

**Quantitative analysis of phase transitions in two-dimensional  $XY$  models using persistent homology**Nicholas Sale<sup>✉,\*</sup>, Jeffrey Giansiracusa<sup>✉,†</sup> and Biagio Lucini<sup>✉,‡</sup>*Department of Mathematics, Swansea University, Bay Campus, SA1 8EN, Swansea, Wales, United Kingdom*

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We use persistent homology and persistence images as an observable of three variants of the two-dimensional  $XY$  model to identify and study their phase transitions. We examine models with the classical  $XY$  action, a topological lattice action, and an action with an additional nematic term. In particular, we introduce a way of computing the persistent homology of lattice spin model configurations and, by considering the fluctuations in the output of logistic regression and  $k$ -nearest neighbor models trained on persistence images, we develop a methodology to extract estimates of the critical temperature and the critical exponent of the correlation length. We put particular emphasis on finite-size scaling behavior and producing estimates with quantifiable error. For each model we successfully identify its phase transition(s) and are able to get an accurate determination of the critical temperatures and critical exponents of the correlation length.

DOI: [10.1103/PhysRevE.105.024121](https://doi.org/10.1103/PhysRevE.105.024121)**I. INTRODUCTION**

An emerging body of work is exploring the use of machine learning and other data analysis methods to detect and classify phase transitions in statistical physics systems (see, among others, Refs. [1–14]). One of the motivations of this approach is to develop methodologies which require minimal *a priori* knowledge about the systems in question. The hope then is that these data-centric methods will be able to offer new insights into those models at the forefront of physics which seem to defy analytical methods [15]. Much of the work in this area makes use of neural network models which, while unparalleled in machine learning tasks, are generally hard to interpret. But recently, among other geometric and topological approaches [16–19], there has been an interest in using persistent homology, a tool from the new field of topological data analysis (TDA), to produce interpretable features which are inherently sensitive to topological objects. These can then be compared in their own right, or fed into a machine learning model [20–25].

There are at least two paradigms for using persistent homology to study phase transitions of a given statistical physics model. The first can be called *persistent homology in configuration/data space*, where the topology of the high-dimensional space of model configurations is probed from samples. This approach is based on the *topology hypothesis* for the origin of phase transitions [26,27] and is the approach used in [25]. The idea here is that a thermodynamic phase transition necessarily coincides with a change in the topology of the energy level set, although such a change does not turn

out to be a sufficient condition [28]. In the present work, however, we shall make use of a more recent paradigm, investigated also in [20–24], which we call *persistent homology as an observable*. Given a sampled configuration of a model, we construct a sequence of geometric complexes based on that configuration. This sequence of topological spaces is known as a filtration. Applying persistent homology to the filtration yields a collection of points called a persistence diagram, which represents this configuration. We can think of this process as a means to reduce the degrees of freedom of the model and produce nonlinear summaries of configurations. Statistics of these persistence diagrams are then analyzed as the system undergoes a phase transition. Previous works have focused on identifying the different phases in various models in a mostly qualitative manner. While Ref. [24] makes some steps towards obtaining quantitative measurements of the multiscale structure of the Ising model at criticality, a framework for using persistent homology observables to make rigorous numerical estimates of critical temperatures and exponents with quantified error has not yet been explored in the literature to the best of our knowledge.

While the existing works on the *persistent homology as an observable* paradigm share the same underlying idea, the approaches seen so far have differed significantly, both in how filtrations have been constructed and in how the resulting persistence diagrams have been analyzed. Tran, Chen, and Hasegawa investigated phase transitions in the two-dimensional (2D)  $XY$  model, one-dimensional (1D) transverse-field Ising, and 1D Bose-Hubbard models [22]. They computed the Vietoris-Rips persistence of point clouds of lattice sites with interpoint distances given by a linear combination of the Euclidean distance in the lattice and the difference in the spins. They show that clustering configurations based on the persistence Fisher kernel [29], persistence entropy, and second moment of persistence of the  $H_1$  diagrams identifies the different phases. They demonstrate that increasing the lattice size produces sharper estimates of the critical temperature. This approach is extended to the XXZ model

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on a pyrochlore lattice by Olsthoorn, Hellsvik, and Balatsky, approximately separating the six different phases of the model [23].

Cole, Loges, and Shiu apply a different methodology to the previous works. Looking at the 2D Ising, square-ice,  $XY$ , and fully frustrated  $XY$  models, they introduce general constructions of filtrations for configurations of discrete-valued and circle-valued spin models [24]. In particular, configurations of circle-valued models are given a sublevel set filtration of the map  $f: \Lambda \rightarrow (-\pi, \pi]$  which assigns each site  $i \in \Lambda$  in the lattice a parametrization of its spin  $f(i) \in (-\pi, \pi]$ . This filtration yields cubical subcomplexes of the lattice. They make use of persistence images [30] to vectorize persistence diagrams, allowing the application of a logistic regression model to separate the phases. They relate some quantitative aspects of the persistence diagrams to the estimation of critical exponents in the case of the Ising model. For discrete models they construct  $\alpha$ -complexes on subsets of the lattice sites with the same spin. This is similar to the approach used by Hirakida *et al.* [21], who look at the effective Polyakov line model.

Comparing the approach in [22,23] to that in [24] makes it clear that there is a significant degree of choice in picking the filtration used to compute the persistent homology of a given lattice configuration. We will demonstrate that this choice is an important factor in determining what information about phase transitions one can derive from the persistence. In particular, we investigate an  $XY$  model with a nematic interaction term and find that using two different filtrations is required to detect and analyze the two different phase transitions undergone by the system.

Our main contributions are as follows:

(1) We introduce a class of filtrations on lattice spin systems which, while general, allow persistent homology to easily detect topological defects.

(2) Extending the approach of using logistic regression on persistence images introduced in [24], we investigate the applicability of finite-size scaling analysis. In particular, we apply the standard statistical tools of histogram reweighting and bootstrapping to obtain estimates of the critical temperature and the critical exponent of the correlation length with quantified error.

(3) Finding inadequacies with using logistic regression for precise estimates of the critical temperature, we introduce a nonparametric method using  $k$ -nearest-neighbor classification as a tool to estimate the critical temperature of phase transitions from persistence images. This yields improved results.

(4) We consider a model with both an Ising-type and Berezinskii-Kosterlitz-Thouless (BKT) transition (the nematic  $XY$  model) and find that two different filtrations are required to capture the two transitions. Each filtration sees one transition, but neither is able to capture information about both transitions. We take this as evidence that the technique is not applicable entirely unsupervised; rather, care must be taken to design a filtration tuned for the problem.

The rest of the paper is organized as follows. In Sec. II we give a brief review of the techniques we use, covering persistent homology, supervised classification, finite-size scaling analysis, histogram reweighting, and bootstrapping. At the end we detail the steps of the data generation and analysis

pipeline. In Sec. III we look at the three models under consideration. In each case we give a brief review of the model and its phase transition(s) before discussing the analysis and results using logistic regression and then  $k$ -nearest-neighbors. In Sec. IV we discuss our findings and identify potential directions for future work. The Appendixes contain more detailed reviews of some of the tools we use as well as the argument demonstrating the stability of the persistence diagrams obtained using our filtrations.

## II. METHOD

### A. Background on persistent homology and persistence images

Persistent homology is a computational topology tool introduced in its modern form in [31] and popularized in [32]. It is one of the main tools of the emerging TDA field. We shall give a brief overview here, but for a more complete review of persistent homology useful references are [33–36].

Given a topological space, such as a manifold or a simplicial or cubical complex, homology can intuitively be thought of as an algebraic way of describing the “holes” in the space. In particular, the spaces we consider will be cubical complexes. A very brief technical introduction to cubical complexes and their homology can be found in Appendix A. But in general terms, given a cubical complex  $C$ , its  $k$ th cubical homology  $H_k(C)$  is a vector space which has a basis in 1-1 correspondence with the  $k$ -dimensional holes in  $C$ . Moreover, given a map of cubical complexes  $f: C \rightarrow C'$ , we obtain induced linear maps  $f_k: H_k(C) \rightarrow H_k(C')$ . The rank of  $f_k$  tells us how many of the holes survived after being mapped into  $C'$ , i.e., how many *persisted*. Given some data  $D$ , the idea of (cubical) persistence then is to construct a sequence

$$F_1(D) \rightarrow F_2(D) \rightarrow \dots \rightarrow F_N(D)$$

of cubical complexes called a filtration using the data. Typically the  $F_i(D)$  are each subcomplexes of the final complex  $F_N(D)$ ; for each cell we specify the index  $i$  at which it appears, and then  $F_i$  is the subcomplex consisting of all cells that have appeared at or before  $i$ . The maps  $F_i(D) \rightarrow F_{i+1}(D)$  are simply the inclusions.

We then apply homology to obtain a sequence of linear maps

$$H_k(F_1(D)) \rightarrow H_k(F_2(D)) \rightarrow \dots \rightarrow H_k(F_N(D)).$$

Using the ranks of these maps we can track the birth of new holes, their persistence through the filtration, and their deaths. We summarize this information as a multiset called a persistence diagram  $PH_k(F(D)) \subset \{(a, b) \in \mathbb{R}^2 \mid a \leq b\}$  which contains a pair  $(b, d)$  every time a hole is born in  $F_b(D)$  and dies in  $F_d(D)$ . We say that a feature is born at  $b$  and dies at  $d$  and that its persistence is  $d - b$ . This can also be represented as a barcode (a multiset of intervals  $[b, d)$ ). There are a few ways to define distances between persistence diagrams, but those which are most commonly used are the bottleneck and Wasserstein distances. For many typical choices of filtration a small change in the input data  $D$  leads to only a small change in the persistence diagram  $PH_k(F(D))$  as measured by these distances. This property of persistent homology is known as stability and makes persistence a useful tool for dealing with real-world, noisy data.

In the *persistent homology in configuration space* paradigm,  $D$  is the entire collection of sampled configurations, and we obtain a single persistence diagram. However, when we use persistent homology as an observable,  $D$  is a single configuration of the model we are studying. We therefore obtain a persistence diagram for each sampled configuration, and we can consider statistics computed from these diagrams. Unfortunately persistence diagrams in their raw form as multi-sets do not lend themselves to computing the typical statistics of interest such as means and variances. While there has been work developing notions of these quantities as Fréchet means and variances [37], we shall instead prefer to work with a vector representation of the diagrams known as persistence images [30] which preserve stability.

Let  $g_{a,b}: \mathbb{R}^2 \rightarrow \mathbb{R}$  denote a 2D Gaussian of standard deviation  $\sigma$  centered at  $(a, b)$ :

$$g_{a,b}(x, y) = \frac{1}{2\pi\sigma^2} \exp\left[-\frac{(x-a)^2 + (y-b)^2}{2\sigma^2}\right].$$

Given a persistence diagram  $PH_k = \{(b_i, d_i)\}_{i \in I}$ , its persistence surface is the function  $\rho_k: \mathbb{R}^2 \rightarrow \mathbb{R}$  obtained by translating each point  $(b, d) \in PH_k$  into birth-persistence coordinates  $(b, d - b)$ , then placing Gaussians with variance  $\sigma^2$  on them, weighted by the persistence of the point:

$$\rho_k(x, y) = \sum_{(b,d) \in PH_k} (d - b) g_{b,d-b}(x, y).$$

The persistence image  $PI_k$  is obtained by discretizing a rectangular region of the domain of  $\rho_k$  into a collection of  $n_l \times n_l$  pixels  $p_i$  and integrating  $\rho_k$  within each:

$$PI_k^i = \iint_{p_i} \rho_k(x, y) dx dy.$$

In this way we obtain a  $(n_l)^2$ -dimensional vector representing our persistence diagram. See Fig. 1 for an example. So long as we choose the same  $\sigma$  and discretization for each diagram, we can compute averages and variances componentwise. As observed in [38], if we are sampling data from some distribution and the expected persistence diagram has a density with respect to the Lebesgue measure on  $\{(a, b) \in \mathbb{R}^2 \mid a \leq b\}$ , then the average of the persistence images can be thought of as an estimator for this density, multiplied by an additional weighting equal to the persistence. Besides emphasizing high-persistence points, the linear weighting by the persistence ensures the stability of the persistence image. Finally we note that, as discussed in [30], machine learning models trained on persistence images are generally insensitive to the resolution and variance parameters  $n_l$  and  $\sigma$ . Therefore in this work, we shall fix the parameters with a resolution of  $30 \times 30$  and  $\sigma$  equal to 10% of a pixel. However, as a check we also performed one of the later experiments with a  $15 \times 15$  resolution, finding no significant change in the results or estimated errors.

## B. Filtrations

In this work we will be working with finite 2D lattices with a circle valued spin at each lattice site. To apply persistent homology we must choose how to define a filtration for a given configuration  $\theta = \{\theta_i\}$ , where  $\theta_i$  represents the spins as angles. Our idea is to filter the square tiling of the plane

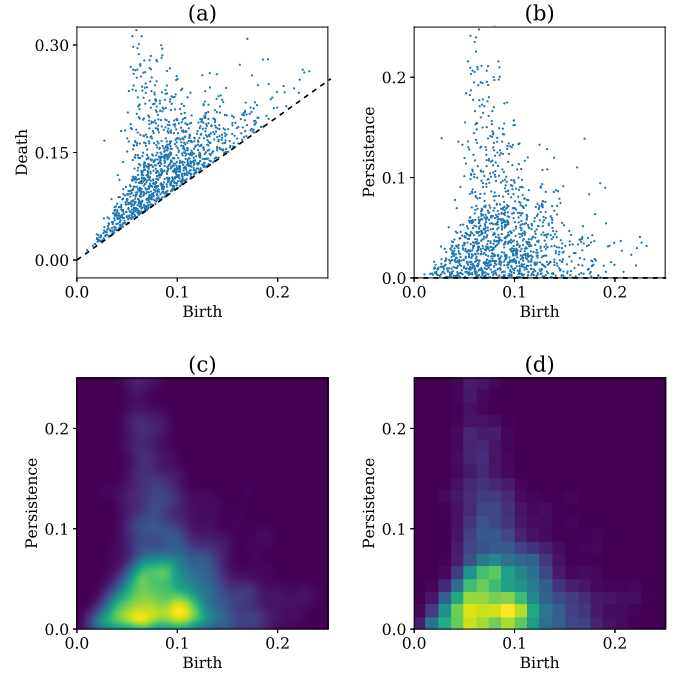


FIG. 1. An illustration of how the persistence image is obtained from a persistence diagram (a). It is first transformed into birth-persistence coordinates (b), then the persistence surface (c) is computed before discretization, yielding the persistence image (d).

corresponding to the lattice  $\Lambda$  according to the differences in neighboring spins. For each cell in this cubical complex, we will specify a time at which it appears, and then  $F_t$  is the subcomplex of the plane consisting of all cells that have appeared by time  $t$ . Denote the smallest angle between spins  $\theta_i$  and  $\theta_j$  by  $d_{ij}$ . This can also be seen as the length of the shortest arc between  $\theta_i$  and  $\theta_j$  on the unit circle. Then taking the lattice as a 2D cubical complex, we introduce each vertex  $i$  at time 0, each edge  $\langle ij \rangle$  at time  $\frac{1}{2\pi} d_{ij}$ , and each plaquette  $\square$  at time  $\max_{i,j \in \square} \frac{1}{2\pi} d_{ij}$ . We will call this the angle difference filtration. We will also introduce another similar filtration to use with the nematic XY model in Sec. III C. This will instead use a nematic angle difference  $d_{ij}^n$  which denotes the smallest angle between the spins  $\theta_i$  and  $\theta_j$  considered as directionless rods. We can think of this as the length of the shortest arc connecting the head of one spin to either the head or tail of the other spin; that is  $d_{ij}^n = \min(d_{ij}, \pi - d_{ij})$ . We will call this the nematic angle difference filtration.

The intuition behind these filtrations originally came from considering the 2D XY model. Regions of the lattice where spins vary slowly will be introduced in the angle difference filtration early, while regions containing rapidly varying spins, such as at the center of vortices, will enter the filtration later. We should expect then, at least at low temperatures, that each vortex will be manifested as a hole in the filtered lattice which is formed early on in the filtration and which gets filled in only much later: i.e., a persistent  $H_1$  class. Figure 2 shows an example of this. However, we will see that this kind of filtration can capture other structure such as spin waves or half-vortices and domain walls when we look at the nematic XY model. Moreover, compared to the point cloud filtrations used in [22,23] this class of filtrations has the computational

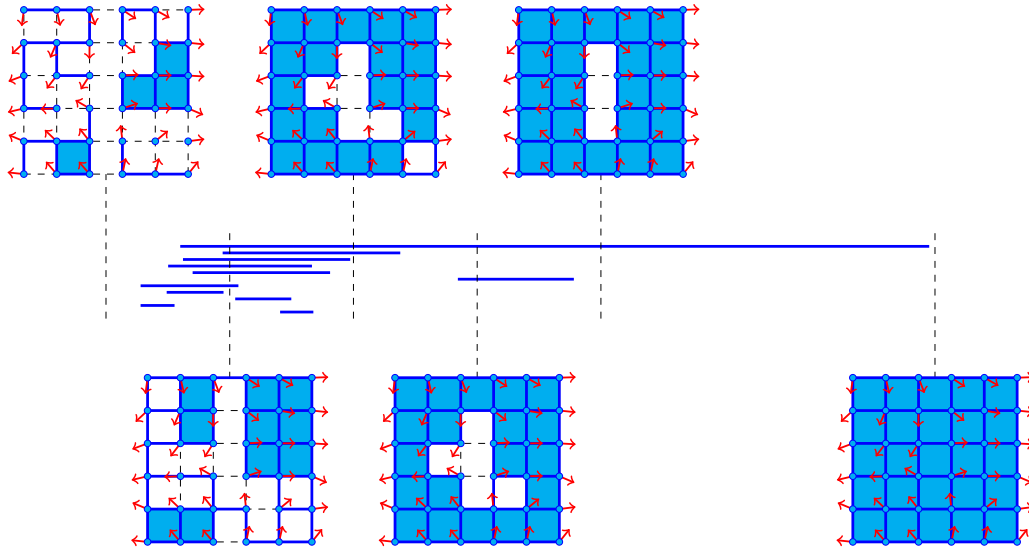


FIG. 2. An illustration of the angle difference filtration for a configuration of the  $XY$  model with an antivortex. The filtration parameter increases from left to right, and the state of the filtration is shown at six different stages. On the left-hand side only those neighboring spins which do not differ too much are connected by edges and plaquettes. As we move towards the right, more and more edges are introduced between more disparate spins. Note the correspondence between the bars and the holes in the filtration. For example, the longest bar corresponds to the hole around the antivortex in the center of the configuration. This hole is formed early as the spins far from the center vary slowly but survives until the central plaquette is added to the filtration.

benefit that edges are introduced only between neighboring lattice sites and only elementary cubes up to dimension 2 are included, greatly speeding up the computation of persistent homology. In this case the filtrations consist of subcomplexes of the plane and so contain cubes of dimension at most 2 anyway. But note that for models on higher dimensional lattices, including cubes of higher dimension in the filtration would not have any effect on  $H_1$  which is the only homological degree we use in our analysis. As discussed in Appendix B, the persistence diagrams obtained using these filtrations are stable with respect to small perturbations to the spins, in contrast to the sublevel set filtration used in [24].

**C. Logistic regression**

Following the approach introduced in [24], we will train a logistic regression model to map the persistence images obtained from configurations onto phases. Recall that logistic regression is a generalized linear model which models a binary dependent variable  $y(\mathbf{x}) \in \{0, 1\}$ . For input  $\mathbf{x} \in \mathbb{R}^N$ , a logistic regression model is parameterized by a weight vector  $\mathbf{w} = (w_1, \dots, w_N)^T \in \mathbb{R}^N$  and intercept  $b \in \mathbb{R}$ . Its output is a logistic function

$$p_{\mathbf{w},b}(\mathbf{x}) = \frac{1}{1 + e^{\mathbf{x}^T \mathbf{w} + b}} \in (0, 1),$$

which can be interpreted as the probability that  $y(\mathbf{x}) = 1$ , with  $1 - p(\mathbf{x})$  giving the probability that  $y(\mathbf{x}) = 0$ . Given training data  $\{(\mathbf{x}_i, y_i)\}$ , the weights  $\mathbf{w}$  and intercept  $b$  are learnt by minimizing a cross-entropy loss function

$$J(\mathbf{w}, b) = - \sum_i \{y_i \log(p_{\mathbf{w},b}(\mathbf{x}_i)) + (1 - y_i) \log[1 - p_{\mathbf{w},b}(\mathbf{x}_i)]\} + \frac{1}{C} (\mathbf{w}^T \mathbf{w} + b^2).$$

The first term penalizes misclassifications with the penalty increasing as the confidence in the incorrect classification increases. The second term implements  $\ell_2$  regularization, reducing overfitting by preventing the weights from becoming too large, where  $C$  is a hyperparameter controlling the amount of regularization.

In our case,  $\mathbf{x}$  will be a persistence image,  $y(\mathbf{x}) = 0$  will indicate the low-temperature phase, and  $y(\mathbf{x}) = 1$  will indicate the high-temperature phase. As in [24] we will train the model using data drawn in the low- and high-temperature phases. However, since we are interested in making a precise estimate of the critical temperature, we will use data closer to the critical region. After successful training, the weights will indicate features in the persistence image characteristic of each phase. Weights  $w_j < 0$  will indicate features of the low-temperature phase, and weights  $w_j > 0$  will indicate features of the high-temperature phase. In the intermediate range of temperatures where there is no training data, the logistic regression model will output an estimated classification  $O_{LR} \in \{0, 1\}$  depending on whether  $p$  is less than or greater than 0.5. Notice that we clamp the output to 0 or 1 rather than using the direct output of the logistic function. We find that this leads to better finite-size scaling behavior later. We may then treat  $\langle O_{LR} \rangle$  as a phase indicator, if not a true (dis)order parameter. In this work we shall be interested in the distribution  $O_{LR}$  at different temperatures and different lattice sizes.

We note that training the logistic regression model directly on raw configurations is ineffective due to the highly nonlinear nature of the system.

**D.  $k$ -nearest-neighbor classification**

We will also make use of  $k$ -nearest-neighbor ( $k$ -NN) classification to map persistence images onto phases. This is a

nonparametric model which models a categorical dependent variable  $y(\mathbf{x}) \in \mathbb{N}$ , where  $\mathbf{x} \in \mathbb{R}^N$ . The behavior of the model is determined by the training data  $\{(\mathbf{x}_i, y_i)\}$  and a choice of the hyperparameter  $k \in \mathbb{N}$ . Given new input  $\mathbf{x}$ , it finds the  $k$  indices  $i_{\mathbf{x}}^1, \dots, i_{\mathbf{x}}^k$  which minimize the Euclidean distance  $\|\mathbf{x} - \mathbf{x}_i\|_2$ . It then outputs the most common label among the  $y_{i_{\mathbf{x}}^1}, \dots, y_{i_{\mathbf{x}}^k}$ .

As in the case of logistic regression,  $\mathbf{x}$  will be a persistence image,  $y(\mathbf{x}) = 0$  will indicate the low-temperature phase, and  $y(\mathbf{x}) = 1$  will indicate the high-temperature phase. We will train the model using data drawn in the low- and high-temperature phases close to the critical region. In the intermediate range of temperatures where there is no training data, the  $k$ -NN model will output an estimated classification  $O_{kNN} \in \{0, 1\}$ . We may then treat  $\langle O_{kNN} \rangle_T$  as a phase indicator.

We note that training the classifier directly on raw configurations is not computationally feasible; doing so would require a vastly larger number of samples to sufficiently fill out the configuration space, and the computational cost of the  $k$ -NN method would consequently grow too large. The mapping from configurations to persistence images concentrates the distribution near a low-dimensional subspace, and hence  $k$ -NN becomes effective with far fewer samples.

### E. Finite-size scaling analysis

A typical approach to extracting the critical temperature and critical exponents of continuous phase transitions in spin systems is a finite-size scaling analysis of quantities such as the magnetic susceptibility

$$\chi(T) = \frac{L^2}{T} [\langle |M|^2 \rangle_T - \langle M \rangle_T^2],$$

which diverges at the critical temperature in the thermodynamic limit, where  $M = L^{-2} \sum_i (\cos \theta_i, \sin \theta_i)$  is the magnetization vector. On a finite lattice of length  $L$  this quantity will remain analytic, instead displaying a pronounced peak at a pseudocritical temperature somewhere above or below the true critical temperature  $T_c$ . As  $L \rightarrow \infty$  this peak grows taller and moves closer towards  $T_c$ . For a second-order phase transition, like that in the Ising model, the way in which the susceptibility scales close to  $T_c$  can be described by the form

$$\chi(L, t) = L^{\gamma/\nu} \hat{\chi}(L^{1/\nu} t), \quad (1)$$

where  $\hat{\chi}$  is a dimensionless function,  $t = \frac{T - T_c}{T_c}$  is the reduced temperature, and  $\gamma$  and  $\nu$  are the critical exponents for the susceptibility and correlation length, respectively. For a BKT transition, like that in the 2D XY model, it scales approximately according to

$$\chi(L, t) \approx L^{\gamma/\nu} \hat{\chi}[L \exp(-bt^{-\nu})], \quad (2)$$

where we have ignored some small logarithmic corrections. By simulating close to the phase transition on different lattice sizes  $L$  we can extract the heights and locations of the different peaks then fit these to Eq. (1) or Eq. (2) as appropriate to estimate  $T_c$ ,  $\gamma$  and  $\nu$ . Note that the logarithmic corrections we ignored in the BKT case mean that this method is not typically used for high-precision studies, where approaches based on the spin stiffness are more common.

Analogously, we might expect the persistent homology of a configuration to demonstrate large variations at criticality. We quantify this by looking at the fluctuations in the output  $O_{LR}$  and  $O_{kNN}$  of the trained logistic regression and  $k$ -NN models, measuring the variance

$$\chi_{LR}(T) = \langle O_{LR}^2 \rangle_T - \langle O_{LR} \rangle_T^2 = \langle O_{LR} \rangle_T (1 - \langle O_{LR} \rangle_T). \quad (3)$$

Note that the second equation follows since  $O_{LR}$  takes values in  $\{0, 1\}$ . This will display a peak, indicating the temperature at which the model is least certain about which phase configurations are from, when  $\langle O_{LR} \rangle_T$  crosses 0.5.  $\chi_{kNN}$  is defined similarly. We find evidence that these quantities may also display finite-size scaling behavior similar to Eqs. (1) and (2), which we will use to estimate the critical temperature  $T_c$  and the critical exponent of correlation length  $\nu$ .

We will initially assume that  $\nu$  is known and estimate the critical temperature  $T_c$  by fitting the peak temperatures  $T_c(L)$  of  $\chi_{LR}$  and  $\chi_{kNN}$  obtained from multiple lattice sizes to the ansatz

$$T_c(L) - T_c(\infty) \propto \frac{1}{L^{1/\nu}} \quad (4)$$

in the case of a second-order transition, or

$$T_c(L) - T_c(\infty) \propto \frac{1}{\log(L)^{1/\nu}} \quad (5)$$

for a BKT transition.

To estimate  $\nu$  (as well as  $T_c$ ), we will use a curve collapse approach, plotting  $y = \chi_{LR}$  or  $y = \chi_{kNN}$  for multiple lattice sizes simultaneously against  $x = L^{1/\nu} t$  (second order) or  $x = L \exp(-bt^{-\nu})$  (BKT) and finding values of  $\nu$  and  $T_c$  which minimize the distance between the curves using the Nelder-Mead method, as in the procedure described in [39].

### F. Statistical analysis

The use of histogram reweighting to extrapolate estimates of ensemble averages to an interval of temperatures around the critical temperature [40,41] and the use of bootstrap or jackknife analysis to obtain error estimates [42] are standard in quantitative investigations of phase transitions. To provide a full demonstration of a quantitative analysis based on persistent homology we will make use of both techniques, which are briefly reviewed in Appendixes C and D.

In particular, we use histogram reweighting to interpolate the outputs of our models  $\langle O_{LR} \rangle_T$  and  $\langle O_{kNN} \rangle_T$ . This allows us to obtain interpolated values of the variances  $\chi_{LR}$  and  $\chi_{kNN}$ . Assuming the sampling temperatures are close enough, this allows us to obtain a better estimate of the height and location of peaks of each quantity.

We estimate the sampling error in the training data and the sampling error in the data in the critical region independently. We do this by performing two bootstrap procedures: the first by resampling the training data, and the second by resampling the data in the critical region. In both cases we resample the data from each temperature individually. The two bootstrap procedures yield approximate sampling distributions of the quantity we are measuring, which we then turn into an error by combining the standard deviations treating the distributions as independent.

### G. Analysis pipeline

Combining the previous sections we arrive at the procedure for our analysis of each model at each lattice size.

(1) We sample the model on the given lattice size using the Wolff cluster algorithm [43] at a range of temperatures spanning the phase transition(s). We perform 50 000 Wolff cluster flips to properly thermalize the model, and 100 cluster flips between samples to ensure that the autocorrelation is negligible.

(2) For each sample, we compute persistence images with  $30 \times 30$  resolution and  $\sigma$  equal to 10% of a pixel.

(3) We use persistence images from the low- and high-temperature phases to train the logistic regression and  $k$ -NN models.

(4) Using the trained classification models, we assign a predicted phase to each sample from the critical region.

(5) Close to the peaks in the variances  $\chi_{LR}$  and  $\chi_{kNN}$  of the classifier we apply multiple histogram reweighting to obtain an interpolated curve and a more precise estimate of the location of the peak.

Once we have the interpolated variance curve and peak temperature for each of the lattice sizes, we estimate  $T_c$  and  $\nu$  by fitting the peak temperatures to the appropriate finite-size scaling ansatz [Eqs. (4) and (5)] and optimizing the data collapse of the variance curves. For each lattice size we perform two bootstraps: first by resampling the training samples, and second by resampling the samples in the critical region. In each case we resample 500 times, obtaining bootstrap distributions for the estimates of  $T_c$  and  $\nu$ . We estimate the error in these quantities by taking the square root of the sum of the variances of the two bootstrap distributions.

## III. ANALYSIS

We analyze three different variants of the 2D  $XY$  model; each undergoes a Berezinskii-Kosterlitz-Thouless (BKT) phase transition. One of the variants also exhibits a second-order transition in the Ising universality class, and it presents an interesting challenge to classify both transitions. For each model, we considered square lattices with periodic boundary conditions and linear sizes  $L = 30, 40, 50, 60, 70, 80, 100, 120, 140$ .

### A. $XY$ model

The 2D  $XY$  model is defined on an  $L \times L$  square lattice  $\Lambda$  by assigning an angle  $\theta_i \in S^1$  to each lattice site  $i \in \Lambda$ . The energy of a given configuration of spins  $\theta = \{\theta_i\}_{i \in \Lambda}$  is given by the Hamiltonian

$$H(\theta) = -J \sum_{\langle ij \rangle} \cos(\theta_i - \theta_j),$$

where  $\langle ij \rangle$  ranges over neighboring lattice sites and  $J$  is a coupling parameter we shall set equal to 1. At low temperatures spins tend to align with their neighbors but collectively twist in spin waves preventing true long-range order. Moreover a small number of vortices and antivortices, where spins twist round the full circle, may be found in bound pairs. As the temperature increases, the model undergoes a BKT transition driven by the unbinding of these vortex-antivortex pairs, so

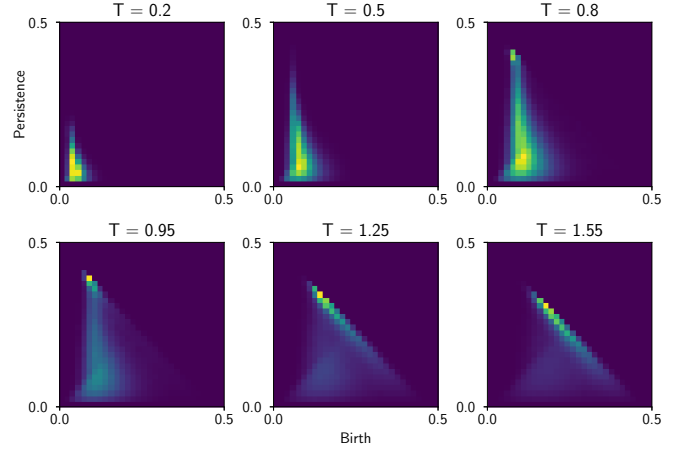


FIG. 3. The average  $H_1$  persistence images in birth-persistence coordinates at different temperatures for the  $XY$  model with  $L = 30$ .

that at high temperatures lone (anti)vortices proliferate. The critical temperature is approximately  $T = 0.8929$  [44], and the critical exponent of correlation length is  $\nu = \frac{1}{2}$ .

Using the angle difference filtration described in Sec. II B we obtain average persistence images as shown in Fig. 3. At low temperatures we see that most points in the persistence diagrams are concentrated in the lower left corner. These come from the presence of spin waves: spins tend to differ more with those in the opposite corner of a plaquette than with their immediate neighbors, producing a short-lived cycle. As the temperature increases we observe that the spin-wave cycles persist longer and longer. At around  $T = 0.8, 0.95$ , close to the downwards diagonal  $persistence = 0.5 - birth$ , or equivalently  $death = 0.5$ . These represent (anti)vortices: they are born reasonably early, as spins far away from the center vary slowly, but die much later due to the large difference in spins at the vortex core. In fact, we can check that the sum of the components of the persistence image lying on the diagonal and the two immediate subdiagonals correlates well with the absolute vorticity (the total count of vortices and antivortices) of the configurations. For example, computing the Pearson correlation coefficient on 2000 configurations at  $T = 1.0$  for  $L = 140$  yields a correlation coefficient of  $r = 0.70$ ,  $p < 0.001$ . At high temperatures we see this concentration of cycles on the diagonal increase and shift rightwards, indicating a disordered phase with many vortices.

#### 1. Logistic regression analysis

We trained logistic regression models on samples drawn from  $T = 0.85, 0.86, \text{ and } 0.87$  in the low-temperature phase, and  $T = 0.91, 0.92, \text{ and } 0.93$  in the high-temperature phase with 10 000 samples from each. The regularization hyperparameter was set to  $C = 0.001$ . We evaluated the models with 10 000 samples from each of  $T = 0.88, 0.89, \text{ and } 0.90$ . A plot of the resulting phase indicators is shown in Fig. 4 and the plot of the pseudocritical temperatures against  $\log(L)^{-2}$  is shown in Fig. 5. We do not observe any significant lattice-size dependence in the pseudocritical temperatures. They instead seem to be distributed close to  $T = 0.89$ , which is the midpoint of the

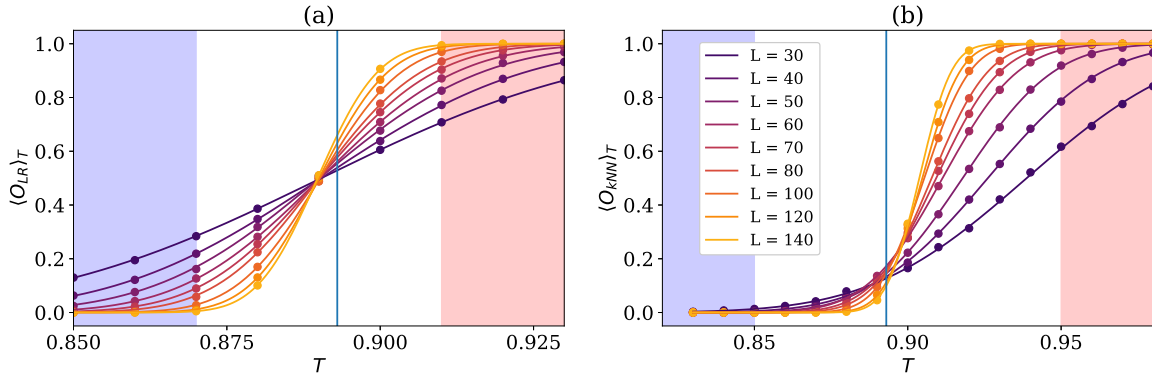


FIG. 4. Plots showing (a)  $\langle O_{LR} \rangle$  and (b)  $\langle O_{kNN} \rangle$  as a function of temperature for each lattice size for the XY model. The shaded regions indicate the temperatures used for the low- and high-temperature training data. The vertical line shows the location of the expected critical temperature  $T_c = 0.8929$ .

training temperatures. A straight line fit yields an extrapolated critical temperature of

$$T_c = 0.8872 \pm 0.0009,$$

well below the expected  $T_c = 0.8929$ . The variance curve of the phase indicator is shown in Fig. 6 and the resulting curve collapse (Fig. 7) procedure gives

$$\begin{aligned} T_c &= 0.8824 \pm 0.0001, \\ \nu &= 0.4968 \pm 0.0055, \\ b &= 0.5098 \pm 0.0068, \end{aligned}$$

not accounting within one standard deviation for the expected values of  $T_c = 0.8929$  and  $\nu = \frac{1}{2}$ .

An advantage of using a generalized linear model like logistic regression, as explored in [24], is that we can easily match the learned weights against the pixels of the persistence images. This allows us to interpret how the classifier distinguishes phases. The weights of the logistic regression model trained on the  $L = 140$  XY model data are shown in Fig. 8. We see that the low-temperature phase is characterized by cycles which are born early and which tend to have low persistence, representing spin waves. The high-temperature phase is indicated by cycles with a later birth time and persistence. In

particular, the most important region in identifying the high-temperature phase is close to  $birth = 0.1$ ,  $persistence = 0.4$ , which detects (anti)vortex cycles beginning to change behavior and move down the diagonal  $persistence = 0.5 - birth$ .

## 2. $k$ -nearest-neighbor analysis

In the case of the XY model, we found that the  $k$ -nearest-neighbor classification worked best when trained on a broad range of temperatures. We trained the models on samples drawn from  $T = 0.20, 0.25, \dots, 0.85$  in the low-temperature phase, and  $T = 0.95, 1.00, \dots, 1.60$  in the high-temperature phase with 2000 samples from each. The neighbor hyperparameter was set to  $k = 30$ . We evaluated the models with 10 000 samples from each of  $T = 0.90, 0.905, \dots, 0.95$ . A plot of the resulting phase indicators is shown in Fig. 4. The plot of the pseudocritical temperatures against  $\log(L)^{-2}$  is shown in Fig. 5. Here we see an asymptotic convergence towards a linear dependence between the pseudocritical temperatures  $T_c(L)$  and  $\log(L)^{-2}$ . Fitting a straight line to the largest three lattice sizes yields

$$T_c = 0.8935 \pm 0.0043,$$

much closer to the expected  $T_c \approx 0.8929$  than the result of the logistic regression approach. The curve collapse (Fig. 9)

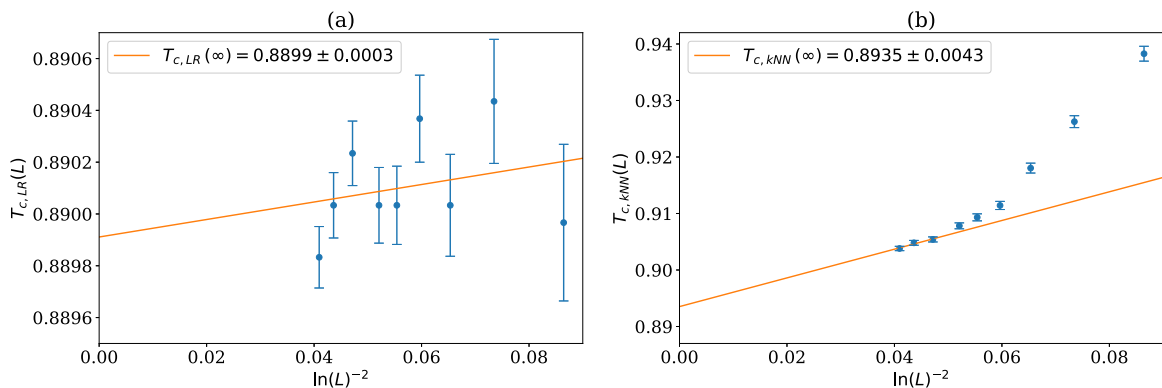


FIG. 5. Estimating the critical temperature for the XY model using (a) logistic regression and (b)  $k$ -nearest neighbors. The pseudocritical temperatures for the different lattice sizes, calculated from finding the peak of  $\chi_p$ , are fitted to the ansatz in Eq. (5). For the logistic regression we use all the lattice sizes in the fit, and for the  $k$ -nearest neighbors we use the largest three lattice sizes. The intercept gives the estimate for  $T_c(\infty)$ . Error bars are estimated by bootstrapping.

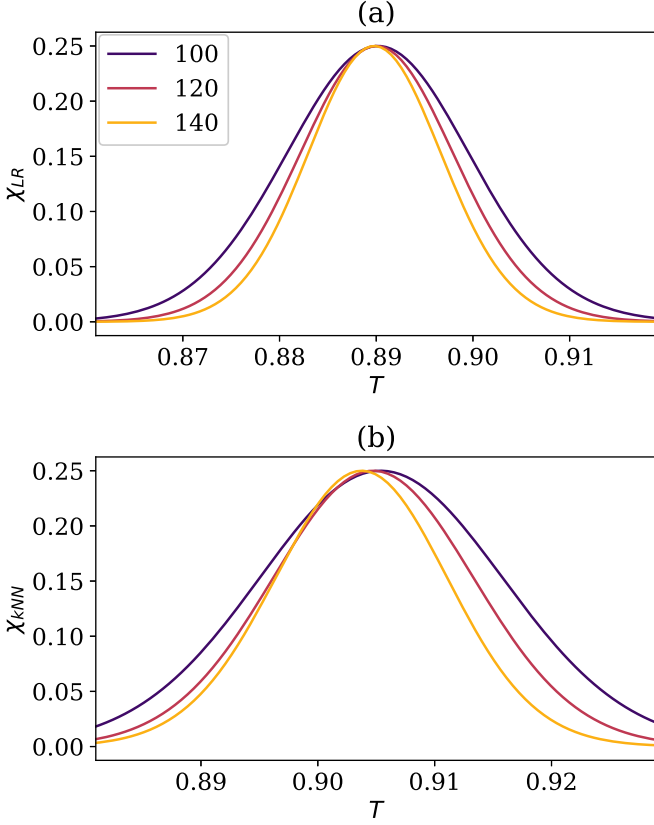


FIG. 6. Plots showing (a)  $\chi_{LR}$  and (b)  $\chi_{kNN}$  as a function of temperature for the largest three lattice sizes of the XY model. These are what we use to perform the curve collapse procedure.

procedure gives

$$\begin{aligned} T_c &= 0.8918 \pm 0.0033, \\ \nu &= 0.4972 \pm 0.0264, \\ b &= 0.5073 \pm 0.0137, \end{aligned}$$

very close to the expected values.

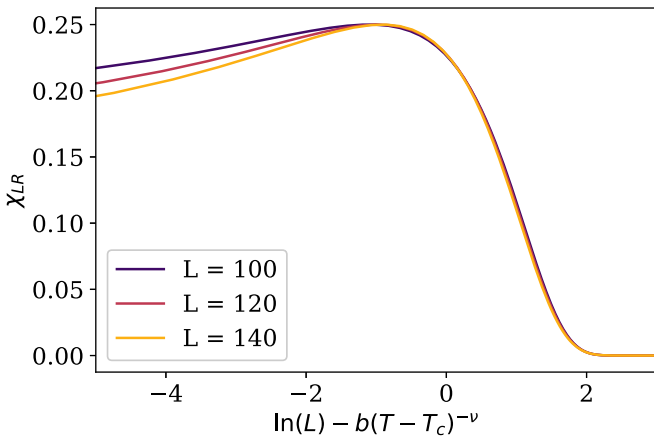


FIG. 7. The curve collapse of  $\chi_{LR}$  for the XY model with  $T_c = 0.8824$ ,  $\nu = 0.4968$ , and  $b = 0.5098$ .

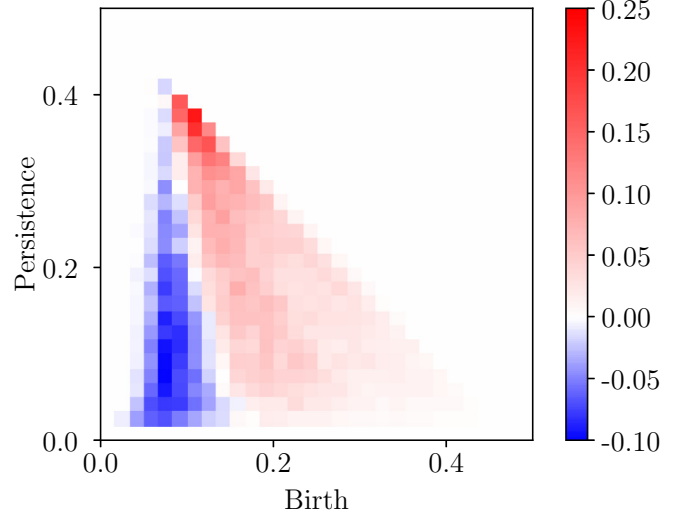


FIG. 8. The weights of the logistic regression model trained on the XY model configurations with  $L = 140$ .

### B. Constrained XY model

What we will refer to as the 2D constrained XY model was introduced and investigated in [45,46] where it is called an XY model with a topological lattice action. It is defined similarly to the classical XY model by assigning an angle  $\theta_i \in S^1$  to each lattice site  $i \in \Lambda$  of an  $L \times L$  square lattice  $\Lambda$ . However, the Hamiltonian is defined as

$$H(\theta) = \begin{cases} 0 & \text{if } \frac{1}{2\pi} |\theta_i - \theta_j| \leq \delta \text{ for all } (i, j) \\ \infty & \text{otherwise.} \end{cases}$$

Therefore all configurations are constrained so that the spins at neighboring sites cannot differ by more than  $\delta$ . Since the partition function does not depend on the thermodynamic temperature, we consider the parameter  $\delta$  as taking on this role instead, and the model undergoes a BKT transition as  $\delta$  increases at approximately  $\delta = 0.2825$  [46] with  $\nu = \frac{1}{2}$ . Notice that while  $\delta < 0.25$  no (anti)vortices may form.

Using the angle difference filtration described in Sec. II B we obtain average persistence images as shown in Fig. 10. We

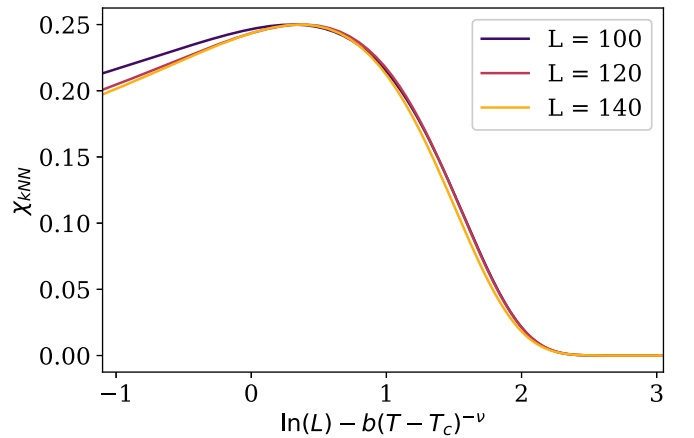


FIG. 9. The curve collapse of  $\chi_{kNN}$  for the XY model with  $T_c = 0.8918$ ,  $\nu = 0.4972$ , and  $b = 0.5073$ .



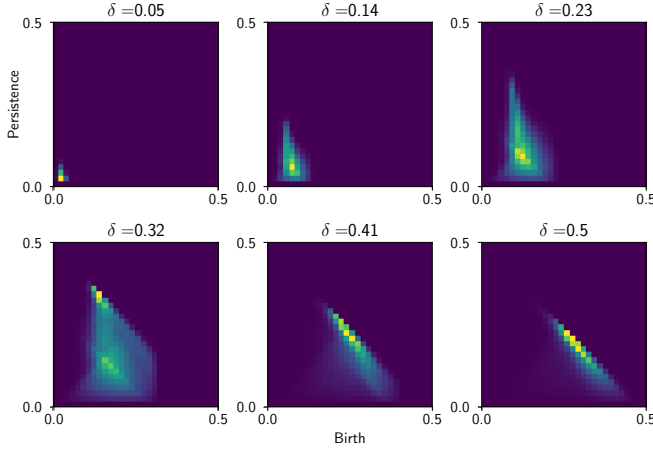


FIG. 10. The average  $H_1$  persistence image in birth-persistence coordinates at different deltas for the constrained XY model with  $L = 30$ .

immediately see a resemblance with the persistence images obtained for the XY model in Fig. 3 except that we see a cutoff effect at  $birth = \delta$ , since by this point all neighboring lattice sites must have been connected in the filtration. For this model we must adjust our methodology slightly since histogram reweighting is not possible. Instead we will sample deltas more densely, then to extract the maximums of  $\chi_{LR}$  and  $\chi_{kNN}$  we will fit a parabola to the three highest points.

### 1. Logistic regression analysis

We trained logistic regression models on samples drawn from  $\delta = 0.27, 0.272, \dots, 0.28$  in the low delta phase, and  $\delta = 0.286, 0.288, \dots, 0.296$  in the high delta phase with 4000 samples from each. The regularization hyperparameter was set to  $C = 0.001$ . We evaluated the models with 4000 samples from each of  $\delta = 0.27, 0.271, \dots, 0.296$ . A plot of the resulting phase indicators is shown in Fig. 11. The plot of the pseudocritical deltas against  $\log(L)^{-2}$  is shown in Fig. 12. We do not observe any significant lattice-size dependence in the pseudocritical deltas. They instead seem to be distributed close to  $\delta = 0.283$  which is the midpoint of the training deltas.

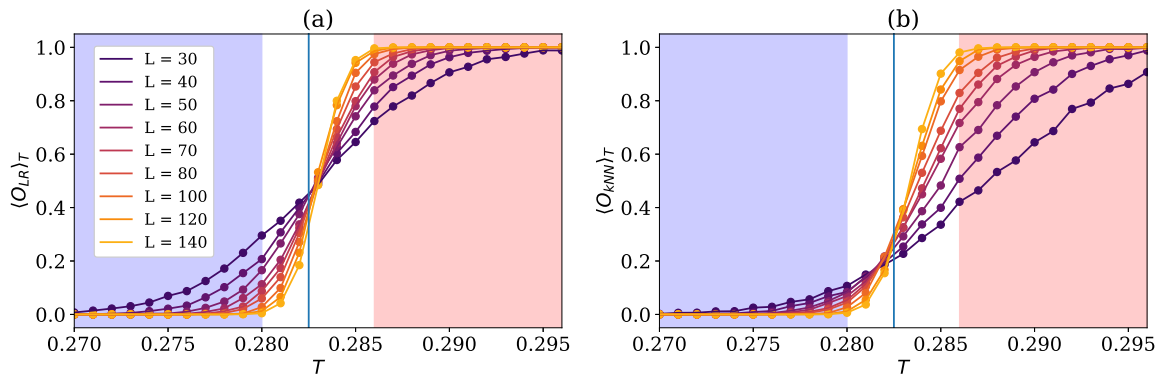


FIG. 11. Plots showing (a)  $\langle O_{LR} \rangle$  and (b)  $\langle O_{kNN} \rangle$  as a function of delta for each lattice size for the constrained XY model. The shaded regions indicate the deltas used for the low and high delta training data. The vertical line shows the location of the expected critical delta  $\delta_c = 0.2825$ . Note that for the  $k$ -NN plot the training regions extend farther away than what is shown.

The curve collapse (Fig. 13) procedure gives

$$\delta_c = 0.2843 \pm 0.0013,$$

$$\nu = 0.4999 \pm 0.0189,$$

$$b = 0.3009 \pm 0.0041,$$

which is not likely to account for the expected value of  $\delta_c = 0.2825$  but does support  $\nu = \frac{1}{2}$ .

The weights of the logistic regression model trained for  $L = 140$  are shown in Fig. 14. We observe a similarity to the weights learnt for the XY model in Fig. 8 although in this case it appears to be more difficult to delineate which regions of the persistence images indicate the two phases.

### 2. $k$ -nearest-neighbor analysis

We trained the  $k$ -nearest-neighbor models on samples drawn from  $\delta = 0.27, 0.272, \dots, 0.28$  in the low delta phase, and  $\delta = 0.286, 0.288, \dots, 0.296$  in the high delta phase with 4000 samples from each. The neighbor hyperparameter was set to  $k = 30$ . We evaluated the models with 4000 samples from each of  $\delta = 0.27, 0.271, \dots, 0.296$ . A plot of the resulting phase indicators is shown in Fig. 11. The plot of the pseudocritical deltas against  $\log(L)^{-2}$  is shown in Fig. 12. Here we see an asymptotic convergence towards a linear dependence between the pseudocritical deltas  $\delta_c(L)$  and  $\log(L)^{-2}$ . Fitting a straight line to the largest three lattice sizes yields

$$\delta_c = 0.2821 \pm 0.0014.$$

The curve collapse (Fig. 15) procedure gives

$$\delta_c = 0.2818 \pm 0.0017,$$

$$\nu = 0.5003 \pm 0.0206,$$

$$b = 0.5022 \pm 0.0048,$$

very close to the expected values.

### C. Nematic XY model

There are a variety of generalized XY models with nematic interactions. We will consider the model with Hamiltonian

$$H(\theta) = - \sum_{\langle ij \rangle} \{ \Delta \cos(\theta_i - \theta_j) + (1 - \Delta) \cos[2(\theta_i - \theta_j)] \},$$

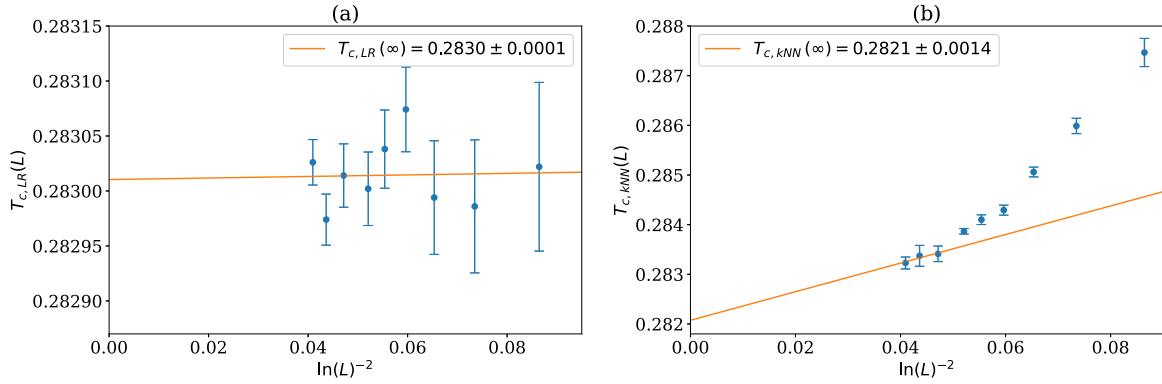


FIG. 12. Estimating the critical delta for the constrained XY model using (a) logistic regression and (b)  $k$ -nearest neighbors. The pseudocritical deltas for the different lattice sizes, calculated from finding the peak of  $\chi_p$ , are fitted to the ansatz in Eq. (5). For the logistic regression we use all the lattice sizes in the fit, and for the  $k$ -nearest neighbors we use the largest three lattice sizes. The intercept gives the estimate for  $\delta_c(\infty)$ . Error bars are estimated by bootstrapping.

where we will fix  $\Delta = 0.15$ . The first term is the usual XY interaction, but the second term is a nematic interaction which remains invariant when any individual spin is rotated 180 degrees. We can imagine this as an interaction between the spins considered as headless rods: spins which are parallel contribute less energy, even if they point in opposite directions. The  $T$ - $\Delta$  phase diagram of this model is explored in [47–49], and we see that at our chosen  $\Delta = 0.15$ , it undergoes two phase transitions as temperature increases. The first is an Ising-type transition from a magnetic phase to a nematic phase at  $T \approx 0.3314$  (as estimated using the magnetic susceptibility) resulting in (anti)vortices (which remain bound into vortex-antivortex pairs) stretching into domain walls with a half-(anti)vortex at each end; across the wall the spins flip by  $\pi$ . See Fig. 16 for an example. The second is a BKT transition to a paramagnetic phase at  $T \approx 0.7808$  (as estimated using the magnetic susceptibility) driven by the unbinding of these pairs of now-elongated vortices and antivortices.

Following the intuition developed in Sec. II B, we consider two different filtrations: The first is the angle difference filtration used for the XY and constrained XY model, where each edge  $(ij)$  of the lattice is added into the filtration at time  $\frac{1}{2\pi}d_{ij}$ ; The second is the nematic angle difference filtration which

uses the nematic distance between spins, adding in edges at time  $\frac{1}{2\pi}d_{ij}^n = \min(\frac{1}{2\pi}d_{ij}, 0.5 - \frac{1}{2\pi}d_{ij})$ . The resulting average persistence images are shown in Figs. 17 and 18, respectively.

From Fig. 17 we see that the magnetic-nematic transition is manifested in the angle difference filtration by the emergence of a cluster in the bottom right of the persistence image and the rightwards movement of the cluster in the top left. These correspond to the appearance of domain walls in configurations. In particular, at a time close to 0.5 in the filtration, the edges which cross domain walls will get added all at once, forming many short-lived cycles. Meanwhile, (anti)vortices get stretched out into strings so that more spins must be connected in the filtration before a hole is formed, generally causing the time at which this happens to increase a little. There is little qualitative difference between the images across the BKT transition, however. In Fig. 18 we see a familiar picture of the BKT transition which is very similar to that observed in the XY model and constrained XY model, while the Ising-type transition is not detectable at all. We also looked at

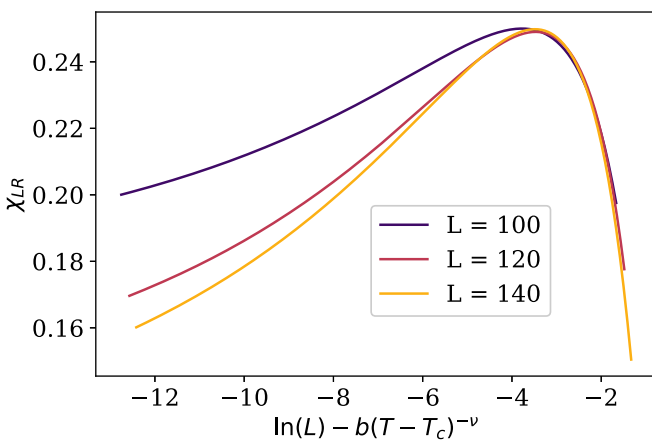


FIG. 13. The curve collapse of  $\chi_{LR}$  for the constrained XY model with  $\delta_c = 0.2843$ ,  $\nu = 0.4999$ , and  $b = 0.3009$ .

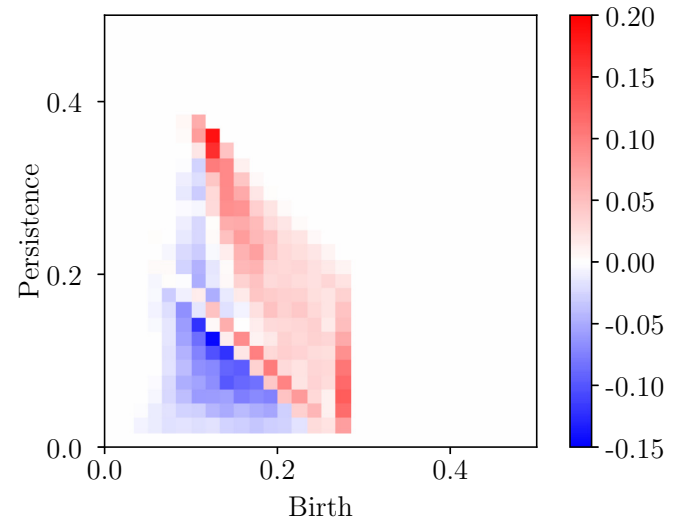


FIG. 14. The weights of the logistic regression model trained on configurations over the BKT transition in the constrained XY model with  $L = 140$ .

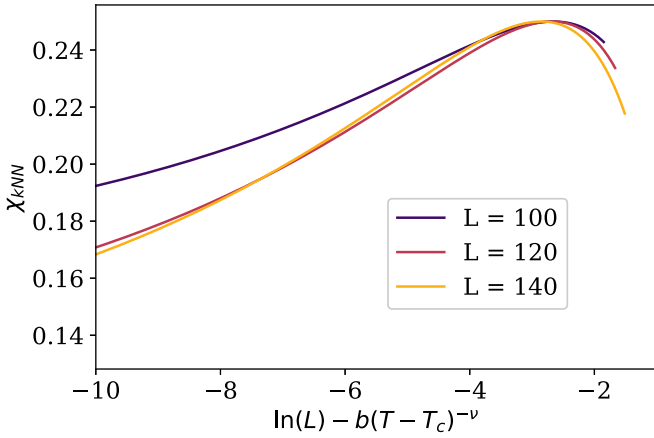


FIG. 15. The curve collapse of  $\chi_{kNN}$  for the constrained XY model with  $\delta_c = 0.2818$ ,  $\nu = 0.5003$ , and  $b = 0.3022$ .

a combined angle difference filtration using  $\frac{\Delta}{2\pi}d_{ij} + \frac{1-\Delta}{2\pi}d_{ij}^n$ , but while this did seem to detect both phase transitions, it was difficult to effectively train the classification models to identify two phases at a time.

**1. Logistic regression analysis of magnetic-nematic transition**

We trained logistic regression models on samples drawn from  $T = 0.32$  and  $0.3225$  in the low-temperature phase, and  $T = 0.3425$  and  $0.345$  in the high-temperature phase with 10 000 samples from each. The regularization hyperparameter was set to  $C = 10^{-6}$ . We evaluated the models with 10 000 samples from each of  $T = 0.33, 0.33125, \dots, 0.335$ . A plot of the resulting phase indicators is shown in Fig. 19. The plot of the pseudocritical temperatures against  $L^{-1}$  is shown in Fig. 20. For the lower lattice sizes  $L < 60$ , we do not observe any significant lattice-size dependence in the pseudocritical temperatures. They instead seem to be distributed close to

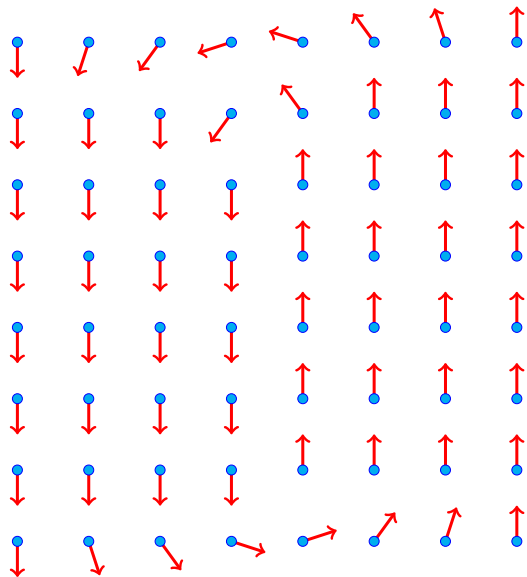


FIG. 16. A configuration with a vortex that has stretched out into two half-vortices separated by a domain wall.

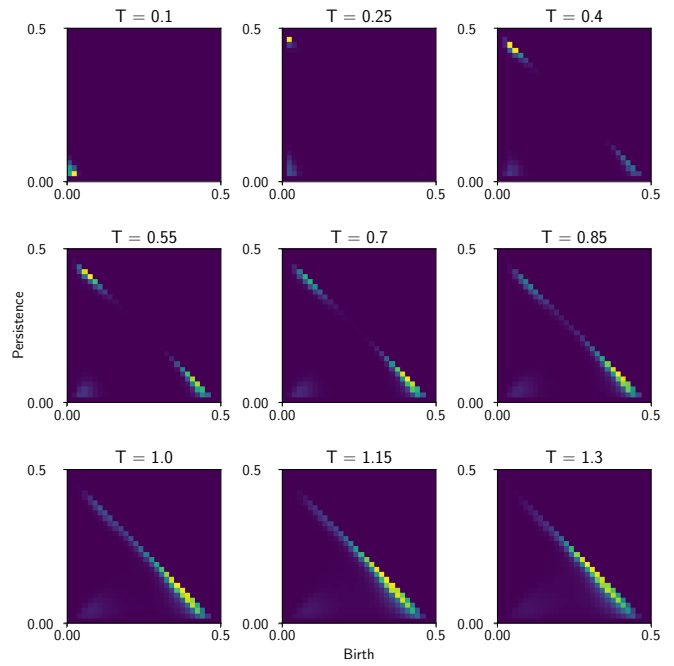


FIG. 17. The average  $H_1$  persistence image in birth-persistence coordinates at different temperatures for the nematic XY model with  $L = 30$  using the angle difference filtration. The magnetic-nematic phase transition occurs between the middle and right images on the top row and the nematic-paramagnetic BKT transition occurs between the middle and right images on the middle row.

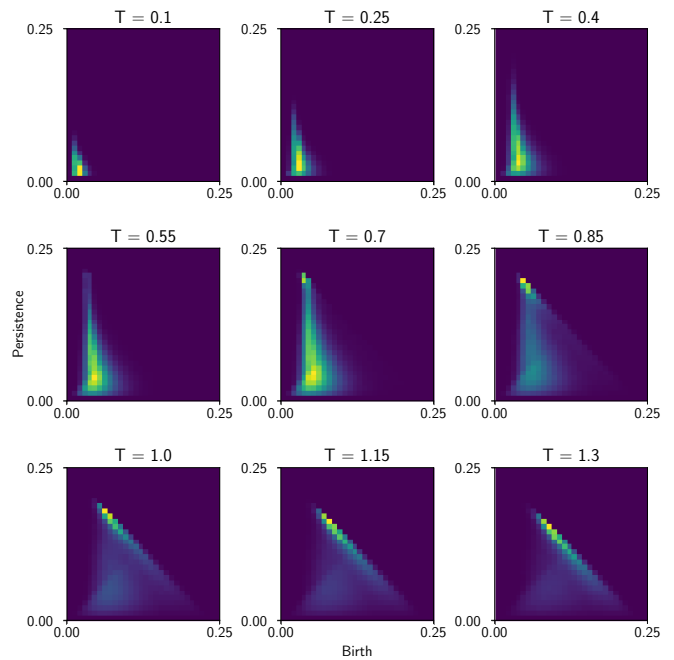


FIG. 18. The average  $H_1$  persistence image in birth-persistence coordinates at different temperatures for the nematic XY model with  $L = 30$  using the nematic angle difference filtration. The magnetic-nematic phase transition occurs between the middle and right images on the top row and the nematic-paramagnetic BKT transition occurs between the middle and right images on the middle row. Note the similarity with Figs. 3 and 10.

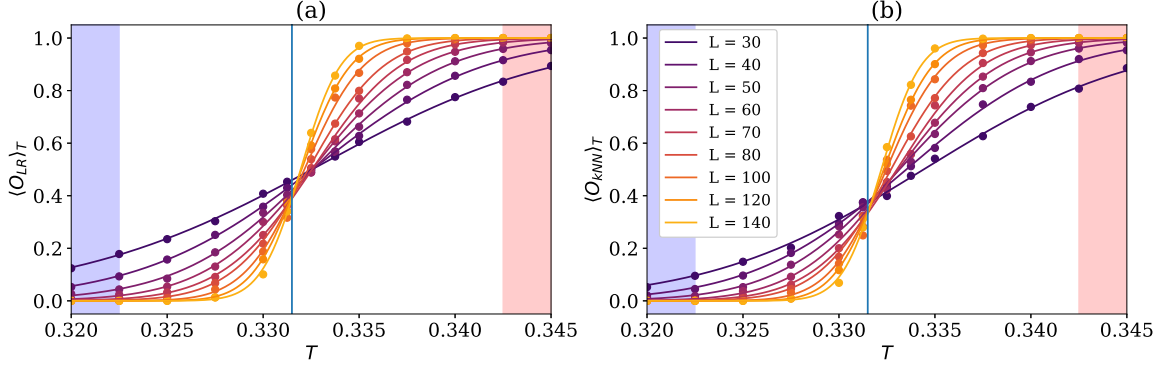


FIG. 19. Plots showing (a)  $\langle O_{LR} \rangle_T$  and (b)  $\langle O_{kNN} \rangle_T$  as a function of temperature for each lattice size for the magnetic-nematic transition in the nematic XY model. The shaded regions indicate the temperatures used for the low- and high-temperature training data. The vertical line shows the location of the expected critical temperature  $T_c = 0.3314$ .

$T = 0.3325$ , which is the midpoint of the training temperatures. At the larger lattice sizes  $L \geq 60$ , a linear dependence on  $L^{-1}$  emerges. Fitting a line to the largest four lattice sizes yields an extrapolated critical temperature of

$$T_c = 0.3314 \pm 0.0001.$$

The curve collapse (Fig. 21) procedure gives

$$\begin{aligned} T_c &= 0.3315 \pm 0.0001, \\ \nu &= 0.8562 \pm 0.0102. \end{aligned}$$

While these estimates of the critical temperature are good, the expected value of  $\nu = 1$  doesn't fall within the error bars estimated with this approach.

The weights of the logistic regression model trained for  $L = 140$  are shown in Fig. 22. We observe that the classifier learns to detect exactly what we saw in Fig. 17, namely, a rightwards shift of the upper left cluster, and the emergence of a cluster in the bottom right, corresponding to domain walls forming in the configurations.

### 2. $k$ -nearest-neighbor analysis of magnetic-nematic transition

We trained the  $k$ -nearest-neighbor models on samples drawn from  $T = 0.32$  and  $0.3225$  in the low-temperature

phase, and  $T = 0.3425$  and  $0.345$  in the high-temperature phase with 4000 samples from each. The neighbor hyperparameter was set to  $k = 30$ . We evaluated the models with 4000 samples from each of  $T = 0.33, 0.33125, \dots, 0.335$ . A plot of the resulting phase indicators is shown in Fig. 19. The plot of the pseudocritical temperatures against  $L^{-1}$  is shown in Fig. 20. Here we see that for  $L \geq 60$ , the pseudocritical temperatures fit reasonably well on a straight line when plotted against  $L^{-1}$ . Fitting a line to the largest four lattice sizes yields

$$T_c = 0.3315 \pm 0.0002.$$

The curve collapse (Fig. 23) procedure gives

$$\begin{aligned} T_c &= 0.3316 \pm 0.0002, \\ \nu &= 0.9551 \pm 0.0196, \end{aligned}$$

very close to the expected value of  $T_c = 0.3314$ , but not quite compatible with  $\nu = 1$  although better than the logistic regression result.

### 3. Logistic regression analysis of nematic-paramagnetic transition

We trained logistic regression models on samples drawn from  $T = 0.74, 0.75$ , and  $0.76$  in the low-temperature phase, and  $T = 0.8, 0.81$  and  $0.82$  in the high-temperature phase

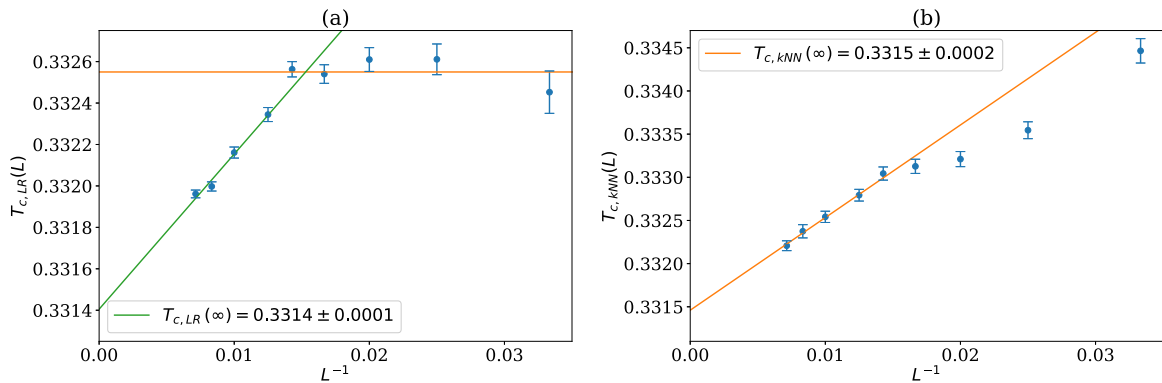


FIG. 20. Estimating the critical temperature for the magnetic-nematic transition in the nematic XY model using (a) logistic regression and (b)  $k$ -nearest neighbors. The pseudocritical temperatures for the different lattice sizes, calculated from finding the peak of  $\chi_p$ , are fitted to the ansatz in Eq. (4). In both cases we use the largest four lattice sizes for the fit. The intercept gives the estimate for  $T_c(\infty)$ . Error bars are estimated by bootstrapping.

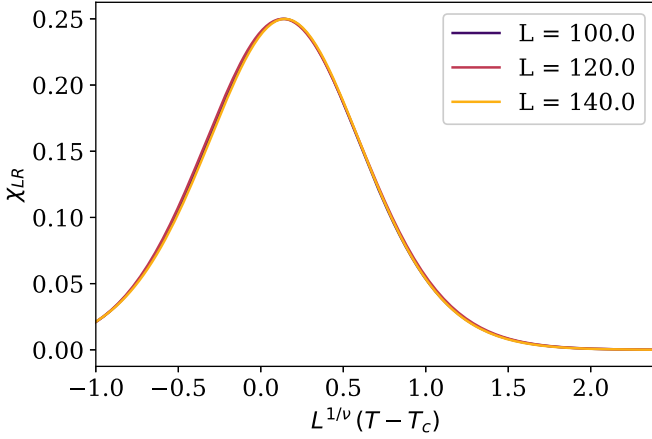


FIG. 21. The curve collapse of  $\chi_{LR}$  for the magnetic-nematic transition in the nematic  $XY$  model with  $T_c = 0.3315$  and  $\nu = 0.8562$ .

with 10 000 samples from each. The regularization hyperparameter was set to  $C = 0.001$ . We evaluated the models with 10,000 samples from each of  $T = 0.74, 0.75, \dots, 0.82$ . A plot of the resulting phase indicators is shown in Fig. 24. The plot of the pseudocritical temperatures against  $\log(L)^{-2}$  is shown in Fig. 25. We do not observe any significant lattice-size dependence in the pseudocritical temperatures. They instead seem to be distributed just above to  $T = 0.78$ , which is the midpoint of the training temperatures. While a straight line fit to all but the smallest lattice size yields an extrapolated critical temperature of

$$T_c = 0.7804 \pm 0.0002,$$

not too far from the expected  $T_c \approx 0.7808$ , Fig. 25 does not suggest that increasing the statistics would lead to increased accuracy. However, the curve collapse (Fig. 26) procedure

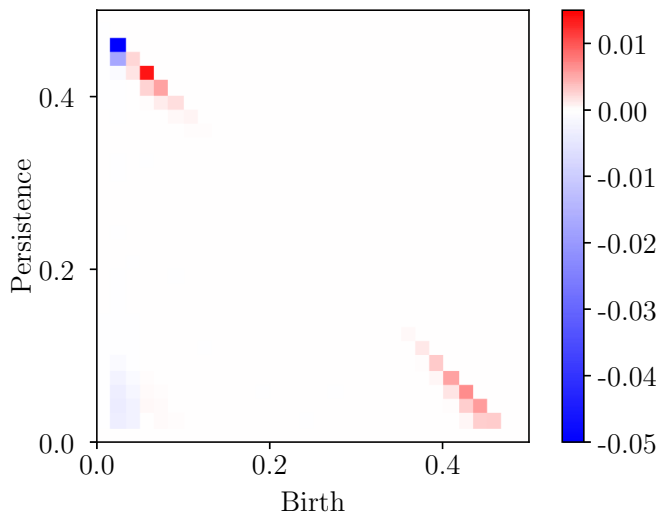


FIG. 22. The weights of the logistic regression model trained on configurations over the magnetic-nematic transition in the nematic  $XY$  model with  $L = 140$ .

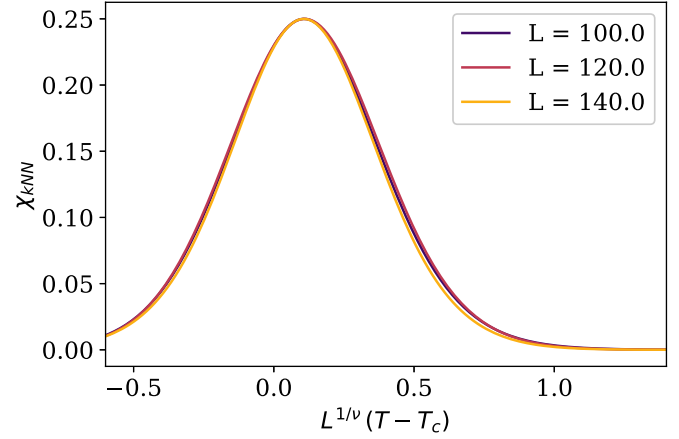


FIG. 23. The curve collapse of  $\chi_{kNN}$  for the magnetic-nematic transition in the nematic  $XY$  model with  $T_c = 0.3316$  and  $\nu = 0.9551$ .

gives

$$\begin{aligned} T_c &= 0.7803 \pm 0.0025, \\ \nu &= 0.5107 \pm 0.0101, \\ b &= 0.3037 \pm 0.0076, \end{aligned}$$

accounting for the expected value of  $T_c = 0.7808$ , but giving a potentially questionable result for  $\nu = \frac{1}{2}$  which lies just outside one standard deviation.

The weights of the logistic regression model trained for  $L = 140$  are shown in Fig. 27. We note the similarity to the weights learnt for the  $XY$  model in Fig. 8 except now the region in the top left represents half-vortices and half-antivortices which change behavior, shifting down to the right as temperature increases and they unbind.

#### 4. $k$ -nearest-neighbor analysis of nematic-paramagnetic transition

Similarly to the case of the  $XY$  model, we found that the  $k$ -nearest-neighbor classification worked best when trained on a broad range of temperatures. We trained the models on samples drawn from  $T = 0.5, 0.55, \dots, 0.7$  in the low-temperature phase, and  $T = 0.85, 0.9, \dots, 1.05$  in the high-temperature phase with 2000 samples from each. The neighbor hyperparameter was set to  $k = 30$ . We evaluated the models with 10 000 samples from each of  $T = 0.74, 0.75, \dots, 0.82$ . A plot of the resulting phase indicators is shown in Fig. 24. The plot of the pseudocritical temperatures against  $\log(L)^{-2}$  is shown in Fig. 25. Here we see an asymptotic convergence towards a linear dependence between the pseudocritical temperatures  $T_c(L)$  and  $\log(L)^{-2}$ . Fitting a straight line to the largest four lattice sizes yields

$$T_c = 0.7766 \pm 0.0034.$$

While this is further from the expected  $T_c \approx 0.7808$  than the result of the logistic regression approach, the approach towards the correct finite-size scaling is much clearer. The curve collapse (Fig. 28) procedure gives

$$\begin{aligned} T_c &= 0.7757 \pm 0.0064, \\ \nu &= 0.4983 \pm 0.0226, \\ b &= 0.3051 \pm 0.0083, \end{aligned}$$

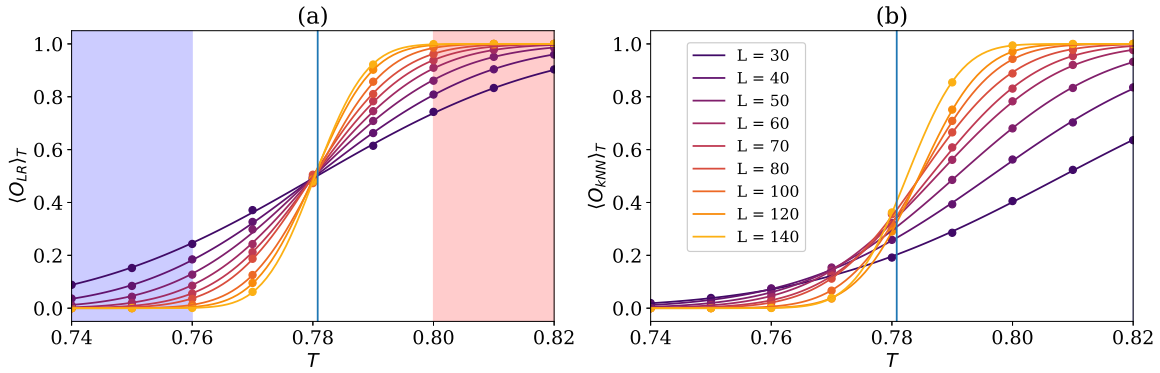


FIG. 24. Plots showing (a)  $\langle O_{LR} \rangle_T$  and (b)  $\langle O_{kNN} \rangle_T$  as a function of temperature for each lattice size for the nematic-paramagnetic transition in the nematic  $XY$  model. The shaded regions indicate the temperatures used for the low- and high-temperature training data. The vertical line shows the location of the expected critical temperature  $T_c = 0.7808$ . Note that for the  $k$ -nearest-neighbor case the training regions lie outside the bounds of the plot.

which is compatible with the expected values of  $T_c = 0.7808$  and  $\nu = \frac{1}{2}$ .

#### IV. CONCLUSIONS AND DISCUSSION

We have introduced a way of applying persistent homology to analyze the configurations of lattice spin models, investigating the phase transitions in the 2D  $XY$  model with three different Hamiltonians: the standard action, a topological lattice action, and a modified standard action with an additional nematic interaction term. In each case we were able to successfully identify the phase transition and estimate its critical temperature and critical exponent of the correlation length by considering the finite-size scaling of observables derived from the persistent homology of configurations. In particular we trained logistic regression and  $k$ -nearest-neighbor classifiers to identify the phases of the models from persistence images. The critical point was estimated as the temperature at which the variance in the classification reached a maximum.

We have found that the previously introduced approach of using logistic regression for classification, while useful for interpreting which regions of the persistence image indicate the different phases, fails to produce accurate estimates of

the critical temperature or exponents in the case of the BKT transitions. Instead it will tend to yield the midpoint between the low- and high-temperature training temperatures as the critical temperature. Indeed, using different temperatures for the training causes the estimated critical temperature to shift accordingly. We believe this failure is because logistic regression is a generalized linear model and the data here are highly nonlinear. On the other hand, the nonparametric  $k$ -nearest-neighbor approach generally produces good results, with a clear asymptotic approach towards the expected finite-size scaling behavior in all cases.

There are a number of interesting questions and directions for further research:

(1) The approach presented in this paper could easily be extended to other lattice spin models, but it would also be interesting to see if the filtration presented in Sec. II B could be adapted to more complex models such as those from lattice gauge theory.

(2) The similarity of the persistence images across the BKT transition in all three models raises the question of the extent to which the persistence is a universal quantity. This could potentially facilitate a transfer learning approach where classifiers trained on one model can identify phase transitions

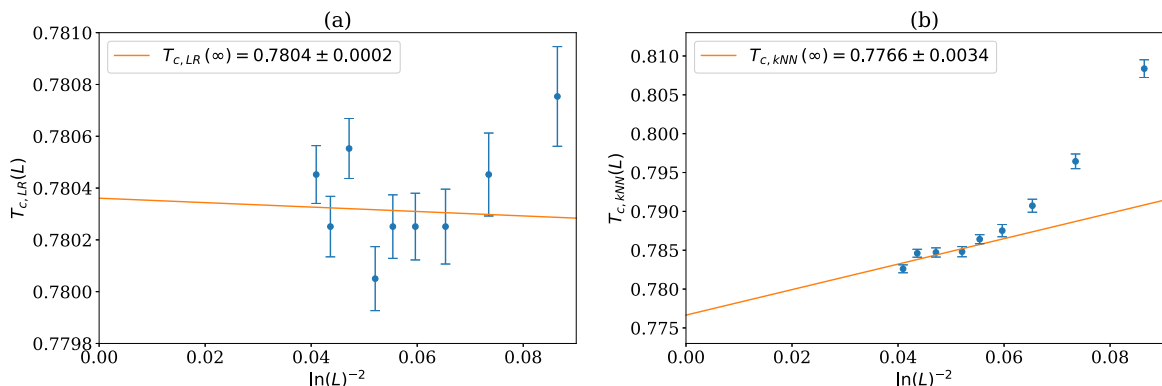


FIG. 25. Estimating the critical temperature for the nematic-paramagnetic transition in the nematic  $XY$  model using (a) logistic regression and (b)  $k$ -nearest neighbors. The pseudocritical temperatures for the different lattice sizes, calculated from finding the peak of  $\chi_p$ , are fitted to the ansatz in Eq. (5). For the logistic regression we use all the lattice sizes except the smallest in the fit, and for the  $k$ -nearest neighbors we use the largest four lattice sizes. The intercept gives the estimate for  $T_c(\infty)$ . Error bars are estimated by bootstrapping.

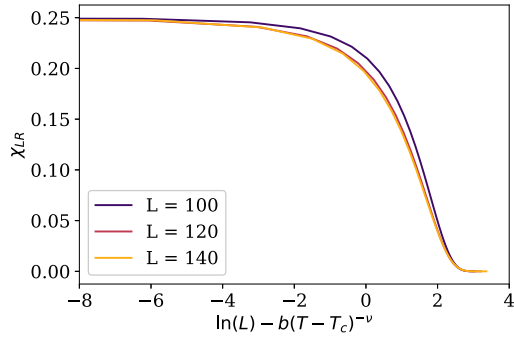


FIG. 26. The curve collapse of  $\chi_{LR}$  for the nematic-paramagnetic transition in the nematic XY model with  $T_c = 0.7803$ ,  $\nu = 0.5107$ , and  $b = 0.3037$ .

of the same universality class in another model (see, e.g., [11]).

(3) It could also be investigated if the use of a vectorization and a classifier is necessary in the first place. There is a notion of variance for persistence diagrams called Fréchet variance [37] which might show finite-size scaling behavior directly. However, this is computationally expensive to measure.

(4) Finally, we note that there have been a variety of different filtrations used to compute the persistent homology of configurations of lattice spin models. It would be interesting to see how these perform and complement one another on a single data set.

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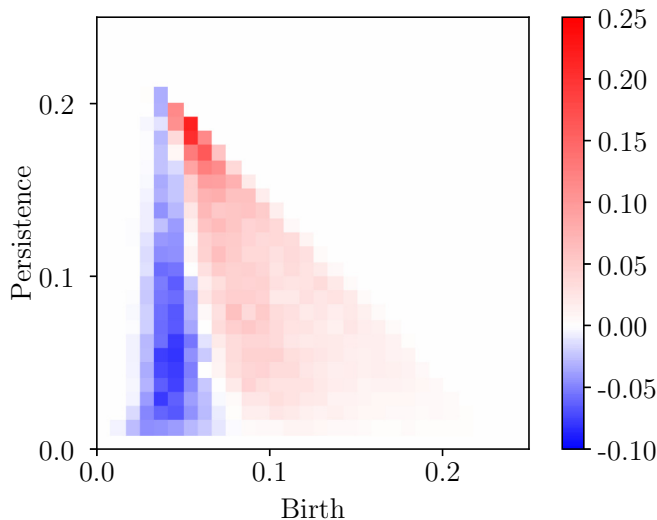


FIG. 27. The weights of the logistic regression model trained on configurations over the nematic-paramagnetic transition in the nematic XY model with  $L = 140$ .

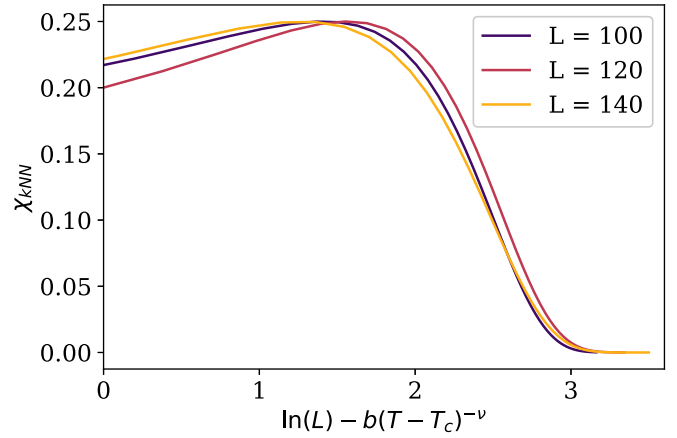


FIG. 28. The curve collapse of  $\chi_{kNN}$  for the nematic-paramagnetic transition in the nematic XY model with  $T_c = 0.7757$ ,  $\nu = 0.4983$ , and  $b = 0.3051$ .

performed using giotto-tda [50]. N.S. has been supported by a Swansea University Research Excellence Scholarship (SURES). J.G. was supported by EPSRC Grant No. EP/R018472/1. B.L. received funding from the European Research Council (ERC) under the European Union's Horizon 2020 research and innovation program under Grant Agreement No. No 813942. The work of B.L. was further supported in part by the UKRI Science and Technology Facilities Council (STFC) Consolidated Grant No. ST/T000813/1, by the Royal Society Wolfson Research Merit Award No. WM170010, and by the Leverhulme Foundation Research Fellowship RF-2020-4619.

#### APPENDIX A: CUBICAL COMPLEXES AND HOMOLOGY

This is a very compressed version of the exposition found in [51]. An *elementary interval* is an interval of the form  $[i, i + 1] \subset \mathbb{R}$  (*nondegenerate*) or  $[i, i] = \{n\}$  (*degenerate*) for some choice of  $i \in \mathbb{Z}$ . An *elementary cube* is a finite product of elementary intervals  $Q = I_1 \times \cdots \times I_n \subset \mathbb{R}^n$ , where  $n$  is some fixed *embedding dimension*. Its *dimension*  $\dim Q$  is the number of nondegenerate intervals in the product. A *cubical complex*  $C$  is a subset of  $\mathbb{R}^n$  which is a union of elementary cubes. Specifying a field  $\mathbf{F}$ , we define  $\mathbf{F}$ -vector spaces  $C_k = \{\sum \alpha_i Q_i \mid Q_i \subseteq C, \dim Q_i = k, \alpha_i \in \mathbf{F}\}$  for each  $k \in \mathbb{N}$ , consisting of finite formal sums of elementary cubes. The *boundary* of a nondegenerate elementary interval is given by the formal sum  $\partial[i, i + 1] = [i + 1, i + 1] - [i, i]$ . For a degenerate elementary interval the boundary is zero. The boundary of an elementary cube  $Q = (I_1 \times \cdots \times I_n)$  is a formal sum

$$\partial Q = \sum_{j=1}^n (-1)^{\sum_{i=1}^{j-1} \dim Q_i} Q_i (I_1 \times \cdots \times \partial I_j \times \cdots \times I_n), \quad (\text{A1})$$

where we consider  $\times$  as distributing over the formal summation. We can see that for  $\dim Q \geq 1$  we have  $\dim \partial Q = \dim Q - 1$ . Therefore we can extend  $\partial$  to linear maps  $\partial_k : C_k \rightarrow C_{k-1}$  via the mapping  $\sum \alpha_i Q_i \mapsto \sum \alpha_i (\partial Q_i)$ . Since  $\partial \partial I = 0$  for any elementary interval  $I$ , we also see that  $\partial_k \circ \partial_{k+1} = 0$  for all  $k \in \mathbb{N}$ , so that  $\text{im } \partial_{k+1} \subseteq \ker \partial_k$ . See Fig. 29

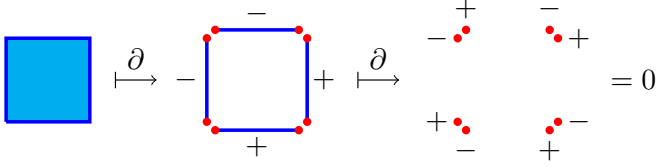


FIG. 29. Example of how the boundary operator  $\partial$  acts on a simple cubical complex consisting of a single 2D cube. Note how the sum in Eq. (A1) being alternating ensures that  $\partial\partial = 0$ .

for an example. A sequence of linear maps

$$\dots \rightarrow C_3 \xrightarrow{\partial_3} C_2 \xrightarrow{\partial_2} C_1 \xrightarrow{\partial_1} C_0 \xrightarrow{\partial_0} 0$$

with this property is called a *chain complex*. The  $k$ th cubical homology of  $C$  over  $\mathbf{F}$  is defined to be the quotient vector space

$$H_k(C; \mathbf{F}) = \frac{\ker \partial_k}{\text{im } \partial_{k+1}}.$$

This construction is functorial: given a suitable definition of a *cubical map*  $f: C \rightarrow D$  between cubical complexes, there is an induced map  $f_k: H_k(C; \mathbf{F}) \rightarrow H_k(D; \mathbf{F})$  for each  $k \in \mathbb{N}$ . We will not introduce these in general, but will note that given  $C \subseteq D$ , the inclusion map  $C \hookrightarrow D$  is cubical and hence induces maps on homology.

## APPENDIX B: STABILITY OF FILTRATION

We briefly show that the filtrations introduced in Sec. II B yields persistent homology which is stable under small perturbations of spins with respect to the bottleneck distance between persistence diagrams. We make use of the following concept:

*Definition (Interleaving).* Given filtrations

$$F, G : (\mathbb{R}, \leq) \rightarrow \text{CubicalComplex},$$

we say that  $F$  and  $G$  are  $\epsilon$ -interleaved if for all  $t \in \mathbb{R}$  we have that  $F(t) \subseteq G(t + \epsilon)$  and  $G(t) \subseteq F(t + \epsilon)$ . We say that the interleaving distance between  $F$  and  $G$  is

$$d_I(F, G) = \inf\{\epsilon \mid F \text{ and } G \text{ are } \epsilon\text{-interleaved}\}.$$

Then from [52] we have the following theorem.

**Theorem** Given filtrations  $F$  and  $G$  as before, and  $k \in \mathbb{N}$ , we have that

$$d_B(PH_k(F), PH_k(G)) \leq d_I(F, G),$$

where  $d_B$  is the bottleneck distance.

Thus we just need to show that the filtration we assign to a configuration  $F_\theta$  are  $\epsilon$ -interleaved with the filtration obtained after a perturbation of the configuration for some  $\epsilon$  bounded by the changes in the spins. Take  $F$  to be the angle difference filtration introduced in Sec. II B and suppose we have a configuration  $\theta$  and change spin  $\theta_i$  to  $\theta_i + \Delta\theta$  to obtain a configuration  $\theta'$ . Then given a neighboring spin  $\theta_j$ , denote the length of the smallest arc between  $\theta_i$  and  $\theta_j$  by  $d_{ij}$  and the length of the arc between  $\theta_i + \Delta\theta$  and  $\theta_j$  by  $d'_{ij}$ . Then we have

that

$$d_{ij} - \Delta\theta \leq d'_{ij} \leq d_{ij} + \Delta\theta.$$

Hence for all filtration values  $t \in \mathbb{R}$  we have inclusions  $F_\theta(t) \hookrightarrow F_{\theta'}(t + \Delta\theta/2\pi)$  and  $F_{\theta'}(t) \hookrightarrow F_\theta(t + \Delta\theta/2\pi)$  so that the filtrations are  $\frac{\Delta\theta}{2\pi}$ -interleaved. By the theorem above, we have

$$d_B(PH_k(F_\theta), PH_k(F_{\theta'})) \leq \frac{\Delta\theta}{2\pi}$$

for all  $k \in \mathbb{N}$ . A straightforward application of the triangle inequality shows that if multiple spins are altered then the bottleneck distance is bounded by the sum of the alterations. The same argument applies in the case of the nematic angle difference filtration.

On the other hand, we observe that the sublevel set filtration for  $\mathbb{S}^1$ -valued spins introduced in [24] does not have this stability. An arbitrarily small perturbation  $\epsilon$  to one of the spins can change its angle from  $\pi$  to  $-\pi + \epsilon$ , potentially introducing or removing a high persistence point in the persistence diagram. Although this is mitigated to an extent in practice by choosing angle 0 to be in the direction of the magnetization.

## APPENDIX C: HISTOGRAM REWEIGHTING

Histogram reweighting allows us to express the ensemble average of an observable  $O$  at temperature  $T'$  in terms of averages at any other temperature  $T$  according to the equation

$$\langle O \rangle_{T'} = \frac{\langle O e^{-(\beta' - \beta)E} \rangle_T}{\langle e^{-(\beta' - \beta)E} \rangle_T}, \quad (\text{C1})$$

where  $\beta = 1/T$ ,  $\beta' = 1/T'$ , and  $E$  is the energy of a configuration [40]. However, in practice we can only reweight so far, so that the energy distributions for  $T$  and  $T'$  have a sizable overlap. To reliably extrapolate to a wider region we can make use of multiple histogram reweighting [41] where we sample at multiple temperatures  $T_1, \dots, T_R$  (with corresponding inverses  $\beta_1, \dots, \beta_R$ ). Suppose we sample  $N_i$  configurations at temperature  $T_i$ , then we can iterate the equation

$$e^{-f_\beta} = \sum_{i=1}^R \sum_{a=1}^{N_i} \frac{g_i^{-1} e^{-\beta E_i^a}}{\sum_{j=1}^R N_j g_j^{-1} e^{-\beta_j E_i^a + f_j}}$$

to estimate the free energies  $f_i = f_{\beta_i}$  at the temperatures  $T_i$  up to an additive constant, where each  $g_i$  is a quantity related to the integrated autocorrelation of the samples in run  $i$ . Given the  $f_i$  we can estimate

$$\langle O \rangle_{T'} = \sum_{i=1}^R \sum_{a=1}^{N_i} \frac{O_i^a g_i^{-1} e^{-\beta_k E_i^a + f_{\beta'}}}{\sum_{j=1}^R N_j g_j^{-1} e^{-\beta_j E_i^a + f_j}}.$$

## APPENDIX D: BOOTSTRAP ERROR ESTIMATION

To make any reasonable conclusions from the results of our analysis we need to be able to estimate the error in any numerical values obtained. While the error in ensemble averages can be directly estimated from the sample, we also calculate various fits to the data. The way in which error propagates here is not necessarily easy to calculate directly. Recall that the idea of bootstrap analysis is to sidestep these concerns by estimating the sampling distribution of a statistic directly. Suppose



we obtain  $N$  sampled configurations  $S = \{\theta_1, \dots, \theta_N\}$  and calculate some numerical statistic  $f(S)$  from the data. Given some preset integer  $N_B$ , bootstrap analysis proceeds by the following:

- (1) Resampling  $S$  with replacement  $N_B$  times to obtain samples  $S_1, \dots, S_{N_B}$  each of size  $N$ ; then
- (2) Computing  $f(S_i)$  for each  $i \in \{1, \dots, N_B\}$ .

For large enough  $N_B$ , the distribution of the  $f(S_i)$  approximates the sampling distribution of  $f$  and we can estimate the standard error

$$\sigma_f \approx \sqrt{\frac{1}{N_B - 1} \sum_i [f(S_i) - \overline{f(S)}]^2}.$$

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