## Supplement

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## Contents

1 Supplement A. Arcs update ..... 4
1.1 Arcs update version A ..... 4
1.2 Arcs update version B ..... 4
1.3 Arcs update version C ..... 5
1.4 Arcs update version D ..... 5
1.5 Arcs update: Version E ..... 6
1.6 Multiple arcs connected to the same node ..... 7
2 Supplement B: updating nodes given fixed arcs ..... 8
2.1 Node update version A ..... 8
2.2 Node update version B ..... 9
2.3 Node update version C ..... 9
3 Supplement C: enumeration examples ..... 10
3.1 Independent arcs ..... 10
3.2 Linear example ..... 10
3.3 X-shaped example ..... 11
3.3.1 Independent arcs ..... 12
3.3.2 Updating both nodes and arcs ..... 14
4 Supplement D. Graph support ..... 15
4.1 List of Nodes ..... 15
4.2 List of Arcs ..... 15
4.3 List of Arc Types ..... 15
4.4 List of Brain Tissue Names ..... 15

## List of Figures

1 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A, arc update from priors).
2 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A, arc update version C-detailed numerical information corresponding to this figure is shown in Tables 4 and 5).
3 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A , arc update version D (equivalent to version E with $\beta=1$ )—detailed numerical information corresponding to this figure is shown in Tables 6 and 7).
4 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B , arc update version B - detailed numerical information corresponding to this figure is shown in Tables 8 and 9).
5 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B, arc update version C-detailed numerical information corresponding to this figure is shown in Tables 10 and 11).
6 Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B , arc update version E (with $\beta=0.2$ )—detailed numerical information corresponding to this figure is shown in Tables 12 and 13).

## List of Tables

$$
\begin{array}{ll}
1 \text { Marginals for the nodes exact and estimated with the Gibbs sampler: node } \\
\text { update version A. (Gibbs sampler: 50, } 000 \text { different starting points, } 100 \text { iter- } \\
\text { ations for each chain.) . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . }
\end{array}
$$

2 Marginals for the nodes - comparison of the exact computation with the Gibbs sampler: node update version B.(Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.) ..... 13
3 Versions of update depicted in the supplement. ..... 14
4 Marginals for the nodes estimated with the Gibbs sampler: node update model A, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.) ..... 14
5 Marginals for the arcs estimated with the Gibbs sampler: node update model A, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.) ..... 15
6 Marginals for the nodes estimated with the Gibbs sampler: node update model A, arc update model D. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.) ..... 15
7 Marginals for the arcs estimated with the Gibbs sampler: node update model A, arc update model D. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.) ..... 16
8 Marginals for the nodes estimated with the Gibbs sampler: node update model B, arc update model B.(Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.) ..... 17
9 Marginals for the arcs estimated with the Gibbs sampler: node update model B, arc update model B. (Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.) ..... 18
10 Marginals for the nodes estimated with the Gibbs sampler: node update model B, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.) ..... 19
11 Marginals for the arcs estimated with the Gibbs sampler: node update modelB, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100iterations for each chain.)1912 Marginals for the nodes estimated with the Gibbs sampler: node update modelB , arc update model E (with $\beta=0.2$ ). (Gibbs sampler: 10,000 differentstarting points, 100 iterations for each chain.)19
13 Marginals for the arcs estimated with the Gibbs sampler: node update modelB , arc update model E (with $\beta=0.2$ ), (Gibbs sampler: 10,000 differentstarting points, 100 iterations for each chain.)20

## 1 Supplement A. Arcs update

In this section, we consider different schemes of conditional probabilities for updating one arc, $A_{i j}$ given its adjacent nodes, $V_{i}$ and $V_{j}$.

### 1.1 Arcs update version A

$$
\begin{align*}
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=1\right)= \begin{cases}1-P_{0}\left(A_{i j}=0\right) & \text { if } a=1, \\
0 & \text { if } a=-1, \\
P_{0}\left(A_{i j}=0\right) & \text { if } a=0 .\end{cases}  \tag{1}\\
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=0\right)= \begin{cases}0 & \text { if } a=1, \\
1-P_{0}\left(A_{i j}=0\right) & \text { if } a=-1, \\
P_{0}\left(A_{i j}=0\right) & \text { if } a=0 .\end{cases}  \tag{2}\\
& P\left(A_{i j}=a \mid V_{i}=0, V_{j}=*\right)= \begin{cases}P_{0}\left(A_{i j}=1\right) & \text { if } a=1, \\
P_{0}\left(A_{i j}=-1\right) & \text { if } a=-1, \\
P_{0}\left(A_{i j}=0\right) & \text { if } a=0 .\end{cases} \tag{3}
\end{align*}
$$

where $\operatorname{symbol} *$ is a "wildcard" character representing either 0 or 1 , and $P_{0}(\cdot)$ represents the prior value on a given event.

This setup corresponds to an assumption that the states of the arc-adjacent nodes are insufficient to increase the certainty that the nodes interact. We are not altering the probability that two nodes have connection no effect, but can informatively re-distribute probability between activation and inhibition.

An assumption common across versions is that if the upstream node, $V_{i}$ for an arc, $A_{i j}$, is inactive $\left(V_{i}=0\right)$, the value for $A_{i j}$ is sampled from the prior distribution for the arc, $P\left(A_{i j}=a_{i j}\right)$. An inactive upstream node $\left(V_{i}\right)$ does not provide evidence on its "effects" on the downstream node $\left(V_{j}\right)$.

### 1.2 Arcs update version B

$$
\begin{align*}
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=1\right)= \begin{cases}1 & \text { if } a=1, \\
0 & \text { if } a=-1, \\
0 & \text { if } a=0,\end{cases}  \tag{4}\\
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=0\right)= \begin{cases}0 & \text { if } a=1, \\
1 & \text { if } a=-1, \\
0 & \text { if } a=0 .\end{cases} \tag{5}
\end{align*}
$$

$$
P\left(A_{i j}=a \mid V_{i}=0, V_{j}=*\right)= \begin{cases}P_{0}\left(A_{i j}=1\right) & \text { if } a=1  \tag{6}\\ P_{0}\left(A_{i j}=-1\right) & \text { if } a=-1 \\ P_{0}\left(A_{i j}=0\right) & \text { if } a=0\end{cases}
$$

This version corresponds to an assumption that the states of the arc-adjacent nodes can be used as direct evidence in support of existence of an interaction.

### 1.3 Arcs update version C

Consider the conditional probability

$$
\begin{equation*}
P\left(A_{i j}=a_{i j} \mid V_{i}=v_{i}, V_{j}=v_{j}\right)=\frac{P\left(a_{i j}, v_{i}, v_{j}\right)}{P\left(v_{i}, v_{j}\right)} \propto p\left(a_{i j} \mid v_{i}\right) p\left(v_{j} \mid a_{i j}, v_{i}\right) \tag{7}
\end{equation*}
$$

To simplify the computation, we assume that $A_{i j}$ is independent from the incoming node $V_{i}$ and we only consider inputs from $V_{i}$ and $A_{i j}$ when deciding the output at $V_{j}$. Then, from (7), the conditional probabilities for updating $A_{i j}$ are derived as follows:

$$
\begin{gather*}
P\left(A_{i j}=a \mid V_{i}=1, V_{j}=1\right)= \begin{cases}\frac{P_{0}\left(A_{i j}=1\right)}{P_{0}\left(A_{i j}=1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=1\right)} & \text { if } a=1, \\
0 & \text { if } a=-1, \\
\frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=1\right)}{P_{0}\left(A_{i j}=1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=1\right)} & \text { if } a=0 .\end{cases}  \tag{8}\\
P\left(A_{i j}=a \mid V_{i}=1, V_{j}=0\right)= \begin{cases}0 & \text { if } a=1, \\
\frac{P_{0}\left(A_{i j}=-1\right)}{P_{0}\left(A_{i j}=-1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=-1, \\
\frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=0\right)}{P_{0}\left(A_{i j}=-1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=0 .\end{cases}  \tag{9}\\
P\left(A_{i j}=a \mid V_{i}=0, V_{j}=*\right)= \begin{cases}P_{0}\left(A_{i j}=1\right) & \text { if } a=1, \\
P_{0}\left(A_{i j}=-1\right) & \text { if } a=-1, \\
P_{0}\left(A_{i j}=0\right) & \text { if } a=0 .\end{cases} \tag{10}
\end{gather*}
$$

This update uses Bayesian probability calculation and allows for a more intuitive redistribution of probabilities. It can also be readily extended to more sophisticated updating schemes by incorporating more complex models for multiple inputs at $V_{j}$.

### 1.4 Arcs update version D

A slightly modified version.

$$
P\left(A_{i j}=a \mid V_{i}=1\right) \propto\left\{\begin{array}{cc}
P_{0}\left(A_{i j}=1\right) / P_{0}\left(V_{i}=1\right) & a=1 \\
P_{0}\left(A_{i j}=-1\right) / P_{0}\left(V_{i}=1\right) & a=-1 \\
P_{0}\left(A_{i j}=0\right) & a=0
\end{array}\right.
$$

$$
P\left(A_{i j}=a \mid V_{i}=1, V_{j}=1\right)= \begin{cases}\frac{P_{0}\left(A_{i j}=1\right)}{\frac{P_{0}\left(A_{i j}=1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)}{}} & \text { if } a=1  \tag{11}\\ 0 \quad & \text { if } a=-1 \\ \frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)}{P_{0}\left(A_{i j}=1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)} & \text { if } a=0\end{cases}
$$

$$
P\left(A_{i j}=a \mid V_{i}=1, V_{j}=0\right)= \begin{cases}0 & \text { if } a=1,  \tag{12}\\ \frac{P_{0}\left(A_{i j}=-1\right)}{P_{0}\left(A_{i j}=-1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=-1, \\ \frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)}{P_{0}\left(A_{i j}=-1\right)+P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=0\end{cases}
$$

$$
P\left(A_{i j}=a \mid V_{i}=0, V_{j}=*\right)= \begin{cases}P_{0}\left(A_{i j}=1\right) & \text { if } a=1  \tag{13}\\ P_{0}\left(A_{i j}=-1\right) & \text { if } a=-1 \\ P_{0}\left(A_{i j}=0\right) & \text { if } a=0\end{cases}
$$

This arc update version differs in that when there is an input from $V_{i}$ (i.e., $V_{i}=1$ ), the arc action likelihood is updated. The alteration maintains the ratio of the probabilities on activation and inhibition, and down-weights the probability on no effect by an artificial factor $P_{0}\left(V_{i}=1\right)$.

### 1.5 Arcs update: Version E

One last proposal; a more generalized of version D.

$$
\begin{align*}
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=1\right)= \begin{cases}\frac{P_{0}\left(A_{i j}=1\right)}{P_{0}\left(A_{i j}=1\right)+\beta P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)} & \text { if } a=1, \\
0 & \text { if } a=-1, \\
\frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)}{P_{0}\left(A_{i j}=1\right)+\beta P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=1\right)} & \text { if } a=0,\end{cases}  \tag{14}\\
& P\left(A_{i j}=a \mid V_{i}=1, V_{j}=0\right)= \begin{cases}0 & \text { if } a=1, \\
\frac{P_{0}\left(A_{i j}=-1\right)}{P_{0}\left(A_{i j}=-1\right)+\beta P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=-1, \\
\frac{P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)}{P_{0}\left(A_{i j}=-1\right)+\beta P_{0}\left(A_{i j}=0\right) P_{0}\left(V_{i}=1\right) P_{0}\left(V_{j}=0\right)} & \text { if } a=0 .\end{cases} \tag{15}
\end{align*}
$$

$$
P\left(A_{i j}=a \mid V_{i}=0, V_{j}=*\right)= \begin{cases}P_{0}\left(A_{i j}=1\right) & \text { if } a=1  \tag{16}\\ P_{0}\left(A_{i j}=-1\right) & \text { if } a=-1, \\ P_{0}\left(A_{i j}=0\right) & \text { if } a=0 .\end{cases}
$$

Here the probability on no effect is down-weighted by a factor $\beta P_{0}\left(V_{i}=1\right)$ where an arbitrary $(0 \leq \beta \leq 1)$ can be named. When $\beta=0)$ this equation is equivalent to version C . When $\beta=1$ ) this equation is equivalent to version D .

### 1.6 Multiple arcs connected to the same node

The scenario gets more complicated when you consider a situation where there are multiple adjacent arcs to one node. One possible way to proceed is to update each arc independently (as outlined above). We favor the simplicity of this solution. The one by one idea behind our "independent arc update" is in accordance with the way in which most interactions are discovered and published in the literature ${ }^{1}$. Since these interactions are independently derived, they can be tested for consistency with node-related data in the same fashion.

An alternate solution would be to update values for all the arcs adjacent to the same node simultaneously. We are not addressing this in detail in this article because of the difficulty of finding a set of assumptions that we can plausibly justify by biological data. Another complication of multiple-arc updating is the possibility of parental nodes to the given node which are not included in the data. Such a complicated update would not give good probability estimates while adding computational burden.

[^0]
## 2 Supplement B: updating nodes given fixed arcs

In this section we describe different options for updating a node given fixed incoming arcs. Our topology is a directed graph to incorporate causal knowledge ${ }^{2}$. This allows us to calculate the joint distribution for all nodes by decomposing the computation into the product of the conditional probabilities of nodes, their parental nodes and associated arcs.

We begin with the input nodes (nodes with no parent nodes) and follow the graph until all sink nodes (nodes with no offspring nodes) are updated.

$$
G \longrightarrow B \longrightarrow C \longrightarrow D
$$

Node $G$ is the only input node, and node $D$ is the only sink node. The joint distribution of node values given arcs for this simple graph has the following form:

$$
\begin{align*}
P\left(V_{G}\right. & \left.=v_{G}, V_{B}=v_{B}, V_{C}=v_{C}, V_{D}=v_{D} \mid A_{G B}=a_{G B}, A_{B C}=a_{B C}, A_{C D}=a_{C D}\right) \\
& =P\left(V_{G}=v_{G}\right) P\left(V_{B}=v_{B} \mid V_{G}=v_{G}, A_{G B}=a_{G B}\right) \\
& \times P\left(V_{C}=v_{C} \mid V_{B}=v_{B}, A_{B C}=a_{B C}\right) \\
& \times P\left(V_{D}=v_{D} \mid V_{C}=v_{C}, A_{C D}=a_{C D}\right) . \tag{17}
\end{align*}
$$

Therefore, to arrive at a sample from the full joint distribution of nodes given arcs, we first update the value for the input node $(G)$-sampling from its prior distribution. For all non-input nodes ( $B, C$, and $D$ ) the value is sampled from the appropriate conditional distributions (given the values of arcs and parental nodes for the node being updated). Each child node can be updated only after all parental nodes are updated.

A more general case of node updating concerns sampling of values for a node, $V$, that has multiple incoming edges with possibly conflicting arc values (both inhibition and activation values). We need to define the conditional probability that $V_{i}=v_{i}$ given the values of upstream nodes and arcs-assuming that node $V_{i}$ has $n$ parents, with values $\left\{V_{j}=v_{j}\right\}_{j=1, \ldots, n}$ at the parental nodes, and the values $\left\{A_{i j}=a_{i j}\right\}_{j=1, \ldots, n}$ at the direct incoming arcs of $V_{i}$. Different definitions of this conditional probability actually represent different assumptions and models at the local network of $V_{j}$.

### 2.1 Node update version A

$$
P\left(V_{i}=1 \left\lvert\, \begin{array}{ll}
\left\{V_{j}=v_{j}\right\}_{j=1, \ldots, n}  \tag{18}\\
\left\{A_{i j}=a_{i j}\right\}_{j=1, \ldots, n}
\end{array}\right., j \in \operatorname{par}\left(V_{i}\right)\right)= \begin{cases}\phi\left(I_{i}^{+} \triangleright I_{i}^{-}\right) & \text {if } I_{i}^{+}>0, \text { and } I_{i}^{-}>0 \\
1 & \text { if } I_{i}^{+}>0, \text { and } I_{i}^{-}=0 \\
0 & \text { if } I_{i}^{+}=0, \text { and } I_{i}^{-}>0 \\
P_{0}\left(V_{i}=1\right) & \text { if } I_{i}^{+}=0, \text { and } I_{i}^{-}=0\end{cases}
$$

[^1]where $I_{i}^{+}=\sum_{\left\{j: a_{i j}=1\right\}} v_{j}$ and $I_{V}^{-}=\sum_{\left\{j: a_{i j}=-1\right\}} v_{j}$ are combined incoming activating and inhibiting signals upon $V_{i}$. Here $\phi\left(I_{i}^{+} \triangleright I_{i}^{-}\right)$is a function that resolves a conflict between activating and inhibiting signals - in the simplest case it can be a single probability parameter common for the whole network. In a more sophisticated setup, the value of the parameter can vary for different nodes. In even more complex case, function $\phi\left(I_{i}^{+} \triangleright I_{i}^{-}\right)$can explicitly depend on values of $I_{i}^{+}$and $I_{i}^{-}$and have different a form for different nodes in the network.

### 2.2 Node update version B

$$
\begin{gather*}
P\left(V_{i}=1 \left\lvert\, \begin{array}{l}
\left\{V_{j}=v_{j}\right\}_{j=1, \ldots, n} \\
\left\{A_{i j}=a_{i j}\right\}_{j=1, \ldots, n}
\end{array}\right., j \in \operatorname{par}\left(V_{i}\right)\right)= \begin{cases}P_{0}\left(V_{i}=1\right) & \text { if } I_{i}^{+}>0, \text { and } I_{i}^{-}>0 \\
1 & \text { if } I_{i}^{+}>0, \text { and } I_{i}^{-}=0 \\
0 & \text { if } I_{i}^{+}=0, \text { and } I_{i}^{-}>0 \\
P_{0}\left(V_{i}=1\right) & \text { if } I_{i}^{+}=0, \text { and } I_{i}^{-}=0\end{cases}  \tag{20}\\
P\left(V_{i}=0 \left\lvert\, \begin{array}{l}
\left\{V_{j}=v_{j}\right\}_{j=1, \ldots, n} \\
\left\{A_{i j}=a_{i j}\right\}_{j=1, \ldots, n}
\end{array}\right., j \in \operatorname{par}\left(V_{i}\right)\right)=1-P\left(V_{i}=1 \left\lvert\, \begin{array}{l}
\left\{V_{j}=v_{j}\right\}_{j=1, \ldots, n} \\
\left\{A_{i j}=a_{i j}\right\}_{j=1, \ldots, n}
\end{array}\right., j \in \operatorname{par}\left(V_{i}\right)\right) \tag{21}
\end{gather*}
$$

This update version resolves conflicting inputs by an assumption that every conflict in input signals brings us back to the prior distribution for the node.

### 2.3 Node update version C

In this version, we assume that state of node $V_{i}$ is determined by a Boolean function over values of $V_{j} \in \operatorname{par}\left(V_{i}\right)$ and values of the corresponding edges, $A_{i j}$. For example, if vertex $V_{1}$ has just two parental nodes, $V_{2}$ and $V_{3}$, as shown below:

we can assume that node $V_{1}$ is active only if both parental nodes are active, $V_{2}=V_{3}=1$, and both arcs are in the state activation, $A_{21}=A_{31}=1$. Similarly, we can assume that to inhibit $V_{1}$ we need both incoming edges in the state inactivation, $A_{21}=A_{31}=-1$, and both parental nodes active. Then the probability to find node $V_{1}$ activated given states of parental nodes and arc is as follows.

$$
P\left(V_{1}=1 \left\lvert\, \begin{array}{cl}
V_{2}=v_{2}, V_{3}=v_{3}  \tag{23}\\
A_{21}=a_{21}, A_{31}=a_{31}
\end{array}\right.\right)= \begin{cases}1 & \text { if }\left(v_{2}, v_{3}, a_{21}, a_{31}\right)=(1,1,1,1) \\
0 & \text { if }\left(v_{2}, v_{3}, a_{21}, a_{31}\right)=(1,1,-1,-1) \\
P_{0}\left(V_{1}=1\right) & \text { otherwise }\end{cases}
$$

Given $n$ inputs to a node $V_{i}$, we can define the state of the node as a probabilistic function of all possible $2^{n} \times 3^{n}$ states of the parental nodes and arcs. Unfortunately, it would be very hard to find biological data to estimate parameters for such modeling at the present state of knowledge about biological circuitry.

## 3 Supplement C: enumeration examples

In this section, we give numerical illustrations of different arc update and node update versions. For comparison, in addition to the arc updates version considered in supplement A, we also include examples where the arcs are assumed to be independent with the nodes.

### 3.1 Independent arcs

Consider a special case when distribution values of the arcs are fixed despite the values of incoming nodes, that is,

$$
\begin{equation*}
P(\mathbf{A} \mid \mathbf{V})=P(\mathbf{A})=\prod_{j, i} P\left(A_{i j}=a_{i j}\right) \tag{24}
\end{equation*}
$$

Then, we can compute the marginal probabilities for nodes through direct summation over the appropriate joint distribution for nodes and arcs. Below we illustrate computation of the joint and marginal distributions for a simple linear network example.

### 3.2 Linear example

$$
\begin{equation*}
G \longrightarrow B \longrightarrow C \longrightarrow D \tag{25}
\end{equation*}
$$

Priors for the nodes:
$P_{0}\left(V_{G}=1\right)=0.9, P_{0}\left(V_{G}=0\right)=0.1$,
$P_{0}\left(V_{B}=1\right)=0.9, P_{0}\left(V_{B}=0\right)=0.1$,
$P_{0}\left(V_{C}=1\right)=0.9, P_{0}\left(V_{C}=0\right)=0.1$,
$P_{0}\left(V_{D}=1\right)=0.9, P_{0}\left(V_{C}=0\right)=0.1$.
Priors for arcs:
$P_{0}\left(A_{G B}=[1,-1,0]\right) \rightarrow(0.9,0.05,0.05)$
$P_{0}\left(A_{B C}=[1,-1,0]\right) \rightarrow(0.05,0.9,0.05)$
$P_{0}\left(A_{C D}=[1,-1,0]\right) \rightarrow(0.7,0.2,0.1)$.

For this simple example, the exact enumeration of the joint probabilities and marginal probabilities with independent arcs may be carried out as follows.

$$
\begin{gather*}
P\left(V_{G}=v_{G}, V_{B}=v_{B}, V_{C}=v_{C}, V_{D}=v_{D}, A_{G B}=a_{G B}, A_{B C}=a_{B C}, A_{C D}=a_{C D}\right) \\
=P_{0}\left(V_{G}=v_{G}\right) P_{0}\left(A_{G B}=a_{G B}\right) P\left(V_{B}=v_{B} \mid V_{G}=v_{G}, A_{G B}=a_{G B}\right) \\
\times P_{0}\left(A_{B C}=a_{B C}\right) P\left(V_{C}=v_{C} \mid V_{B}=v_{B}, A_{B C}=a_{B C}\right) \\
\times P_{0}\left(A_{C D}=a_{C D}\right) P\left(V_{D}=v_{D} \mid V_{C}=v_{C}, A_{C D}=a_{C D}\right) .  \tag{26}\\
P\left(V_{G}=v_{G}\right) \stackrel{\text { def }}{=} \sum_{v_{B}=0,1} \sum_{v_{C}=0,1} \sum_{v_{D}=0,1} \sum_{a_{G B}=-1,0,1} \sum_{a_{B C}=-1,0,1} \\
\sum_{a_{C D}=-1,0,1} P\left(V_{G}=v_{G}, V_{B}=v_{B},\right. \\
V_{C}=v_{c}, V_{D}=v_{D}, A_{G B}=a_{G B}, \\
\left.A_{B C}=a_{B C}, A_{C D}=a_{C D}\right) .  \tag{27}\\
P\left(A_{i j}=a_{i j}\right) \equiv P_{0}\left(A_{i j}=a_{i j}\right)(\text { prior values }), \forall i, j . \tag{28}
\end{gather*}
$$

Enumerated marginals for the nodes can be easily calculated:
$P\left(V_{G}=1\right)=0.9, P\left(V_{G}=0\right)=0.1$,
$P\left(V_{B}=1\right)=0.9405, P\left(V_{B}=0\right)=0.0595$,
$P\left(V_{C}=1\right)=0.1429, P\left(V_{C}=0\right)=0.8571$,
$P\left(V_{D}=1\right)=0.8843, P\left(V_{D}=0\right)=0.1157$.

### 3.3 X-shaped example

In this section, we use a more general example to illustrate different arc and node update versions, starting with simpler computations where independent arcs are assumed.


Priors for the nodes:
$P_{0}\left(V_{G}=1\right)=0.7, P_{0}\left(V_{G}=0\right)=0.3$,
$P_{0}\left(V_{B}=1\right)=0.8, P_{0}\left(V_{B}=0\right)=0.2$,
$P_{0}\left(V_{C}=1\right)=0.9, P_{0}\left(V_{C}=0\right)=0.1$,
$P_{0}\left(V_{D}=1\right)=0.35, P_{0}\left(V_{D}=0\right)=0.65$,
$P_{0}\left(V_{E}=1\right)=0.75, P_{0}\left(V_{E}=0\right)=0.25$.
Priors for arcs:
$P_{0}\left(A_{G C}=[1,-1,0]\right) \rightarrow(0.9,0.05,0.05)$,
$P_{0}\left(A_{B C}=[1,-1,0]\right) \rightarrow(0.05,0.85,0.1)$,
$P_{0}\left(A_{C D}=[1,-1,0]\right) \rightarrow(0.8,0.11,0.09)$,
$P_{0}\left(A_{C E}=[1,-1,0]\right) \rightarrow(0.1,0.1,0.8)$.

### 3.3.1 Independent arcs

Assuming independent arcs, the probabilities can be enumerated as follows.

$$
\begin{align*}
P\left(V_{G}\right. & =v_{G}, V_{B}=v_{B}, V_{C}=v_{C}, V_{D}=v_{D}, V_{E}=v_{E}, \\
A_{G C} & \left.=a_{G C}, A_{B C}=a_{B C}, A_{C D}=a_{C D}, A_{C E}=a_{C E}\right) \\
& =P_{0}\left(V_{G}=v_{G}\right) P_{0}\left(A_{G C}=a_{G C}\right) \\
& \times P_{0}\left(V_{B}=v_{B}\right) P_{0}\left(A_{B C}=a_{B C}\right) \\
& \times P\left(V_{C}=v_{C} \mid V_{G}=v_{G}, V_{B}=v_{B}, A_{G C}=a_{G C}, A_{B C}=a_{B C}\right) \\
& \times P_{0}\left(A_{C D}=a_{C D}\right) P\left(V_{D}=v_{D} \mid V_{C}=v_{C}, A_{C D}=a_{C D}\right) \\
& \times P_{0}\left(A_{C E}=a_{C E}\right) P\left(V_{E}=v_{E} \mid V_{C}=v_{C}, A_{C E}=a_{C E}\right) . \tag{30}
\end{align*}
$$

We implemented this example with two of our node update version. Table 1 shows the results for using node update version $\mathbf{A}$ with $\phi\left(I_{i}^{+} \triangleright I_{i}^{-}\right)=\frac{1}{2}$ (Equation (18)). The enumerated marginals (using exact probability computation) on the nodes are compared with both the priors and the caluculated values using the proposed numerical updates with Gibbs sampler. Since we are assuming independent arcs for this computation, the marginals on the arcs are unchanged from the prior values. The resulting network is compared with the network based on prior values in Figure 1.

| Node | Priors |  | Exact marginals |  | Gibbs sampler |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7 | 0.3 | 0.7 | 0.3 | 0.6996 | 0.30048 |
| $V_{B}$ | 0.8 | 0.2 | 0.8 | 0.2 | 0.7999 | 0.2001 |
| $V_{C}$ | 0.9 | 0.1 | 0.5143 | 0.4857 | 0.5140 | 0.4860 |
| $V_{D}$ | 0.35 | 0.65 | 0.5976 | 0.4024 | 0.5974 | 0.4026 |
| $V_{E}$ | 0.75 | 0.25 | 0.7243 | 0.2757 | 0.7243 | 0.2757 |

Table 1: Marginals for the nodes exact and estimated with the Gibbs sampler: node update version A. (Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.)


Figure 1: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A, arc update from priors).

For comparison, we also implemented the enumeration and numerical updating using the node update version B. The results are listed in Table 2.

| Node | Priors |  | Exact marginals |  | Gibbs sampler |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7 | 0.3 | 0.7 | 0.3 | 0.7002 | 0.2998 |
| $V_{B}$ | 0.8 | 0.2 | 0.8 | 0.2 | 0.8001 | 0.1999 |
| $V_{C}$ | 0.9 | 0.1 | 0.6862 | 0.3138 | 0.6864 | 0.3136 |
| $V_{D}$ | 0.35 | 0.65 | 0.6804 | 0.3196 | 0.6802 | 0.3198 |
| $V_{E}$ | 0.75 | 0.25 | 0.7157 | 0.2843 | 0.7156 | 0.2844 |

Table 2: Marginals for the nodes - comparison of the exact computation with the Gibbs sampler: node update version B.(Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.)

Apparently from Tables 1 and 2, the Gibbs sampler gives acceptable estimates of the marginals (precision can be improved by increasing the number of sampling iterations).

These examples also illustrate the difference between node update versions A and B. In the case of conflicting inputs for node $C$, version A assumes complete lack of information about the state of $V_{c}\left(\phi\left(I_{i}^{+} \triangleright I_{i}^{-}\right)=\frac{1}{2}\right.$, Equation 18), while version B samples values from the prior distribution for node $V_{C}$.

### 3.3.2 Updating both nodes and arcs

In this section, we apply different versions of arc and node update combinations to the same numerical x-shape example, with exactly the same prior values.

| Node update | Arc update | Numerical results | Graph |
| :--- | :--- | :--- | :--- |
| Version A | Version C | Tables 4 and 5 | Figure 2 |
| Version A | Version D | Tables 6 and 7 | Figure 3 |
| Version B | Version B | Tables 8 and 9 | Figure 4 |
| Version B | Version C | Tables 10 and 11 | Figure 5 |
| Version B | Version E (with $\beta=0.2)$ | Tables 12 and 13 | Figure 6 |

Table 3: Versions of update depicted in the supplement.

| Node | Priors |  | Marginals <br> (Gibbs sampler) |  |
| :---: | :--- | :--- | :--- | :--- |
|  | Gion |  |  |  |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7000 | 0.3000 | 0.7005 | 0.2995 |
| $V_{B}$ | 0.8000 | 0.2000 | 0.8000 | 0.2000 |
| $V_{C}$ | 0.9000 | 0.1000 | 0.5147 | 0.4853 |
| $V_{D}$ | 0.3500 | 0.6500 | 0.5979 | 0.4021 |
| $V_{E}$ | 0.7500 | 0.2500 | 0.7243 | 0.2757 |

Table 4: Marginals for the nodes estimated with the Gibbs sampler: node update model A, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)

| Arc | Priors |  |  | Marginals (Gibbs sampler) |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $a=1$ | $a=-1$ | $a=0$ | $a=1$ | $a=-1$ | $a=0$ |
| $A_{G C}$ | 0.9000 | 0.0500 | 0.0500 | 0.8998 | 0.0500 | 0.0501 |
| $A_{B C}$ | 0.0500 | 0.8500 | 0.1000 | 0.0499 | 0.8499 | 0.1002 |
| $A_{C D}$ | 0.8000 | 0.1100 | 0.0900 | 0.8007 | 0.1096 | 0.0897 |
| $A_{C E}$ | 0.1000 | 0.1000 | 0.8000 | 0.1002 | 0.0999 | 0.7999 |

Table 5: Marginals for the arcs estimated with the Gibbs sampler: node update model A, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)

| Node | Priors |  | Marginals (Gibbs sampler) |  |
| :---: | :---: | :---: | :---: | :---: |
| $V_{G}$ | 0.7000 | 0.3000 | 0.6994 | 0.3006 |
| $V_{B}$ | 0.8000 | 0.2000 | 0.8004 | 0.1996 |
| $V_{C}$ | 0.9000 | 0.1000 | 0.5762 | 0.4238 |
| $V_{D}$ | 0.3500 | 0.6500 | 0.6215 | 0.3785 |
| $V_{E}$ | 0.7500 | 0.2500 | 0.7199 | 0.2801 |

Table 6: Marginals for the nodes estimated with the Gibbs sampler: node update model A, arc update model D. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)

## 4 Supplement D. Graph support

In this section, we included the data supporting the graph depicting the genes assoiciated with the four neurological diseases.

### 4.1 List of Nodes

### 4.2 List of Arcs

### 4.3 List of Arc Types

### 4.4 List of Brain Tissue Names



Figure 2: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A, arc update version C-detailed numerical information corresponding to this figure is shown in Tables 4 and 5).

| Arc | Priors |  |  | Marginals (Gibbs sampler) |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $a=1$ | $a=-1$ | $a=0$ | $a=1$ | $a=-1$ | $a=0$ |
| $A_{G C}$ | 0.9000 | 0.0500 | 0.0500 | 0.6766 | 0.2757 | 0.0477 |
| $A_{B C}$ | 0.0500 | 0.8500 | 0.1000 | 0.1833 | 0.5438 | 0.2729 |
| $A_{C D}$ | 0.8000 | 0.1100 | 0.0900 | 0.7941 | 0.1179 | 0.0880 |
| $A_{C E}$ | 0.1000 | 0.1000 | 0.8000 | 0.1056 | 0.1046 | 0.7898 |

Table 7: Marginals for the arcs estimated with the Gibbs sampler: node update model A, arc update model D. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)


Figure 3: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version A , arc update version D (equivalent to version E with $\beta=1$ )—detailed numerical information corresponding to this figure is shown in Tables 6 and 7).

| Node | Priors |  | Marginals <br> (Gibbs sampler) |  |
| :--- | :--- | :--- | :--- | :--- |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7 | 0.3 | 0.6999 | 0.3001 |
| $V_{B}$ | 0.8 | 0.2 | 0.8000 | 0.2000 |
| $V_{C}$ | 0.9 | 0.1 | 0.8455 | 0.1545 |
| $V_{D}$ | 0.35 | 0.65 | 0.7241 | 0.3199 |
| $V_{E}$ | 0.75 | 0.25 | 0.7105 | 0.2895 |

Table 8: Marginals for the nodes estimated with the Gibbs sampler: node update model B, arc update model B.(Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.)

| Arc | Priors |  |  | Marginals (Gibbs sampler) |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $a=1$ | $a=-1$ | $a=0$ | $a=1$ | $a=-1$ | $a=0$ |
| $A_{G C}$ | 0.9 | 0.05 | 0.05 | 0.8962 | 0.0887 | 0.0151 |
| $A_{B C}$ | 0.05 | 0.85 | 0.1 | 0.6745 | 0.3055 | 0.02 |
| $A_{C D}$ | 0.8 | 0.11 | 0.09 | 0.7913 | 0.1948 | 0.0139 |
| $A_{C E}$ | 0.1 | 0.1 | 0.8 | 0.61052 | 0.2662 | 0.1233 |

Table 9: Marginals for the arcs estimated with the Gibbs sampler: node update model B, arc update model B. (Gibbs sampler: 50, 000 different starting points, 100 iterations for each chain.)


Figure 4: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B, arc update version B-detailed numerical information corresponding to this figure is shown in Tables 8 and 9).

| Node | Priors |  | Marginals <br> (Gibbs sampler) |  |
| :---: | :---: | :---: | :---: | :--- |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7000 | 0.3000 | 0.7001 | 0.2999 |
| $V_{B}$ | 0.8000 | 0.2000 | 0.7998 | 0.2002 |
| $V_{C}$ | 0.9000 | 0.1000 | 0.8057 | 0.1943 |
| $V_{D}$ | 0.3500 | 0.6500 | 0.7306 | 0.2694 |
| $V_{E}$ | 0.7500 | 0.2500 | 0.7102 | 0.2898 |

Table 10: Marginals for the nodes estimated with the Gibbs sampler: node update model B, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)

| Arc | Priors |  |  | Marginals (Gibbs sampler) |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $a=1$ | $a=-1$ | $a=0$ | $a=1$ | $a=-1$ | $a=0$ |
| $A_{G C}$ | 0.9000 | 0.0500 | 0.0500 | 0.8455 | 0.1023 | 0.0522 |
| $A_{B C}$ | 0.0500 | 0.8500 | 0.1000 | 0.2339 | 0.3411 | 0.4249 |
| $A_{C D}$ | 0.8000 | 0.1100 | 0.0900 | 0.7911 | 0.1161 | 0.0928 |
| $A_{C E}$ | 0.1000 | 0.1000 | 0.8000 | 0.1001 | 0.0999 | 0.7999 |

Table 11: Marginals for the arcs estimated with the Gibbs sampler: node update model B, arc update model C. (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)

| Node | Priors |  | Marginals <br> (Gibbs sampler) |  |
| :---: | :--- | :--- | :--- | :--- |
|  | $v=1$ | $v=0$ | $v=1$ | $v=0$ |
| $V_{G}$ | 0.7000 | 0.3000 | 0.6999 | 0.3001 |
| $V_{B}$ | 0.8000 | 0.2000 | 0.8001 | 0.1999 |
| $V_{C}$ | 0.9000 | 0.1000 | 0.8331 | 0.1669 |
| $V_{D}$ | 0.3500 | 0.6500 | 0.7441 | 0.2559 |
| $V_{E}$ | 0.7500 | 0.2500 | 0.6454 | 0.3546 |

Table 12: Marginals for the nodes estimated with the Gibbs sampler: node update model B, arc update model E (with $\beta=0.2$ ). (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)


Figure 5: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B, arc update version C-detailed numerical information corresponding to this figure is shown in Tables 10 and 11).

| Arc | Priors |  |  | Marginals (Gibbs sampler) |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | $a=1$ | $a=-1$ | $a=0$ | $a=1$ | $a=-1$ | $a=0$ |
| $A_{G C}$ | 0.9000 | 0.0500 | 0.0500 | 0.8854 | 0.0941 | 0.0204 |
| $A_{B C}$ | 0.0500 | 0.8500 | 0.1000 | 0.5173 | 0.3168 | 0.1659 |
| $A_{C D}$ | 0.8000 | 0.1100 | 0.0900 | 0.8123 | 0.1549 | 0.0329 |
| $A_{C E}$ | 0.1000 | 0.1000 | 0.8000 | 0.2672 | 0.2462 | 0.4867 |

Table 13: Marginals for the arcs estimated with the Gibbs sampler: node update model B, arc update model E (with $\beta=0.2$ ), (Gibbs sampler: 10, 000 different starting points, 100 iterations for each chain.)


Figure 6: Update of both arcs and nodes of a hypothetical five-node and four-arc network (node update version B , arc update version E (with $\beta=0.2$ ) -detailed numerical information corresponding to this figure is shown in Tables 12 and 13).


[^0]:    ${ }^{1}$ A notable exception is associated with so-called high-throughput experiments, such as yeast two-hybrid experiments aiming at discovering protein-protein interactions.

[^1]:    ${ }^{2}$ This first version is a directed acyclic graph - we hope to handle cycles in the next version.

