallelization of the dipole sum. Furthermore, the concept of simultaneous flipping might be particularly useful in three dimensions where the computational issues are enhanced, the caveat of course being that care must be taken with parameter choice since the number of simultaneous spins is greatly increased. This technique has then been used to examine the effects of higher order anisotropy in striped systems where we have shown that the anisotropy order can suppress or enhance the SRT in cases with strong out of plane anisotropy.

TABLE I. $\mathcal{T} = 3.5$.

Equilibrium Steps	$a = -1$			$a = 0$			$a = 2$		
	$M_\parallel$ (0.001)	$\mathcal{O}$ (0.002)	η (0.001)	$M_\parallel$ (0.003)	$\mathcal{O}$ (0.002)	η (0.003)	$M_\parallel$ (5×10^{-4})	$\mathcal{O}$ (0.004)	η (0.001)
24 900	0.517	0.024	0.724	0.397	0.026	0.863	0.0092	0.582	1.646
50 000	0.518	0.023	0.724	0.400	0.026	0.862	0.0092	0.583	1.646
75 200	0.518	0.023	0.723	0.402	0.025	0.861	0.0091	0.584	1.647
100 000	0.517	0.025	0.723	0.400	0.027	0.861	0.0092	0.584	1.646

When comparing our simulations to analytic results, we note several discrepancies. Although we were able to observe the coexistence of stripes and magnetic order, we did not observe domain formation of the cone angle as suggested by Abanov.[41] We also did not observe any magnetic order due to correlations in the in-plane magnetization of domain walls in the wall segments dividing stripes. Whitehead *et al.* also noted the absence of such correlations,[43] however, Yafet and Gyorgy have argued that correlations between domain walls leading to ferromagnetic order are possible.[65] The discrepancy between analytic and computational results might largely be a result of limited system size. In particular, it is not possible to make a fine grained examination of wall profiles in these small systems. The parallel algorithm presented here reduces the computational cost of scaling system size. In the future, it would be interesting to use the algorithm to simulate far bigger systems, in which domain wall structure could be more reasonably simulated, providing more insight into correlations that occur on finer scales.

APPENDIX A: CONVERGENCE MEASURES

Here, we give the results of calculating order parameters with variable equilibrium times in Tables I–IV.[67] In each case, the uncertainty is estimated by considering the standard error based on the standard deviation calculated from the longest equilibrium time,[68] this uncertainty is indicated in

brackets at the top of each column. The system is considered equilibrated when values agree within the uncertainty for three subsequent simulation times. For the $a = -1$ case at temperatures below $\mathcal{T} = 1$, the uncertainty due to finite ensemble size is smaller than the uncertainty introduced from a single pass of the multiple spin-flip algorithm (sample variance decreases with temperature, while the multiple spin-flip error increases). In this cases, we assume convergence when the calculated values have converged to three significant figures.

APPENDIX B: PSEUDOCODE

In order to implement the parallel algorithm described, one needs to address the potential update sites and the subset of sites with which they interact as a function of the thread identifier. Here, we give the pseudocode addressing this for the subsystems described above. The state is initialized on the host memory and consists of two one-dimensional arrays of size L^2. These arrays store the azimuthal (ϕ) and zenith (θ) angles. In what follows, we will use $\vec{s}_j$ to represent values of both ϕ and θ at some position j in these arrays. A number of potential update sites are selected separated horizontally and vertically by a fixed number of sites, which we denote l, i.e., $j = (al,bl) + j1$. The interactions between array elements is calculated according to the algorithm below in which blockIdx.x, blockDim.x, and threadIdx.x are system integers that identify a thread address. In the following, $a \backslash b$ is defined as $a \backslash b = \text{floor}(a/b)$. For example, $7\backslash 3 = 2$ and $2 \times (7\backslash 3) + 7 \bmod 3 = 7$.

ON HOST
S_1 = Random Integer $\in [1, l]$
S_2 = Random Integer $\in [1, l]$
$N = L \backslash l$
for $j = 1$; $j < N$ do
Select a series of new states:
$\theta[j]$ = Random Real $\in [0, 2\pi]$
$\phi[j]$ = Arccos(Random Real $\in [-1, 1]$)

TABLE II. $\mathcal{T} = 6$.

Equilibrium Steps	$a = -1$			$a = 0$			$a = 2$		
	$M_\parallel$ (0.006)	$\mathcal{O}$ (0.001)	η (0.001)	$M_\parallel$ (0.009)	$\mathcal{O}$ (9×10^{-4})	η (0.002)	$M_\parallel$ (0.002)	$\mathcal{O}$ (8×10^{-4})	η (0.002)
6300	0.134	0.013	0.719	0.166	0.0121	0.704	0.533	0.0105	0.480
12 500	0.137	0.013	0.719	0.166	0.0121	0.704	0.532	0.0105	0.480
24 900	0.135	0.013	0.719	0.168	0.0120	0.704	0.532	0.0105	0.480
50 000	0.133	0.013	0.719	0.172	0.0120	0.703	0.532	0.0105	0.480

TABLE III. $\mathcal{T} = 4$.

Equilibrium Steps	$a = -1$			$a = 0$			$a = 2$		
	$M_\parallel$ (0.002)	$\mathcal{O}$ (0.002)	η (0.001)	$M_\parallel$ (0.003)	$\mathcal{O}$ (0.001)	η (0.003)	$M_\parallel$ (7×10^{-4})	$\mathcal{O}$ (8×10^{-4})	η (8×10^{-4})
3200	0.480	0.018	0.708	0.455	0.016	0.740	0.6486	0.0099	0.4381
6300	0.488	0.018	0.708	0.455	0.016	0.740	0.6576	0.0098	0.4235
12 500	0.495	0.017	0.708	0.454	0.016	0.740	0.6764	0.0098	0.3936
24 900	0.495	0.018	0.707	0.455	0.016	0.739	0.7159	0.0098	0.3310
50 000	0.496	0.018	0.708	0.454	0.016	0.740	0.8094	0.0099	0.1832
75 200	0.495	0.018	0.708	0.457	0.016	0.739	0.8141	0.0098	0.1762
100 000	0.495	0.019	0.707	0.457	0.016	0.738	0.8146	0.0096	0.1760
125 000	...	...	...	...	...	...	0.8146	0.0099	0.1761
150 000	...	...	...	...	...	...	0.8148	0.0106	0.1762

Copy all variables arrays to GPU
end for
ON GPU: Exchange Interaction
T_{id} = blockIdx.x *blockDim.x + threadIdx.x;
if $T_{id} \leqslant L^2 \backslash l^2$ then
$S_x = S_1 + l(T_{id} \bmod N)$
$S_y = [S_2 + l(T_{id} \backslash N)]$
$j = S_y L + S_x$
calculate the exchange coupling between $\vec{s}_j$ and its immediate neighbors
calculate the exchange coupling between $(\theta[j], \phi[j])$ and $\vec{s}_j$'s immediate neighbors
end if
ON GPU: Dipole Interaction
T_{id} = blockIdx.x*blockDim.x + threadIdx.x
$S_{id} = T_{id} \backslash P$
if $S_{id} \leqslant L^2 \backslash l^2$ then
$S_x = S_1 + l(S_{id} \bmod N)$
$S_y = [S_2 + l(S_{id} \backslash N)]$
$j = S_y L + S_x$
$\sigma = (L + 1) \backslash P$
$\rho = [(L + 1) \bmod P] - 1$
for $k = -L \backslash 2$; $k \leqslant L \backslash 2$ do
for $i = -L \backslash 2 + \sigma T_{id} \bmod P)$; $i < -L \backslash 2 + \sigma(T_{id} \bmod P + 1)$ do
$j' = \{[(S_y + k) \bmod L] L\} + (S_x + i) \bmod L$
Calculate the dipole coupling between $\vec{s}_j$ and $\vec{s}_{j'}$
Calculate the dipole coupling between $(\theta[j], \phi[j])$ and $\vec{s}_{j'}$
end for

end for
end if
ON GPU: Single Site Energies and Spin Flips
T_{id} = blockIdx.x*blockDim.x + threadIdx.x;
if $T_{id} \leqslant L^2 \backslash l^2$ then
$S_x = S_1 + l(T_{id} \bmod N)$
$S_y = [S_2 + l(T_{id} \backslash N)]$
$j = S_y L + S_x$
calculate anisotropy and Zeeman energies for $\vec{s}_j$
calculate anisotropy and Zeeman energies $(\theta[j], \phi[j])$
replace $\vec{s}_j$ with $(\theta[j], \phi[j])$ or $\vec{s}_j$ according to the Boltzmann probability
end if

APPENDIX C: DIPOLE ISING MODEL

In Sec. III B, it was argued that the error introduced by a single pass of the proposed GPU algorithm is bounded. However, this does not ensure that the error is not compounded over a large number of passes leading to large systematic errors. In order to investigate this possibility, we consider the result of applying the algorithm when every accepted spin update corresponds to a complete reversal: $\vec{s}_i = (0, 0, s_i^z) = (0, 0, \pm 1)$. We select the same algorithm parameters that are used in Sec. IV: $L = 64$ and $l = 32$. This corresponds to four simultaneous attempted spin flips for each cycle of the algorithm. Since each site is restricted to only two states the energy can be

TABLE IV. $\mathcal{T} = 0.5$.

Equilibrium Steps	$a = -1$			$a = 0$			$a = 2$		
	$M_\parallel$	$\mathcal{O}$	η	$M_\parallel$ (4×10^{-4})	$\mathcal{O}$ (0)	η (5×10^{-4})	$M_\parallel$ (1×10^{-4})	$\mathcal{O}$ (0)	η
24 900	...	...	...	0.0075	1	2.0180	0.0017	1	2.203
50 000	...	...	...	0.0076	1	2.0179	0.0017	1	2.203
75 200	...	...	...	0.0074	1	2.0177	0.0017	1	2.203
100 000	0.622	1	0.764	0.0076	1	2.0178	0.0017	1	2.203
125 000	0.625	1	0.764	...	...	...	...	...	...
150 000	0.625	1	0.764	...	...	...	...	...	...
175 000	0.625	1	0.764	...	...	...	...	...	...

TABLE V. Various parameters calculated using the GPU algorithm and conventional techniques[74].

Parameter	GPU algorithm	Conventional MC
$\mathcal{T}_c$	0.85	0.82
γ	1.62	1.75
β	0.08	0.08

written as

$$H = \frac{J}{2} \sum_{\langle i,j \rangle} s_i s_j + \frac{C_D}{2} \sum_{i,j} \frac{s_i s_j}{r_{ij}^3}. \tag{C1}$$

In the presence of a sufficiently strong dipole interaction, the ground state of the system will form a striped pattern of alternating spins. In order to ensure that dipole coupling contributes significantly to the total energy, (increasing the error) parameters are selected such that $J/C_D = 1.7$, which corresponds to the thinnest possible stable stripe width $h = 1$. The order parameter for such a system is formed by considering a series of sublattices in the manner described by Binder and Landau.[69] For $h = 1$, the system is broken into four sublattices m_λ, horizontal stripes are described by $m_h = m_1 + m_2 - m_3 - m_4$ and vertical stripes by $m_v = m_1 + m_4 - m_3 - m_2$. The order parameter is then the staggered magnetization $m_{st} = \langle (m_h^2 + m_v^2)^{1/2} \rangle$. In Fig. 17, this staggered magnetization is shown as a function of a normalized temperature $\mathcal{T} = k_B T C_D^{-1}$. The staggered susceptibility $\chi_{st} = \frac{L^2}{k_B T}(\langle m_{st}^2 \rangle - \langle m_{st} \rangle^2)$ is also calculated. By determining the location of the peak in Fig. 18, the critical temperature of the phase transition $\mathcal{T}_c$ can be determined, the result is shown in Table V. The peak lacks the δ-like structure associated with a first-order transition, instead displaying an exponential decay consistent with a continuous phase transition,[70,71] this is consistent with previous simulations with strong dipole coupling.[72] In Fig. 19, we show the probability distribution of the average energy per spin for various temperatures near the transition. At each temperature, the distribution displays a single turning point, this is also consistent with the expected continuous transition.[72] Near the critical temperature, the system displays critical behavior. After rescaling temperature as $t = L^{\frac{1}{\nu}}|1 - T/T_c|$, one expects the following scaling relations: $m_{st} = L^{-\frac{\beta}{\nu}} m_0(t)$ and $\chi_{st} = \frac{L^{\frac{\gamma}{\nu}}}{t} \chi_0(t)$, where $m_0(t)$ and $\chi_0(t)$ are universal scaling functions.[73,74] Close to the
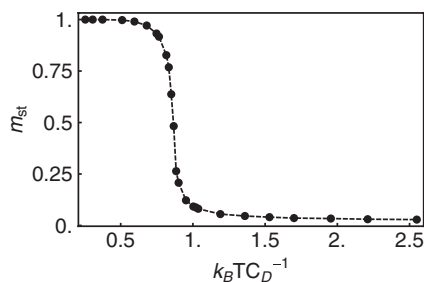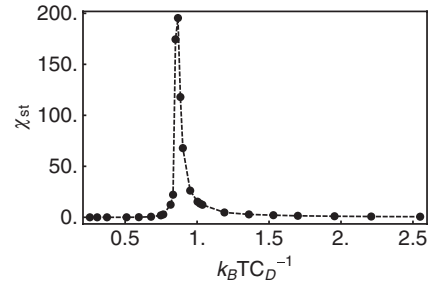
critical point, when $|1 - T/T_c|$ is small and L large, these universal scaling functions are expected to show a power-law behavior $m_0(t) \to Bt^\beta$ and $\chi_0(t) \to At^{-\gamma}$. In order to extract parameters from simulation, one usually makes use of the universality of $m_0(t)$ and $\chi_0(t)$ and simulates the system for a number of different sizes before varying the parameters until the results of all simulations lie on a single curve.[74,75] However, performing such analysis would not be a suitable test of the algorithm, for any choice $L < l$, there is no simultaneous flipping and hence no approximation is being made. If one were to change the size of l to simulate smaller systems, the degree of approximation would be changed (the error will be increased for decreasing l). Instead, we use the value of the scaling constant ν calculated by Rastelli *et al.* to rescale temperature and then use the power-law dependence of the system near T_c to extract β and γ. A comparison of the critical properties is given in Table V. The use of the GPU has introduced some error in the critical parameters of the system, the largest error being slightly less than 8%. Current generation GPU cards use lower than the IEEE recommended precision on certain functions (notably trigonometric and exponential functions required here).[76,77] In previous simulations of the Heisenberg model, the discrepancy between CPU and GPU results of up to 5% have been attributed to different levels of numerical precision.[54] Since this previous implementation did not use Ewald summation, the number of floating point operations performed on the GPU is larger than in our work. It is likely then that the error introduced by our approximation is largest source of discrepancy, indeed the errors introduced in our work are larger than those described by Campos *et al.*

FIG. 18. The staggered susceptibility as a function of $\mathcal{T}$.

FIG. 17. The staggered magnetization as a function of $\mathcal{T}$ showing the order-disorder transition.
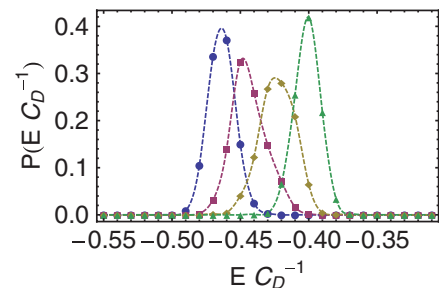

FIG. 19. (Color online) Probability distributions of energy near the critical point: $\mathcal{T} = 0.833$ (blue circles), 0.85 (red squares), 0.867 (yellow diamonds), 0.884 (green triangles).

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