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Published in:
Physical Review E

DOI:
[10.1103/PhysRevE.92.010102](https://doi.org/10.1103/PhysRevE.92.010102)

Published: 01/01/2015

Document Version
Publisher's PDF, also known as Version of record

Please cite the original version:
Del Ferraro, G., & Aurell, E. (2015). Dynamic message-passing approach for kinetic spin models with reversible dynamics. *Physical Review E*, 92(1), 1-7. [010102]. <https://doi.org/10.1103/PhysRevE.92.010102>

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Dynamic message-passing approach for kinetic spin models with reversible dynamics

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(Received 17 November 2014; published 21 July 2015)

A method to approximately close the dynamic cavity equations for synchronous reversible dynamics on a locally treelike topology is presented. The method builds on (a) a graph expansion to eliminate loops from the normalizations of each step in the dynamics and (b) an assumption that a set of auxiliary probability distributions on histories of pairs of spins mainly have dependencies that are local in time. The closure is then effectuated by projecting these probability distributions on n -step Markov processes. The method is shown in detail on the level of ordinary Markov processes ($n = 1$) and outlined for higher-order approximations ($n > 1$). Numerical validations of the technique are provided for the reconstruction of the transient and equilibrium dynamics of the kinetic Ising model on a random graph with arbitrary connectivity symmetry.

DOI: [10.1103/PhysRevE.92.010102](https://doi.org/10.1103/PhysRevE.92.010102)

PACS number(s): 02.50.-r, 89.20.-a

I. INTRODUCTION

Disordered spin systems are an important class of models able to catch and reproduce a large range of phenomena from phase transitions in magnets and amorphous systems [1] to protein folding in biology [2], social media [3,4], epidemic spreading [5], immune and neural networks [6], and applications in finance and optimization problems [7–9]. In the thermodynamic limit these systems have rich and fascinating repertoires of static and dynamic behavior, including the clustering [8] or shattering [10] transition, ergodicity breaking, and ageing [11]. To systematically describe their static properties the replica method (for fully connected systems) and the cavity method (for dilute systems) were developed [12]. General techniques to systematically study the dynamics of single finite systems in this class have, however, been less developed and in practice mostly limited to dynamic mean-field theories [13,14], path integral techniques [15–18], large deviation approaches [19], and, above all, numerical simulation [20].

The cavity method [12,21] here holds a special place as it has become the method of choice to solve the statics of models on sparse networks, while for dynamics it was long restricted to dynamics on fully asymmetric graphs [22–24]. The main problem is that while the cavity technique reduces the complexity “in space” (number of terms in an approximate computation of marginals), there remains a complexity “in time” (cardinality of each term). This is so because the natural variables of the dynamic cavity method are probabilities of spin histories, of which there are exponentially many (2^t for synchronous updates over time $[0, t]$ with time constant one). Therefore, although the dynamic cavity equations themselves only involve a finite number of terms, summing them nevertheless (in general) entails a number of operations which is exponentially large in t . Fully asymmetric networks is a special case since the cavity equations then can be marginalized over time with no loss of information, and the complexity “in time” disappears. Alternative “ways out” investigated in the literature use additional assumptions on the evolution

law such as majority dynamics [25] (i.e., linear dynamics with thresholding) or, more recently, unidirectional dynamics. Models of this latter type, where after a variable makes a transition from one state to another it can never go back, are represented by the zero-temperature random-field Ising model (RFIM) [26], cascade processes [27], spread optimization problems [28], and several epidemic models as, for instance, the susceptible-infected-recovered (SIR) model [29,30]. The main peculiarity of such models is that the dynamics can be parametrized in terms of the time(s) at which the transition from one state of the variable to another occurs, which again eliminates time complexity.

Another approach was taken in Ref. [31], where a simplifying “one-time” assumption was introduced, which for fully asymmetric networks reduces to (exact) marginalization over time. For the kinetic Ising model with synchronous and asynchronous updates this was later shown to be considerably more accurate than dynamic mean field, not only for asymmetric networks but also for partly symmetric networks [32,33]. A different approach, based on variational approximations, was very recently proposed in Ref. [34] where the author shows better performances in recovering stationary states compared to existing methods. Unfortunately, except for fully asymmetric networks, all these approaches are limited to steady states and hence cannot handle dynamic phenomena.

In this contribution we present a method to approximately close the dynamic cavity equations for synchronous updates with no assumptions on the underlying network and evolution law beyond that the network is locally treelike and the evolution law is Markov. Unlike other methods already present in the literature, our approach is built not only to recover stationary states but also potentially the transient, i.e., the out-of-equilibrium dynamics. The method is built on two ingredients. The first, which already appeared in this context in Ref. [28], is the use of the graph expansion technique of Ref. [12] to rewrite the probabilistic model in a way such that the underlying graph is explicitly locally treelike, and the standard cavity equations can be used. The second ingredient

graph, as in the static case (see Eq. (14.15) in Ref. [12]) as:

$$m_{i \rightarrow (ij)}(X_i, X_j) \propto \sum_{\{X_k\}} \Phi(X_i, X_j, \{X_k\}) \prod_{k \in \partial i \setminus j} m_{k \rightarrow (ik)}(X_k, X_i), \quad (3)$$

where $\Phi(X_i, X_j, \{X_k\}) = P_0 \prod_{s=1}^t w_i(\sigma_i^s | \sigma_i^{s-1}, \{\sigma_k^{s-1}\}_{k \in \partial i \setminus j}, \sigma_j^{s-1})$ is analogous to a potential sitting in the factor nodes which transfers the dynamics to its neighbor variable nodes and the normalization factor can be computed by the condition that $\sum_{X_i, X_j} m_{i \rightarrow (ij)}(X_i, X_j) = 1$. Above we have shortened the notation to $X_i = X_i^{(ij)}$, $X_k = X_k^{(ik)}$ and the same for X_j and all the σ variables contained in w_i . Let us note that we have fewer distinct messages with respect to the static BP formulation since messages from factor to variable nodes are, in this case, the same as messages from variable to factor nodes, i.e., $m_{j \rightarrow (ij)}(X_j, X_i) = m_{(ij) \rightarrow i}(X_j, X_i)$. Hence, for the topology shown in Fig. 2, the message on the left-hand side of (3) is illustrated in black, whereas the messages on the right-hand side are those two in red coming from the right side of the picture. To simplify notation, from now on variables with no apex refer to variables at time t , i.e., $\sigma_i = \sigma_i(t)$, whereas variables with apex(es) refer respectively to previous time(s), i.e., $\sigma_i' = \sigma_i(t-1)$, $\sigma_i'' = \sigma_i(t-2)$, ..., and so X_i' is the spin history of spin i up to time $t-1$. Let us observe that, while notationally compact, Eq. (3) is, except for short times, computationally intractable as the right-hand side involves sums over complete spin histories.

III. MARKOVIAN CLOSURE OF THE DYNAMIC CAVITY EQUATIONS

We first observe that, given the Markovianity of the dynamics contained in $\Phi(X_i, X_j, \{X_k\})$, the messages $m_{i \rightarrow (ij)}(X_i, X_j)$ in (3) actually do not depend on the spin variable σ_j at time t but only at time $t-1$ and earlier times. The messages $m_{i \rightarrow (ij)}(X_i, X_j)$ are hence always uniform distributions on $\sigma_j(t)$ and, since $\sigma_j(t)$ is therefore a kind of dummy argument, we may simplify the notation by writing $m_{i \rightarrow (ij)}(X_i, X_j')$. Furthermore, if we use the same simplification on the right-hand side of (3), then we have incoming messages $m_{k \rightarrow (ik)}(X_k, X_i')$ which can be marginalized to $m_{k \rightarrow (ik)}(X_k', X_i')$ as there is no other dependence on σ_k , the value of spin k at time t . Using Markovianity again we can simplify further to $m_{k \rightarrow (ik)}(X_k', X_i'')$ and write (3) as

$$m_{i \rightarrow (ij)}(X_i, X_j') \propto \sum_{\{X_k'\}} \Phi(X_i, X_j', \{X_k'\}) \prod_{k \in \partial i \setminus j} m_{k \rightarrow (ik)}(X_k', X_i''). \quad (4)$$

Equation (4) shows that the dynamic cavity equations have a different structure and are actually, in some respects, simpler than standard BP updates, since if all messages up to some time $t-1$ are known, then messages up to time t can be evaluated directly without the need to iterate to a fixed point.

A. Fully asymmetric graph

It is worth observing how Eq. (4) simplifies when referred to a fully asymmetric graph, with interaction couplings between node i and j such that $J_{ij} \neq 0$ and $J_{ji} = 0$. Under this

assumption the interaction function Φ no longer depends on the spin history X_j' and, as a consequence, the message on the left-hand side of the equation does not depend on that history either, since such dependence is carried in only through the function Φ . For consistency, we can apply the same argument to the messages on the right-hand side and conclude that they only depend on the spin history X_k' and no longer on X_i'' . Then, remembering that Φ is a normalized function with respect to the variable σ_i^s , as shown in the text below Eq. (3), we can sum both sides of (4) over the spin history X_i' and then make use of the sum over $\{X_k'\}$ on the right-hand side to obtain the simplified version of the dynamic message-passing equation for a fully asymmetric graph,

$$m_{i \rightarrow (ij)}(\sigma_i^t) = \sum_{\{\sigma_k^{t-1}\}_{k \in \partial i \setminus j}} w_i(\sigma_i^t | \{\sigma_k^{t-1}\}_{k \in \partial i \setminus j}) \times \prod_{k \in \partial i \setminus j} m_{k \rightarrow (ik)}(\sigma_k^{t-1}). \quad (5)$$

This equation is in agreement with the literature for the same graph topology [24,31,32].

B. Graph with arbitrary connectivity symmetry

In what follows, we propose a Markovian closure of the dynamic *BP update* equation (4) for a network with arbitrary connectivity symmetry. In the next section the same closure is used to derivate the dynamic *BP output* equations. In a closure in the class of n -th order Markov processes we assume $m_{i \rightarrow (ij)}(X_i, X_j') = \prod_{s=n}^t T_{i \rightarrow (ij)}^{(n)}(\sigma_i^s | \sigma_i^{s-1}, \sigma_j^{s-1}, \dots, \sigma_i^{s-n}, \sigma_j^{s-n})$ and solve (4) iteratively. We here consider $n=1$. The marginalizations over the last and last two times of the variable node (i, j) are

$$P_{i \rightarrow (ij)}^{(t-1)}(\sigma_i', \sigma_j') = \sum_{X_i'', X_j''} m_{i \rightarrow (ij)}(X_i', X_j''), \quad (6)$$

$$P_{i \rightarrow (ij)}^{(t, t-1)}(\sigma_i, \sigma_i', \sigma_j') = \sum_{X_i'', X_j''} m_{i \rightarrow (ij)}(X_i, X_j'), \quad (7)$$

by assumption linked by

$$P^{(t, t-1)}(\sigma_i, \sigma_i', \sigma_j') = T(\sigma_i | \sigma_i', \sigma_j') P^{(t-1)}(\sigma_i', \sigma_j'), \quad (8)$$

where we omitted the subscript $i \rightarrow (ij)$ and the superscript time dependence of $T^{(t, t-1)}$ for readability. We note that, by the above, $P^{(t-1)}(\sigma_i', \sigma_j')$ actually does not depend on its second argument and we will therefore from now on simplify to $P^{(t-1)}(\sigma_i')$. Closure means to make the same assumptions for the upstream messages $m_{k \rightarrow (ik)}(X_k, X_i')$, use (4) to compute $P_{i \rightarrow (ij)}^{(t-1)}$ and $P_{i \rightarrow (ij)}^{(t, t-1)}$ in (6) and (7), and then take (8) to define $T(\sigma_i | \sigma_i', \sigma_j')$. This can be done by introducing an auxiliary function $\mathcal{F}$,

$$\begin{aligned} \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma_i', \{\sigma_k'\}_{k \in \partial i \setminus j}) &= \sum_{\{X_k'', X_i'', X_j''\}} \prod_{k \in \partial i \setminus j} m_{k \rightarrow (ik)}(X_k', X_i'') \\ &\times \prod_{s=1}^{t-1} w_i(\sigma_i^s | \sigma_i^{s-1}, \{\sigma_k^{s-1}\}_{k \in \partial i \setminus j}, \sigma_j^{s-1}) P_0, \end{aligned} \quad (9)$$

which is strictly a specific marginalization of the right-hand side of (4). From now on, we use the notation $\{\sigma_k\}_\setminus = \{\sigma_k\}_{k \in \partial i \setminus j}$ and $\prod_{k \setminus} = \prod_{k \in \partial i \setminus j}$. In terms of (9) we have

$$P_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i) \propto \sum_{\{\sigma'_k\}} \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i, \{\sigma'_k\}_\setminus), \quad (10)$$

$$P_{i \rightarrow (ij)}^{(t, t-1)}(\sigma_i, \sigma'_i, \sigma'_j) \propto \sum_{\{\sigma'_k\}} w_i(\sigma_i | \sigma'_i, \{\sigma'_k\}_\setminus, \sigma'_j) \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i, \{\sigma'_k\}_\setminus). \quad (11)$$

On the other hand, by the (assumed) Markovianity of the upstream messages we can write an iterative equation in time for $\mathcal{F}$:

$$\begin{aligned} \mathcal{F}_{i \rightarrow (ij)}^{(t)}(\sigma_i, \{\sigma_k\}_\setminus) &= \sum_{\{\sigma'_k\}, \sigma'_i, \sigma'_j} \prod_{k \setminus} T_{k \rightarrow (ik)}(\sigma_k | \sigma'_k, \sigma'_i) w_i(\sigma_i | \sigma'_i, \{\sigma'_k\}_\setminus, \sigma'_j) \\ &\times \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i, \{\sigma'_k\}_\setminus). \end{aligned} \quad (12)$$

Whereas the iterative equation for T , using (10) and (11), reads as

$$\begin{aligned} T_{i \rightarrow (ij)}(\sigma_i | \sigma'_i, \sigma'_j) &= \frac{\sum_{\{\sigma'_k\}} w_i(\sigma_i | \sigma'_i, \{\sigma'_k\}_\setminus, \sigma'_j) \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i, \{\sigma'_k\}_\setminus)}{\sum_{\sigma_i, \{\sigma'_k\}} w_i(\sigma_i | \sigma'_i, \{\sigma'_k\}_\setminus, \sigma'_j) \mathcal{F}_{i \rightarrow (ij)}^{(t-1)}(\sigma'_i, \{\sigma'_k\}_\setminus)}, \end{aligned} \quad (13)$$

where the denominator takes care of the normalization. Equations (12) and (13), which solve the dynamic cavity equations under the assumption that messages are Markovian, are the first result of this paper and represent the ingredients to solve the BP update equations at any time. The entire procedure clearly takes polynomial time instead of exponential time as the original formulation and, as noted above, does not involve iteration to a fixed point.

IV. MARKOVIAN CLOSURE OF THE BP OUTPUT EQUATIONS

We now turn to the *BP output equations*, i.e., the equations for the actual marginal probability both on the auxiliary and on the original factor graph. The marginal probability of one variable node (i, j) in the auxiliary graph is simply given by the product of the incoming messages to the node (i, j) :

$$P_{(ij)}(X_i, X_j) \propto m_{i \rightarrow (ij)}(X_i, X'_j) m_{j \rightarrow (ij)}(X_j, X'_i). \quad (14)$$

We are, however, interested in the single-site one-time marginal probability on the original graph, $P_i[\sigma_i(t)]$, from which we can compute physical observables of interest such as, for instance, the magnetization at any given time. We can get this marginal starting from the one-site marginal probability on a spin history, $P_i(X_i)$. In terms of the auxiliary probability distribution on the expanded graph this reads as

$$\begin{aligned} P_i(X_i) &\propto \sum_{X_j} \prod_{j \in \partial i} m_{j \rightarrow (ij)}(X_j, X'_i) \\ &\times \left[\prod_{s=1}^t w_i(\sigma_i^s | \sigma_i^{s-1}, \{\sigma_j^{s-1}\}_{j \in \partial i}) P_0 \right], \end{aligned} \quad (15)$$

where spins j 's are the neighbors of site i in the original and expanded graph (see Fig. 2). To solve these equations on the same level of approximation as the update equation we define a new auxiliary function:

$$\begin{aligned} \mathcal{G}^{(t-1)}(\sigma'_i, \{\sigma'_j\}_{j \in \partial i}) &= \sum_{X''_i, \{X''_j\}_{j \in \partial i}} \prod_{j \in \partial i} m_{j \rightarrow (ij)}(X'_j, X''_i) \\ &\times \left[\prod_{s=1}^{t-1} w_i(\sigma_i^s | \sigma_i^{s-1}, \{\sigma_j^{s-1}\}_{j \in \partial i}) P_0 \right]. \end{aligned} \quad (16)$$

The single-site one-time and two-time marginal probabilities on the original graph then follow from (15):

$$P_i^{(t-1)}(\sigma'_i) \propto \sum_{\{\sigma'_j\}_{j \in \partial i}} \mathcal{G}^{(t-1)}(\sigma'_i, \{\sigma'_j\}_{j \in \partial i}), \quad (17)$$

$$P_i^{(t, t-1)}(\sigma_i, \sigma'_i) \propto \sum_{\{\sigma'_j\}_{j \in \partial i}} w_i(\sigma_i | \sigma'_i, \{\sigma'_j\}_{j \in \partial i}) \mathcal{G}^{(t-1)}(\sigma'_i, \{\sigma'_j\}_{j \in \partial i}). \quad (18)$$

In analogy to $\mathcal{F}$ above, we can write a recursive equation for $\mathcal{G}$ by using the Markovian assumption for messages:

$$\begin{aligned} \mathcal{G}^{(t)}(\sigma_i, \{\sigma_j\}_{j \in \partial i}) &= \sum_{\sigma'_i, \{\sigma'_j\}_{j \in \partial i}} \prod_{j \in \partial i} T_{j \rightarrow (ij)}(\sigma_j | \sigma'_j, \sigma'_i) \\ &\times w_i(\sigma_i | \sigma'_i, \{\sigma'_j\}_{j \in \partial i}) \mathcal{G}^{(t-1)}(\sigma'_i, \{\sigma'_j\}_{j \in \partial i}). \end{aligned} \quad (19)$$

Hence starting with an initial value for the functions T , $\mathcal{F}$, and $\mathcal{G}$, Eqs. (12), (13), and (19) can be iterated up to the desired time and Eq. (17) can be used to compute the time-dependent marginal probability of site i . We highlight that, unlike the original formulation, which takes an exponential time to compute marginals, the entire scheme presented here has a polynomial computational cost.

V. RESULTS

In this section we test the accuracy of our dynamic message-passing (DMP) approach on a statistical physics model often chosen as a case of study to investigate dynamics of complex systems: the kinetic Ising model.

Case study: The kinetic Ising model with arbitrary connectivity symmetry

We compare the performance of DMP to Monte Carlo Markov-chain simulations (MC) with Glauber dynamics on the kinetic Ising model on a random diluted graph with arbitrary connectivity symmetry: fully symmetric, partially asymmetric, and fully asymmetric networks. The comparison is performed computing the behavior of the magnetization of the model both during the transient and at equilibrium (or at the stationary states when detailed balance does not hold). Following Ref. [35], we introduce a connectivity matrix c_{ij} , where $c_{ij} = 1$ if there is a link from vertex i to vertex j , $c_{ij} = 0$ otherwise, and matrix elements c_{ij} and c_{kl} are independent unless $\{kl\} = \{ji\}$. The following distributions then specify

FIG. 3. (Color online) Comparison between dynamic message passing algorithm (solid lines) and Monte Carlo simulations (dashed lines) for the evolution of the magnetization $m(t)$ in an Erdős-Rényi network of $N = 5000$ nodes and average connectivity $c = 3$. Different colors represent different initial magnetizations $m(0)$, from the largest (upper line) to the smallest (lower line) according to the legend. (a) Fully asymmetric networks at high temperatures. [(b) and (c)] Partially asymmetric network ($\epsilon = 0.5$) respectively at high and low temperatures. [(d) and (e)] Fully symmetric network respectively at high and low temperatures. In MC simulations $m(t)$ is averaged over 5000 samples.

the graph topology: the marginal one-link distribution

$$p(c_{ij}) = \frac{c}{N} \delta_{1,c_{ij}} + \left(1 - \frac{c}{N}\right) \delta_{0,c_{ij}} \quad (20)$$

and the conditional distribution

$$p(c_{ij}|c_{ji}) = \epsilon \delta_{c_{ij},c_{ji}} + (1 - \epsilon) p(c_{ij}). \quad (21)$$

Above, N is the size of the network, c the average connectivity, and $\epsilon \in [0, 1]$ a parameter which controls the asymmetry. The value $\epsilon = 0$ gives a fully asymmetric network, whereas $\epsilon = 1$ gives a fully symmetric one.

In the kinetic Ising model the transition probability rate w_i , which appears above, is given by $w_i(\sigma_i^s | \{\sigma_j^{s-1}\}) = \exp[\beta \sigma_i^s J_{ij} \sigma_j^{s-1} - \log 2 \cosh(\beta \sum_j J_{ij} \sigma_j^{s-1})]$, where the j 's variable are neighbors of spin i , J_{ij} are the interaction strengths between sites i and j , and β is the inverse temperature. We observe that this model belongs to a subclass of the models considered within the general formulation above as the transition rate does not depend explicitly on σ_i^{s-1} . Since we here want to mainly investigate the effect of the asymmetry on the performances of our algorithm, we restrict our numerical analysis to ferromagnetic models, i.e., interaction strength $J_{ij} = J > 0$ for every pair of sites i, j . For simplicity, all the spins on the graph are chosen to be independent at the initial time although correlated initial conditions could also be considered in the above formulation. Results are shown in Fig. 3 for several values of the initial magnetization at different temperatures and for various values of the asymmetry parameter ϵ . Numerical results for fully asymmetric networks are obtained by using the simplified version (5) of the dynamic message-passing equation and, as expected from Refs. [31–33], they show a perfect agreement with Monte Carlo simulations both for the computation of the transient and the stationary state [see Fig. 3(a)]. When the full asymmetry is broken and a network with feedback is considered, numerics is obtained through the DMP scheme presented above. Comparison with MC shows that the results provided by DMP are still very good for high-enough temperature (above the critical transition) both for partially and fully symmetric connectivities. Indeed, as it is possible to note from Fig. 3 [partially asymmetric network in Fig. 3(b) and the fully

symmetric one in Fig. 3(d)], the transient regime is very well reproduced by DMP and the stationary (or equilibrium) state is the same as MC. When temperature is decreased below the critical ferromagnetic transition the performances of DMP begin to worsen [Figs. 3(c) and 3(e)]. In this regime, the transient is very well recovered only for the first few initial steps of the dynamics and progressively worsen for longer times. Nevertheless, for the partially asymmetric network considered ($\epsilon = 0.5$) the stationary state reached by the DMP and MC simulations coincide, although DMP has a faster convergence to that [Fig. 3(c)]. For fully symmetric networks similar considerations follow for the transient regime although the equilibrium state reached by the DMP and MC simulations at this temperature slightly differ [Fig. 3(e)]. This is not always the case; indeed, we noticed numerically that, for some of the temperatures even below the critical transition, DMP reaches the same equilibrium state as MC [see, for instance, Fig. 4(c)]. However, for fully symmetric networks, it is known that the stationary solutions of the one-time approximation (OTA) method presented in Refs. [24,31] are also the solution of the static BP equations. Therefore the agreement between the MC simulations and the one-time approximation for the equilibrium state of fully symmetric networks is expected to be near perfect, whereas, for some temperatures, the DMP approach proposed here presents small differences with the equilibrium MC solution.

In order to investigate the performances of DMP with respect to the one-time approximation, we compared both algorithms with MC simulations for different temperatures and for various values of the asymmetry parameter ϵ . The numerical investigation, partially illustrated in Fig. 4, shows that, regardless, the network connectivity symmetry DMP always outperforms OTA for the transient dynamic regime. For temperatures above the critical transition (T_c) both algorithms converge to the same stationary or equilibrium state, in agreement with the MC simulations [see Figs. 4(a) and 4(d)] and, surprisingly, the one-time approximation recovers well the MC stationary solution also for partially asymmetric networks [Fig. 4(d)]. For temperatures lower than T_c the following picture emerges. For fully symmetric networks OTA always reaches the same equilibrium state as MC, whereas DMP in some cases does [see Fig. 4(c)] and in some cases

FIG. 4. (Color online) Comparison among the dynamic message passing algorithm (solid lines with $\bullet$ dots), the one-time approximation (solid lines with $\blacklozenge$ dots), and the Monte Carlo simulations (solid lines) for the evolution of the magnetization $m(t)$ in an Erdős-Rényi network of $N = 5000$ nodes and average connectivity $c = 3$. [(a)–(c)] Fully symmetric network at high and low temperatures. [(d) and (e)] Partially symmetric networks with different connectivity symmetry at high and low temperatures. In the MC simulations $m(t)$ is averaged over 5000 samples.

does not [see Fig. 4(b)], depending on the temperature. For partially asymmetric networks, the performances of DMP get better and, surprisingly, OTA also gives fairly good results for the stationary states [see Fig. 4(d)]. The differences in the reconstruction of the transient and stationary regime of the DMP and the one-time approach can be understood, remembering that the two algorithms do not belong to the same class of approximations.

We conclude this section by underlining that we expect improvement of the results provided by the dynamic message-passing scheme presented, both for the transient and for the stationary or equilibrium states, when a higher-order closure of the DMP equations is made (see the next section).

VI. HIGHER-ORDER CLOSURES

To close (3) in the class of n -th order Markov processes, we consider, instead of $P_{i \rightarrow (ij)}^{(t-1)}$ and $P_{i \rightarrow (ij)}^{(t,t-1)}$ in (10) and (11), the marginalizations $P_{i \rightarrow (ij)}^{(t-1,t-2,\dots,t-n)}$ and $P_{i \rightarrow (ij)}^{(t,t-1,t-2,\dots,t-n)}$ and

auxiliary functions $F^{(n)}$ and $G^{(n)}$ which depend on n earlier times. The time iteration of $F^{(n)}$, $G^{(n)}$ and the iterative solution of the n th-order kernel $T_{i \rightarrow (ij)}^{(n)}[\sigma_i | \sigma'_i, \sigma'_j, \dots, \sigma_i(t-n), \sigma_j(t-n)]$ then proceed analogously as above. The computational cost obviously increases quickly with n and the value of higher-order closures will depend on the application and the model. This issue will be addressed in future contributions.

ACKNOWLEDGMENTS

The authors acknowledge valuable discussions with Federico Ricci-Tersenghi, Silvio Franz, Haiping Huang, Andrey Y. Lokhov, Yoshiyuki Kabashima, and Gabriele Perugini. This work was finalized during the program “Collective Dynamics in Information System” at Kavli Institute for Theoretical Physics China (KITPC) in Beijing, China. It has been funded under FP7/2007-2013/Grant No. 290038 (GDF) and supported by the Swedish Science Council through Grant No. 621-2012-2982 and by the Academy of Finland through its Center of Excellence COIN (E.A.).

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