bindsnet Documentation

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Contents:

1	Installation	3
	1.1 Pip install	3
	1.2 Installing from source	
	1.3 Running the tests	3
2	Quickstart	5
3	BindsNET User Manual	7
	3.1 Part I: Creating and Adding Network Components	7
	3.2 Part II: Creating and Adding Learning Rules	
4	bindsnet package	17
	4.1 Subpackages	17
	4.2 Module contents	40
5	Indices and tables	41
Ру	thon Module Index	43
In	dex	45

BindsNET is built on top of the PyTorch deep learning platform. It is used for the simulation of spiking neural networks (SNNs) and is geared towards machine learning and reinforcement learning.

BindsNET takes advantage of the torch.Tensor object to build spiking neurons and connections between them, and simulate them on CPUs or GPUs (for strong acceleration / parallelization) without any extra work. Recently, torchvision.datasets has been integrated into the library to allow the use of popular vision datasets in training SNNs for computer vision tasks. Neural network functionality contained in torch.nn.functional module is used to implement more complex connections between populations of spiking neurons.

Spiking neural networks are sometimes referred to as the third generation of neural networks. Rather than the simple linear layers and nonlinear activation functions of deep learning neural networks, SNNs are composed of neural units which more accurately capture properties of their biological counterparts. An important difference between spiking neurons and the artificial neurons of deep learning are the former's integration of input *in time*; they are naturally short-term memory devices by their maintenance of a (possibly decaying) membrane voltage. As a result, some have argued that SNNs are particularly well-suited to model time-varying data.

Neurons are connected together with directed edges (*synapses*) which are (in general) plastic. Synapses may have their own dynamics as well, which may or may not *depend on pre- and post-synaptic neural activity https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3395004/* or *other biological signals https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4717313/*. The modification of synaptic strengths is thought to be an important mechanism by which organisms learn. Accordingly, BindsNET provides a module (**bindsnet.learning**) which contains functions used for the updating of synapse weights.

At its core, BindsNET provides software objects and methods which support the simulation of groups of different types of neurons (**bindsnet.network.nodes**), as well as different types of connections between them (**bindsnet.network.topology**). These may be arbitrarily combined together under a single **bind-snet.network.Network** object, which is responsible for the coordination of the simulation logic of all underlying components. On creation of a network, the user can specify a simulation timestep constant, dt, which determines the granularity of the simulation. Choosing this parameter induces a trade-off between simulation speed and numerical precision: large values result in fast simulation, but poor simulation accuracy, and vice versa. Monitors (**bindsnet.network.monitors**) are available for recording state variables from arbitrary network components (e.g., the voltage v of a group of neurons).

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Installation

1.1 Pip install

Issue:

```
pip install git+https://github.com/BindsNET/bindsnet.git
```

1.2 Installing from source

On *nix systems, issue one of the following in a shell:

```
git clone https://github.com/Hananel-Hazan/bindsnet.git # HTTPS
git clone git@github.com:Hananel-Hazan/bindsnet.git # SSH
```

Change directory into bindsnet and issue one of the following:

```
pip install . # Typical install
pip install -e . # Editable mode (package code can be edited without reinstall)
```

This will install bindsnet and all its dependencies.

1.3 Running the tests

If BindsNET is installed from source, install pytest and issue the following from BindsNET's installation directory:

python -m pytest test

Quickstart

Check out some example use cases for BindsNET in the examples/ folder (link). For example, changing directory to *[bindsnet-root]/examples/mnist* and running the following will result in a near-replication of the architecture of Diehl & Cook 2015:

python eth_mnist.py [options]

The token [options] should be replaced with any command-line arguments you'd like to use to modify the behavior of the program.

BindsNET User Manual

Welcome to BindsNET's user manual! To get started, click on one of the links below.

3.1 Part I: Creating and Adding Network Components

3.1.1 Creating a Network

The bindsnet.network.Network object is BindsNET's main offering. It is responsible for the coordination of simulation of all its constituent components: neurons, synapses, learning rules, etc. To create one:

from bindsnet.network import Network
network = Network()

The bindsnet.network.Network object accepts optional keyword arguments dt, batch_size, learning, and reward_fn.

The dt argument specifies the simulation time step, which determines what temporal granularity, in milliseconds, simulations are solved at. All simulation is done with the Euler method for the sake of computational simplicity. If instability is encountered in your simulation, use a smaller dt to resolve numerical instability.

The batch_size argument specifies the expected minibatch size of the input data. However, since BindsNET supports dynamics minibatch size, this argument can safely be ignored. It is used to initialize stateful neuronal and synaptic variables, and may provide a small speedup if specified beforehand.

The learning argument acts to enable or disable updates to adaptive parameters of network components; e.g., synapse weights or adaptive voltage thresholds. See 'Using Learning Rules'_ for more details.

The reward_fn argument takes in class that specifies how a scalar reward signal will be computed and fed to the network and its components. Typically, the output of this callable class will be used in certain "reward-modulated", or "three-factor" learning rules. See **'Using Learning Rules'** for more details.

3.1.2 Adding Network Components

BindsNET supports three categories of network component: *layers* of neurons (nodes), *connections* between them (*bindsnet.network.topology*), and *monitors* for recording the evolution of state variables (*bindsnet.network.monitors*).

Note: Names of components in a network are arbitrary, and need only be unique within their component group (layers, connections, and monitors) in order to address them uniquely. We encourage our users to develop their own naming conventions, using whatever works best for them.

Creating and adding layers

To create a layer (or *population*) of nodes (in this case, leaky integrate-and-fire (LIF) neurons:

```
from bindsnet.network.nodes import LIFNodes
# Create a layer of 100 LIF neurons with shape (10, 10).
layer = LIFNodes(n=100, shape=(10, 10))
```

Each *bindsnet.network.nodes* object has many keyword arguments, but one of either n (the number of nodes in the layer, or shape (the arrangement of the layer, from which the number of nodes can be computed) is required. Other arguments for certain nodes objects include thresh (scalar or tensor giving voltage threshold(s) for the layer), rest (scalar or tensor giving resting voltage(s) for the layer), traces (whether to keep track of "spike traces" for each neuron in the layer), and tc_decay (scalar or tensor giving time constant(s) of the layer's neurons' voltage decay).

To add a layer to the network, use the add_layer function, and give it a name (a string) to call it by:

```
network.add_layer(
    layer=layer, name="LIF population"
)
```

Such layers are kept in the dictionary attribute network.layers, and can be accessed by the user; e.g., by network.layers['LIF population'].

Other layer types include *bindsnet.network.nodes.Input* (for user-specified input spikes), *bindsnet.network.nodes.McCullochPitts* (the McCulloch-Pitts neuron model), *bindsnet.network.nodes.AdaptiveLIFNodes* (LIF neurons with adaptive thresholds), and *bindsnet.network.nodes.IzhikevichNodes* (the Izhikevich neuron model). Any number of layers can be added to the network.

Custom nodes objects can be implemented by sub-classing *bindsnet.network.nodes.Nodes*, an abstract class with common logic for neuron simulation. The functions forward(self, x: torch. Tensor) (computes effects of input data on neuron population; e.g., voltage changes, spike occurrences, etc.), reset_state_variables(self) (resets neuron state variables to default values), and _compute_decays(self) must be implemented, as they are included as abstract functions of *bindsnet. network.nodes.Nodes*.

Creating and adding connections

Connections can be added between different populations of neurons (a *projection*), or from a population back to itself (a *recurrent* connection). To create an all-to-all connection:

```
from bindsnet.network.nodes import Input, LIFNodes
from bindsnet.network.topology import Connection
# Create two populations of neurons, one to act as the "source"
# population, and the other, the "target population".
source_layer = Input(n=100)
target_layer = LIFNodes(n=1000)
# Connect the two layers.
connection = Connection(
    source=source_layer, target=target_layer
)
```

Like nodes, each connection object has many keyword arguments, but both source and target are required. These must be objects that subclass *bindsnet.network.nodes.Nodes*. Other arguments include w and b (weight and bias tensors for the connection), wmin and wmax (minimum and maximum allowable weight values), update_rule (bindsnet.learning.LearningRule; used for updating connection weights based on pre- and post-synaptic neuron activity and / or global neuromodulatory signals), and norm (a floating point value to normalize weights by).

To add a connection to the network, use the add_connection function, and pass the names given to source and target populations as source and target arguments. Make sure that the source and target neurons are added to the network as well:

```
network.add_layer(
    layer=source_layer, name="A"
)
network.add_layer(
    layer=target_layer, name="B"
)
network.add_connection(
    connection=connection, source="A", target="B"
)
```

Connections are kept in the dictionary attribute network.connections, and can be accessed by the user; e.g., by network.connections['A', 'B']. The layers must be added to the network with matching names (respectively, A and B) in order for the connection to work properly. There are no restrictions on the directionality of connections; layer "A" may connect to layer "B", and "B" back to "A", or "A" may connect directly back to itself.

Custom connection objects can be implemented by sub-classing *bindsnet.network.topology*. *AbstractConnection*, an abstract class with common logic for computing synapse outputs and updates. This includes functions compute (for computing input to downstream layer as a function of spikes and connection weights), update (for updating connection weights based on pre-, post-synaptic activity and possibly other signals; e.g., reward prediction error), normalize (for ensuring weights incident to post-synaptic neurons sum to a pre-specified value), and reset_state_variables (for re-initializing stateful variables for the start of a new simulation).

Specifying monitors

bindsnet.network.monitors.AbstractMonitor objects can be used to record tensor-valued variables over the course of simulation in certain network components. To create a monitor to monitor a single component:

```
from bindsnet.network import Network
from bindsnet.network.nodes import Input, LIFNodes
from bindsnet.network.topology import Connection
from bindsnet.network.monitors import Monitor
```

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```
network = Network()
source_layer = Input(n=100)
target_layer = LIFNodes(n=1000)
connection = Connection(
    source=source_layer, target=target_layer
)
# Create a monitor.
monitor = Monitor(
    obj=target_layer,
    state_vars=("s", "v"), # Record spikes and voltages.
    time=500, # Length of simulation (if known ahead of time).
)
```

The user must specify a Nodes or AbstractConnection object from which to record, attributes of that object to record (state_vars), and, optionally, how many time steps the simulation(s) will last, in order to save time by pre-allocating memory.

To add a monitor to the network (thereby enabling monitoring), use the add_monitor function of the bindsnet. network.Network class:

```
network.add_layer(
    layer=source_layer, name="A"
)
network.add_layer(
    layer=target_layer, name="B"
)
network.add_connection(
    connection=connection, source="A", target="B"
)
network.add_monitor(monitor=monitor, name="B")
```

The name given to the monitor is not important. It is simply used by the user to select from the monitor objects controlled by a Network instance.

One can get the contents of a monitor by calling network.monitors [<name>].get (<state_var>), where <state_var> is a member of the iterable passed in for the state_vars argument. This returns a tensor of shape (time, n_1 , ..., n_k), where (n_1 , ..., n_k) is the shape of the recorded state variable.

The bindsnet.network.monitors.NetworkMonitor is used to record from many network components at once. To create one:

```
from bindsnet.network.monitors import NetworkMonitor
network_monitor = NetworkMonitor(
    network: Network,
    layers: Optional[Iterable[str]],
    connections: Optional[Iterable[Tuple[str, str]]],
    state_vars: Optional[Iterable[str]],
    time: Optional[int],
)
```

The user must specify the network to record from, an iterable of names of layers (entries in network.layers), an iterable of 2-tuples referring to connections (entries in network.connections), an iterable of tensor-valued state

variables to record during simulation (state_vars), and, optionally, how many time steps the simulation(s) will last, in order to save time by pre-allocating memory.

Similarly, one can get the contents of a network monitor by calling network.monitors[<name>].get(). Note this function takes no arguments; it returns a dictionary mapping network components to a sub-dictionary mapping state variables to their tensor-valued recording.

3.1.3 Running Simulations

After building up a Network object, the next step is to run a simulation. Here, the function Network.run comes into play. It takes arguments inputs (a dictionary mapping names of layers subclassing AbstractInput to input data of shape [time, batch_size, *input_shape], where input_shape is the shape of the neuron population to which the data is passed), time (the number of simulation timesteps, generally thought of as milliseconds), and a number of keyword arguments, including clamp (and unclamp), used to force neurons to spike (or not spike) at any given time step, reward, for supplying to reward-modulated learning rules, and masks, a dictionary mapping connections to boolean tensors specifying which synapses weights to clamp to zero.

Building on the previous parts of this guide, we present a simple end-to-end example of simulating a two-layer, inputoutput spiking neural network.

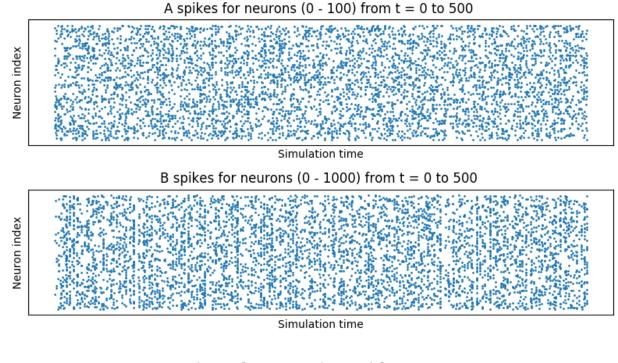
```
import torch
import matplotlib.pyplot as plt
from bindsnet.network import Network
from bindsnet.network.nodes import Input, LIFNodes
from bindsnet.network.topology import Connection
from bindsnet.network.monitors import Monitor
from bindsnet.analysis.plotting import plot_spikes, plot_voltages
# Simulation time.
time = 500
# Create the network.
network = Network()
# Create and add input, output layers.
source_layer = Input (n=100)
target_layer = LIFNodes(n=1000)
network.add_layer(
   layer=source_layer, name="A"
)
network.add laver(
    layer=target_layer, name="B"
)
# Create connection between input and output layers.
forward_connection = Connection(
    source=source layer,
   target=target_layer,
   w=0.05 + 0.1 * torch.randn(source_layer.n, target_layer.n), # Normal(0.05, 0.01).
\rightarrow weights.
)
network.add_connection(
    connection=forward connection, source="A", target="B"
)
```

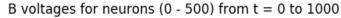
(continues on next page)

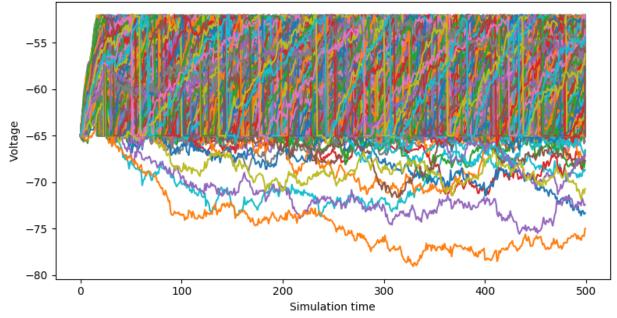
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```
# Create recurrent connection in output layer.
recurrent_connection = Connection(
   source=target_layer,
   target=target_layer,
   w=0.025 * (torch.eye(target_layer.n) - 1), # Small, inhibitory "competitive"
\rightarrow weights.
)
network.add_connection(
   connection=recurrent_connection, source="B", target="B"
)
# Create and add input and output layer monitors.
source_monitor = Monitor(
   obj=source_layer,
   state_vars=("s",), # Record spikes and voltages.
   time=time, # Length of simulation (if known ahead of time).
target_monitor = Monitor(
   obj=target_layer,
    state_vars=("s", "v"), # Record spikes and voltages.
   time=time, # Length of simulation (if known ahead of time).
)
network.add_monitor(monitor=source_monitor, name="A")
network.add_monitor(monitor=target_monitor, name="B")
# Create input spike data, where each spike is distributed according to Bernoulli(0.
\hookrightarrow 1
input_data = torch.bernoulli(0.1 * torch.ones(time, source_layer.n)).byte()
inputs = {"A": input_data}
# Simulate network on input data.
network.run(inputs=inputs, time=time)
# Retrieve and plot simulation spike, voltage data from monitors.
spikes = {
   "A": source_monitor.get("s"), "B": target_monitor.get("s")
voltages = {"B": target_monitor.get("v")}
plt.ioff()
plot_spikes(spikes)
plot_voltages(voltages, plot_type="line")
plt.show()
```

This script will result in figures that looks something like this:







Notice that, in the voltages plot, no voltage goes above -52mV, the default threshold of the LIFNodes object. After hitting this point, neurons' voltage is reset to -64mV, which can also be seen in the figure.

3.1.4 Simulation Notes

The simulation of all network components is *synchronous* (*clock-driven*); i.e., all components are updated at each time step. Other frameworks use *event-driven* simulation, where spikes can occur at arbitrary times instead of at regular

multiples of dt. We chose clock-driven simulation due to ease of implementation and for computational efficiency considerations.

During a simulation step, input to each layer is computed as the sum of all outputs from layers connecting to it (weighted by synapse weights) from the *previous* simulation time step (implemented by the _get_inputs method of the bindsnet.network.Network class). This model allows us to decouple network components and perform their simulation separately at the temporal granularity of chosen dt, interacting only between simulation steps.

This is a strict departure from the computation of *deep neural networks* (DNNs), in which an ordering of layers is supposed, and layers' activations are computed *in sequence* from the shallowest to the deepest layer in a single time step, with the exclusion of recurrent layers, whose computations are still ordered in time.

3.2 Part II: Creating and Adding Learning Rules

3.2.1 What is considered a learning rule?

Learning rules are necessary for the automated adaption of network parameters during simulation. At present, BindsNET supports two different categories of learning rules:

- Two factor: Associative learning takes place based on pre- and post-synaptic neural activity. Examples include:
 - The typical example is Hebbian learning, which may be summarized as "Cells that fire together wire together." That is, co-active neurons causes their connection strength to increase.
 - Spike-timing-dependent plasticity (STDP) stipulates that the ordering of pre- and post-synaptic spikes
 matters. A synapse is strengthened if the pre-synaptic neuron fires *before* the post-synaptic neuron,
 and, conversely, is weakened if it fires *after* the post-synaptic neuron. The magnitude of these updates
 is a decreasing function of the time between pre- and post-synaptic spikes.
- Three factor: In addition to associating pre- and post-synaptic neural activity, a third factor is introduced which modulate
 - (Reward, error, attention)-modulated (Hebbian learning, STDP): The same learning rules described above are modulated by the presence of global signals such as reward, error, or attention, which can be variously defined in machine learning or reinforcement learning contexts. These signals act to gate plasticity, turning it on or off and switching its sign and magnitude, based on the task at hand.

The above are examples of local learning rules, where the information needed to make updates are thought to be available at the synapse. For example, pre- and post-synaptic neurons are adjacent to synapses, rendering their spiking activity accessible, whereas chemical signals like dopamine (hypothesized to be a reward prediction error (RPE) signal) are widely distributed across certain neuron populations; i.e., they are *globally* available. This is in contrast to learning algorithms like back-propagation, where per-synapse error signals are derived by computing backwards from a loss function at the network's output layer. Such error derivation is thought to be biologically implausible, especially compared to the two- and three-factor rules mentioned above.

3.2.2 Creating a learning rule in BindsNET

At present, learning rules are attached to specific Connection objects. For example, to create a connection with a STDP learning rule on the synapses:

```
from bindsnet.network.nodes import Input, LIFNodes
from bindsnet.network.topology import Connection
from bindsnet.learning import PostPre
```

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```
# Create two populations of neurons, one to act as the "source"
# population, and the other, the "target population".
# Neurons involved in certain learning rules must record synaptic
# traces, a vector of short-term memories of the last emitted spikes.
source_layer = Input(n=100, traces=True)
target_layer = LIFNodes(n=1000, traces=True)
# Connect the two layers.
connection = Connection(
    source=source_layer, target=target_layer, update_rule=PostPre, nu=(1e-4, 1e-2)
)
```

The connection may be added to a Network instance as usual. The Connection object takes arguments update_rule, of type bindsnet.learning.LearningRule, as well as nu, a 2-tuple specifying pre- and post-synaptic learning rates; i.e., multiplicative factors which modulate how quickly synapse weights change.

Learning rules also accept arguments reduction, which specifies how parameter updates are aggregated across the batch dimension, and weight_decay, which specifies the time constant of the rate of decay of synapse weights to zero. By default, parameter updates are averaged across the batch dimension, and there is no weight decay.

Other supported learning rules include Hebbian, WeightDependentPostPre, MSTDP (reward-modulated STDP), and MSTDPET (reward-modulated STDP with eligibility traces).

Custom learning rules can be implemented by subclassing bindsnet.learning.LearningRule and providing implementations for the types of AbstractConnection objects intended to be used. For example, the Connection and LocalConnection objects rely on the implementation of a private method, _connection_update, whereas the Conv2dConnection object uses the _conv2d_connection_update version.

bindsnet package

4.1 Subpackages

4.1.1 bindsnet.analysis package

Submodules

bindsnet.analysis.plotting module

<pre>bindsnet.analysis.plotting.plot_assignments</pre>	(assign	ments:	torch.Tenso	or, in	m:	Op-
	tional[matplotli	b.image.Axes	sImage	2]	=
	None,	figsize:	Tuple[int,	int]	= (5,	5),
	classes	: Opt	ional[Sized]	= 1	Vone)	\rightarrow
	matplo	tlib.imag	e.AxesImage	e		

Plot the two-dimensional neuron assignments.

Parameters

- **assignments** Vector of neuron label assignments.
- **im** Used for re-drawing the assignments plot.
- **figsize** Horizontal, vertical figure size in inches.
- **classes** Iterable of labels for colorbar ticks corresponding to data labels.

Returns Used for re-drawing the assignments plot.

bindsnet.analysis.plotting.plot_conv2d_weight	s (weights: torch.Tensor, wmin: float
	= 0.0, wmax: float = 1.0, im: Op-
	tional[matplotlib.image.AxesImage]
	= None, figsize: Tuple[int, int] =
	$(5, 5), cmap: str = 'hot_r') \rightarrow mat-$
	plotlib.image.AxesImage

Plot a connection weight matrix of a Conv2dConnection.

Parameters

- weights Weight matrix of Conv2dConnection object.
- wmin Minimum allowed weight value.
- wmax Maximum allowed weight value.
- im Used for re-drawing the weights plot.
- **figsize** Horizontal, vertical figure size in inches.
- **cmap** Matplotlib colormap.

Returns Used for re-drawing the weights plot.

<pre>bindsnet.analysis.plotting.plot_input</pre>	(image:	torch.Ter	ısor, in	pt:	torch.	Tensor,
	label:	Optiona	l[int]	= Λ	one,	axes:
	List[ma	tplotlib.axes	axes.Axes	1	=	None,
	ims:	List[mat	plotlib.im	ige.Axe	sImage] =
	None,	figsize: T	Tuple[int,	int]	= (8	3, 4))
	\rightarrow	Tuple[L	.ist[matplc	tlib.axe	esaxes	s.Axes],
	List[ma	tplotlib.image	.AxesImag	[[e]		

Plots a two-dimensional image and its corresponding spike-train representation.

Parameters

- **image** A 2D array of floats depicting an input image.
- inpt A 2D array of floats depicting an image's spike-train encoding.
- **label** Class label of the input data.
- **axes** Used for re-drawing the input plots.
- **ims** Used for re-drawing the input plots.
- **figsize** Horizontal, vertical figure size in inches.

Returns Tuple of (axes, ims) used for re-drawing the input plots.

bindsnet.analysis.plotting.plot_locally_connected_weights	s (weights:	torch.T	ensor,
	n_filters:	int,	ker-
	nel_size:	Unio	on[int,
	Tuple[int,		int]],
	conv_size.	: Unio	on[int,
	Tuple[int,	int]],	loca-
	tions:	torch.T	ensor,
	input_sqr	t: Unio	on[int,
	Tuple[int,	int]], v	wmin:
	float =	0.0, w	vmax:
	float = 1	.0, im:	Op-
	tional[ma	tplotlib.i	mage.AxesImage]
	= None,	lines: b	ool =
	True, figst	ize: Tupi	le[int,
	int] = (.1)	5, 5), a	cmap:
	str = 'ho	$t_r') \rightarrow$	mat-
	plotlib.im	age.Axes	sImage
Plot a connection weight matrix of a Connection with locally connected s	structure <ht< td=""><td>tn•//vann</td><td>lecun com/</td></ht<>	tn•//vann	lecun com/

Plot a connection weight matrix of a Connection with 'locally connected structure <http://yann.lecun.com/ exdb/publis/pdf/gregor-nips-11.pdf>_.

Parameters

• weights – Weight matrix of Conv2dConnection object.

- n_filters No. of convolution kernels in use.
- **kernel_size** Side length(s) of 2D convolution kernels.
- **conv_size** Side length(s) of 2D convolution population.
- locations Indices of input receptive fields for convolution population neurons.
- **input_sqrt** Side length(s) of 2D input data.
- wmin Minimum allowed weight value.
- wmax Maximum allowed weight value.
- im Used for re-drawing the weights plot.
- lines Whether or not to draw horizontal and vertical lines separating input regions.
- figsize Horizontal, vertical figure size in inches.
- **cmap** Matplotlib colormap.

Returns Used for re-drawing the weights plot.

```
bindsnet.analysis.plotting.plot_performance (performances: Dict[str, List[float]], ax: Op-
tional[matplotlib.axes._axes.Axes] = None,
figsize: Tuple[int, int] = (7, 4)) → mat-
plotlib.axes._axes.Axes
```

Plot training accuracy curves.

Parameters

- **performances** Lists of training accuracy estimates per voting scheme.
- **ax** Used for re-drawing the performance plot.
- figsize Horizontal, vertical figure size in inches.

Returns Used for re-drawing the performance plot.

bindsnet.analysis.plotting.plot_spikes (spikes: Dict[str, torch.Tensor], time: Of	р-
tional[Tuple[int, int]] = None, n_neurons: O	р-
tional[Dict[str, Tuple[int, int]]] = None, ims: O	р-
tional[List[matplotlib.collections.PathCollection]]	
= None, axes: Union[matplotlib.axesaxes.Axe	?S,
List[matplotlib.axesaxes.Axes], None]	=
None, figsize: Tuple[float, float] = (8.0, 4.5)))
\rightarrow Tuple[List[matplotlib.image.AxesImage]	*],
List[matplotlib.axesaxes.Axes]]	

Plot spikes for any group(s) of neurons.

- **spikes** Mapping from layer names to spiking data. Spike data has shape [time, n_1, ..., n_k], where [n_1, ..., n_k] is the shape of the recorded layer.
- time Plot spiking activity of neurons in the given time range. Default is entire simulation time.
- n_neurons Plot spiking activity of neurons in the given range of neurons. Default is all neurons.
- **ims** Used for re-drawing the plots.
- **axes** Used for re-drawing the plots.
- figsize Horizontal, vertical figure size in inches.

Returns ims, axes: Used for re-drawing the plots.

bindsnet.analysis.plotting.plot_voltages	(voltages: Dict[str, torch.Tensor], ims: Op-
	tional[List[matplotlib.image.AxesImage]]
	= None, axes: Op-
	tional[List[matplotlib.axesaxes.Axes]] =
	None, time: Tuple[int, int] = None, n_neurons:
	Optional[Dict[str, Tuple[int, int]]] = None,
	cmap: Optional[str] = 'jet', plot_type: str =
	'color', thresholds: Dict[str, torch.Tensor] =
	<i>None, figsize: Tuple[float, float] = (8.0, 4.5))</i>
	\rightarrow Tuple[List[matplotlib.image.AxesImage],
	List[matplotlib.axesaxes.Axes]]

Plot voltages for any group(s) of neurons.

Parameters

- voltages Contains voltage data by neuron layers.
- **ims** Used for re-drawing the plots.
- **axes** Used for re-drawing the plots.
- time Plot voltages of neurons in given time range. Default is entire simulation time.
- **n_neurons** Plot voltages of neurons in given range of neurons. Default is all neurons.
- **cmap** Matplotlib colormap to use.
- **figsize** Horizontal, vertical figure size in inches.
- plot_type The way how to draw graph. 'color' for pcolormesh, 'line' for curved lines.
- thresholds Thresholds of the neurons in each layer.

Returns ims, axes: Used for re-drawing the plots.

```
bindsnet.analysis.plotting.plot_weights (weights: torch.Tensor, wmin: Optional[float]
= 0, wmax: Optional[float] = 1, im: Op-
tional[matplotlib.image.AxesImage] = None, fig-
size: Tuple[int, int] = (5, 5), cmap: str = 'hot_r')
→ matplotlib.image.AxesImage
```

Plot a connection weight matrix.

Parameters

- weights Weight matrix of Connection object.
- wmin Minimum allowed weight value.
- wmax Maximum allowed weight value.
- im Used for re-drawing the weights plot.
- **figsize** Horizontal, vertical figure size in inches.
- **cmap** Matplotlib colormap.

Returns AxesImage for re-drawing the weights plot.

bindsnet.analysis.visualization module

```
bindsnet.analysis.visualization.plot_spike_trains_for_example(spikes:
```

torch.Tensor, n_ex : Optional[int] = None, top_k: Optional[int] = None, indices: Optional[List[int]] = None) \rightarrow None

Plot spike trains for top-k neurons or for specific indices.

Parameters

- **spikes Spikes** for one simulation run of shape (n_examples, n_neurons, time).
- **n_ex** Allows user to pick which example to plot spikes for.
- top_k Plot k neurons that spiked the most for n_ex example.
- **indices** Plot specific neurons' spiking activity instead of top_k.

bindsnet.analysis.visualization.**plot_voltage** (voltage: torch.Tensor, n_ex: int = 0, n_neuron: int = 0, time: Optional[Tuple[int, int]] = None, threshold: float = None) \rightarrow None

Plot voltage for a single neuron on a specific example.

Parameters

- voltage Tensor or array of shape [n_examples, n_neurons, time].
- **n_ex** Allows user to pick which example to plot voltage for.
- **n_neuron** Neuron index for which to plot voltages for.
- time Plot spiking activity of neurons between the given range of time.
- threshold Neuron spiking threshold.

bindsnet.analysis.visualization.plot_weights_movie(ws: numpy.ndarray, sample_every: int = 1) \rightarrow None

Create and plot movie of weights.

Parameters

- ws Array of shape [n_examples, source, target, time].
- **sample_every** Sub-sample using this parameter.

Module contents

4.1.2 bindsnet.datasets package

Submodules

bindsnet.datasets.preprocess module

```
class bindsnet.datasets.preprocess.BoundingBox (x1, y1, x2, y2)
     Bases: object
     compute_output_height()
     compute_output_width()
     edge_spacing_x()
     edge_spacing_y()
     get_bb_list()
     get_center_x()
     get_center_y()
     get_height()
     get_width()
     print_bb()
     recenter (search_loc, edge_spacing_x, edge_spacing_y, bbox_gt_recentered)
     scale(image)
     shift (image, lambda_scale_frac, lambda_shift_frac, min_scale, max_scale, shift_motion model,
            bbox rand)
     uncenter (raw_image, search_location, edge_spacing_x, edge_spacing_y)
     unscale(image)
class bindsnet.datasets.preprocess.NormalizeToTensor
     Bases: object
     Returns torch tensor normalized images.
class bindsnet.datasets.preprocess.Rescale (output size)
     Bases: object
     Rescale image and bounding box. Args:
         output_size (tuple or int): Desired output size. If int, square crop is made.
bindsnet.datasets.preprocess.bgr2rgb(image)
bindsnet.datasets.preprocess.binary_image(image: numpy.ndarray) → numpy.ndarray
     Converts input image into black and white (binary)
         Parameters image – Gray-scaled image.
         Returns Black and white image.
bindsnet.datasets.preprocess.computeCropPadImageLocation(bbox_tight, image)
```

bindsnet.datasets.preprocess.crop (*image: numpy.ndarray*, x1: int, x2: int, y1: int, y2: int) \rightarrow numpy.ndarray

Crops an image given coordinates of cropping box.

- **image** 3-dimensional image.
- **x1** Left x coordinate.

- **x2** Right x coordinate.
- **y1** Bottom y coordinate.
- **y2** Top y coordinate.

Returns Image cropped using coordinates (x1, x2, y1, y2).

bindsnet.datasets.preprocess.cropPadImage(bbox_tight, image)

bindsnet.datasets.preprocess.crop_sample(sample)

Given a sample image with bounding box, this method returns the image crop at the bounding box location with twice the width and height for context.

bindsnet.datasets.preprocess.gray_scale(*image: numpy.ndarray*) \rightarrow numpy.ndarray Converts RGB image into grayscale.

Parameters image – RGB image.

Returns Gray-scaled image.

bindsnet.datasets.preprocess.sample_exp_two_sides(lambda_)

```
bindsnet.datasets.preprocess.sample_rand_uniform()
```

bindsnet.datasets.preprocess.shift_crop_training_sample (sample, bb_params)
Given an image with bounding box, this method randomly shifts the box and generates a training example. It
returns current image crop with shifted box (with respect to current image).

```
bindsnet.datasets.preprocess.subsample(image: numpy.ndarray, x: int, y: int) \rightarrow numpy.ndarray
```

Scale the image to (x, y).

Parameters

- **image** Image to be rescaled.
- **x** Output value for image's x dimension.
- **y** Output value for image's y dimension.

Returns Re-scaled image.

Module contents

4.1.3 bindsnet.encoding package

Module contents

4.1.4 bindsnet.environment package

Module contents

4.1.5 bindsnet.evaluation package

Module contents

4.1.6 bindsnet.learning package

Module contents

4.1.7 bindsnet.network package

Submodules

bindsnet.network.monitors module

class bindsnet.network.monitors.AbstractMonitor Bases: abc.ABC

Abstract base class for state variable monitors.

class bindsnet.network.monitors.**Monitor** (*obj*: Union[bindsnet.network.nodes.Nodes, bindsnet.network.topology.AbstractConnection], state_vars: Iterable[str], time: Optional[int] = *None, batch_size: int* = 1) Bases: bindsnet.network.monitors.AbstractMonitor

Records state variables of interest.

Constructs a Monitor object.

Parameters

- **obj** An object to record state variables from during network simulation.
- **state_vars** Iterable of strings indicating names of state variables to record.
- time If not None, pre-allocate memory for state variable recording.

get (*var: str*) \rightarrow torch.Tensor

Return recording to user.

Parameters var – State variable recording to return.

Returns Tensor of shape [time, n_1, \ldots, n_k], where $[n_1, \ldots, n_k]$ is the shape of the recorded state variable.

record() \rightarrow None

Appends the current value of the recorded state variables to the recording.

$\texttt{reset_state_variables()} \rightarrow None$

Resets recordings to empty "torch.Tensor"s.

class	bindsnet.network.monitc	rs.NetworkMonitor	(networ	k: Network,	layers:	Op-
			tional[Iterable[str]] =	None, co	onnec-
			tions:	Optional[Iterabl	e[str]] =	None,
			state_v	ars: Optional[Iterable[st	[r]] =
			None, i	time: Optional[in	t] = None)

Bases: bindsnet.network.monitors.AbstractMonitor

Record state variables of all layers and connections.

Constructs a NetworkMonitor object.

- **network** Network to record state variables from.
- **layers** Layers to record state variables from.
- connections Connections to record state variables from.

- **state_vars** List of strings indicating names of state variables to record.
- time If not None, pre-allocate memory for state variable recording.

Returns Dictionary of dictionary of all layers' and connections' recorded state variables.

$\texttt{record}\,(\,)\,\rightarrow None$

Appends the current value of the recorded state variables to the recording.

$\texttt{reset_state_variables()} \rightarrow None$

Resets recordings to empty torch. Tensors.

save (*path: str, fmt: str* = '*npz*') \rightarrow None Write the recording dictionary out to file.

Parameters

- path The directory to which to write the monitor's recording.
- fmt Type of file to write to disk. One of "pickle" or "npz".

bindsnet.network.nodes module

```
class bindsnet.network.nodes.AbstractInput
    Bases: abc.ABC
```

Abstract base class for groups of input neurons.

class bindsnet.network.nodes.AdaptiveLIFNode	s (n: Optional[int] = None, shape: Op-
	tional[Iterable[int]] = None, traces: bool
	= False, traces_additive: bool = False,
	tc_trace: Union[float, torch.Tensor]
	= 20.0, trace_scale: Union[float,
	torch.Tensor] = 1.0, sum_input: bool =
	False, rest: Union[float, torch.Tensor] =
	-65.0, reset: Union[float, torch.Tensor] =
	-65.0, thresh: Union[float, torch.Tensor]
	= -52.0, refrac: Union[int, torch.Tensor]
	= 5, tc_decay: Union[float, torch.Tensor]
	= 100.0, theta_plus: Union[float,
	torch.Tensor] = 0.05, tc_theta_decay:
	Union[float, torch.Tensor] = 10000000.0,
	<i>lbound: float = None</i> , ** <i>kwargs</i>)

Bases: bindsnet.network.nodes.Nodes

Layer of leaky integrate-and-fire (LIF) neurons with adaptive thresholds. A neuron's voltage threshold is increased by some constant each time it spikes; otherwise, it is decaying back to its default value.

Instantiates a layer of LIF neurons with adaptive firing thresholds.

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- **traces_additive** Whether to record spike traces additively.

- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **rest** Resting membrane voltage.
- **reset** Post-spike reset voltage.
- **thresh** Spike threshold voltage.
- **refrac** Refractory (non-firing) period of the neuron.
- tc_decay Time constant of neuron voltage decay.
- theta_plus Voltage increase of threshold after spiking.
- tc_theta_decay Time constant of adaptive threshold decay.
- **lbound** Lower bound of the voltage.

```
forward (x: torch.Tensor) \rightarrow None
Runs a single simulation step.
```

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size (*batch_size*) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

class bindsnet.network.nodes.CurrentLIFNodes	(n: Optional[int] = None, shape: Op-
	tional[Iterable[int]] = None, traces: bool
	= False, traces_additive: bool = False,
	<i>tc_trace: Union[float, torch.Tensor] = 20.0,</i>
	trace_scale: Union[float, torch.Tensor]
	= 1.0, sum_input: bool = False, thresh:
	Union[float, torch.Tensor] = -52.0, rest:
	Union[float, torch.Tensor] = -65.0, reset:
	<i>Union[float, torch.Tensor] = -65.0, refrac:</i>
	<i>Union[int, torch.Tensor] = 5, tc_decay:</i>
	Union[float, torch.Tensor] = 100.0,
	tc_i_decay: Union[float, torch.Tensor] =
	2.0, <i>lbound: float = None</i> , **kwargs)

Bases: bindsnet.network.nodes.Nodes

Layer of current-based leaky integrate-and-fire (LIF) neurons. Total synaptic input current is modeled as a decaying memory of input spikes multiplied by synaptic strengths.

Instantiates a layer of synaptic input current-based LIF neurons. :param n: The number of neurons in the layer. :param shape: The dimensionality of the layer. :param traces: Whether to record spike traces. :param traces_additive: Whether to record spike traces additively. :param tc_trace: Time constant of spike trace decay. :param trace_scale: Scaling factor for spike trace. :param sum_input: Whether to sum all inputs. :param thresh: Spike threshold voltage. :param rest: Resting membrane voltage. :param reset: Post-spike reset voltage. :param refrac: Refractory (non-firing) period of the neuron. :param tc_decay: Time constant of neuron voltage decay. :param tc_i_decay: Time constant of synaptic input current decay. :param lbound: Lower bound of the voltage.

forward (*x: torch.Tensor*) \rightarrow None Runs a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size ($batch_size$) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

class bindsnet.network.nodes.**DiehlAndCookNodes** (n: Optional[int] = None, shape: Optional[Iterable[int]] = None, traces: bool = False, traces_additive: bool = False, tc_trace: Union[float, torch.Tensor] = 20.0, trace scale: Union[float, torch.Tensor] = 1.0,sum input: bool = False, thresh: *Union[float, torch.Tensor] = -52.0, rest:* Union[float, torch.Tensor] = -65.0, reset: Union[float, torch.Tensor] = -65.0, refrac: Union[int, torch.Tensor] = 5, tc_decay: Union[float, torch.Tensor] = 100.0, theta plus: Union[float, *torch.Tensor*] = 0.05, *tc_theta_decay:* Union[float, *torch.Tensor*] 10000000.0, lbound: float = None, *one_spike: bool = True*, ***kwargs*) Bases: bindsnet.network.nodes.Nodes

Layer of leaky integrate-and-fire (LIF) neurons with adaptive thresholds (modified for Diehl & Cook 2015 replication).

Instantiates a layer of Diehl & Cook 2015 neurons.

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- traces_additive Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- thresh Spike threshold voltage.
- **rest** Resting membrane voltage.
- **reset** Post-spike reset voltage.
- **refrac** Refractory (non-firing) period of the neuron.
- tc_decay Time constant of neuron voltage decay.

- theta_plus Voltage increase of threshold after spiking.
- tc_theta_decay Time constant of adaptive threshold decay.
- **lbound** Lower bound of the voltage.
- **one_spike** Whether to allow only one spike per timestep.

sets the relevant decays.

forward (*x: torch.Tensor*) \rightarrow None Runs a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size (*batch_size*) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

class bindsnet.network.nodes.IFNodes (n: Optional[int] = None, shape: Optional[Iterable[int]] = None, traces: bool = False, traces_additive: bool = False, tc_trace: Union[float, torch.Tensor] = 20.0, trace_scale: Union[float, torch.Tensor] = 1.0, sum_input: bool = False, thresh: Union[float, torch.Tensor] = -52.0, reset: Union[float, torch.Tensor] = -65.0, refrac: Union[int, torch.Tensor] = 5, lbound: float = None, **kwargs)

Bases: bindsnet.network.nodes.Nodes

Layer of integrate-and-fire (IF) neurons.

Instantiates a layer of IF neurons.

Parameters

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- traces_additive Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **thresh** Spike threshold voltage.
- **reset** Post-spike reset voltage.
- **refrac** Refractory (non-firing) period of the neuron.
- **lbound** Lower bound of the voltage.

forward (*x: torch.Tensor*) \rightarrow None Runs a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size (*batch_size*) \rightarrow None

Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

Bases: bindsnet.network.nodes.Nodes, bindsnet.network.nodes.AbstractInput

Layer of nodes with user-specified spiking behavior.

Instantiates a layer of input neurons.

Parameters

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- traces Whether to record decaying spike traces.
- **traces_additive** Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.

forward (*x: torch.Tensor*) \rightarrow None

On each simulation step, set the spikes of the population equal to the inputs.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

class bindsnet.network.nodes.IzhikevichNodes	s (n: Optional[int] = None, shape: Op-
	tional[Iterable[int]] = None, traces: bool
	= False, traces_additive: bool = False,
	tc_trace: Union[float, torch.Tensor] = 20.0,
	<pre>trace_scale: Union[float, torch.Tensor] =</pre>
	1.0, sum_input: bool = False, excitatory:
	<pre>float = 1, thresh: Union[float, torch.Tensor]</pre>
	= 45.0, rest: Union[float, torch.Tensor] = -
	65.0, lbound: float = None, **kwargs)
Bases: bindsnet.network.nodes.Nodes	
Layer of Izhikevich neurons.	

Instantiates a layer of Izhikevich neurons.

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.

- **traces_additive** Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **excitatory** Percent of excitatory (vs. inhibitory) neurons in the layer; in range [0, 1].
- **thresh** Spike threshold voltage.
- **rest** Resting membrane voltage.
- **lbound** Lower bound of the voltage.

```
forward (x: torch.Tensor) \rightarrow None
Runs a single simulation step.
```

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size ($batch_size$) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

class bindsnet.network.nodes.**LIFNodes**(*n*: *Optional[int]* = *None*, shape: Op*tional*[*Iterable*[*int*]] = *None*, *traces*: bool =False, traces_additive: bool = False, tc_trace: *Union[float, torch.Tensor] = 20.0, trace_scale:* Union[float, torch.Tensor] = 1.0, sum_input: bool = False, thresh: Union[float, torch.Tensor] = -52.0, rest: Union[float, torch.Tensor] = -65.0, reset: Union[float, torch.Tensor] = -65.0, refrac: Union[int, torch.Tensor] = 5, tc_decay: Union[float, torch.Tensor] = 100.0, lbound: float = None, **kwargs) Bases: bindsnet.network.nodes.Nodes

Layer of leaky integrate-and-fire (LIF) neurons.

Instantiates a layer of LIF neurons.

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- traces_additive Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **thresh** Spike threshold voltage.
- **rest** Resting membrane voltage.

- reset Post-spike reset voltage.
- **refrac** Refractory (non-firing) period of the neuron.
- tc_decay Time constant of neuron voltage decay.
- **lbound** Lower bound of the voltage.

forward (*x: torch.Tensor*) \rightarrow None Runs a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size (*batch_size*) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

class bindsnet.network.nodes.McCullochPitts (n: Optional[int] = None, shape: Optional[Iterable[int]] = None, traces: bool = False, traces_additive: bool = False, tc_trace: Union[float, torch.Tensor] = 20.0, trace_scale: Union[float, torch.Tensor] = 1.0, sum_input: bool = False, thresh: Union[float, torch.Tensor] = 1.0, **kwargs)

Bases: bindsnet.network.nodes.Nodes

Layer of McCulloch-Pitts neurons.

Instantiates a McCulloch-Pitts layer of neurons.

Parameters

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- traces_additive Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- trace_scale Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **thresh** Spike threshold voltage.

```
forward (x: torch.Tensor) \rightarrow None
Runs a single simulation step.
```

Parameters \mathbf{x} – Inputs to the layer.

- **reset_state_variables** () \rightarrow None Resets relevant state variables.
- $\texttt{set_batch_size}(batch_size) \rightarrow \text{None}$

Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size – Mini-batch size.

Bases: torch.nn.modules.module.Module

Abstract base class for groups of neurons.

Abstract base class constructor.

Parameters

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record decaying spike traces.
- **traces_additive** Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **learning** Whether to be in learning or testing.
- **compute_decays** (dt) \rightarrow None

Abstract base class method for setting decays.

forward (*x: torch.Tensor*) \rightarrow None

Abstract base class method for a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

```
\texttt{reset\_state\_variables()} \rightarrow None
```

Abstract base class method for resetting state variables.

```
set_batch_size (batch_size) \rightarrow None
Sets mini-batch size. Called when layer is added to a network.
```

Parameters batch_size – Mini-batch size.

train (*mode: bool* = True) \rightarrow bindsnet.network.nodes.Nodes Sets the layer in training mode.

Parameters mode (*bool*) – Turn training on or off

Returns self as specified in *torch.nn.Module*

class bindsnet.network.nodes.**SRMONodes**(*n*: Optional[int] = None,shape: Optional[Iterable[int]] = None, traces: bool = *False*, *traces_additive: bool* = *False*, *tc_trace: Union[float, torch.Tensor] = 20.0, trace_scale:* Union[float, torch.Tensor] = 1.0, sum_input: bool = False, thresh: Union[float, torch.Tensor] = -50.0, rest: Union[float, torch.Tensor] = -70.0, reset: Union[float, torch.Tensor] = -70.0, re*frac:* Union[int, torch.Tensor] = 5, tc decay: Union[float, torch.Tensor] = 10.0, lbound: float = None, eps_0: Union[float, torch.Tensor] = 1.0, *rho_0: Union[float, torch.Tensor] = 1.0, d_thresh: Union[float, torch.Tensor] = 5.0, **kwargs)*

Bases: bindsnet.network.nodes.Nodes

Layer of simplified spike response model (SRM0) neurons with stochastic threshold (escape noise). Adapted from (Vasilaki et al., 2009).

Instantiates a layer of SRM0 neurons.

Parameters

- **n** The number of neurons in the layer.
- **shape** The dimensionality of the layer.
- **traces** Whether to record spike traces.
- **traces_additive** Whether to record spike traces additively.
- tc_trace Time constant of spike trace decay.
- **trace_scale** Scaling factor for spike trace.
- **sum_input** Whether to sum all inputs.
- **thresh** Spike threshold voltage.
- **rest** Resting membrane voltage.
- **reset** Post-spike reset voltage.
- **refrac** Refractory (non-firing) period of the neuron.
- tc_decay Time constant of neuron voltage decay.
- **lbound** Lower bound of the voltage.
- eps_0 Scaling factor for pre-synaptic spike contributions.
- **rho_0** Stochastic intensity at threshold.
- **d_thresh** Width of the threshold region.

compute_decays $(dt) \rightarrow$ None Sets the relevant decays.

forward (*x: torch.Tensor*) \rightarrow None Runs a single simulation step.

Parameters \mathbf{x} – Inputs to the layer.

reset_state_variables () \rightarrow None Resets relevant state variables.

set_batch_size ($batch_size$) \rightarrow None Sets mini-batch size. Called when layer is added to a network.

Parameters batch_size - Mini-batch size.

bindsnet.network.topology module

class bindsnet.network.topology.AbstractConnection (source: bindsnet.network.nodes.Nodes, target: bindsnet.network.nodes.Nodes, nu: Union[float, Sequence[float], None] = None, reduction: Optional[callable] = None, weight_decay: float = 0.0, **kwargs)
Bases: abc.ABC,torch.nn.modules.module.Module

Abstract base method for connections between Nodes.

Constructor for abstract base class for connection objects.

Parameters

- **source** A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- **nu** Learning rate for both pre- and post-synaptic events.
- **reduction** Method for reducing parameter updates along the minibatch dimension.
- weight_decay Constant multiple to decay weights by on each iteration.

Keyword arguments:

Parameters

- update_rule (LearningRule) Modifies connection parameters according to some rule.
- wmin (float) The minimum value on the connection weights.
- wmax (float) The maximum value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization.

compute (s: torch.Tensor) \rightarrow None

Compute pre-activations of downstream neurons given spikes of upstream neurons.

Parameters s – Incoming spikes.

```
\begin{array}{l} \textbf{reset\_state\_variables} () \rightarrow None \\ Contains resetting logic for the connection. \end{array}
```

update (**kwargs) \rightarrow None Compute connection's update rule.

Keyword arguments:

Parameters

- **learning** (*bool*) Whether to allow connection updates.
- **mask** (*ByteTensor*) Boolean mask determining which weights to clamp to zero.

```
class bindsnet.network.topology.Connection (source: bindsnet.network.nodes.Nodes, tar-
get: bindsnet.network.nodes.Nodes, nu:
Union[float, Sequence[float], None] = None,
reduction: Optional[callable] = None,
weight_decay: float = 0.0, **kwargs)
```

Bases: bindsnet.network.topology.AbstractConnection

Specifies synapses between one or two populations of neurons.

Instantiates a Connection object.

Parameters

- **source** A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- nu Learning rate for both pre- and post-synaptic events.
- reduction Method for reducing parameter updates along the minibatch dimension.
- weight_decay Constant multiple to decay weights by on each iteration.

Keyword arguments:

Parameters

- update_rule (LearningRule) Modifies connection parameters according to some rule.
- w (torch. Tensor) Strengths of synapses.
- **b** (torch. Tensor) Target population bias.
- wmin (float) Minimum allowed value on the connection weights.
- wmax (float) Maximum allowed value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization constant.

$\texttt{compute} (\textit{s: torch.Tensor}) \rightarrow \texttt{torch.Tensor}$

Compute pre-activations given spikes using connection weights.

Parameters s – Incoming spikes.

Returns Incoming spikes multiplied by synaptic weights (with or without decaying spike activation).

$\texttt{normalize()} \rightarrow None$

Normalize weights so each target neuron has sum of connection weights equal to self.norm.

$\texttt{reset_state_variables()} \rightarrow None$

Contains resetting logic for the connection.

update (**kwargs) \rightarrow None Compute connection's update rule.

class bindsnet.network.topology.Conv2dConnection (source: bindsnet.network.nodes.Nodes, target: bindsnet.network.nodes.Nodes, kernel size: Union[int, Tuple[int, int]], stride: Union[int, Tuple[int, int]] = 1, padding: Union[int, Tuple[int, int]] = 0, dilation: Union[int, Tuple[int, int]] = 1, nu: Union[float, Sequence[float], None] = None, reduction: Optional[callable] =None, weight_decay: float = 0.0, **kwargs) Bases: bindsnet.network.topology.AbstractConnection

Specifies convolutional synapses between one or two populations of neurons.

Instantiates a Conv2dConnection object.

Parameters

- **source** A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- **kernel size** Horizontal and vertical size of convolutional kernels.
- **stride** Horizontal and vertical stride for convolution.
- padding Horizontal and vertical padding for convolution.
- dilation Horizontal and vertical dilation for convolution.
- **nu** Learning rate for both pre- and post-synaptic events.
- **reduction** Method for reducing parameter updates along the minibatch dimension.
- weight_decay Constant multiple to decay weights by on each iteration.

Keyword arguments:

Parameters

- update rule (LearningRule) Modifies connection parameters according to some rule.
- w (torch. Tensor) Strengths of synapses.
- **b** (*torch*. *Tensor*) Target population bias.
- wmin (float) Minimum allowed value on the connection weights.
- wmax (float) Maximum allowed value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization constant.

compute (*s: torch.Tensor*) \rightarrow torch.Tensor

Compute convolutional pre-activations given spikes using layer weights.

Parameters s – Incoming spikes.

Returns Incoming spikes multiplied by synaptic weights (with or without decaying spike activation).

normalize () \rightarrow None

Normalize weights along the first axis according to total weight per target neuron.

$\texttt{reset_state_variables()} \rightarrow None$

Contains resetting logic for the connection.

```
update (**kwargs) \rightarrow None
     Compute connection's update rule.
```

```
bind-
class bindsnet.network.topology.LocalConnection (source:
                                                                snet.network.nodes.Nodes,
                                                                                             target:
                                                                bindsnet.network.nodes.Nodes,
                                                                                                ker-
                                                                nel_size: Union[int, Tuple[int, int]],
                                                                stride: Union[int, Tuple[int, int]],
                                                                n_filters:
                                                                                        Union[float,
                                                                            int, nu:
                                                                Sequence[float], None] = None, re-
                                                                duction: Optional[callable] = None,
                                                                weight decay: float = 0.0, **kwargs)
```

Bases: bindsnet.network.topology.AbstractConnection

Specifies a locally connected connection between one or two populations of neurons.

Instantiates a LocalConnection object. Source population should be two-dimensional.

Neurons in the post-synaptic population are ordered by receptive field; that is, if there are n_{conv} neurons in each post-synaptic patch, then the first n_{conv} neurons in the post-synaptic population correspond to the first receptive field, the second n_{conv} to the second receptive field, and so on.

Parameters

- source A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- kernel_size Horizontal and vertical size of convolutional kernels.
- **stride** Horizontal and vertical stride for convolution.
- n_filters Number of locally connected filters per pre-synaptic region.
- nu Learning rate for both pre- and post-synaptic events.
- **reduction** Method for reducing parameter updates along the minibatch dimension.
- weight_decay Constant multiple to decay weights by on each iteration.

Keyword arguments:

Parameters

- update_rule (LearningRule) Modifies connection parameters according to some rule.
- w (torch. Tensor) Strengths of synapses.
- **b** (torch. Tensor) Target population bias.
- wmin (float) Minimum allowed value on the connection weights.
- wmax (float) Maximum allowed value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization constant.
- int] input_shape (Tuple[int,) Shape of input population if it's not [sqrt, sqrt].

compute (*s: torch.Tensor*) \rightarrow torch.Tensor

Compute pre-activations given spikes using layer weights.

Parameters s – Incoming spikes.

Returns Incoming spikes multiplied by synaptic weights (with or without decaying spike activation).

```
normalize() \rightarrow None
```

Normalize weights so each target neuron has sum of connection weights equal to self.norm.

```
reset_state_variables () \rightarrow None
Contains resetting logic for the connection.
```

update (***kwargs*) \rightarrow None

Compute connection's update rule.

Keyword arguments:

Parameters mask (*ByteTensor*) – Boolean mask determining which weights to clamp to zero.

class bindsnet.network.topology.MaxPool2dConnection(sou	urce:	bind-
Sne	et.network.nodes.Nodes,	
tar	rget:	bind-
sne	et.network.nodes.Nodes,	
ker	rnel_size: Union[int,	Tu-
ple	e[int, int]], stride: Unic	on[int,
Тир	ple[int, int]] = 1, pac	lding:
Un	nion[int, Tuple[int, int]]	= <i>0</i> ,
dile	lation: Union[int, Tup	le[int,
int	[] = 1, **kwargs)	
Bases: bindsnet.network.topology.AbstractConnection		

Specifies max-pooling synapses between one or two populations of neurons by keeping online estimates of maximally firing neurons.

Instantiates a MaxPool2dConnection object.

Parameters

- source A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- **kernel_size** Horizontal and vertical size of convolutional kernels.
- **stride** Horizontal and vertical stride for convolution.
- **padding** Horizontal and vertical padding for convolution.
- dilation Horizontal and vertical dilation for convolution.

Keyword arguments:

Parameters decay – Decay rate of online estimates of average firing activity.

compute (*s: torch.Tensor*) \rightarrow torch.Tensor

Compute max-pool pre-activations given spikes using online firing rate estimates.

Parameters s – Incoming spikes.

Returns Incoming spikes multiplied by synaptic weights (with or without decaying spike activation).

normalize () \rightarrow None No weights -> no normalization.

reset state variables() \rightarrow None

Contains resetting logic for the connection.

update (***kwargs*) \rightarrow None Compute connection's update rule.

class bindsnet.network.topology.MeanFieldConnection (source: bindsnet.network.nodes.Nodes, target: bindsnet.network.nodes.Nodes, nu: Union[float, Sequence[float], None] = None, weight_decay: float = 0.0, **kwargs) Bases: bindsnet.network.topology.AbstractConnection

A connection between one or two populations of neurons which computes a summary of the pre-synaptic population to use as weighted input to the post-synaptic population.

Instantiates a MeanFieldConnection object. :param source: A layer of nodes from which the connection originates. :param target: A layer of nodes to which the connection connects. :param nu: Learning rate for both pre- and post-synaptic events. :param weight_decay: Constant multiple to decay weights by on each iteration. Keyword arguments: :param LearningRule update_rule: Modifies connection parameters according to

some rule.

Parameters

- w (torch. Tensor) Strengths of synapses.
- wmin (float) Minimum allowed value on the connection weights.
- wmax (float) Maximum allowed value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization constant.

compute (*s: torch.Tensor*) \rightarrow torch.Tensor

Compute pre-activations given spikes using layer weights. :param s: Incoming spikes. :return: Incoming spikes multiplied by synaptic weights (with or without

decaying spike activation).

```
\texttt{normalize()} \rightarrow None
```

Normalize weights so each target neuron has sum of connection weights equal to self.norm.

```
\texttt{reset\_state\_variables()} \rightarrow None
```

Contains resetting logic for the connection.

```
update (**kwargs) \rightarrow None
Compute connection's update rule.
```

<pre>lass bindsnet.network.topology.SparseConnection(source:</pre>	bind-
snet.network.nodes.Nodes,	target:
bindsnet.network.nodes.No	des,
nu: Union[float, Seque	nce[float],
None] = None,	reduction:
<i>Optional[callable]</i> =	None,
weight_decay: float	= None,

**kwargs)

Bases: bindsnet.network.topology.AbstractConnection

Specifies sparse synapses between one or two populations of neurons.

Instantiates a Connection object with sparse weights.

Parameters

- source A layer of nodes from which the connection originates.
- target A layer of nodes to which the connection connects.
- nu Learning rate for both pre- and post-synaptic events.
- reduction Method for reducing parameter updates along the minibatch dimension.
- weight_decay Constant multiple to decay weights by on each iteration.

Keyword arguments:

Parameters

- w (torch. Tensor) Strengths of synapses.
- **sparsity** (*float*) Fraction of sparse connections to use.

- **update_rule** (*LearningRule*) Modifies connection parameters according to some rule.
- wmin (float) Minimum allowed value on the connection weights.
- wmax (float) Maximum allowed value on the connection weights.
- **norm** (*float*) Total weight per target neuron normalization constant.

compute (*s: torch.Tensor*) \rightarrow torch.Tensor

Compute convolutional pre-activations given spikes using layer weights.

Parameters s – Incoming spikes.

Returns Incoming spikes multiplied by synaptic weights (with or without decaying spike activation).

 $normalize() \rightarrow None$

Normalize weights along the first axis according to total weight per target neuron.

$\texttt{reset_state_variables}() \rightarrow None$

Contains resetting logic for the connection.

update (***kwargs*) \rightarrow None Compute connection's update rule.

Module contents

4.1.8 bindsnet.pipeline package

Module contents

4.2 Module contents

CHAPTER 5

Indices and tables

- genindex
- search

Python Module Index

b

```
bindsnet,40
bindsnet.analysis,21
bindsnet.analysis.plotting,17
bindsnet.analysis.visualization,21
bindsnet.datasets,23
bindsnet.datasets.preprocess,22
bindsnet.encoding,23
bindsnet.environment,23
bindsnet.evaluation,23
bindsnet.learning,24
bindsnet.network,40
bindsnet.network.monitors,24
bindsnet.network.nodes,25
bindsnet.network.topology,34
bindsnet.pipeline,40
```

Index

Α

AbstractConnection	a (class	in	bind-
snet.network.top	ology), 34		
AbstractInput (class	in bindsnet.ne	etwork.ne	odes), 25
AbstractMonitor	(class	in	bind-
snet.network.mo	nitors), 24		
AdaptiveLIFNodes	(class	in	bind-
snet.network.nod	les), 25		

В

```
bgr2rgb() (in module bindsnet.datasets.preprocess),
       22
binary_image()
                      (in
                              module
                                         bind-
       snet.datasets.preprocess), 22
bindsnet (module), 40
bindsnet.analysis (module), 21
bindsnet.analysis.plotting (module), 17
bindsnet.analysis.visualization (module),
       21
bindsnet.datasets (module), 23
bindsnet.datasets.preprocess(module), 22
bindsnet.encoding (module), 23
bindsnet.environment (module), 23
bindsnet.evaluation (module), 23
bindsnet.learning(module), 24
bindsnet.network (module), 40
bindsnet.network.monitors (module), 24
bindsnet.network.nodes (module), 25
bindsnet.network.topology (module), 34
bindsnet.pipeline (module), 40
BoundingBox (class in bindsnet.datasets.preprocess),
       22
С
```

method), 36

A	compute() (bindsnet.network.topology.LocalConnection
AbstractConnection (class in bind-	method), 37
snet.network.topology), 34	<pre>compute() (bindsnet.network.topology.MaxPool2dConnection</pre>
AbstractInput (class in bindsnet.network.nodes), 25	method), 38
AbstractMonitor (class in bind-	<pre>compute() (bindsnet.network.topology.MeanFieldConnection</pre>
snet.network.monitors), 24	method), 39
AdaptiveLIFNodes (class in bind-	<pre>compute() (bindsnet.network.topology.SparseConnection</pre>
snet.network.nodes), 25	method), 40
D	compute_decays() (bind-
В	snet.network.nodes.AdaptiveLIFNodes
bgr2rgb() (in module bindsnet.datasets.preprocess),	method), 26
22	compute_decays() (bind-
binary_image() (in module bind-	snet.network.nodes.CurrentLIFNodes method),
snet.datasets.preprocess), 22	26
bindsnet (module), 40	compute_decays() (bind-
bindsnet.analysis(module),21	snet.network.nodes.DiehlAndCookNodes
<pre>bindsnet.analysis.plotting(module), 17</pre>	method), 28
bindsnet.analysis.visualization (module),	compute_decays() (bind-
21	snet.network.nodes.LIFNodes method), 31
bindsnet.datasets(module),23	compute_decays() (bindsnet.network.nodes.Nodes
bindsnet.datasets.preprocess(module),22	method), 32
bindsnet.encoding (module), 23	compute_decays() (bind-
bindsnet.environment (module), 23	snet.network.nodes.SRM0Nodes method),
bindsnet.evaluation (module), 23	33 compute output height() (bind-
bindsnet.learning(module),24	
bindsnet.network (module), 40	snet.datasets.preprocess.BoundingBox
bindsnet.network.monitors (module), 24	<pre>method), 22 compute_output_width() (bind-</pre>
bindsnet.network.nodes(module), 25	
bindsnet.network.topology(module), 34	snet.datasets.preprocess.BoundingBox method), 22
bindsnet.pipeline (module), 40	computeCropPadImageLocation() (in module
BoundingBox (class in bindsnet.datasets.preprocess),	bindsnet.datasets.preprocess), 22
22	Connection (<i>class in bindsnet.network.topology</i>), 34
С	Conv2dConnection (class in bind-
<pre>compute() (bindsnet.network.topology.AbstractConnect method), 34</pre>	crop() (in module bindsnet.datasets.preprocess), 22
compute() (bindsnet.network.topology.Connection	
method), 35	snet.datasets.preprocess), 23
compute () (bindsnet.network.topology.Conv2dConnection	
method) 36	snet datasets preprocess) 23

CurrentLIFNodes (*class in bindsnet.network.nodes*), gray_scale() 26 snet.dat

D

DiehlAndCookNodes (class in snet.network.nodes), 27

Ε

edge_spacing_x()	(bind-
snet.datasets.preprocess.BoundingBox	
method), 22	
edge_spacing_y()	(bind-
snet.datasets.preprocess.BoundingBox	
method), 22	

F

forward()(<i>bi</i>	indsnet.network.nodes.AdaptiveLIFNodes
method	<i>d</i>), 26
forward() (b	indsnet.network.nodes.CurrentLIFNodes
method	<i>d</i>), 27
forward()(<i>bi</i>	ndsnet.network.nodes.DiehlAndCookNode
method	<i>l</i>), 28
forward() (b	indsnet.network.nodes.IFNodes method),
28	
forward()(<i>bi</i>	indsnet.network.nodes.Input method), 29
forward() (bindsnet.network.nodes.IzhikevichNodes
method	<i>d</i>), 30
forward()	(bindsnet.network.nodes.LIFNodes
method	<i>d</i>), 31
forward()	(bindsnet.network.nodes.McCullochPitts
method	<i>l</i>), 31
forward() (<i>l</i>	bindsnet.network.nodes.Nodes method),
32	
forward()	(bindsnet.network.nodes.SRM0Nodes
method	<i>1</i>), 33
G	

G

get() (bindsnet.network.monitors.Monitor	method), 24
get() (bindsnet.network.monitors.Network.monitors	vorkMonitor
method), 25	
get_bb_list()	(bind-
snet.datasets.preprocess.Bounding.	Box
method), 22	
get_center_x()	(bind-
snet.datasets.preprocess.Bounding.	Box
method), 22	
get_center_y()	(bind-
snet.datasets.preprocess.Bounding.	Box
method), 22	
get_height()	(bind-
snet.datasets.preprocess.Bounding.	Box
method), 22	
get_width() (bindsnet.datasets.preproce.	ss.BoundingBo
method), 22	-

gray_scale() (in module bindsnet.datasets.preprocess), 23

L

bind-

IFNodes (class in bindsnet.network.nodes), 28
Input (class in bindsnet.network.nodes), 29
IzhikevichNodes (class in bindsnet.network.nodes),
29

L

LIFNodes (class in bindsnet.network.nodes), 30 LocalConnection (class in bindsnet.network.topology), 36

Μ

```
MaxPool2dConnection (class in bind-
snet.network.topology), 37
McCullochPitts (class in bindsnet.network.nodes),
31
MeanFieldConnection (class in bind-
es snet.network.topology), 38
Monitor (class in bindsnet.network.monitors), 24
```

Ν

NetworkMonitor snet.network.mo		in	bind-	
Nodes (class in bindsnet.		es) 31		
•			maation	
normalize() (bindsne method), 35	21.11011011.10	pology.Co	nnection	
normalize() (bindsnet	.network.top	ology.Con	w2dConnectio	on
method), 36				
normalize()(bindsnet	.network.top	ology.Loc	alConnection	
method), 37				
normalize()(bindsnet	.network.top	ology.Max	Pool2dConne	ection
method), 38				
normalize() (bindsnet	.network.top	ology.Med	ınFieldConne	ction
method), 39				
normalize()(bindsnet	.network.top	ology.Spa	rseConnection	п
method), 40				
NormalizeToTensor	(class	in	bind-	
snet.datasets.pre	eprocess), 22			
Ρ				

plot_assignments() (in module bindsnet.analysis.plotting), 17

plot_conv2d_weights() (in module bindsnet.analysis.plotting), 17

plot_performance() (in module bindox snet.analysis.plotting), 19

plot_spike_trains	_for_	example() (in	n mod-				
ule bindsnet.an	ule bindsnet.analysis.visualization), 21						
plot_spikes()	(in	module	bind-				
snet.analysis.pl	otting),	19					
<pre>plot_voltage()</pre>	(in	module	bind-				
snet.analysis.vi	sualizat	tion), 21					
<pre>plot_voltages()</pre>	(in	module	bind-				
snet.analysis.pl	lotting),	20					
plot_weights()	(in	module	bind-				
snet.analysis.pl	lotting),	20					
plot_weights_movi	e()	(in module	bind-				
snet.analysis.vi	sualizat	tion), 21					
print_bb() (bindsnet.	dataset	s.preprocess.Bour	ndingBox				
method), 22		• •					

R

recent	er() (bindsnet.datasets.preprocess.Ba	oundingBox
	method), 22	
record	d() (bindsnet.network.monitor.	s.Monitor
	method), 24	
record	d() (bindsnet.network.monitors.Networ	<i>kMonitor</i>
	method), 25	
Rescal	e (class in bindsnet.datasets.preproces	s), 22
reset_	_state_variables()	(bind-
	snet.network.monitors.Monitor	method),
	24	
reset_	_state_variables()	(bind-
	snet.network.monitors.NetworkMonite	or
	method), 25	
reset_	_state_variables()	(bind-
	snet.network.nodes.AdaptiveLIFNode.	\$
	method), 26	
reset_	_state_variables()	(bind-
	snet.network.nodes.CurrentLIFNodes	method),
	27	
reset_	_state_variables()	(bind-
	snet.network.nodes.DiehlAndCookNod	des
	method), 28	
reset_	_state_variables()	(bind-
	snet.network.nodes.IFNodes method),	
reset_	_state_variables()	(bind-
	snet.network.nodes.Input method), 29	
reset_	_state_variables()	(bind-
	snet.network.nodes.IzhikevichNodes	method),
	30	(1 • 1
reset_	_state_variables()	(bind-
	snet.network.nodes.LIFNodes method	
reset_	_state_variables()	(bind-
	snet.network.nodes.McCullochPitts	method),
	31	(1 + 1
reset_	_state_variables()	(bind-
	snet.network.nodes.Nodes method), 32	
reset_	_state_variables()	(bind-
	snet.network.nodes.SRM0Nodes	method),

33

	55	
	reset_state_variables()	(bind-
	snet.network.topology.AbstractConnectio	n
	method), 34	
	reset_state_variables()	(bind-
	snet.network.topology.Connection me	ethod),
	35	
	reset_state_variables()	(bind-
	snet.network.topology.Conv2dConnection	1
	method), 36	
	reset_state_variables()	(bind-
	snet.network.topology.LocalConnection	
ĸ	method), 37	
	reset_state_variables()	(bind-
	snet.network.topology.MaxPool2dConnet	ction
	method), 38	
r	reset_state_variables()	(bind-
•	snet.network.topology.MeanFieldConnec	tion
	method), 39	
	reset_state_variables()	(bind-
	snet.network.topology.SparseConnection	
	method), 40	

S

sample	_exp_	_two_	_sides()	(in	module	bind
	snet.a	lataset	s.preprocess), 23		

- sample_rand_uniform() (in module bindsnet.datasets.preprocess), 23

- set_batch_size() (bindsnet.network.nodes.AdaptiveLIFNodes method), 26
- set_batch_size() (bindsnet.network.nodes.CurrentLIFNodes method), 27
- set_batch_size() (bindsnet.network.nodes.DiehlAndCookNodes method), 28
- set_batch_size() (bindsnet.network.nodes.IFNodes method), 29
- set_batch_size() (bindsnet.network.nodes.IzhikevichNodes method), 30
- set_batch_size() (bindsnet.network.nodes.LIFNodes method), 31 set_batch_size() (bind-
- snet.network.nodes.McCullochPitts method), 31

<pre>set_batch_size()</pre>					(bind-
snet.network.nodes.SRM0Nodes				5	method),
33	5				
<pre>shift() (bindsnet.datasets.preprocess.BoundingBox</pre>					
method), 22					
shift_cr	op_train	ing_s	sample()	(in	module
bindsnet.datasets.preprocess), 23					
SparseConnection (class				in	bind-
snet.network.topology), 39					
SRM0Nodes (class in bindsnet.network.nodes), 32					
subsampl	e()	(in	modu	ıle	bind-
snet.datasets.preprocess), 23					

Т

train() (bindsnet.network.nodes.Nodes method), 32

U

- update() (bindsnet.network.topology.Connection method), 35
- update() (bindsnet.network.topology.Conv2dConnection method), 36

- update() (bindsnet.network.topology.SparseConnection method), 40