Inverse problem beyond two-body interaction: The cubic mean-field Ising model

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In this paper, we study the inverse problem for a class of mean-field models in statistical mechanics with cubic interaction. The direct problem of statistical mechanics is to compute macroscopic variables (i.e., the average values of magnetizations and correlations) when the couplings and fields are known. In the inverse problem, the reverse is done: the couplings and fields are computed using the (statistical) datum of the macroscopic quantities. This is often achieved by a technique known as Boltzmann machine learning, a special case of learning in statistical inference theory [1,2] when the probability measure is the Boltzmann-Gibbs one.

The system we consider here is made of Ising spins and, beside a homogeneous magnetic field and a constant two-body interaction, it contains a constant three-body term. One of the peculiarities of this model, which turns out to have a cubic Hamiltonian function, is that it lacks the standard convexity property of its quadratic version and its direct and inverse problems are therefore outside the general methods of convex optimization problems. Taking into account the three-body term, we move from a generic graph (network) structure, where we consider only dyadic or pairwise interactions, into hypergraphs, where faces are also considered [3–5]. This allows for the consideration of a large spectrum of applications that are closely related to real-world phenomena, such as team collaborations rather than collaborations between pairs (see [6]). According to [4,6], the presence of higher-order interactions, such as three- or more-body interactions, may also have a significant impact on the dynamics of interacting networked systems and potentially lead to abrupt transitions between states [7]. Abrupt transitions are a prevalent phenomenon in nature that can be found in areas as diverse as social networks or biology [6,8].

In recent years, studies in deep learning for artificial intelligence have been approached in terms of the inverse problem in statistical mechanics [9–11]. The techniques to study that case are of a very different nature than those we treat in this work because the parameters to be identified are of very high dimension and the involved models concern the theory of disordered systems [12]. Although in this study we are only interested in computing three parameters, we believe that a robust understanding of the statistical mechanics low-dimensional of the inverse problem may shed some light on the general Boltzmann machine learning problem due to the presence of phase transitions for very large systems.

A further reason of interest for the problem we deal with is that in recent times, this method has attracted some attention due to its ability to advance a useful novel approach for several applications such as neural networks, protein structures, computer vision [13–17], and socioeconomic sciences [18–26].

The model we consider is invariant under the permutation group, but its extension to the case in which that symmetry is not present has already been considered in [27] for the two-populated case with the same perspectives of the multi-populated quadratic models [21,28]. An intriguing feature of such model is that it shows a discontinuous first-order phase transition which is not present in the case of the standard quadratic mean-field model.

To solve the inverse problem, we first compute, exploiting the exact solution of the model [27,29], the analytical formulas for the system’s macroscopic variables in the thermodynamic limit where they provide explicit expressions for the interaction couplings (cubic and quadratic) and the magnetic field. It is worth noticing that since the number of necessary relations to compute the free parameters is three, we need to make observations up to the third moment of the probability distribution. To relate the analytical inversion with the (statistical) observations, we use the maximum likelihood criteria and we establish a link between estimated and theoretical values. Finally, we test how well the model’s free parameters are reconstructed using the inversion formulas and how their robustness is affected by both the system size and the number of independent samples simulated from the model’s equilibrium configuration.

The paper is organized as follows. The cubic mean-field model is introduced in Sec. II, where it is shown how to
compute and test the robustness of the analytical inverse formulas using the maximum-likelihood estimation procedure. Section III is devoted to the numerical testing of the robustness of the inversion formulas for unique stable solutions. In Sec. IV, the case of metastable or multiple solutions for finite-size systems is discussed. The final section, Sec. V, provides a general conclusion and the model’s future prospects.

II. INVERSE PROBLEM FOR THE CUBIC MEAN-FIELD ISING MODEL

Let us consider the Hamiltonian of an Ising model on $N$ spin configurations, $\Omega_N = \{-1, +1\}^N$, with cubic interaction and spin moments $\sigma_i = \pm 1$, $i = 1, \ldots, N$, defined as

$$H_N(\sigma) = -\sum_{i,k=1}^{N} K_{i,k} \sigma_i \sigma_k - \sum_{i,j=1}^{N} J_{i,j} \sigma_i \sigma_j - \sum_{i=1}^{N} h_i \sigma_i.$$  

(1)

Assuming mean-field interaction, we set $K_{i,k} = K N$, $J_{i,j} = J N$, and $h_i = h$, where $K, J$ are the cubic and binary spin coupling and $h$ is the external magnetic field. Hence, the Hamiltonian per particle is

$$H_N(\sigma) = -N \left[ K \frac{1}{3} m_N^3(\sigma) + \frac{J}{2} m_N^2(\sigma) + h m_N(\sigma) \right].$$

(2)

where

$$m_N(\sigma) = \frac{1}{N} \sum_{i=1}^{N} \sigma_i$$

(3)

is the magnetization per particle of the configuration $\sigma$. The Boltzmann-Gibbs state on a configuration $\sigma$ is given by

$$P_{N,K,J,h}(\sigma) = \frac{e^{-H_N(\sigma)}}{Z_N},$$

(4)

where $Z_N = \sum_{\sigma \in \Omega_N} e^{-H_N(\sigma)}$ is the partition function of the system. As a result, we obtain the Gibbs free energy associated with the thermodynamic system as

$$p_N = \frac{1}{N} \ln Z_N.$$  

(5)

For a given observable $f(\sigma)$, the Boltzmann-Gibbs expectation $\omega_N(f(\sigma))$ is defined as follows:

$$\omega_N(f(\sigma)) = \frac{\sum_{\sigma \in \Omega_N} f(\sigma) e^{-H_N(\sigma)}}{Z_N}.$$  

(6)

Furthermore, the Gibbs free energy (5) can be used to generate the moments of the system with respect to the Boltzmann-Gibbs measure. Hence, one obtains the following finite-size quantities:

$$\frac{\partial p_N}{\partial h} = \omega_N(m_N(\sigma)),$$

(7)

$$\frac{\partial^2 p_N}{\partial h^2} = \chi_N = N \left[ \omega_N(m_N^2(\sigma)) - \omega_N^2(m_N(\sigma)) \right],$$

(8)

and

$$\frac{\partial^3 p_N}{\partial h^3} = \psi_N = N^2 \left[ \omega_N(m_N^3(\sigma)) - 3 \omega_N(m_N) \omega_N(m_N^2) + 2 \omega_N^3(m_N) \right].$$  

(9)

where $\omega_N(m_N(\sigma))$, $\chi_N$, and $\psi_N$ are the finite-size average magnetization, susceptibility, and third moment, respectively. The considered model can be solved exactly using the large deviations technique, which was proposed in [30]. The thermodynamic limit of (5) admits the following variational representation:

$$p = \lim_{N \to \infty} p_N = \sup_{m \in \{-1,1\}} \phi(m),$$

(10)

where $\phi(m) = U(m) - I(m)$, and

$$U(m) = K \frac{1}{3} m^3 + J \frac{1}{2} m^2 + hm,$$

(11)

is the energy contribution and

$$I(m) = \frac{1}{2} m \ln \left( \frac{1 - m}{2} \right) + 1 + m \ln \left( \frac{1 + m}{2} \right)$$

(12)

is the entropy contribution. Notice that $p$ is the Legendre transform of $\phi(m) - hm$. The stationarity condition, which acts as a consistency equation, gives

$$m = \tanh(K m^3 + J m + h),$$

(13)

and must be satisfied by the solutions of the variational principle (10). In order to solve the inverse problem analytically for a given configuration of spin particles, we first find the relation between the model parameters and the variational principle (10). Observe that

$$\frac{\partial^2 \phi}{\partial h^2} = \chi = \left( 1 - m^2 \right)(1 - (1 - m^2)(J + 2Km),$$

(14)

and

$$\frac{\partial^3 \phi}{\partial h^3} = \psi = \chi^3 \left( 2K - \frac{2m}{1 - m^2} \right).$$

(15)

The peculiar feature of the cubic mean-field model is the presence of three distinct stable phases in the magnetic order parameter $m$. Unlike the usual quadratic model, here an unpolarized stable phase close to $m = 0$ appears beyond the usual two phases of positive and negative magnetization. From Fig. 1, one can observe a triple point $(K, J, h) = (0, 1, 0)$ where all three phases meet [27]. Let us consider the model in its simplest form with zero quadratic coupling and magnetic field, i.e., when $J = h = 0$ and only the cubic coupling in (2) is present. It is worth mentioning that when $J = h = 0$ and $K$ is progressively increased from negative to positive, one encounters two transitions: from a negatively polarized phase to an unpolarized one and from an unpolarized phase to a positively polarized one (see Fig. 1 and Fig. 1 of [27]). In Fig. 2, we illustrate an example of critical behavior for our model with the presence of phase transitions occurring at $J = h = 0$ when $K$ is varied.

The quantities, $m, \chi, \text{ and } \psi$ are the infinite-volume limit average magnetization, susceptibility, and third moment corresponding to the finite-size quantities $\omega_N, \chi_N$, and $\psi_N$, respectively, in the thermodynamic limit. The system of equations (14)–(16) has three unknowns $K, J,$ and $h$ for which one can solve. Having knowledge of $m, \chi, \text{ and } \psi$, one can con-
FIG. 1. Phase diagram of the stable solutions of (13) showing the coexistence curves. For $J < 1$, three distinct phases are observed: the negatively polarized phase (yellow), the zero or unpolarized phase (orange), and the positively polarized phase (brown). As a result, in that region, a progressive increase in $K$ from negative to positive values encounters two consecutive jumps.

pute the parameters (i.e., $K$, $J$, and $h$) of the model through the following equations:

$$K = \frac{m}{(1 - m^2)^2} + \frac{\psi}{2\chi},$$
$$J = \frac{1}{1 - m^2} - \frac{1}{\chi} - 2Km,$$

and the external magnetic field is then obtained from (13) as

$$h = \text{arctanh}(m) - Km^2 - Jm.$$  

Let us observe that in the region of the parameter space where the consistency equation (13) has a unique stable solution, the following holds:

$$\lim_{N \to \infty} \omega_N(m_N(\sigma)) = m.$$  

In analogy to the behavior of the quadratic case [31], the Boltzmann-Gibbs measure (4) may be multimodal for some $(K, J, h)$ in the parameter space for both the finite-size system and in the thermodynamic limit. In this case, Eq. (20) fails to hold. We will discuss later how to handle such a case, following the work done in [31,32]. The procedure discussed so far deals with the analytical inverse problem. The remainder of this section will be devoted to the statistical procedure required to compute the estimators of $K$, $J$, and $h$.

We start by generating $M$ independent configurations $\sigma^{(1)}, \ldots, \sigma^{(M)}$ distributed according to (4) from the model’s equilibrium configuration. Notice that the analytical inverse formulas of $K$, $J$, and $h$ in Eqs. (17), (18), and (19), respectively, are valid on the infinite-volume limit of the observables, i.e., $m$, $\chi$, and $\psi$. Hence, to compute the estimates of the model parameters $K$, $J$, and $h$, the maximum-likelihood estimation procedure will be adopted having knowledge of real data. This procedure ensures that the estimated model

FIG. 2. $J = 0$, $h = 0$. First three moments of the model as a function of $K$: In (a) the total magnetization shows indications of phase transitions occurring at a critical point around $\pm 2$. At the critical point, the (b) susceptibility and (c) third moment have a jump to 1 and a jump to around $\pm 4$, respectively, parameters maximize the probability of getting the given sample of spin configurations from the distribution. Furthermore, the analytical inverse procedure requires statistical approximation of the infinite-volume limit quantities (i.e., $m$, $\chi$, and
The function $h$ for each of the have that

\[
\psi
\]

are the total magnetizations of the $M$ sampled spin configurations. Further, we have that

\[
\ln L(K, J, h) = \frac{1}{M} \sum_{i=1}^{M} \left( -H_N(\sigma^{(i)}) \right) - \ln \sum_{\sigma \in \Omega_N} e^{-H_N(\sigma)}. \tag{21}
\]

Requiring the derivatives with respect to the parameters $K$, $J$, and $h$ to vanish amounts to

\[
\frac{\partial}{\partial h} \ln L(K, J, h) = N \sum_{i=1}^{M} [m_N(\sigma^{(i)}) - \omega_N(\sigma^{(i)})] = 0,
\]

\[
\frac{\partial}{\partial J} \ln L(K, J, h) = \frac{N}{2} \sum_{i=1}^{M} [m_N^2(\sigma^{(i)}) - \omega_N^2(\sigma^{(i)})] = 0,
\]

\[
\frac{\partial}{\partial K} \ln L(K, J, h) = \frac{N}{3} \sum_{i=1}^{M} [m_N^3(\sigma^{(i)}) - \omega_N(\sigma^{(i)})] = 0.
\]

Therefore,

\[
\omega_N(\sigma^{(i)}) = \frac{1}{M} \sum_{i=1}^{M} m_N(\sigma^{(i)}),
\]

\[
\omega_N(\sigma^{(i)}) = \frac{1}{M} \sum_{i=1}^{M} m_N^2(\sigma^{(i)}),
\]

\[
\omega_N(\sigma^{(i)}) = \frac{1}{M} \sum_{i=1}^{M} m_N^3(\sigma^{(i)}). \tag{22}
\]

The function $L(K, J, h)$ is at its stationary points when the first, second, and third moments of the magnetization in Eq. (22) are obtained. It is worth noticing that

\[
m_N(\sigma^{(i)}) = \frac{1}{N} \sum_{i=1}^{N} \sigma^{(i)} \quad \text{for} \quad l = 1, \ldots, M \tag{23}
\]

are the total magnetizations of the $M$ sample spin configurations. Let us note that $L(K, J, h)$ and its derivatives are only used to solve the forward problem, but not the inverse problem. Now, the inverse problem can be solved when we make use of (17)–(19) and (22). The maximum-likelihood procedure computes the estimators of the infinite-volume quantities $m$, $\chi$, and $\psi$ from a sample data set through the following:

\[
\tilde{m} = \frac{1}{M} \sum_{i=1}^{M} m_N(\sigma^{(i)}), \tag{24}
\]

\[
\tilde{\chi} = N \left[ \frac{1}{M} \sum_{i=1}^{M} m_N^2(\sigma^{(i)}) - \tilde{m}^2 \right], \tag{25}
\]

and

\[
\tilde{\psi} = N^2 \left[ \frac{1}{M} \sum_{i=1}^{M} m_N^3(\sigma^{(i)}) - 3\tilde{m} \frac{1}{M} \sum_{i=1}^{M} m_N^2(\sigma^{(i)}) + 2\tilde{m}^3 \right]. \tag{26}
\]

We now define the estimators of the three parameters of the cubic mean-field model using the statistical estimators for the magnetization, susceptibility, and third moment (24)–(26) in the infinite-volume limit relations among those quantities (17)–(19),

\[
\tilde{K} = \frac{\tilde{m}}{(1 - \tilde{m}^2)^2} + \frac{\tilde{\psi}}{2\tilde{\chi}^3}, \tag{27}
\]

\[
\tilde{J} = \frac{1}{1 - \tilde{m}^2} - \frac{1}{\tilde{\chi}} - 2\tilde{K}\tilde{m}, \tag{28}
\]

and

\[
\tilde{h} = \arctanh(\tilde{m}) - \tilde{K}\tilde{m} - \tilde{J}\tilde{m}. \tag{29}
\]

At the critical point $(K, J, h) = (0, 1, 0)$ where all the three phases meet, the magnetization is zero and the infinite-volume magnetic susceptibility $\chi$ and the third moment $\psi$ defined by Eqs. (15) and (16), respectively, diverge. Hence, the inversion formulas (17)–(19) do not hold, as will be illustrated at the end of the next section. We do not include the inversion formulas at the critical point in this work, but the problem will be considered in future work.

## III. TEST FOR THE CASE OF UNIQUE SOLUTION

In this section, we are going to examine how the inversion equations perform for different and increasing choices of $N$ and $M$, respectively, the number of particles and sampled configurations. The specific case we consider is the inversion problem for those values of the triple $(K, J, h)$ where there is a unique stable solution of (13). In this case, the Boltzmann-Gibbs distribution of the total magnetization has a unique peak always centered around the analytic solution $m$: some examples are shown in Fig. 3 for fixed $N$. The accuracy of the estimation increases as $N$ and $M$ increase.

The parameters $K$, $J$, and $h$ are obtained from the computation of the finite-size quantities $m_N$, $\chi_N$, and $\psi_N$ using configurations extracted from the Boltzmann-Gibbs distribution of the data. Estimation of $m_N$, $\chi_N$, and $\psi_N$ for fixed triples of the parameters $(K, J, h)$ and varying $N \in [500, 10000]$ are shown in Fig. 4. In the same figure, the thermodynamic limits of those quantities are also shown.

From Fig. 4, we can observe the monotonic behavior of $m_N$, $\chi_N$, and $\psi_N$ as $N$ increases. In Fig. 5, we study the
relationship between the absolute difference of the finite-size quantities and their corresponding thermodynamic values as a function of the system size \( N \). We find evidence that the finite-size quantities \( m_N, \chi_N, \) and \( \psi_N \) converge to their true values with a power-law behavior as \( N \) increases. The obtained results indicate that using \( N = 10000 \), one can estimate the infinite-volume magnetization, susceptibility, and third moment with vanishing error. We will proceed to use \( N = 10000 \) as the size for each of the \( M \) independent spin configurations \( \sigma^{(1)}, \ldots, \sigma^{(M)} \). Further numerical tests will be performed to determine a suitable number of sample configurations \( M \) that can be used for reconstructing the model parameters using the inversion formulas.

To obtain the statistics associated to the reconstruction of the estimators, we simulate from the model’s equilibrium configuration 50 different instances of the \( M = \) iid sample configurations, i.e., \( (\sigma^{(1)}, \ldots, \sigma^{(M)}) \), apply the maximum-likelihood estimation procedure to each of them separately, solve the inverse problem using (27)–(29), and then average the inferred values over the 50 different \( M \) samples. The mean value of the estimators \( \bar{m}, \bar{\chi}, \bar{\psi}, \) and \( (\bar{K}, \bar{J}, \bar{h}) \) over the 50 different \( M \) samples of spin configurations is denoted by \( \bar{m}, \bar{\chi}, \bar{\psi}, \) and \( (\bar{K}, \bar{J}, \bar{h}) \), respectively. The results are shown in Figs. 6 and 7.

Figures 6 and 7 illustrate that at \( M = 20000 \), we get smaller error bounds for the reconstruction as compared to lesser values of \( M \).

In the sequel, we study the behavior of the reconstructed parameter for fixed values of \( J \) and \( h \) and varying \( K \) (Figs. 8 and 9), and also for fixed values of \( K \) and \( h \) and varying \( J \) (Figs. 10 and 11). The simulations are performed using \( M = 20000, N = 10000 \), and error bars are standard deviations on 50 different \( M \) samples of the same system. We find all the reconstructed parameter values in good agreement with the exact ones. We can observe that as the intensity of the cubic and quadratic coupling increases, the error bars associated to the reconstructed parameters grow, as we can expect since in that region of the parameter space the system is more disordered, due to the presence of multiple local stable states, and the fluctuations are greater. Furthermore, Fig. 12
shows the reconstructed parameters as a function of $N$ at the critical point ($K = 0$, $J = 1$, $h = 0$). We notice that the reconstruction at the critical point for $K$ and $h$ agrees with their exact values with only a small percentage of error and that of $J$ is underestimated. The monotonic behavior observed for the reconstruction of $J$ relates to higher-order corrections of the free energy from the variational principle with respect to the magnetization.

It is worth observing that when $K = h = 0$ and $J > 1$, the consistency equation (13) has two stable solutions. In such a case, the inverse problem procedure discussed in Sec. II cannot be used for the reconstruction of the model parameters. We refer readers to [28], where this case has been studied using the spin-flip approach due to

FIG. 6. Reconstructed average magnetization $\bar{m}$, susceptibility $\bar{\chi}$, and third moment $\bar{\psi}$ (given as blue crosses in each panel) as a function of $M$ with standard deviation on 50 different $M$ samples and $N = 10^4$. The continuous red line corresponds to $m$, $\chi$, and $\psi$ in the thermodynamic limit.

FIG. 7. $\bar{K}$, $\bar{J}$, and $\bar{h}$ as a function of $M$ for $N = 10^4$. The blue crosses are the estimation of $\bar{K}$, $\bar{J}$, and $\bar{h}$ with standard deviations on 50 different $M$ samples of configurations of the same system. The horizontal red line in each panel corresponds to the exact values of $K$, $J$, and $h$.

FIG. 8. $\bar{K}$ as a function of $K$ for $N = 10^4$ and $M = 20^4$. $J = 0.3$, $h = 0.1$ in the left panel and $J = 0.4$, $h = 0.3$ in the right panel. The estimations of $\bar{K}$ are given as the blue crosses in both panels, with standard deviations on 50 different $M$ samples of configurations of the same system. The red continuous line represents $\bar{K} = K$.

FIG. 9. $\bar{J}$ and $\bar{h}$ as a function of $K$ for $N = 10^4$ and $M = 20^4$. $J = 0.3$, $h = 0.1$ in the left panels and $J = 0.4$, $h = 0.3$ in the right panels. The estimates of $\bar{J}$ and $\bar{h}$ are given as the blue crosses in all the panels, with standard deviations on 50 different $M$ samples of configurations of the same system. The red continuous lines represent the exact values of $J$ and $h$. 
symmetry of the solution in both finite-size and infinite-volume systems for the quadratic mean-field model. The clustering algorithm to be outlined in the next section provides a more general approach to handle the reconstruction of the model parameters when the phase space has multiple locally stable solutions.

FIG. 10. \( \overline{J} \) as a function of \( J \) for \( N = 10,000 \) and \( M = 20,000 \). \( K = 0.05, h = 0 \) in the left panel and \( K = 0.05, h = -0.02 \) in the right panel. The blue crosses are the reconstructed values of \( J \) in both panels, with standard deviations on 50 different \( M \) samples of configurations of the same system. The red continuous line represents the exact value \( \overline{J} = J \).

FIG. 11. \( \overline{K} \) and \( \overline{h} \) as a function of \( J \) for \( N = 10,000 \) and \( M = 20,000 \). \( K = 0.05, h = 0 \) in the left panels and \( K = 0.05, h = 0.02 \) in the right panels. The estimates of \( \overline{K} \) and \( \overline{h} \) are given as the blue crosses in all the panels, with standard deviations on 50 different \( M \) samples of configurations of the same system. The red continuous lines represent the exact values of \( K \) and \( h \).

FIG. 12. \( K = 0, J = 1, h = 0 \). \( \overline{K}, \overline{J}, \) and \( \overline{h} \) as a function of \( N \) for \( M = 20,000 \). The reconstructed estimates of \( K, J, \) and \( h \) are given as the blue crosses on statistical error bars of 50 different \( M \) samples. The red continuous line is the exact value of the parameters \( K, J, \) and \( h \) in the respective panels.

IV. CLUSTERING ALGORITHM FOR METASTABLE STATE SOLUTIONS

Here, we focus on cases where Eq. (13) has a metastable solution. This corresponds to the case where there is more than one locally stable solution of the consistency equation (13). For this model, Eq. (13) can have, at most, three solutions and \( \phi \) has, at most, two local maxima for fixed \((K, J, h)\). The existence of the metastable solution in the infinite-volume limit is represented at finite \( N \) by the occurrence of an extra peak in the distribution. Therefore, while in the thermodynamic limit the Boltzmann-Gibbs distribution of the magnetization is unimodal with the peak corresponding to the stable solution, in the finite-size case also the peak corresponding to the metastable one is present and the distribution is bimodal. Hence, in this case, the inversion problem cannot be studied globally, as done in the previous section. Instead, the procedure has to be applied locally, that is, to each subset of configurations clustered around the two local maxima. Given \( M \) spin configurations, \( \sigma^{(1)}, \ldots, \sigma^{(M)} \), we perform the reconstruction by first partitioning the \( M \) configurations in clusters according to their local densities around each local maximum. More precisely, using the clustering algorithm discussed in [32–36], we divide the \( M \) configurations into different clusters using the mutual distances between their magnetizations of each configuration. Configurations form a cluster if the magnetization distances are less than a fixed threshold \( d_c \). The choice of the optimal threshold is obviously crucial: a too small threshold will produce too many clusters, while a too large one will give only one cluster. Given \( d_c \), for each configuration \( l \) the algorithm computes two quantities: the local density \( \rho_l \), defined as the number of magnetizations within the given distance \( d_c \) to the magnetization of \( \sigma^{(l)} \), and
the minimum distance $\delta_l$ between the magnetization of configuration $l$ and any other configuration with a higher density.

The algorithm is based on the assumptions that the cluster centers are surrounded by points with a lower density and that the centers are at a relatively large distance from each other. For each configuration, plotting the minimum distance $\delta$ as a function of the local density $\rho$ provides a decision graph that gives the cluster centers; the cluster centers are the outliers in the graph. Finally, each remaining configuration is assigned to the same cluster of its nearest neighbor of higher density. In this study, we identify two clusters $C_1, k = 1, 2$, using the optimal threshold $d_c = 0.001$. Notice that it is not possible to observe three clusters in the inverse problem due to the analytical properties of the consistency equation (13).

Then, for each cluster $C_k$, $k = 1, 2$, we compute the estimates of the finite-size quantities, $\hat{m}, \hat{\xi},$ and $\hat{\psi}$, and the corresponding $\hat{K}, \hat{J}, \hat{h}$. More precisely, we can define the estimators of the finite-size quantities with reference to the clusters as follows:

$$\hat{m}_{C_k} = \frac{1}{M_k} \sum_{l \in C_k} m_N(\sigma^{(l)}),$$

(30)

$$\hat{\xi}_{C_k} = N^2 \left[ \frac{1}{M_k} \sum_{l \in C_k} m_{N}^{2}(\sigma^{(l)}) - \hat{m}_{C_k}^{2} \right],$$

(31)

and

$$\hat{\psi}_{C_k} = N^2 \left[ \frac{1}{M_k} \sum_{l \in C_k} m_{N}^{3}(\sigma^{(l)}) - 3 \hat{m}_{C_k} \frac{1}{M_k} \sum_{l \in C_k} m_{N}^{2}(\sigma^{(l)}) + 2 \hat{m}_{C_k}^{3} \right],$$

(32)

where $M_k$ is the size of the cluster $C_k$, $k = 1, 2$, such that $M_1 + M_2 = M$. After obtaining the above quantities, we now compute the estimated values, $\hat{K}, \hat{J}, \hat{h}$, using Eqs. (27)–(29) for each cluster, and compute the final estimates of the parameters $K, J,$ and $h$ as the weighted averages,

$$\hat{K} = \frac{1}{M} \sum_{k=1}^{2} M_k \hat{K}_{C_k},$$

(33)

$$\hat{J} = \frac{1}{M} \sum_{k=1}^{2} M_k \hat{J}_{C_k},$$

(34)

and

$$\hat{h} = \frac{1}{M} \sum_{k=1}^{2} M_k \hat{h}_{C_k}.$$ (35)

Observe that if a point $(K, J, h)$ in the parameter space corresponds to a metastable solution (at finite volume) and it is sufficiently distant from the coexistence curve, we can expect a better reconstruction of the parameters by applying Eqs. (27)–(29) to the configurations in the largest cluster. However, if the point $(K, J, h)$ is close to the coexistence curve, a better reconstruction can be expected using the density clustering algorithm, i.e., by using (33)–(35).

Figure 13 illustrates how the Boltzmann-Gibbs measure of the magnetization is changing with varying $K, J,$ and $h$ in each column, starting from the left, respectively.
where the two peaks are centered around crosses on statistical error bars on 50 different \( M \) samples of configurations of the same system. The horizontal red lines in each panel correspond to the exact values of \( K, J, \) and \( h \).

\[ M = 20000 \text{ and } N = 3000. \] The distribution of the magnetization at this point is given as the blue dashed curve in Fig. 14, where the two peaks are centered around \( m_1 = 0.1311 \) and \( m_2 = 0.8973 \), the stable solution and the metastable solution of the consistency equation (13), respectively.

As is evident from Fig. 14, the cluster centered around \( m_1 \) (i.e., \( C_1 \)) has more configurations as compared to the other cluster centered around \( m_2 \) (i.e., \( C_2 \)). We get the following reconstructed estimates for the parameter values by applying Eqs. (27)–(29) just to the configurations in both clusters (i.e., \( C_1 \) and \( C_2 \)) according to formulas (33)–(35),

\[ (\vec{K}, \vec{J}, \vec{h}) = (1.76 \pm 0.67, -0.11 \pm 1.11, 0.15 \pm 0.49). \]

Instead, we obtain the following reconstructed parameter values by applying Eqs. (27)–(29) to the setups in both clusters (i.e., \( C_1 \) and \( C_2 \)) according to formulas (33)–(35),

\[ (\bar{K}, \bar{J}, \bar{h}) = (1.69 \pm 0.23, 0.01 \pm 0.06, 0.10 \pm 0.004). \]

Note that the reconstructed parameters using only the configurations in the more dense cluster are in better agreement with the exact ones when compared to the reconstructed parameters on both clusters. This is an indication that the point \((K, J, h) = (1.67, 0.01, 0.1)\) is sufficiently distant from the coexistence curve. Observe that if two clusters have the same density, we do not choose between them and the clustering algorithm provides an optimal reconstruction.

Now, we perform reconstruction of the parameters using the cluster with the largest size for fixed values of the model parameters and observe its performance for varying \( M \) in Fig. 15. It can be observed that the reconstructed parameters are in good agreement with their corresponding exact values.

As a last remark, note that given a point \((K, J, h)\) in a neighborhood of the coexistence curve, one can observe a metastable state when the number of particles, \( N \), is not large enough. In this case, the clustering algorithm is useful to reconstruct the parameters, but it has a high computational cost. This is easily overcome by using a large number of particles, which causes the metastable state to vanish (see Fig. 14) and the inversion formulas in Eqs. (27)–(29) become efficient.

V. CONCLUSION

In this work, we consider a mean-field statistical mechanics model with three-body interaction displaying a first-order phase transition. The findings of this paper can be used to infer interaction couplings and fields from real-world datasets, particularly in network structures where long-range interactions are a realistic description of the phenomenon, such as the protein structure [37] and human genome [38].

We studied and solved the inverse problem and tested the statistical robustness of the inversion method. We numerically tested the inversion method for cases where the consistency equation (13) has a unique stable solution as well as more than one locally stable solution. For the case where the consistency equation (13) has multiple locally stable solutions, we used the clustering algorithm to reconstruct the model parameters. Robustness was tested for different values of the number of particles, \( N \), and samples \( M \) and reached the precision of a few percent for \( M = 2 \times 10^4 \).

We plan to investigate, in the future, two extensions of the inverse problem: first to the critical point where some of the observables such as \( \chi \) and \( \psi \) diverge, and second to the multi-populated version of the model that found applications in the description of human-AI (Artificial Intelligence) ecosystems [27].

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