Jointly Learning Network Connections and Link Weights in Spiking Neural Networks

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Abstract
Spiking neural networks (SNNs) are considered to be biologically plausible and power-efficient on neuromorphic hardware. However, unlike the brain mechanisms, most existing SNN algorithms have fixed network topologies and connection relationships. This paper proposes a method to jointly learn network connections and link weights simultaneously. The connection structures are optimized by the spike-timing-dependent plasticity (STDP) rule with timing information, and the link weights are optimized by a supervised algorithm. The connection structures and the weights are learned alternately until a termination condition is satisfied. Experiments are carried out using four benchmark datasets. Our approach outperforms classical learning methods such as STDP, Tempotron, SpikeProp, and a state-of-the-art supervised algorithm. In addition, the learned structures effectively reduce the number of connections by about 24%, thus facilitate the computational efficiency of the network.

1 Introduction
Artificial neural networks (ANNs) have obtained great success in the computer vision and pattern recognition fields in recent years. Although ANNs are inspired by brain, most studies only imitate layer-like brain structures roughly [Hinton and Salakhutdinov, 2006]. Spiking neural networks (SNNs), known as the third generation of artificial neural network, work in the manner of spiking neurons in brain, thus are more biologically plausible and have potential to approach the intelligence of brain [Tsukada and Pan, 2005] [Ghoshr-Dastidar and Adeli, 2009] [Maass, 1997]. However, existing SNN algorithms usually have fixed network topologies and connection relationships. According to the Hebbian theory [Hebb, 1949], only a few cells would strongly connect together when brain transfers information, e.g. the persistently co-active cells. Therefore, how to build a spiking neural network with both optimal connection and link weights is an important problem.

Much effort has been made on structure design and learning of SNN. Various spiking neuron models, such as the leaky integrate-and-fire (LIF) model [Gerstner and Kistler, 2002] and the Hodgkin-Huxley type model [Hodgkin and Huxley, 1952], have been proven to be capable of encoding temporal information in synapses between neurons. To improve the learning capacity in SNN, different approaches including unsupervised rules and supervised algorithms were proposed.

- **Unsupervised learning methods.** The unsupervised approaches are mostly biomimetic methods inspired by the mechanism in biological neurons. The Hebbian theory, which is often summarized as ‘Cells that fire together wire together’ [Shatz, 1992], is considered to be the foundation of connection learning in biological neurons. According to the Hebbian theory, the spike-timing-dependent plasticity (STDP) rule has been proposed which exploits the timing and the emitting order of the spikes [Bi and Poo, 2001]. The STDP algorithm strengthens a weight when the presynaptic neuron emits spikes preceding the postsynaptic one, and weakens it when the order is reversed. Recently, lots of STDP-based approaches are proposed with promising results [Diehl and Cook, 2015] [Masquelier et al., 2009]. Despite the effectiveness of the unsupervised methods, the convergence of these approaches can not be guaranteed due to the difficulty in setting appropriate parameters.

- **Supervised learning methods.** Some supervised rules were proposed in order to accelerate the learning progress. With the help of instructor signals, supervised algorithms can potentially improve the learning speed. Classical supervised learning algorithms are mostly based on gradient descent methods. SpikeProp [Bohte et al., 2002] is one of the representative learning algorithms to process spatiotemporal information in SNN. The Tempotron rule [Güttig and Sompolinsky, 2006] then extends the ability of SNN to binary classification tasks. Another type of supervised training approach uses learning windows. Unlike the gradient descent methods that minimize the cost functions, these methods such as the remote supervised learning method (ReSûMe) [Filip, 2005] and the precise-spike-driven (PSD) [Yu et al., 2013a] are motivated by the Widrow-Hoff rule, and use the Hebbian and anti-
Hebbian learning windows to drive training. The training window-based algorithms are commonly more efficient than methods with gradient descent rules.

Although various learning methods have been proposed, joint optimization of SNN connections and link weights has not been well addressed. In this study, we propose a method to jointly learn network connections and link weights simultaneously in SNNs. To enable structure learning, we propose a spiking neuron with connection gate (SNN-CG) to control whether a connection should be established. A connection weight is effective only if the gate is open. In SNN-CG, the connection gates are learned based on the Hebbian theory and the STDP rule. The link weights are updated using a supervised algorithm. The structure learning and weight updating work alternately until a termination condition is satisfied. Experiments are conducted using four benchmark datasets. Our approach outperforms classical learning methods such as STDP, Tempotron, SpikeProp, and a state-of-the-art work.

In order to enable structural learning of connections between neurons, we assemble connection gates to the SNN model. Then we elaborate the STDP rule-based learning and supervised link weight updating algorithms respectively. Finally, we present the iterative training scheme to jointly learn both connection structures and the corresponding weights.

2 Method

In this section, we present the SNN-CG algorithm in details. We first introduce the formulations of the spiking neural networks with connection gates. Then we elaborate the STDP rule-based learning and supervised link weight updating algorithms respectively. Finally, we present the iterative training scheme to jointly learn both connection structures and the corresponding weights.

2.1 SNN with Connection Gates

In order to enable structural learning of connections between neurons, we assemble connection gates to the SNN model. We start from the classical leaky integrate-and-fire (LIF) neuron model and then describe the model with connection gates.

**LIF neuron model.** Given an LIF neuron $i$, suppose there are $N$ presynaptic afferents contributing to it. Neuron $i$ is driven by exponential decaying synaptic currents generated by its $N$ presynaptic neurons. Then the subthreshold membrane voltage of neuron $i$ is a weighted sum of postsynaptic potentials (PSPs) contributed by all incoming spikes:

$$ V_j(t) = \sum_{i=1}^{N} W_{ij} \sum_{t_i < t} K(t-t_i) + V_{rest}, \quad (1) $$

where $W_{ij}$ is the synaptic efficacy between postsynaptic neuron $j$ and presynaptic afferent $i$, $t_i$ and $V_{rest}$ denote the firing time of presynaptic afferent $i$, and the rest potential of postsynaptic neuron $j$, respectively. A normalized PSP kernel $K$ vanishing for $t > t_i$ is as follows:

$$ K(t-t_i) = V_0 (\exp(-\frac{t-t_i}{\tau_m}) - \exp(-\frac{t-t_i}{\tau_s})), \quad (2) $$

where $V_0$ is used to normalize the maximum of kernel to be 1.0, which ensures unitary PSP amplitudes to be given by synaptic efficacies $W_{ij}$. The parameters $\tau_m$ and $\tau_s$ denote the decay time constants of membrane integration and synaptic currents, respectively. The postsynaptic neuron $j$ fires a spike once its voltage $V_j$ crosses the firing threshold $V_{thr}$. That is, neuron $j$ generates an output spike at that time. Since we only consider the situation that postsynaptic neuron is fired only once in this paper, the voltages of that fired neuron smoothly decline to $V_{rest}$ by shutting down all the following incoming spikes.

**Connection gates.** Based on LIF, we enable structural learning by using connection gates in the links between neurons, called spiking neural network with connection gates (SNN-CG). The connection gates control whether two neurons are linked with each other. With the connection gate $G_{ij}$ between neurons $i$ and $j$, (1) can be rewritten as follows:

$$ V_j(t) = \sum_{i=1}^{N} W_{ij} * G_{ij} \sum_{t_i < t} K(t-t_i) + V_{rest}. \quad (3) $$

With a proper connection adjusting rule, the SNN-CG model establishes connections between neurons only if they are related. Therefore, the neurons become more sensitive to related neurons, and thus are able to transmit information more effectively and robustly than fully connected networks.

**Gate functions.** The connection gates in SNN-CG model can be implemented using different functions. We apply the sigmoid function as the gate function in this work:

$$ G_{ij} = \frac{1}{1 + \exp(-b_{ij})}. \quad (4) $$

where $b_{ij}$ is the connecting coefficient between neuron $i$ and neuron $j$. A higher $b_{ij}$ indicates stronger connections.

2.2 Connection Learning

In biological systems, the links between neurons are adjusted using the timing information in spikes. According to the Hebbian theory, repeatedly and persistently co-active cells would increase connective synaptic efficacy among populations of interconnected neurons [Hebb, 1949].

Motivated by the biological mechanism, we adopt the STDP rule to learn the connections in SNN neurons. The STDP rule provides powerful guidance for establishing or eliminating the connections between neurons by considering both the timing and the order of the spikes. The plasticity in STDP depends on the intervals between presynaptic and postsynaptic spikes. As illustrated in Fig. 1 (a), a connection is strengthened (long-term potentiation, LTP) when the presynaptic neuron emits spikes preceding the postsynaptic one and weakened (long-term depression, LTD) when the order is reversed. The STDP rule used is as follows:

$$ \Delta W_{ij}^{stdp} = \begin{cases} A_+ \exp(\frac{t_j-t_i}{\tau^+}), & \text{if } t_j < t_i (\text{LTP}), \\ A_- \exp(\frac{t_j-t_i}{\tau^-}), & \text{if } t_j < t_i (\text{LTD}), \end{cases} \quad (5) $$

where the parameters $\tau^+$ and $\tau^-$ determine the ranges of pre-to-postsynaptic interspike intervals on which synaptic strengthening and weakening occur. $A_+$ and $A_-$ refer to the maximum amounts of synaptic modification for LTP and LTD, respectively. The ratio between the $A_+$ and $A_-$ balances the magnitudes of strengthening and weakening of the connections in STDP.
According to the STDP rule, the connection gate can be updated based on $W_{stdp}$:

$$b_{ij} = b_{ij} + \lambda_c \cdot W_{stdp}$$  \hspace{1cm} (6)

where $\lambda_c$ is the learning rate for connections, and $b_{ij}$ is the connecting coefficient between neuron $i$ and neuron $j$. With the STDP-based connection learning, the connections between related neurons are strengthened toward one and the connections between unrelated neurons decay to zero.

### 2.3 Weight Learning

The link weights in SNN can be learned with general supervised learning methods suitable for temporal coding. In this study, we employ Tempotron rule due to its good ability for classification.

In binary classification, the input patterns to the neurons belong to one of two types of $\oplus$ and $\ominus$. When a $\oplus$ is presented to the neuron, it fires a spike, and when a $\ominus$ appears, the neuron does not fire. Tempotron rule learns the synaptic weights of $W_{ij}$ with gradient descent to minimize the error signals:

$$E_j = \begin{cases} V_{thr} - V_j(t_{max}) & \text{if } \oplus \text{ error}, \\ V_j(t_{max}) - V_{thr} & \text{if } \ominus \text{ error}, \end{cases}$$  \hspace{1cm} (7)

where $t_{max}$ is the time point that the neuron reaches its maximum voltage, and $V_{thr}$ is the threshold for neurons to fire a spike. The gradients of parameter $W_{ij}$ are computed follows:

$$\begin{align*}
\Delta W_{ij}^+ &= \lambda_w \sum_{t_i < t_{max}} G_{ij} K(t_{max} - t_i) & \text{if } \oplus \text{ error}, \\
\Delta W_{ij}^- &= -\lambda_w \sum_{t_i < t_{max}} G_{ij} K(t_{max} - t_i) & \text{if } \ominus \text{ error},
\end{align*}$$

8

where $\lambda_w$ is the weight learning rate. $\oplus$ error means the error that the neuron should emit but it does not, and $\ominus$ error is the error that the neuron should not emit but it does. The learning window of the Tempotron rule is illustrated in Fig 1 (b). According to the kernel shape, the spikes near $t_{max}$ change more than those far away from it.

### 2.4 Iterative Training

To jointly learn the connections and the link weights, we propose an iterative learning scheme. In the connection learning stage, the weights of the links are fixed and the connection strengths are tuned by the STDP rule. In the weight learning stage, the connections are fixed and the weights are updated. The connection learning and the weight learning stages are conducted alternately until a termination condition is satisfied. The details of the iterative training process are presented in Algorithm 1. The key parameters in the iterative training algorithm are $M$ and $N$, which indicate how many epochs to train in the weight learning stage and connection learning stage respectively.

### 2.5 Network Architectures

The network includes an encoding layer and a decoding layer. The encoding layer converts the real-valued features into spike trains. The decoding layer determines the classification results using the population decoding method [Yu et al., 2013b].

**Encoding layer.** The continuous feature values are encoded into spike trains using the Gaussian receptive field method [Bohle et al., 2002] [Eurich and Wilke, 2000]. Suppose a feature value $x$ is in range $[x_{min}, x_{max}]$, and it is encoded by different $N_{encode}$ Gaussian receptive fields. The center and width of the $k_{th}$ Gaussian kernel are defined as $\mu_k = x_{min} + (2k - 3)/2(x_{max} - x_{min})/(N_{encode} - 2)$, and $\sigma_k = 1/1.5(x_{max} - x_{min})/(N_{encode} - 2)$, respectively. With the Gaussian kernels, a feature value $x$ is encoded to spike arrays of $N_{encode}$ dimensions. Because the Tempotron model only permits postsynaptic neuron to fire only once,
there are $N_{\text{encode}}$ neurons for each attribute. With the encoding method, the higher activated neurons fire earlier, and vice versa. Suppose there are a total of $N_{\text{feature}}$ features. Then there are $N_{\text{encode}} \times N_{\text{feature}}$ neurons in the input layer of the network.

**Decoding layer.** The binary outputs of networks predict the class labels by population decoding [Yu et al., 2013b], according to the biological mechanisms in neurons [Ranhe, 2012]. For classification, each class is represented by a group of $N_{\text{decode}}$ neurons. The decisions are made by majority voting, and the incoming samples are classified into the class with maximum numbers of firing neurons. Suppose there are $N_{\text{class}}$ classes. The total number of output neurons is $N_{\text{decode}} \times N_{\text{class}}$. The architecture of the SNN-CG method is illustrated in Fig. 2.

**Network.** We use a single-layer SNN model. The neurons of the encoding layer are fully connected with neurons in the decoding layer by gated links. The connections and the link weights can be jointly adjusted in training. Fig. 2 shows a simple example of the network with three features and two classes.

![The spiking neural network with connection gates.](image)

**3 Experiments**

In this section, experiments are carried out to evaluate the performance of our method. Firstly, we give the detail of the evaluation datasets and settings. Then we assess the influence of the model settings in SNN-CG, including parameters in connection learning and iterative training. Finally, we compare our approach with existing methods to demonstrate the strength of the SNN-CG model.

**3.1 Datasets**

Four datasets are used in our experiments. The datasets are selected from the OpenML and the UCI repositories [Dheeru and Karra Taniskidou, 2017]. The detail of the datasets is as follows.

- **Iris.** A benchmark dataset for plant classification. It contains three types of iris plants with four-dimensional features. There are a total of 150 instances in the dataset.
- **Breast Cancer.** A binary classification dataset for breast cancer recognition. It contains 295 samples, and each sample has nine features.

- **Liver Disorder.** A binary classification dataset for liver disease detection. There are a total of 345 samples in the dataset. For each sample, the dimension of feature is six.
- **Pima Indians Diabetes.** The dataset includes 500 instances of two classes. The instances are described by eight-dimensional features.

**3.2 Settings**

Here we give the detail of the configurations of our model, including the model settings and the parameters in training.

For the network architecture, a one-layer SNN-CG model is adopted. For the LIF neuron model, the parameters are set as follows: $V_{\text{th}} = 1.0mV$, $V_{\text{rest}} = 0mV$, $V_0 = 0.4725mV$, $\tau_m = 4\tau_s = 15ms$. For each feature, the encoding neuron number $N_{\text{encode}}$ is set to 12, and the number of decoding neurons $N_{\text{decode}}$ for each class is set to 8. In the initialization, the connection coefficients $b$ are set to $0.5 \times I$ for the connections gates. The link weights in the network are randomly initialized in the range of [-0.01, 0.01].

For connection learning, the time constants are set to $\tau^+ = 16.8ms$ and $\tau^- = 33.7ms$ in accordance to the observed measurements experimentally [Bi and Poo, 2001]. The learning rate of connection gates is $\lambda_c = 1.0$.

The performance is evaluated by the classification accuracy. For each dataset, we split the data to ten parts randomly. The experimental result is the average accuracy of the ten parts with cross-validation.

**3.3 Influence of the Parameters**

In this section, we assess the influence of the parameters in the SNN-CG model. Firstly, we compare the models with different parameters in connection learning. Then we evaluate the parameters in iterative training of connection learning and weight updating.

**Connection learning.** Here we test the influence of the learning parameters in STDP-based connection training. We tune the synaptic modification magnitude of $A_+$ and $A_-$ in (5) to change the balance between strengthening and weakening of the connections. In the STDP rule, the ratio between the two parameters is considered to be important. In the experiment, we set the $A_+ = 0.0156$ and $A_- = 0.02656$. Then we tune the ratio between them by multiplying the value of $A_-$ with a coefficient $C$ in the range of 0.001 to 0.5. The criteria of performance are the classification accuracy and the connection sparsity. The connection sparsity is computed as the proportion of disconnected links. The links with connection weights lower than 1e-8 are considered to be disconnected. In training, for each iteration, we use $M = 100$ epochs for weight learning and $N = 10$ epochs for connection learning.

Fig. 3 illustrates the connection sparsity under different settings. As shown in Fig. 3 (a), the heights of the bars indicate the proportions of disconnected links. With a larger parameter $C$, the connections in the networks are more sparse. For the Iris dataset, when $C = 0.001$, the network is almost fully connected. When we tune $C$ to 0.5, 97% of the connections
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Table 1: Comparison of parameter $C$ in connection learning.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>$C=0.001$</th>
<th>$C=0.01$</th>
<th>$C=0.1$</th>
<th>$C=0.5$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Iris</td>
<td>0.9467</td>
<td>0.9667</td>
<td>0.9133</td>
<td>0.7600</td>
</tr>
<tr>
<td>Breast Cancer</td>
<td>0.9542</td>
<td>0.9571</td>
<td>0.9270</td>
<td>0.8470</td>
</tr>
<tr>
<td>Liver Disorder</td>
<td>0.6162</td>
<td>0.6214</td>
<td>0.6210</td>
<td>0.5429</td>
</tr>
<tr>
<td>Pima Indians Diabetes</td>
<td>0.7238</td>
<td>0.7304</td>
<td>0.6966</td>
<td>0.6353</td>
</tr>
</tbody>
</table>

Table 2: Comparison of parameter $R$ in iterative training.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>$R=0.05$</th>
<th>$R=0.1$</th>
<th>$R=0.2$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Iris</td>
<td>0.9667</td>
<td>0.9533</td>
<td>0.9533</td>
</tr>
<tr>
<td>Breast Cancer</td>
<td>0.9571</td>
<td>0.9600</td>
<td></td>
</tr>
<tr>
<td>Liver Disorder</td>
<td>0.6214</td>
<td>0.6529</td>
<td>0.6024</td>
</tr>
<tr>
<td>Pima Indians Diabetes</td>
<td>0.7304</td>
<td>0.7342</td>
<td>0.7368</td>
</tr>
</tbody>
</table>

Figure 3: The connection sparsity of SNN-CG. The connection sparsity is the percentage of unconnected links. (a) is the connection sparsity with different $C$ on the datasets. With a bigger $C$, the links are prone to be weakened in learning, thus higher sparsity is shown. (b) is the sparsity of the decoding neurons with $C=0.01$.

3.4 Comparison with Other Methods

In this section, experiments are conducted to compare our approach with existing methods. Firstly, we compare our model with STDP and Tempotron separately, to evaluate the effectiveness of the joint learning scheme. Then we assess the structure learning ability of our method in comparison with Tempotron with L1-norm regularization. Finally, the SNN-CG model is compared with other supervised learning approaches in SNN and the SVM classifier.

The methods to be compared in this experiment are configured as follows:

- **Tempotron.** The classical Tempotron method [Gütig and Sompolinsky, 2006]. The LIF neurons are applied in this model, and the network structure is the same as that in the SNN-CG model.
- **STDP.** The classical STDP method [Bi and Poo, 2001]. The LIF neurons are applied in this model, and the network structure is the same as the SNN-CG model.
- **Tempotron-L1.** The settings of this method are the same as that of the Tempotron. In order to constrain the sparsity of the weights, we add the L1 norm of the weights in the cost function with a coefficient of $\lambda$. The parameter $\lambda$ is selected from $\{0.1, 0.01, 0.001\}$ by cross-validation. In Tempotron-L1, we remove the connections with top $K\%$ smallest weights, and the $K\%$ equals to the proportion of links cut in the SNN-CG model.
- **SpikeProp.** A multiple-layer SNN based on the gradient descent rule [Bohte et al., 2002]. We employ a three-layer network containing $N_{\text{feature}} \times N_{\text{encode}}$ input neurons, 20 hidden neurons, and $N_{\text{class}}$ output neurons. Each neuron is modeled as Spike Response Model (SRM). The time constant is set to 7ms. The firing threshold is set to 1.0 which is the same as that in the above LIF neuron model. The networks are trained for 100 iterations and the learning rate is set to 0.01.

of the parameters $M$ and $N$ in the iterative training. $M$ and $N$ denote how many epochs are applied in the weight learning and connection learning stages, respectively. Here we tune the ratio $R$ between $M$ and $N$, with $R = M/N$. The ratio $R$ balances the update frequencies between the connections and link weights. Here we tune the parameter $R$ from 0.05 to 0.2.

As illustrated in Table 2, the best performance is achieved with different $R$ on different datasets. For the Iris dataset, the highest accuracy of 96.67% is obtained when $R$ is set to 0.05. While for the Breast Cancer and the Liver Disorder, the best accuracies of 96.86% and 65.29% are obtained with $R = 0.1$. For the Pima Indians Diabetes, the best accuracy of 73.68% is achieved when $R = 0.2$. The results also indicate that, a larger dataset may require a bigger $R$ in the iterative training.

Iterative training. In this section, we assess the influence
ASA. A supervised learning algorithm with the accurate synaptic-efficiency adjustment method [Xie et al., 2017]. By focusing on the main contents in the target spike trains and ignoring other neuron states, this algorithm converges rapidly. The cost function is based on the voltage difference between the potential of the output neuron and the firing threshold. The accuracies of this method are quoted from [Xie et al., 2017].

SVM. The support vector machines. The standard libSVM toolbox with a radial-basis-function (RBF) kernel is applied to this classification.

SNN-CG. Our method. The settings of the model are the same as those described in Section 3.2. The parameter of $C$ is set to 0.01, and the parameter of $R$ is selected according to the model selection in 3.3.

<table>
<thead>
<tr>
<th>Accuracy</th>
<th>Iris</th>
<th>Breast Cancer</th>
<th>Liver Disorder</th>
<th>Pima Indians Diabetes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tempotron</td>
<td>0.96</td>
<td>0.96</td>
<td>0.61</td>
<td>0.73</td>
</tr>
<tr>
<td>STDP</td>
<td>0.79</td>
<td>0.86</td>
<td>0.58</td>
<td>0.65</td>
</tr>
<tr>
<td>Tempotron-L1</td>
<td>0.95</td>
<td>0.96</td>
<td>0.61</td>
<td>0.73</td>
</tr>
<tr>
<td>SpikeProp</td>
<td>0.96</td>
<td>0.97</td>
<td>0.61</td>
<td>0.72</td>
</tr>
<tr>
<td>ASA</td>
<td>0.95</td>
<td>0.95</td>
<td>0.60</td>
<td>0.72</td>
</tr>
<tr>
<td>SVM</td>
<td><strong>0.99</strong></td>
<td><strong>0.96</strong></td>
<td>0.61</td>
<td>0.71</td>
</tr>
<tr>
<td>SNN-CG (ours)</td>
<td>0.97</td>
<td>0.97</td>
<td><strong>0.65</strong></td>
<td><strong>0.74</strong></td>
</tr>
</tbody>
</table>

Table 3: Comparison with other methods.

Effectiveness of joint learning. In SNN-CG, network connections and link weights are jointly learned in the iterative training scheme. To verify the benefit of joint learning, we compare SNN-CG with single Tempotron and STDP-based learning methods. The results are shown in Table 3. Overall, the SNN-CG method outperforms both of Tempotron and STDP algorithms. Compared with the STDP method, the SNN-CG model achieves significantly better performances with 7%-18% improvement in accuracies. Compared with the classical Tempotron, our method obtains slightly better accuracies with much lower computational costs. As shown in Fig. 4, the SNN-CG method outperforms the classical Tempotron with better accuracies and lower computational costs. Since the SNN-CG reduces the number of connections in the network, the feedforward computation becomes more efficient.

Effectiveness of connection learning. To demonstrate the strength of STDP-based connection learning, we compare SNN-CG with Tempotron using L1 norm regularization, denoted as Tempotron-L1. With the L1-regularization, the weights are more prone to zeros. As illustrated in Fig. 3, compared with the Tempotron-L1 method, the SNN-CG obtains better performance. We further compare the SNN-CG and Tempotron-L1 with different sparsity. As shown in Fig. 5, compared with Tempotron-L1, the SNN-CG method achieves better performance, especially in the high connection sparsity.

Comparison with existing methods. To evaluate the capability of SNN-CG, we compare our method with SVM, SpikeProp [Bohte et al., 2002], and ASA [Xie et al., 2017]. As illustrated in Table 3, for the Iris dataset, we achieves 97% which is higher than SpikeProp (96%) and ASA (95%), and slightly lower than SVM (99%). For the Liver Disorder and the Pima Indians Diabetes datasets, SNN-CG achieves higher accuracies of 65% and 74%. For the Breast Cancer dataset, both SNN-CG and SpikeProp obtain the highest accuracy of 97%, while those of the SVM and ASA are 96% and 95%.

4 Conclusions

This paper proposed a spiking neural network with connection gates (SNN-CG) to jointly learn the connections and the link weights in SNN. The connection structures were optimized by the STDP rule with timing information, and the link weights were optimized by a supervised algorithm. Our method was evaluated on four benchmark datasets. The SNN-CG method outperformed other supervised learning rules in SNN, and the SVM classifier. Experimental results demonstrated that, the joint learning of the connections and link weights improved the strength of SNN models with higher performance and lower computational costs.

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