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Spiking neural network's neuron model describe how the membrance potential of a neuron change over the time. In this chapter, we dive into the biological neuron's dynamics and exhibit several examples neuron models in the spiking neural network.

#### 2.1 Biological Neuron

#### 2.2 Abstract of Neuron Models

In spiking neural network, Equation 2.2.1 to Equation 2.2.4 in the form of **state-space models** (**SMMs**). provide a highly abstract description for describe a biological neuron.

In these equation,  $\mathbf{x}(t)$  is the state vector at time t,  $\dot{x}(t)$  is the state variation at time t,  $\mathbf{I}(t)$  is the input current at time t, V is the membrane potential of the neuron,  $\mathbf{y}(t)$  is the neuron's output at time t, and  $V_{out}$  is the output voltage tha will be sent to the synapses that depature from this neuron.  $(\mathbf{x}(t))_i$  is the i-th element of the state vector.  $\mathcal{P}$  is the model parameters.

$$\dot{\mathbf{x}}(t) = f(\mathbf{x}(t), \mathbf{I}(t), t, \mathscr{P}) \tag{2.2.1}$$

$$\mathbf{y}(t) = g(\mathbf{x}(t), \mathbf{I}(t), t, \mathscr{P}) \tag{2.2.2}$$

$$s.t., (\exists i \in [0, |\mathbf{x}(t)|])(\mathbf{x}(t))_i = V(t)$$
 (2.2.3)

$$(\exists i \in [0, |\mathbf{y}(t)|])_i \mathbf{y}(t) = V_{out}(t)$$
(2.2.4)

Equation 2.2.3 indicates that, a neuron should maintain a membrane potential V, and Equation 2.2.4 indicates that, the neuron's output should contains a voltage  $V_{out}$ .

A discretization version present in Equation 2.2.5 to Equation 2.2.8.

$$\mathbf{x}_{t+1} = f(\mathbf{x}_t, \mathbf{I}_t, t, \mathscr{P}) \tag{2.2.5}$$

$$\mathbf{y}_t = g(\mathbf{x}_t, \mathbf{I}_t, t, \mathscr{P}) \tag{2.2.6}$$

$$s.t., (\exists i \in [0, |\mathbf{x}_t|])(\mathbf{x}_t)_i = V_t$$
 (2.2.7)

$$(\exists i \in [0, |\mathbf{y}_t|])_i \mathbf{y} = V_{out_t} \tag{2.2.8}$$

#### 2.3 Neuron Model Examples

#### 2.3.1 Hodgkin-Huxley (HH) Model

The Hodgkin-Huxley (HH) model can be utilize to accurately reproduce the bio-neuron's dynamics. Its mathematical formulation is presented in Equations 2.3.1 to 2.3.5. By combining all these equations, we arrive at Equation 2.3.6.

$$I(t) = C_m \frac{dV_m(t)}{dt} + I_i(t)$$
(2.3.1)

$$I_i(t) = I_{Na}(t) + I_K(t) + I_l(t)$$
(2.3.2)

$$I_{Na}(t) = g_{Na}(t)(V_m(t) - V_{Na}(t))$$
(2.3.3)

$$I_K(t) = g_K(t)(V_m(t) - V_K(t))$$
(2.3.4)

$$I_l(t) = \bar{g}_l(t)(V_m(t) - V_l(t)) \tag{2.3.5}$$

$$I(t) = C_m \frac{dV_m(t)}{dt} + g_{Na}(t)(V_m(t) - V_{Na}(t)) + g_K(V_m(t) - V_K(t)) + \bar{g}_l(t)(V_m(t) - V_l(t))$$
 (2.3.6)

Ion channel function g are function respect to time t and membrance potential V. Specifically, Equation 2.3.7 is held.

$$g_{Na}(t) = \bar{g}_{Na}m^3(t)h(t)$$
  $g_K(t) = \bar{g}_K n^4(t)$   $g_I(t) = \bar{g}_I$  (2.3.7)

By combining Equations 2.3.1 to 2.3.7, we arrive at Equation 2.3.8.

$$I = C_m \frac{dV_m(t)}{dt} + \bar{g}_{Na}(t)m^3(t)h(t)(V_m(t) - V_{Na}(t)) + \bar{g}_K(t)n^4(t)(V_m(t) - V_K(t)) + \bar{g}_l(t)(V_m(t) - V_l(t))$$
(2.3.8)

 $\frac{d\cdot}{dt} = \alpha \cdot (V_m)(1-\cdot) - \beta \cdot (V_m) \cdot$  is held. In which,  $\cdot$  is a placeholder for m(t), n(t) and h(t). As such, Equation 2.3.9 to Equation 2.3.11 are held.

$$\frac{dn(t)}{dt} = \alpha_n(V_m(t))(1 - n(t)) - \beta_n(V_m(t))n(t)$$
(2.3.9)

$$\frac{dm(t)}{dt} = \alpha_m(V_m(t))(1 - m(t)) - \beta_m(V_m(t))m(t)$$
(2.3.10)

$$\frac{dh(t)}{dt} = \alpha_h(V_m(t))(1 - h(t)) - \beta_h(V_m(t))h(t)$$
(2.3.11)

(2.3.12)

From experiment, Equation 2.3.13 to Equation 2.3.18 were obtained.

$$\alpha_n(V_m(t)) = \frac{0.01(10 - V_m(t))}{exp(\frac{10 - V_m(t)}{10}) - 1}$$
(2.3.13)

$$\alpha_m(V_m(t)) = \frac{0.1(25 - V_m(t))}{exp(\frac{25 - V_m(t)}{10}) - 1}$$
(2.3.14)

$$\alpha_h(V_m(t)) = 0.07exp(-\frac{V_m(t)}{20})$$
 (2.3.15)

$$\beta_n(V_m(t)) = 0.125 exp(-\frac{V_m(t)}{80})$$
(2.3.16)

$$\beta_m(V_m(t)) = 4exp(-\frac{V_m(t)}{18})$$
(2.3.17)

$$\beta_h(V_m(t)) = \frac{1}{exp(\frac{30 - V_m(t)}{10}) + 1}$$
(2.3.18)

Finally, we write the state-space model representation from Equation 2.3.19 to Equation 2.3.25 for Hodgkin-Huxley model.

$$x(t) = \begin{bmatrix} V_m(t) & n(t) & m(t) & h(t) \end{bmatrix}$$
 (2.3.19)

$$\dot{x}(t) = \begin{bmatrix} \dot{V}_m(t) & \dot{n}(t) & \dot{m}(t) & \dot{h}(t) \end{bmatrix}$$
 (2.3.20)

$$\dot{V}_m(t) = -\frac{1}{C_m} (\bar{g}_K n^4(t) (V_m(t) - V_K) + \bar{g}_{Na} m^3(t) h(t) + \bar{g}_l (V_m(t) - V_l(t)) - I(t))$$
(2.3.21)

$$\dot{n}(t) = \alpha_n(V_m(t))(1 - n(t)) - \beta_n(V_m(t))n(t) \tag{2.3.22}$$

$$\dot{m}(t) = \alpha_m(V_m(t))(1 - m(t)) - \beta_m(V_m(t))m(t) \tag{2.3.23}$$

$$\dot{h}(t) = \alpha_h(V_m(t))(1 - h(t)) - \beta_h(V_m(t))h(t)$$
 (2.3.24)

$$V_o ut(t) = V_m(t)$$
 (2.3.25)

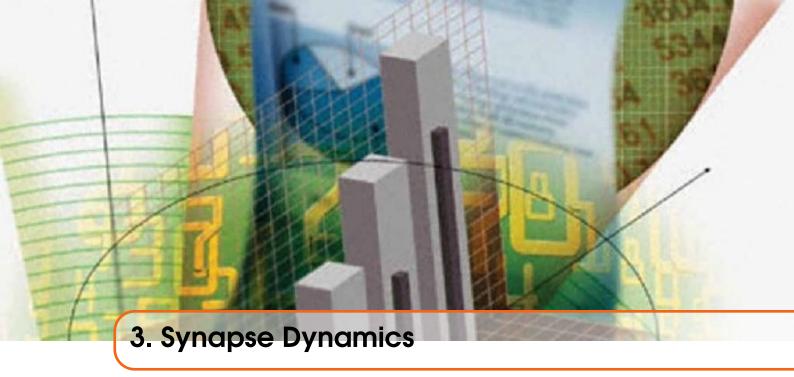
#### 2.3.2 Leaky Integrate-and-fire Model

Leaky Integrate-and-fire model is a computational effective model, in which a threshold is set, when membrance potential cross the threshold, the neuro emit a spike. In implementation, we may set the voltage output at time t,  $V_{out}$  be 1 mV. The scale problem has the potential to be solved by automatically adjusting the synapses' weights in training. A LIF neuron i's output form a **spike train**  $S_i(t) = \sum_m \delta(t - t_m)$ .

$$\dot{V}_t = \frac{1}{\tau_{tot}} \left( -(V_t - V_{rest}) + g_s yn(t) (E_{syn} - V_t) \right)$$
(2.3.26)

- 2.3.3 Izhikevich Model
- 2.3.4 FitzHugh-Nagumo Model
- 2.3.5 Morris-Lecar Model
- 2.3.6 Hindmarsh-Rose Model
- 2.3.7 Cable theory
- 2.3.8 Perfect Integrate-and-fire
- 2.3.9 Adaptive Integrate-and-fire
- 2.3.10 Fring Rate Model
- 2.3.11 Discussion

**Spike Representation** You may note that, although different neuron models have different dynamics and spike representations, they can still communicate with each other through synapses. Some neuron models require current input, while others do not. Nonetheless, spikes can transmit over synapses and cause current variations across them by utilizing a common output element  $V_{out}$ . One challenge that may arise is the normalization of spike representations. To address this issue, one approach is to standardize the spike events by converting all spike representations into binary form. Another approach can involve adjusting the synaptic weights. Although we may face scaling issues with different types of spike representations, these can be mitigated during training by appropriately adjusting the synapse weights.



#### 3.1 Biological Synapse

#### 3.2 Synapse Abstract and Taxonomy

Similar to neuron dynamics, the synapse dynamics could be modeled a state space model, and the current I is included in the state  $\mathbf{x}_t$ . We only need modified the constraint term, as shown in Equation ?? to Equation ??.

The synapse model can be categorized to current-based synapses and conductance-based synapses. Both of them can be seems as a convolution operations on the generalized weighted preneuron spike train. Specifically, we may write  $I(t) = (\theta \cdot S(t)) * h(t)$ , where  $\theta \triangleq \{w_i, g_{max}\}$ , h(t) is a convolution kernel, and S(t) is the pre-synapse neuron's spike train.

- 3.2.1 Current-based Synapse
- 3.2.2 Conductance-based Synapse
- 3.2.3 Chemical Synapse
  Current-based Synapse
  Conductance-based Synapse

#### 3.3 Discussion



### 4.1 Taxonomy of Training Algorithms

From the form of data utilized in the training, we could categories training algorithms of spike neural network into largely two categories: unsupervior learning and supervise learning. While unsupervior learning not explictly specify the expect outcomes in the dataset and align the model to these explicit-specified outcomes, supervior learning explictly specify the expect outcomes in the dataset.

From the theory basis, in this book, we categories the training algorithms into 4 categories:

- surrogate gradient descending methods: The basic idea of surrogate gradient descending methods are utilize surrogate methods to solve the undifferntial term  $\frac{\partial S(t)}{\partial w_{ij}}$  appear in the gradient descending-based loss optimizations. S(t) is a spike train of a neuron, and  $w_{ij}$  is the synapse weight from neuron j to neuron i.
- bio-phenomena-based methods: The bio-phenomena-based inspired by bio-phenomena to adjust the weights dynamically. One kind of these methods is the spike-timing-dependent plasticity (STDP)-based methods, which inspired by the *long-term potentiation (LTP)* and the *long-term depression (LTD)* in the bioneural networks. Base on the difference in time of the the spike emit from pre-post neurons, STDP-based methods adjust the synapse weights.
- statistical mechanism-based methods: Statistical mechanism-based methods tackle the whole networks as a statistical mechanism system. Statistical mechanism methods are used to model the property (e.g., system energy), and the interaction between neurons. With optimize on the system energy, we adjust the weights of synapses.
- **observation methods**: rather than well-train the neural network, observation methods focus on design and training observation models on the neural network. As the system property of neural network at a certain time *t*, contains rich information about the dynamics of input data, by well-design an observation method, we has the potential to obtain the outcomes we expect.

#### 4.2 Unsupervior Learning

- 4.2.1 Spike-timing-dependent plasticity (STDP)
- 4.2.2 Growing Spiking Neural Networks
- 4.2.3 Artola, Bröcher, Singer (ABS) rule
- 4.2.4 Bienenstock, Cooper, Munro (BCM) rule
- 4.2.5 Relationship between BCM and STDP rules
- 4.3 Supervised Learning

#### 4.3.1 STDP-based Methods

Supervised STDP (SSTDP)

Spike-Timing-Dependent Plasticity (STDP) with Supervision

- 4.3.2 Spike-Timing Dependent Backpropagation (STDBP)
- 4.3.3 Liquid State Machine (LSM) and Readout Training

#### 4.3.4 SpikeProp

**Extension** (McKennoch et al., 2006; Booij and tat Nguyen, 2005; Shrestha and Song, 2015; de Montigny and Mâsse, 2016; Banerjee, 2016; Shrestha and Song, 2017).

spike timing based methods is that they cannot learn starting from a quiescent state of no spiking. Bohte (2011)

Huh and Sejnowski (2017)

#### 4.3.5 ReSuMe

**Related Work** (Sporea and Grüning, 2013) Pfister et al. (2006) Gardner et al. (2015) Fremaux et al. (2010)

#### 4.3.6 SuperSpike

SuperSpike [super-spike] is a supervised learning algorithm dedicated to deterministic Leaky Intergrate-and-Fire neuron model. While the backpropagation algorithm used in traditional neural network cannot directly be used in the training of spiking neural network, the author provide a surrogate gradient-based method to tackling with the problems facing in solving the  $S_i/\partial w_{ij}$ , where  $S_i$  is the *i*-th neuron's spike train, and  $w_{ij}$  is the connection weight from neuron *j* to neuron *i*. Specifically,  $S_i(t) = \sum_k \delta(t - t_k)$ , where  $t_k$  is the *k*-th spike emission time, and  $\delta(\cdot)$  is the dirac delta function.

**Approaches** approximate the partial derivative of the hidden unit output by  $f(S_{pre}, g(V_{post}))$ . Let  $\hat{S}_i$  be the target spike train of neuron i. The cost model for optimization that make  $\hat{S}_i$  approach the real  $S_i$  hold the form:  $L = \frac{1}{2} \int_{-\infty}^{t} ds [(\alpha * \hat{S}_i - \alpha * S_i)(s)]^2$ .

 $\alpha$  is a normalized smooth temporal convolution kernel. The original SuperSPike use *double* exponential causal kernel.

$$\partial L/\partial w_{ij} = -\int_{-\infty}^{t} ds [(\alpha * \hat{S}_{i} - \alpha * S_{i})(s)](\alpha * \frac{\partial S_{i}}{\partial w_{ij}})(s)$$

Some existing methods for tackling the term  $\frac{\partial S_i}{\partial w_{ij}}$ : (1) making derivation directly to the membrance voltage, (2) introducing noisy which render the likelihood of  $\langle S_i \rangle$  a smooth function of the membrance potential.

The superspike convert calculation of  $\frac{\partial S_i}{\partial w_{ij}} \to \sigma'(U_i) \frac{\partial U_i}{\partial w_{ij}}$ . In which  $U_i$  is the membrance voltage. Original superspike choose  $\sigma(U)$  be the negative side of a fast sigmoid. This function is objective to increase steeply and peak at the spiking threshold. Other monotonic functions may also work.

For current-based LIF models the membrane potential  $U_i(t)$  can be written in integral form as a spike response model (SRM0 (Gerstner et al., 2014)):  $U_i(t) = \sum_j w_{ij} (\varepsilon * S_j(t)) + (\eta * S_i(t))$  In which,  $\varepsilon$  corresponds to the postsynaptic potential (PSP) shape,  $\eta$  captures spike dynamics and reset.

The existence of term  $(\eta * S_i(t))$  make us difficult to perform derivation. Now,  $U_i(t) \approx (\varepsilon * S_j(t))$ . The gradient calculation for a weight become:

$$\frac{\partial w_{ij}}{\partial t} = r \int_{-\infty}^{t} ds e_i(s) \alpha * (\sigma'(U_i(s))(\varepsilon * S_j)(s))$$

r is the learning rate,  $e_i(s) \equiv \alpha * (\hat{S}_i - S_i)$ ,  $\lambda_{ij} = \alpha * (\sigma'(U_i(s))(\varepsilon * S_j)(s))$  is the eligibility trace. The form above is also known as *non-vanishing surrogate gradient*.

The neuron model utilized by the SuperSpike is

$$\tau^{mem} \frac{dU_i}{dt} = (U^{rest} - U_i) + I_i^{syn}(t)$$

. And synapse evolution model is

$$\frac{d}{dt}I_i^{syn}(t) = -\frac{I_i^{syn}(t)}{\tau^{syn}} + \sum_{j \in pre} w_{ij}S_j(t)$$

- .
- 4.3.7 SPAN (Mohemmed et al., 2012)
- 4.3.8 Remote Supervised Method (ReSuMe)
- 4.3.9 FreqProp
- 4.3.10 Local error-driven associative biologically realistic algorithm (LEABRA)
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  - 4.4 Reinforcement Learning
- 4.4.1 Spiking Actor-Critic method
- 4.4.2 STDP-based Methods
  - 4.5 Convert Trandictional ANN to SNN



#### **5.1** Liquid Neural Network

**State Space Models (SSM)** 

**Linear State Space Models** 

Structured State Space Model (S4) [lnn-s4]

Liquid Time Constant (LTC) Model [ltc]

**Linear Liquid Time Constant Model** 

**High-order Polynomial Projection Operator (HiPPO)** 

Normal Plus Low-Rank (NPLR)

Scaled Legendre Measure (LegS)

**HiPPO Matrix** 

black-box Cauchy Kernel

**Coupled Bilinear** 

**Dynamics Causal Model** 

Liquid structure state space model [lnn-4s] introduce the liquid time constant into the structure state model. Specifically it hold the form in Equation 5.1.1.

$$\frac{d\mathbf{x}(t)}{dt} = -[\mathbf{A} + \mathbf{B} \otimes f(\mathbf{x}(t), \mathbf{u}(t), t, \theta)] \otimes \mathbf{x}(t) + \mathbf{B} \otimes f(\mathbf{x}(t), \mathbf{u}(t), t, \theta)$$
(5.1.1)

A linear version could be expressed by the form of Equation 5.1.2.

$$\dot{\mathbf{x}}(t) = [\mathbf{A} + \mathbb{I}_N \mathbf{B} u(t)] \mathbf{x}(t) + \mathbf{B} u(t), \ y(t) = \mathbf{C} \mathbf{x}(t)$$
(5.1.2)

The output  $y_k$  for input  $u_0$  to  $u_k$  can be represented by  $y = \bar{K} * u + \bar{K}_{liquid} * u_{correlation}$ . In which the \* is the convolution operator. This equation divide the calculation of  $y_k$  to the calculation of the kernel  $\bar{K}$ , and the convolution between the kernal and input, and the convolution between the kernal and input correlation.

- **5.2** Feedforward Neural Network
- **5.3** Recurrent Neural Network
- **5.4** Synfire Chain
- 5.5 Reservoir Computing
- **5.5.1** Liquid State Machine

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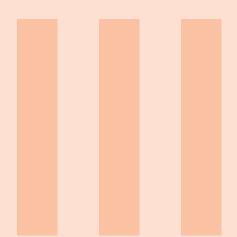


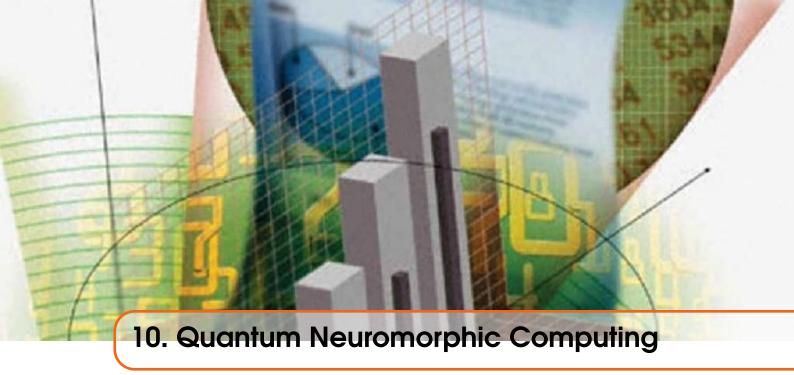
- **6.1** Dynamical System Structures
- 6.1.1 Topology and Geometry
- 6.1.2 Flow and Manifold











- 10.1 System Dynamics-Based Methods
- **10.2** Neuron Simulation Methods







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