Convergence of a Belief Propagation Algorithm for Biological Networks

Sang-Mok Choo and Young-Hee Kim

1Department of Mathematics, University of Ulsan, Ulsan 44610, Republic of Korea
2Ingenium College of Liberal Arts, Kwangwoon University, Seoul 01897, Republic of Korea

Correspondence should be addressed to Young-Hee Kim; yhkim@kw.ac.kr

Received 2 April 2019; Revised 7 June 2019; Accepted 24 June 2019; Published 24 July 2019

Academic Editor: Maria Alessandra Ragusa

Constructing network models of biological systems is important for effective understanding and control of the biological systems. For the construction of biological networks, a stochastic approach for link weights has been recently developed by using experimental data and belief propagation on a factor graph. The link weights were variable nodes of the factor graph and determined from their marginal probability mass functions which were approximated by using an iterative scheme. However, there is no convergence analysis of the iterative scheme. In this paper, at first, we present a detailed explanation of the complicated multistep process step by step with a network of small size and artificial experimental data, and then we show a sufficient condition for the convergence of the iterative scheme. Numerical examples are given to illustrate the whole process and to verify our result.

1. Introduction

Systems of mathematical equations have been used for modeling interactions of genes or proteins in biological systems. The modeling consists of two parts: one is to construct a network of nodes and links, where nodes usually represent genes or proteins and links denote interactions of nodes. The second is to build a set of mathematical formulas for governing the network dynamics, which might be expressed as a system of differential equations or Boolean logic functions [1, 2]. Modeling of a biological network has been focused on the construction of a directed network for a given biological system and its mathematical formulas, with which biological phenomena and mechanisms have been explained and controlled [3] and, furthermore, modeling of a biological network is a useful tool for understanding and explaining the distribution of the biological communities too [4]. However, there are few researches on how to determine an optimal set of nodes and links which are key components of networks for biological phenomena.

There could be multiple networks for explaining a given biological system. So, it would be meaningful to use a probabilistic modeling approach which could yield multiple networks for the biological system. One approach was recently developed in [5, 6] for the construction of network models of a melanoma cell line (SK-MEL-133), the most serious skin cancer, in which each link weight was considered as a discrete random variable and its marginal probability mass function (PMF) was constructed by using a factor graph and a belief propagation (BP) algorithm. The marginal PMFs were used to choose multiple weights of high-probability values and then to construct multiple networks for the biological system. The initial step of the approach was to determine a prior knowledge network (PKN) for melanoma and its mathematical formula structure. Nodes were chosen for representing some drugs (drug nodes) for melanoma or proteins (measured nodes) which take important roles in melanoma. The drug nodes had outgoing links to each measured node without incoming links. Each measured node had an outgoing link to each other measured node and an incoming link from each other measured node. Each link weight in the mathematical formulas was not determined in the initial step. Next, each protein concentration was measured at steady state before and after treatment of the
drugs, which produced experimental data for determining the link weights. Then, the joint PMF of link weights was defined as the Boltzmann/Gibbs form [7] for the factorization of the joint PMF. The form contained a cost function of the experimental data and the simulated values of measured nodes at steady state, where the simulated values were not constants for undetermined link weights. So, the factorization of the joint PMF directly led to the factor graph which is a bipartite graph consisting of variable nodes (link weights in the network) and factor nodes (factors in the factorization). Finally, the marginal PMFs were approximated with an iterative scheme constructed from belief (message) propagation on the factor graph.

In general, computation time could be high to directly compute marginal distributions from a joint probability distribution. So, factor graphs and BP have been applied to infer marginals in diverse problems [8–12]. It is known that a marginal obtained from applying BP becomes the true marginal for an acyclic factor graph and BP could be successfully applied to a cyclic graph [13–18]. There have been many researches on conditions for the convergence of beliefs, which can be applied for computing marginals from a Gaussian joint probability density function (PDF) [19, 20]. However, the convergence conditions cannot be applied to the convergence of beliefs (messages) in [5, 6], since the joint random variable is discrete. And the authors [5, 6] did not analyze the convergence. As far as we know, there is no framework for the inference of marginal PMFs based on both BP and perturbation data except the approach in [5, 6]. For the use of the stochastic approach [5] in the construction of networks based on BP as well as perturbation data, a sufficient condition for the convergence of messages is necessary. Under such a sufficient condition marginal PMFs can be identified and, as a result, multiple network models for the given biological system can be constructed.

In this paper, with our simple network, we present a detailed explanation of the long and intricate process of constructing a system of equations with messages from variable nodes to factor nodes, and vice versa for the approximation of marginal PMFs. Since approximate marginal PMFs are defined with the solution of the system of equations, we identify recursive relations for the solution of the system of equations and construct iterative schemes for solving the recursive relations. Finally we show a sufficient condition for the convergence of the schemes by using a Banach fixed-point theorem (see, e.g., [21, 22]). Numerical examples are given to illustrate the process in Section 4 and to verify our results.

2. Preliminaries

For a clear explanation of the convergence of the iterative schemes mentioned in the Introduction section, we make a simple network of three nodes and four links (see Figure 1(a)), which is assumed to be a PKN. Since the simple network has key components necessary for determining approximate marginal PMFs, the simple network can be extended to any network, including the biological network in [5]. So, our convergence analysis in the Results section can be applied for any network. The network has two measured nodes \((x_1, x_2)\) and one drug node \((x_3)\). We consider two treatments as the 1st and 2nd perturbations. Symbols \(x_i^{u} \ (u = 1, 2, \ i = 1, 2, 3)\) in Figure 1(b) denote \(\log P(x_i^{u}, \text{after}/x_i^{u}, \text{before})\), where \(x_i^{u}, \text{before}\) and \(x_i^{u}, \text{after}\) are the concentrations of \(x_i\) at steady state before and after the \(u^{th}\) perturbation, respectively. Since each drug node has an outgoing link to each measured node and no incoming link to any node, drug node \(x_3\) has two outgoing links to \(x_1\) and \(x_2\) with weights \(w_{2,3}\) and \(w_{2,3}\), respectively. Each measured node has an outgoing link to each other measured node and incoming links from each other nodes, and so, measured node \(x_1\) has one outgoing link to \(x_2\) with weight \(w_{1,2}\) and two incoming links from \(x_2\) and \(x_3\) with weights \(w_{1,2}\) and \(w_{1,3}\), respectively. Similarly, measured node \(x_2\) has one outgoing link to \(x_1\) with weight \(w_{2,1}\) and two incoming links from \(x_1\) and \(x_3\) with weights \(w_{2,1}\) and \(w_{2,3}\), respectively.

As in [5], the dynamics of the given situation is modeled with the mathematical formulas as in Figure 1(c). Simulated values \(x_i^{u, s}\) at steady state under the \(u^{th}\) perturbation are assumed to be the form

\[
x_i^{u, s} = \phi \left( \sum_{j=1, j \neq i}^{3} w_{i,j} x_j^{u, s} \right) \quad (i = 1, 2), \quad x_3^{u, s} = x_3^u, \tag{1}
\]

where \(\phi(x) = \tanh(x)\). From now on, weights \(w_{i,j}\) are considered as discrete random variables with three values \(\omega (-1, 0 \text{ and } 1)\) and their PMFs are denoted as follows: for \(\omega \in \{-1, 0, 1\}\)

\[
P \left( w_{1,2} = \omega \right), \quad P \left( w_{1,3} = \omega \right), \quad P \left( w_{2,1} = \omega \right), \quad P \left( w_{2,3} = \omega \right), \tag{2}
\]

which are approximately calculated in the Results section by using a factor graph and BP based on the factor graph. In [3], discretization over the 3 weight values led to a high rate of convergence and did not capture appropriate uncertainty in a probability distribution and then 11 weight values were used. However, our goals in this paper are to find a convergence condition for messages and to explain the complicated multistep process in [3] step by step using a prior knowledge network of small size. So, the three weight values are enough for our goals. For the extension of our results to networks of large size and real perturbation data, some notations and mathematical expressions used in our paper might be rather cumbersome in later sections.

3. Results

In this section, we present a system of equations for each marginal PMF, iterative schemes for the solutions and a sufficient condition for the convergence of the iterative schemes.

3.1. System of Equations for Marginal PMFs. To define a joint PMF of link weights based on the experimental data we
Consider the sum of squared errors between simulated values \( x_i^{\text{obs}} \) and experimental values \( x_i^v \)

\[
\sum_{i=1}^{3} \sum_{v=1}^{2} \left( x_i^{\text{obs}} - x_i^v \right)^2
\]

as well as the penalty

\[
\sum_{i=1}^{3} \sum_{j=1, j\neq i}^{3} \delta(w_{ij}), \quad \delta(w_{ij}) = \begin{cases} 
0 & (w = 0) \\
1 & (w \neq 0)
\end{cases}
\]

which reflects the property that the total number of links tends not to be large. Note that \( w_{3j} = 0 \) due to no incoming link to drug node \( x_3 \). Then the cost function is defined as

\[
\text{Cost} = \beta \sum_{i=1}^{3} \sum_{v=1}^{2} \left( x_i^{\text{obs}} - x_i^v \right)^2 + \lambda \sum_{i=1}^{3} \sum_{j=1, j\neq i}^{3} \delta(w_{ij}),
\]

where the error and penalty are weighted by positive constants \( \beta \) and \( \lambda \), respectively. Using (1), the cost is written as

\[
\text{Cost} = \beta \sum_{i=1}^{3} \sum_{v=1}^{2} \left\{ \frac{3}{j=1, j\neq i} \left[ x_i^{\text{obs}} - x_i^v \right] - x_i^v \right\}^2 + \lambda \sum_{i=1}^{3} \sum_{j=1, j\neq i}^{3} \delta(w_{ij}).
\]

Posteriori probability (joint PMF) \( P(W) = P(w_{1,2}, w_{1,3}, w_{2,1}, w_{2,3}) \) is defined as

\[
P(W) = \frac{1}{Z} \exp \left( -\text{Cost} \right),
\]

where \( Z \) is the constant ensuring that the sum of the probabilities over all model configurations is equal to one. Let \( Z = Z_1 Z_2 \), where \( Z_1 \) and \( Z_2 \) are the constants ensuring that the sums of probabilities \( P(w_{1,2}, w_{1,3}) \) and \( P(w_{2,1}, w_{2,3}) \) over all model configurations are equal to one, respectively. Substituting the cost (6) into \( P(W) \) gives

\[
P(W) = \frac{1}{Z} \exp \left[ \sum_{i=1}^{2} \left( -\beta \sum_{v=1}^{2} \left\{ \frac{3}{j=1, j\neq i} \left[ x_i^{\text{obs}} - x_i^v \right] - x_i^v \right\}^2 - \lambda \sum_{j=1, j\neq i}^{3} \delta(w_{ij}) \right) \right].
\]
\[\begin{align*}
&= \prod_{i=1}^{2} \left( \frac{1}{Z_{i}} \exp \left\{ -\lambda \sum_{j=1,j\neq i}^{3} \delta (w_{i,j}) \right\} \right) \prod_{i=1}^{3} \exp \left[ -\beta \left\{ \phi \left( \sum_{j=1,j\neq i}^{3} w_{i,j}x_{j}^{v} \right) - x_{i}^{v} \right\}^{2} \right] \\
&\equiv P (w_{1,2}, w_{1,3}) P (w_{2,1}, w_{2,3}),
\end{align*}\]

which implies that \( P(w_{1,2}, w_{1,3}) \) and \( P(w_{2,1}, w_{2,3}) \) can be independently determined by using a similar approach. So, we consider \( P(w_{1,2}, w_{1,3}) \) to show the approximation of marginal PMF \( P(w_{1,2}) \):

\[\begin{align*}
P (w_{1,2}) &= \sum_{w_{1,3}} P (w_{1,2}, w_{1,3}) \\
&= \frac{1}{Z_{1,2}} \sum_{w_{1,3}} e^{-\lambda \sum_{j=1}^{3} \delta (w_{1,j})} \prod_{i=1}^{2} e^{-\beta \left( \phi \left( \sum_{j=1,j\neq i}^{3} w_{i,j}x_{j}^{v} \right) - x_{i}^{v} \right)^{2}}.
\end{align*}\] (9)

It is not efficient to calculate the exact marginal with the enumeration in the cases where the numbers of nodes in a network or experimental perturbations become large. Therefore, the process to approximate the marginal is presented step by step by using a factor graph and BP on the factor graph.

**Step 1** (introduction of a factor graph and BP on the factor graph). Using the factorization in (9), factor nodes \( F_{v}^{i} (v = 1, 2) \) are defined as

\[F_{v}^{i} (w_{1,2}, w_{1,3}) = \exp \left[ -\beta \left\{ \phi \left( \sum_{j=1,j\neq i}^{3} w_{i,j}x_{j}^{v} \right) - x_{i}^{v} \right\}^{2} \right]\] (10)

and then marginal \( P(w_{1,2}) \) in (9) can be written as follows:

\[\begin{align*}
P (w_{1,2}) &= \frac{1}{Z_{1,2}} \sum_{w_{1,3}} e^{-\lambda \sum_{j=1}^{3} \delta (w_{1,j})} \prod_{i=1}^{2} F_{v}^{i} (w_{1,2}, w_{1,3}),
\end{align*}\] (11)

which yields the factor graph of two variable nodes \((w_{1,2}, w_{1,3})\) and two factor nodes \((F_{1}^{1}, F_{2}^{2})\) as in Figure 2. Following BP on the factor graph, message \( F_{v}^{i} (w_{1,j}) \) from variable node \( w_{1,j} \) to factor node \( F_{v}^{i} (w_{1,2}, w_{1,3}) \) is defined as

\[\begin{align*}
P_{v}^{i} (w_{1,j}) &= \frac{1}{Z_{1,2}^{v}} e^{-\lambda \delta (w_{1,j})} \prod_{v=1,v\neq v}^{2} P_{v}^{i} (w_{1,j}) \\
&= \prod_{v=1,v\neq v}^{2} P_{v}^{i} (w_{1,j}) \prod_{2 \leq i \leq 3, i\neq j}^{3} P_{v}^{i} (w_{1,j}).
\end{align*}\] (12)

Using message \( \rho^{v} (w_{1,j}) \), marginal PMF \( P(w_{1,j}) \) in (12) can be approximated as

\[\begin{align*}
P (w_{1,j}) &= \frac{1}{Z_{1,2}^{v}} e^{-\lambda \delta (w_{1,j})} \prod_{v=1}^{2} P_{v}^{i} (w_{1,j}),
\end{align*}\] (14)

where \( Z_{1,2}^{v} \) is the normalization constant ensuring that the sum of the probabilities over all model configurations is equal to one.

**Step 2** (approximation of the summation (13)). The process of the approximation used in (5) can be divided into two parts: the first is to change multiple summations into a single summation by introducing a new random variable. The next is to change the single summation into an integral.

**Step 2-1.** Since \( \rho^{v} \) in (13) is a function of \( w_{1,j} \), all random variables in \( F_{v}^{i} (w_{1,2}, \ldots w_{1,n}) \), which are in (13), can be divided into two type of random variables: one is \( w_{1,j} \) and the other is \( s_{1,j}^{v} = \sum_{1<j}^{n} w_{1,k}x_{k}^{v} \) \((n = 3)\), which becomes the linear combination of random variables \( w_{1,j} \) in the cases where the number \( n \) of nodes becomes large. Using the definitions of \( s_{1,j}^{v} \) and \( F_{v}^{i} (w_{1,2}, \ldots w_{1,n}) \) in (10), we can write \( F_{v}^{i} \) in (13) as

\[\begin{align*}
F_{v}^{i} (s_{1,j}^{v}, w_{1,j}) &= \exp \left\{ -\beta \left( s_{1,j}^{v} + w_{1,j}x_{j}^{v} \right) - x_{i}^{v} \right\}^{2}\right),
\end{align*}\] (15)

which is a function of random variables \( s_{1,j}^{v} \) and \( w_{1,j} \). Then (13) becomes

\[\rho^{v} (w_{1,j}) = \sum_{w_{1,j} \in \{1 \leq s_{1,j}^{v} \leq n \} \setminus \{1, j\}} F_{v}^{i} (s_{1,j}^{v}, w_{1,j}) \prod_{2 \leq i \leq 3, i\neq j}^{3} P_{v}^{i} (w_{1,j}).\] (16)

Letting

\[\sum_{1 \leq s_{1,j}^{v} \leq n} \bar{w}_{1,k} x_{k}^{v} \mid \bar{w}_{1,k} \in \{-1, 0, 1\}, 1 < \xi \leq n, \xi \not\in j\}
\]

\[\bar{w}_{1,k} \in \{-1, 0, 1\}, 1 < \xi \leq n, \xi \not\in j\} = \{s_{1,j,k}^{v} \mid 1 \leq k \leq m\}.
\] (17)
for some positive integer \( m \), message \( \rho^v \) in (16) becomes

\[
\rho^v (w_{1,j}) = \sum_{k=1}^{m} \sum_{\{w_{1,k}=w_{1,j}(1 \leq \xi \leq n, \xi \neq j)\}} \left\{ F_{1}^{v} \left( s_{1,j}, w_{1,j} \right) \right\} \prod_{2 \leq \xi \leq m, \xi \neq j} P^v \left( w_{1,\xi} \right)
\]

Note that

\[
\sum_{k=1}^{m} \sum_{\{w_{1,k}=w_{1,j}(1 \leq \xi \leq n, \xi \neq j)\}} \prod_{2 \leq \xi \leq m, \xi \neq j} P^v \left( w_{1,\xi} \right) = 1,
\]

which implies that the following can be a PMF of \( s^v_{1,j} \)

\[
P_s^v \left( s_{1,j} = s^v_{1,j,k} \right) = \sum_{\{w_{1,k}=w_{1,j}(1 \leq \xi \leq n, \xi \neq j)\}} \prod_{2 \leq \xi \leq m, \xi \neq j} P^v \left( w_{1,\xi} \right)
\]

(1 \leq k \leq m).

So, message \( \rho^v \) in (18) can be written as a single summation

\[
\rho^v (w_{1,j}) = \sum_{s_{1,j}^v} F_{1}^{v} \left( s_{1,j}^v, w_{1,j} \right) P^v \left( s_{1,j}^v \right).
\]

Step 2. Note that \( s_{1,j}^v \) is a sum of random variables \( w_{1,\ell} \) (1 < \( \xi \leq n, \xi \neq j \)), which are assumed to be independent. Even though \( s_{1,j}^v \) is not identically distributed, the authors [5] invoked the central limit theorem to approximate the PMF of \( s_{1,j}^v \) as a Gaussian with reference to [23], where there was no explicit justification for the application of this theorem. Since sums of independent random variables converge in distribution to the standard normal as long as some condition (e.g., the Lindeberg Condition [24]) is satisfied, we think that such a condition might be implicitly assumed in [24]. So, the approximate PMF of \( s_{1,j}^v \) becomes

\[
P_s^v \left( s_{1,j}^v \right) = \frac{1}{\sqrt{2\pi \Delta_{1,j}^v}} \exp \left\{ -\frac{(s_{1,j}^v - \overline{s}_{1,j})^2}{2\Delta_{1,j}^v} \right\},
\]

where average \( \overline{s}_{1,j} \) and variance \( \Delta_{1,j}^v \) of \( s_{1,j}^v \) are defined as

\[
\overline{s}_{1,j} = E(\overline{s}_{1,j}) = E \left( \sum_{\ell \neq 1,j} w_{1,\ell} x_{\ell}^v \right) = \sum_{\ell \neq 1,j} \overline{w}_{1,\ell} x_{\ell}^v,
\]

\[
\Delta_{1,j}^v = V(s_{1,j}) = V \left( \sum_{\ell \neq 1,j} w_{1,\ell} x_{\ell}^v \right) = \sum_{\ell \neq 1,j} V(w_{1,\ell}) (x_{\ell}^v)^2.
\]

Symbols \( \overline{w}_{1,\ell} \) and \( \overline{w}_{1,\ell} x_{\ell}^v \) denote the averages of \( w_{1,\ell} \) and \( (w_{1,\ell})^2 \), respectively:

\[
\overline{w}_{1,\ell} = \sum_w w P^v (w_{1,\ell} = w),
\]

\[
\overline{w}_{1,\ell} x_{\ell}^v = \sum_w w^2 P^v (w_{1,\ell} = w)
\]

and then the sum over configurations \( \{w_{1,\ell} | 2 \leq \ell \leq 3, \ell \neq j\} \) in (19) is approximated with a Gaussian integration:

\[
\rho^v (w_{1,j}) \approx \int_{-\infty}^{\infty} F_{1}^{v} \left( s_{1,j}, w_{1,j} \right) P^v \left( s_{1,j} \right) ds_{1,j}^v.
\]

Step 3 (approximation of the improper integral (26)). For the calculation of the improper integral (26), error \( \phi(s_{1,j}^v + w_{1,\ell} x_{\ell}^v) - x_{1}^v \) is linearized with respect to the maximization of the fitness in (15). Note that the equality

\[
\phi \left( s_{1,j} + w_{1,\ell} x_{\ell}^v \right) - x_{1}^v = 0,
\]
can be written as
\[ \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v = 0, \] (28) under the assumption that

experimental data \( x_i^v \) are contained in the codomain of \( \phi \). (29)

Then error \( \phi(s_{1,j}^v + w_{1,j}x_j^v) - x_1^v \) in (15) is approximated by
\[ \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v, \] (30)
which yields
\[ F_i^v(s_{1,j}^v, w_{1,j}) = \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\} \]
\[ \approx \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\}. \] (31)

Hence we have
\[ \rho^v(w_{1,j}) = \int_{-\infty}^{\infty} F_i^v(s_{1,j}^v, w_{1,j}) P_i^v(s_{1,j}^v) \, ds_{1,j}^v \]
\[ = \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\} \cdot \frac{1}{\sqrt{2\pi \Delta_{1,j}^v}} \int_{-\infty}^{\infty} \exp \left\{ -\frac{1}{2} \left( \frac{s_{1,j}^v - \overline{s}_{1,j}^v}{\Delta_{1,j}^v} \right)^2 \right\} \, ds_{1,j}^v \]
\[ = \frac{1}{\left( 1 + 2\beta \Delta_{1,j}^v \right)^{1/2}} \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\}, \] (32)
where the last equality is obtained both by the identity

and the property of a PDF (the integral of a PDF over its domain is equal to 1). Therefore adopting the final approximation of \( \rho^v \) in [5], approximate marginal PMF \( P(w_{1,j} = w) \) in (14) can be obtained by solving the system of equations
\[ \rho^v(w_{1,j} = w) = \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\}, \] (34)
\[ p^v(w_{1,j} = w) = \frac{1}{Z_{2,j}^v} e^{-\lambda \delta(w_i = w)} \rho^{3-v}(w_{1,j} = w), \] (35)
where \( v = 1, 2, j = 2, 3, w = -1, 0, 1, \overline{s}_{1,j} \) and \( \Delta_{1,j}^v \) are in (23)–(25).

**Remark 1.** Since the approximation (34) was used in [5] and our goal in this paper is to find a convergence condition for the algorithm in [5], we also use (34) instead of
\[ \rho^v(w_{1,j}) = \frac{1}{\left( 1 + 2\beta \Delta_{1,j}^v \right)^{1/2}} \exp \left\{ -\beta \left( \phi^{-1}(x_1^v) - w_{1,j}x_j^v - s_{1,j}^v \right)^2 \right\}. \] (36)

Even when using (36) instead of (34), the convergence analysis in the later subsections could be applied (we do not show it in this paper).

**Remark 2.** Similarly approximate marginal PMF \( P(w_{2,j}) \) can be obtained by replacing subscripts 1 and \( j = 2, 3 \), related with node \( x_1 \), in \( w_{1,j}, s_{1,j}^v, \Delta_{1,j}^v \), and \( w_{1,k} \) in (34), (35), and (23)–(25) with 2 and \( j = 1, 3 \), related with node \( x_2 \): for \( v = 1, 2, j = 1, 3; \)
\[ P_{2,j}^v(w_{2,j}) = \frac{1}{Z_{2,j}^v} e^{-\lambda \delta(w_i = w)} \rho^{3-v}(w_{2,j}), \] (37)
\[ p_{2,j}^v(w_{2,j}) = \exp \left\{ -\beta \left( \phi^{-1}(x_2^v) - w_{2,j}x_j^v - s_{2,j}^v \right)^2 \right\}, \] (38)
\[ P_{2,j}^v(w_{2,j}) = \frac{1}{Z_{2,j}^v} e^{-\lambda \delta(w_i = w)} \rho^{3-v}(w_{2,j}), \] (39)
where \( Z_{2,j}^v \) is the normalization constant ensuring that the sum of the probabilities over all model configurations is equal to one and
\[ \overline{x}_{2,j} = \sum_{\ell \neq 2,j} \left\{ \sum_{w} w^3 P_{1,\ell,0}^v (w_{2,\ell} = w) \right\} x_{\ell}^u, \quad (40) \]

\[ \Delta_{2,j}^u = \sum_{\ell \neq 2,j} \left\{ \left( \sum_{w} w^2 P_{1,\ell,0}^v (w_{2,\ell} = w) \right)^2 - \left( \sum_{w} w P_{1,\ell,0}^v (w_{2,\ell} = w) \right)^2 \right\} (x_{\ell}^u)^2. \quad (41) \]

### 3.2. Iterative Schemes for Solving the System of Equations.

In this subsection, we construct sequences \( \{\rho_{1,j,n}^u (w)\} \) and \( \{P_{1,j,n}^v (w)\} \) \((v = 1, 2, j = 2, 3, w = -1, 0, 1)\) by using (34) and (35). In the next subsection we present a sufficient condition for the convergence of the two sequences, which implies that the two limits satisfy (34) and (35). So, the limit of \( \{\rho_{1,j,n}^u (w)\} \) becomes value \( \rho_{1,j}^u (w) \), leading to the construction of approximate marginal PMF \( P(w_{1,j}) \) in (14). We assume that initial terms \( \rho_{1,j,0}^u (w) \) of \( \{\rho_{1,j,n}^u (w)\} \) are given as positive numbers (42)

\[ \rho_{1,j,0}^u (w) = \frac{1}{Z_{1,j,0}} e^{-\lambda \delta (w)} \rho_{1,3,j,0}^u (w), \quad (43) \]

and initial terms of \( \{P_{1,j,n}^v (w)\} \) are defined as

\[ P_{1,j,0}^v (w) = \frac{1}{Z_{1,j,0}} e^{-\lambda (w)} P_{1,3,j,0}^v (w), \]

where \( Z_{1,j,0}^u \) is the normalization constant ensuring that the sum of the probabilities over all model configurations is equal to one. The 1st iterations \( \rho_{1,j,1}^u (w) \) and \( P_{1,j,1}^v (w) \) are defined similarly to (34) and (35) and so we need to define \( s_{1,j,0}^u \) and \( \Delta_{1,j,0}^u \) by using (23) and (24):

\[ s_{1,j,0}^u = \sum_{\ell \neq 1,j} \left\{ \sum_{w} w^3 P_{1,\ell,0}^v (w_{1,\ell} = w) \right\} x_{\ell}^u \]

\[ = \sum_{\ell \neq 1,j} \left\{ \sum_{w} w P_{1,\ell,0}^v (w_{1,\ell} = w) \right\} (x_{\ell}^u)^2 \]

\[ \Delta_{1,j,0}^u = \sum_{\ell \neq 1,j} \left\{ \left( \sum_{w} w^2 P_{1,\ell,0}^v (w_{1,\ell} = w) \right)^2 - \left( \sum_{w} w P_{1,\ell,0}^v (w_{1,\ell} = w) \right)^2 \right\} (x_{\ell}^u)^2 \]

\[ = \frac{e^{-\lambda}}{Z_{1,5-j,0}} \left\{ \rho_{1,5-j,0}^3 (w_{1,5-j,0} (1)) - \rho_{1,5-j,0}^3 (w_{1,5-j,0} (1)) \right\} (x_{5-j}^u)^2. \quad (45) \]
So, the $1^{st}$ iteration is defined as

$$
\rho_{1,j,1}^w (w) = \exp \left[ -\beta \left\{ \phi^{-1} (x_{j}^w) - wx_{j}^w - \frac{\sigma^2}{1 + 2\beta \Delta_{1,j,0}} \right\} \right],
$$

(46)

$$
P_{1,j,1}^w (w) = \frac{1}{Z_{1,j,1}^w} \exp \left[ -\beta \Delta_{1,j,0} \right] \rho_{1,j,1}^w (w),
$$

where $Z_{1,j,1}^w$ is the normalization constant ensuring that the sum of the probabilities over all model configurations is equal to one. Similarly the $(n+1)^{th}$ iteration is defined as

$$
\Phi_{1,j,w}^v \left( \rho_{1,5-j,n}^3 (-1), \rho_{1,5-j,n}^3 (1) \right)
$$

$$
= \exp \left[ -\beta \left\{ \phi^{-1} (x_{j}^w) - wx_{j}^w - \frac{\sigma^2}{1 + 2\beta \Delta_{1,j,0}} \right\} \right].
$$

(49)

Therefore the desired iterative schemes consist of (42), (43), (47), (48), and (49) under the assumption (29).

Remark 3. Since $P_{1,j,n+1}^w (w_{1,j})$ defined in (48) is a PMF due to (47) and (49), we have

$$
0 < \frac{e^{-\lambda}}{Z_{1,5-j,n}^w} \rho_{1,5-j,n}^3 (-1) < 1,
$$

(50)

$$
0 < \frac{(e^{-\lambda})^3}{Z_{1,5-j,n}^w} \rho_{1,5-j,n}^3 (1) < 1,
$$

which gives the positivity of the following term in (49)

$$
\rho_{1,5-j,n}^3 (-1) + \rho_{1,5-j,n}^3 (1) - \frac{e^{-\lambda}}{Z_{1,5-j,n}^w} \left( \rho_{1,5-j,n}^3 (-1) + \rho_{1,5-j,n}^3 (1) \right)^2 > 0.
$$

(51)

So, we obtain a lower bound of $\rho_{1,j,n+1}^w (w)$

$$
\rho_{1,j,n+1}^w (w) \geq \exp \left[ -\beta \left\{ \phi^{-1} (x_{j}^w) + x_{j}^w + 2|x_{5-j}^w| \right\} \right].
$$

(52)
\[
\Phi_{2,j,\omega}^{\nu} \left( \rho_{2,4-j,\nu}^{3-v}(-1), \rho_{2,4-j,\nu}^{3-v}(1) \right) = \exp \left[ -\beta \frac{\left\{ \phi^{-1}(x_2) - wx_2 - (e^{-\lambda}/Z_{2,4-j,\nu}) \left( -\rho_{2,4-j,\nu}^{3-v}(-1) + \rho_{2,4-j,\nu}^{3-v}(1) \right)x_4 \right\}^2}{1 + 2\beta \left( e^{-\lambda}/Z_{2,4-j,\nu} \right) \left( \left( \rho_{2,4-j,\nu}^{3-v}(-1) + \rho_{2,4-j,\nu}^{3-v}(1) \right)x_4 \right)^2} \right].
\]

3.3. A Sufficient Condition for the Convergence of the Iterative Schemes. Let

\[
X_i^{(n)} = \begin{pmatrix}
X_{1,1}^{(n)}, X_{1,2}^{(n)}, X_{1,3}^{(n)} \\
X_{1,4}^{(n)}, X_{1,5}^{(n)}, X_{1,6}^{(n)} \\
X_{1,7}^{(n)}, X_{1,8}^{(n)}, X_{1,9}^{(n)} \\
X_{1,10}^{(n)}, X_{1,11}^{(n)}, X_{1,12}^{(n)}
\end{pmatrix}
\]

\[
= \begin{pmatrix}
\rho_{1,2,\nu}^{1}(-1), \rho_{1,2,\nu}^{1}(0), \rho_{1,2,\nu}^{1}(1) \\
\rho_{1,3,\nu}^{1}(-1), \rho_{1,3,\nu}^{1}(0), \rho_{1,3,\nu}^{1}(1) \\
\rho_{1,2,\nu}^{2}(-1), \rho_{1,2,\nu}^{2}(0), \rho_{1,2,\nu}^{2}(1) \\
\rho_{1,3,\nu}^{2}(-1), \rho_{1,3,\nu}^{2}(0), \rho_{1,3,\nu}^{2}(1)
\end{pmatrix} \in R^{12}
\]

and

\[
\Phi_1 \left( X \right) = \begin{pmatrix}
\Phi_{1,1} \Phi_{1,2} \Phi_{1,3} \\
\Phi_{1,4} \Phi_{1,5} \Phi_{1,6} \\
\Phi_{1,7} \Phi_{1,8} \Phi_{1,9} \\
\Phi_{1,10} \Phi_{1,11} \Phi_{1,12}
\end{pmatrix}
\]

\[
= \begin{pmatrix}
\Phi_{1,2,\nu}^{1}(-1), \Phi_{1,2,\nu}^{1}(0), \Phi_{1,2,\nu}^{1}(1) \\
\Phi_{1,3,\nu}^{1}(-1), \Phi_{1,3,\nu}^{1}(0), \Phi_{1,3,\nu}^{1}(1) \\
\Phi_{1,2,\nu}^{2}(-1), \Phi_{1,2,\nu}^{2}(0), \Phi_{1,2,\nu}^{2}(1) \\
\Phi_{1,3,\nu}^{2}(-1), \Phi_{1,3,\nu}^{2}(0), \Phi_{1,3,\nu}^{2}(1)
\end{pmatrix},
\]

where the subscript 1 of \(X_i\) and \(\Phi_1\) represents node 1. The \((n+1)\)th iteration in (47) is written as

\[
X_i^{(n+1)} = \Phi_1 \left( X_i^{(n)} \right).
\]

It follows from (47), (57), and (58) that \(\Phi_{1,\theta} (1 \leq \theta \leq 12)\) are functions of two independent variables: for \(X \in R^{12}\)

\[
\Phi_1 \left( X \right) = \begin{pmatrix}
\Phi_{1,1} \hspace{1cm} \Phi_{1,2} \hspace{1cm} \Phi_{1,3} \\
\Phi_{1,4} \hspace{1cm} \Phi_{1,5} \hspace{1cm} \Phi_{1,6} \\
\Phi_{1,7} \hspace{1cm} \Phi_{1,8} \hspace{1cm} \Phi_{1,9} \\
\Phi_{1,10} \hspace{1cm} \Phi_{1,11} \hspace{1cm} \Phi_{1,12}
\end{pmatrix} = \begin{pmatrix}
\Phi_{1,2,\nu}^{1}(-1), \Phi_{1,2,\nu}^{1}(0), \Phi_{1,2,\nu}^{1}(1) \\
\Phi_{1,3,\nu}^{1}(-1), \Phi_{1,3,\nu}^{1}(0), \Phi_{1,3,\nu}^{1}(1) \\
\Phi_{1,2,\nu}^{2}(-1), \Phi_{1,2,\nu}^{2}(0), \Phi_{1,2,\nu}^{2}(1) \\
\Phi_{1,3,\nu}^{2}(-1), \Phi_{1,3,\nu}^{2}(0), \Phi_{1,3,\nu}^{2}(1)
\end{pmatrix}.
\]

We use Banach fixed-point theorem [21] for the convergence of the sequence (59) to prove Theorem 9, which is our main result.

**Theorem 6.** Let \(D\) be a closed subset of \(R^m\) for a positive integer \(m\). If a function \(\Psi : D \rightarrow D\) satisfies that for a constant \(k \in (0,1)\) and all \(x, y\) in \(D\)

\[
\| \Psi(x) - \Psi(y) \| \leq k \| x - y \|,
\]

then there exists a unique fixed point \(x^* \in D\) such that \(\Psi(x^*) = x^*, \) which is the limit of sequence \(x^{(n+1)} = \Psi(x^{(n)}) \) \((n \geq 0)\) for any \(x^{(0)} \in D\).

For the application of Theorem 6 to the proof of our main result we need the following lemmas.

**Lemma 7.** Assume that experimental data \(x^{(n)}\) are contained in the codomain of \(\Phi\). Assume that positive constants \(\beta\) and \(\lambda\) satisfy inequalities \(1 - 2\beta(e^{-\lambda}U_{i,\nu,\lambda}^m(x_i^m)^2 > 0\) for \(\nu = 1, 2, j = 2, 3\) and \(U_{i,\nu,\lambda}^m\) defined in (53). Then \(\Phi_1\) defined in (49) and (57)–(60) becomes a function from domain \([0,1]\) to codomain \([0,1]\).

**Proof.** It follows from (49) and (57)–(60) that for \((\theta, w) = (1, -1), (2, 0), (3, 1)\)

\[
\Phi_{1,\theta}(X) = \Phi_{1,2,\omega}^1 \left( X_{1,\omega}, X_{12} \right),
\]

\[
X = (X_1, \ldots, X_{12}) \in [0,1]^{12}
\]

and

\[
\Phi_{1,2,\omega}^1 (X_{10}, X_{12}) = \exp \left[ -\beta \frac{\left\{ \phi^{-1}(x_1) - wx_1 - (e^{-\lambda}/Z_{1,3,\nu}) \left( -x_{10} + x_{12} \right)x_3 \right\}^2}{1 + 2\beta \left( e^{-\lambda}/Z_{1,3,\nu} \right) \left( \left( -x_{10} + x_{12} \right)x_3 \right)^2} \right].
\]
We have that for $\beta > 0$, $\lambda > 0$ and $\{X_{10}, X_{12}\} \subset [0, 1]$

\[
1 + 2\beta \frac{e^{-\lambda}}{Z_{1,3,n}^1} \left( (X_{10} + X_{12}) - \frac{e^{-\lambda}}{Z_{1,3,n}^1} (-X_{10} + X_{12})^2 \right) \\ \cdot (x_3^1) \geq 1 - 2\beta \left( e^{-\lambda} U_{1,3,n}^1 \right) (x_3^1)^2 > 0,
\]

and for $\beta > 0$

\[
-\beta \left\{ \phi^{-1} (x_1^1) - w x_2^1 + \frac{e^{-\lambda}}{Z_{1,3,n}^1} (-X_{10} + X_{12}) x_3^1 \right\}^2 \leq 0,
\]

and so, we obtain

\[
0 < \Phi_{1,\beta} (X) \leq 1 \quad (1 \leq \theta \leq 3).
\]

Similarly, under the same condition, we can obtain

\[
0 < \Phi_{1,\beta} (X) \leq 1 \quad (4 \leq \theta \leq 12),
\]

which are desired results.

**Lemma 8.** Assume that experimental data $x_i^n$ are contained in the codomain of $\Phi$. Assume that positive constants $\beta$ and $\lambda$ satisfy inequalities $1 - 2\beta \left( e^{-\lambda} U_{1,3,n}^1 \right) (x_i^n)^2 > 0$ for $\nu = 1, 2$, $j = 2, 3$ and $U_{1,3,n}^1$ defined in (53). Then for $\Phi_i$ defined in (49) and (57)–(60) and $\{X, Y\} \subset [0, 1]^2$

\[
\|\Phi_i (X) - \Phi_i (Y)\| \leq \sqrt{6} \max \left\{ e_{1,\beta,\lambda} \mid \nu = 1, 2, j = 2, 3 \right\} \beta e^{-\lambda} \|X - Y\|,
\]

where

\[
\begin{align*}
C_{1,\beta,\lambda}^v &= \left\{ \left[ \phi^{-1} (x_i^1) + x_i^2 - \frac{e^{-\lambda}}{Z_{1,3,n}^1} (-X_{10} + X_{12}) x_3^1 \right] \left[ e^{-\lambda} / Z_{1,3,n}^1 \right] x_3^2 \times \left[ 1 + 2\beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) - \left( e^{-\lambda} / Z_{1,3,n}^1 \right) (-\zeta + \eta)^2 \right) \right] \\
&\quad \times \left[ -\phi^{-1} (x_i^1) + x_i^2 - \frac{e^{-\lambda}}{Z_{1,3,n}^1} (-\zeta + \eta) x_3^1 \right]^2 \\
&\quad \times \left[ 1 + 2\beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) - \left( e^{-\lambda} / Z_{1,3,n}^1 \right) (-\zeta + \eta)^2 \right) \right] \\
&\quad \times \left[ 1 - 2\beta \left( e^{-\lambda} U_{1,3,n}^1 \right) (x_i^1)^2 \right].
\end{align*}
\]

**Proof.** Using (62) and (63), we have that for $\{X, Y\} \subset [0, 1]^2$

\[
\Phi_{1,1} (X) - \Phi_{1,1} (Y) = \Phi_{1,2,1} (X_{10}, X_{12}) - \Phi_{1,2,1} (Y_{10}, Y_{12})
\]

\[
= \exp \left[ f (X_{10}, X_{12}) \right] - \exp \left[ f (Y_{10}, Y_{12}) \right],
\]

where $\{X_{10}, X_{12}, Y_{10}, Y_{12}\} \subset [0, 1]$ and function $f : [0, 1]^2 \rightarrow [0, 1]^2$ is defined as follows: for $\zeta, \eta \in [0, 1]$

\[
f (\zeta, \eta) = -\beta \frac{\phi^{-1} (x_i^1) - (\zeta) x_i^2 - \left( (\zeta)^2 / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) x_i^1 \right)^2}{1 + 2\beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) - \left( e^{-\lambda} / Z_{1,3,n}^1 \right) (-\zeta + \eta)^2 \right) (x_i^1)^2}.
\]

Due to the mean value theorem there exists a constant $c$ in $(0, 1)$ such that

\[
\exp \left[ f (X_{10}, X_{12}) \right] - \exp \left[ f (Y_{10}, Y_{12}) \right] = \left[ \nabla \exp \left[ f \left( (1 - c) (X_{10}, X_{12}) + c (Y_{10}, Y_{12}) \right) \right] \right] \cdot (X_{10} - Y_{10}, X_{12} - Y_{12}).
\]

Note that

\[
\left| \frac{\partial}{\partial \zeta} \exp \left[ f (\zeta, \eta) \right] \right| \leq \left| \frac{\partial}{\partial \zeta} \exp \left[ f (\zeta, \eta) \right] \right|,
\]

where

\[
\frac{\partial}{\partial \zeta} f (\zeta, \eta) = -\beta \frac{\phi^{-1} (x_i^1) + x_i^2 - \left( (\zeta)^2 / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) x_i^1 \right)^2}{1 + 2\beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \left( (\zeta + \eta) - \left( e^{-\lambda} / Z_{1,3,n}^1 \right) (-\zeta + \eta)^2 \right) (x_i^1)^2}.
\]
Using \( \{\zeta, \eta\} \in [0, 1] \), we have

\[
\left| \frac{\partial}{\partial \zeta} f (\zeta, \eta) \right| \leq \beta \left[ \frac{2 \left\{ \phi^{-1} (x_1^1) + \frac{e^{-\lambda}}{Z_{1,3,n}^1} x_1^1 \right\} + \left( e^{-\lambda} / Z_{1,3,n}^1 \right) x_1^1 \right]
\times \left[ 1 + 2 \beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \right]
\times \left( 2 + e^{-\lambda} / Z_{1,3,n}^1 \right) (x_1^2)^2
\times \left. \frac{\left\{ \phi^{-1} (x_1^1) + \frac{e^{-\lambda}}{Z_{1,3,n}^1} x_1^1 \right\} + \left( e^{-\lambda} / Z_{1,3,n}^1 \right) x_1^1 \right|}{\left[ 1 + 2 \beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \right]
\times \left( \zeta + \eta \right) \left( -e^{-\lambda} / Z_{1,3,n}^1 \right) \left( -\zeta + \eta \right)^2 (x_1^2)^2. \tag{76}
\]

Using the given condition in this lemma and (53), we have the positive lower bound

\[
1 + 2 \beta \left( e^{-\lambda} / Z_{1,3,n}^1 \right) \left( \zeta + \eta \right) \left( -e^{-\lambda} / Z_{1,3,n}^1 \right) \left( -\zeta + \eta \right)^2 (x_1^2)^2 \geq 1 - 2 \beta \left( e^{-\lambda} U_{1,3,1}^{1,3,1} \right) (x_1^2)^2 > 0,
\tag{77}
\]

which gives that

\[
\left| \frac{\partial}{\partial \zeta} f (\zeta, \eta) \right| \leq \beta e^{-\lambda} C_{1,3,1}^{-1}.
\tag{78}
\]

Similarly,

\[
\left| \frac{\partial}{\partial \eta} f (\zeta, \eta) \right| \leq \beta e^{-\lambda} C_{1,3,1}^{-1}.
\tag{79}
\]

Substituting (73), (74), (78), and (79) into (70), we have

\[
\left| \Phi_{1,1} (X) - \Phi_{1,1} (Y) \right| = \left| \Phi_{1,2,1}^1 (X) - \Phi_{1,2,1}^1 (Y) \right|
= \left| \exp \left[ f (X_{10}, X_{12}) \right] - \exp \left[ f (Y_{10}, Y_{12}) \right] \right|
\leq C_{1,3,1} \beta e^{-\lambda} \left| X_{10} - Y_{10} \right| + \left| X_{12} - Y_{12} \right| \tag{80}
\]

Following the calculations for obtaining (80), we can obtain that for the 1\textsuperscript{st} perturbation

\[
\left| \Phi_{1,\theta} (X) - \Phi_{1,\theta} (Y) \right|
\leq C_{1,3,1} \beta e^{-\lambda} \left| X_{10} - Y_{10} \right| + \left| X_{12} - Y_{12} \right| \tag{81}
\]

where \( C_{1,3,1} \) is defined in Lemma 8.

Similarly we can have that for the 2\textsuperscript{nd} perturbation

\[
\left| \Phi_{1,\theta} (X) - \Phi_{1,\theta} (Y) \right|
\leq C_{2,3,1} \beta e^{-\lambda} \left| X_{4} - Y_{4} \right| + \left| X_{6} - Y_{6} \right| \tag{82}
\]

Similarly,\]

\[
\left| \Phi_{1,\theta} (X) - \Phi_{1,\theta} (Y) \right|
\leq C_{3,3,1} \beta e^{-\lambda} \left| X_{1} - Y_{1} \right| + \left| X_{3} - Y_{3} \right| \tag{83}
\]

Using Theorem 6 and Lemmas 7 and 8, we can obtain our main result.
Theorem 9. Assume that the experimental data $x_v^v$ ($v=1,2$) are contained in the codomain of $\phi$. Suppose that positive constants $\beta$ and $\lambda$ satisfy

$$1 - 2\beta \left( e^{-\beta U_{1,j,\lambda}^v} \right)^2 (x_v^v)^2 > 0$$

and

$$\sqrt{\beta} \max \left\{ C_{j,\beta,\lambda}^v | v = 1, 2, j = 2, 3 \right\} \beta e^{-\lambda} < 1,$$

for $v=1,2$, $j=2,3$, $U_{j,\beta,\lambda}^v$ defined in (53) and $C_{j,\beta,\lambda}^v$ defined in Lemma 8. Then sequence $X_1^{(n+1)} = X_1(X_1^{(n)})$ converges for any $X_1^{(0)} \in [0,1]^2$.

Proof. Let $D$ be the closed subset $[0,1]^2$ of $\mathbb{R}^2$ and

$$k = \sqrt{\beta} \max \left\{ C_{j,\beta,\lambda}^v | v = 1, 2, j = 2, 3 \right\} \beta e^{-\lambda}. \quad (86)$$

Then Lemmas 7 and 8 give that $\Phi_1 : D \rightarrow D$ and

$$\| \Phi_1(X) - \Phi_1(Y) \| \leq k \| X - Y \|. \quad (87)$$

Therefore we can apply Theorem 6 and so we obtain a unique fixed point $X_1^* \in D$ such that $\Phi_1(X_1^*) = X_1^*$, which implies the convergence of sequence $X_1^{(n+1)} = X_1(X_1^{(n)}) \ (n \geq 0)$ for any $X_1^{(0)} \in [0,1]^2$ with limit $X_1^*$. \hfill $\square$

Remark 10. We define

$$\rho^1 (w_{1,2} = -1), \rho^1 (w_{1,2} = 0), \rho^1 (w_{1,2} = 1),
$$

$$\rho^2 (w_{1,2} = -1), \rho^2 (w_{1,2} = 0), \rho^2 (w_{1,2} = 1),
$$

$$\rho^3 (w_{1,3} = -1), \rho^3 (w_{1,3} = 0), \rho^3 (w_{1,3} = 1),
$$

and

$$\left( \begin{array}{ccc}
\rho^1_{1,2,1,-1}, \rho^1_{1,2,0,0}, \rho^1_{1,2,1,1} \\
\rho^1_{1,3,-1}, \rho^1_{1,3,0,0}, \rho^1_{1,3,1,1} \\
\rho^2_{1,2,1,-1}, \rho^2_{1,2,0,0}, \rho^2_{1,2,1,1} \\
\rho^3_{1,3,-1}, \rho^3_{1,3,0,0}, \rho^3_{1,3,1,1}
\end{array} \right), \quad (88)$$

where the vector in the right side denotes limit $X_1^*$ in Theorem 9. Since $X_1^{(n)}$ in (57)–(59) converges, $\{P_{1,j,\alpha}^v(w)\}$ in (48) also converges to limit $e^{-\lambda \delta} P_{1,j,\alpha}^v$. Therefore, $\rho^1(w_{1,j})$ in (88) and $P^v(w_{1,j}) = e^{-\lambda \delta} P_{1,j,\alpha}^v$ satisfy the system of equations in (34) and (35), which give approximate marginal PMFs $P(w_{1,j}) = (1/Z_{1,j}) e^{-\lambda \delta} \prod_{i=1}^2 \rho^v_{1,i,w}$ in (14) as follows:

$$P(w_{1,j} = w) = \frac{1}{Z_{1,j}} e^{-\lambda \delta} \prod_{i=1}^2 \rho^v_{1,i,w}, \quad (89)$$

$$j = 2, 3, \quad w = -1, 0, 1.$$
Step 2 (determination of cost parameters). We find a pair of positive cost parameters \((\beta, \lambda)\) as in Figure 3(b) for each experimental data set, which satisfy the condition on \((\beta, \lambda)\) in Theorem 9.

Step 3 (generation of values of initial terms in sequences). We randomly generate initial values of 24 terms \(p_{i,j,0}(w)\) \((v = 1, 2, i = 1, 2, j \neq i, 1 \leq j \leq 3, w = -1,0,1)\) for each experimental data set, which satisfy the condition on initial values in Theorem 9 (see Figure 4).

Step 4 (convergence of the sequences). We obtain the convergence of all sequences \(p_{i,j,v}(w)\) \((v = 1, 2, i = 1, 2, j \neq i, 1 \leq j \leq 3, w = -1,0,1)\) for each experimental data set in Figure 3(a), \((\beta, \lambda)\) in Figure 3(b), and initial values in Figure 4, where every sequence converges within 14 iterations (see Figure 5). Due to the recursive relations in (47), (48), (49), (53), (54), (55), and (56), we can divide all sequences into four groups \(\{p_{1,2,v}(w), p_{2,1,v}(w)\}, \{p_{1,3,v}(w), p_{3,1,v}(w)\}, \{p_{2,3,v}(w), p_{3,2,v}(w)\}\) and show their convergences as in Figure 5.

Step 5 (approximation of marginal PMFs). Substituting the limits (Figure 6(c)) of the sequences for the 1st experimental data set (Figure 6(a)) and its corresponding cost parameters \((\beta, \lambda)\) (Figure 6(a)) into (89) and (93), we obtain the approximate marginal PMFs \(P(w_{1,2} = w), P(w_{1,3} = w), P(w_{2,1} = w)\), and \(P(w_{2,3} = w)\) as in Figure 6(d), where the maximum value of each PMF is marked as red.

Step 6 (construction of multiple networks and their mathematical formulas from the approximate marginal PMFs). We can use the maximum values of four link weights to make a network and its corresponding mathematical formulas (Figure 6(d), first grey box), which could be considered as the most possible network model of the artificial biological system. In addition, since value \(P(w_{2,1} = 0)\) is almost equal to value \(P(w_{2,1} = 1)\), we can also consider the network with \(w_{2,1} = 0\) and its mathematical formula as in the second grey box in Figure 6(d). We present two networks and their mathematical formulas in the two grey boxes to show that there could be multiple networks for explaining the artificial biological system by using the approach in [5]. Similarly, in cases where a given PKN has more nodes and discrete values of link weights, we could obtain multiple networks and their corresponding mathematical formulas, where probability \(P(w_{1,3} = w)\) is high.

5. Conclusions and Future Work

In this paper, using a simple network with artificial experimental data, we have presented a detailed explanation of the whole process for determining approximate marginal PMFs of link weights in a prior knowledge network based on BP.
Each set contains values of 12 initial terms \( \rho_{\alpha,j}(w) \) for each experimental dataset. (a) Each set contains values of 12 initial terms \( \rho_{1,j,w} \) (\( v = 1, 2, j = 2, 3, w = -1, 0, 1 \)). (b) Each set contains values of 12 initial terms \( \rho_{2,j,w} \) (\( v = 1, 2, j = 1, 3, w = -1, 0, 1 \)). Figure 5: Iteration number for the convergence. (a) The system of recursive relations for the convergences of the two types of these sequences converges within 10 iterations for tolerance \( 10^{-15} \). Symbols and data in Figures 5(b), 5(c), and 5(d) have similar meanings as in Figure 5(a).

Data Availability

The data used to support the findings of this study are available from the corresponding author upon request.

Conflicts of Interest

The authors declare that there are no conflicts of interest regarding the publication of this paper.
Figure 6: Approximate marginal PMFs. (a) The experimental data and cost parameters \((\beta, \lambda)\) are the 61\textsuperscript{th} data set and its corresponding values of \((\beta, \lambda)\) in Figures 3(a) and 3(b), respectively. (b) 12 initial values in the left and right denote the values in the 61\textsuperscript{th} initial data set in Figures 4(a) and 4(b), respectively. (c) 12 limit values in the top and bottom are obtained for Figures 6(a) and 6(b). (d) The approximate marginal PMFs are obtained from (89) and (93). The red numbers denote the maximum values of the marginal PMFs. The first network and its mathematical model in the right denote those in Figures 1(a) and 1(c) with link weights of maximum values. Replacing \(w_{12} = 1\) in the first network with \(w_{12} = 0\) gives the second network and its mathematical model.

Authors’ Contributions

All authors carried out the main results and completed the corresponding proof. All authors read and approved the final manuscript.

Acknowledgments

This work was supported by the 2018 Research Fund of University of Ul산.

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