

#####

Installation

#####

1. Installing PyNN:

\$ pip install pyNN

Or follow the instructions in

<http://neuralensemble.org/docs/PyNN/installation.html>

2. Compile and install PyCARL

2.1 Clone the PyCARL repository to the CARLsim root.

\$ cd CARLsim4/

\$ git clone https://github.com/adarshabalaji/PyCarlsim.git

2.2 Compile and generate the pyNN -> carlsim interface file (carlsim.py)

2.2.1 Install SWIG:

\$ sudo apt update

\$ sudo apt install swig

2.2.2 Compile and link the interface file (carlsim_wrap.cxx), generated by swig, with the libcarlsim.a library.

\$ cd source

\$ source build.sh

This creates a static library `_carlsim.so` and a pyNN -> CARLsim interface (carlsim.py)

****OPTIONAL**** If you want to compile and link the interface file manually, then

3. 1 Compile the carlsim.i (interface file) using SWIG

```
$ cd source
```

```
$ swig -c++ -python carlsim.i
```

The output of the swig build is a wrapper file (carlsim_wrap.cxx) and the pyNN -> carlsim interface file (carlsim.py).

3. 2 In the source folder, follow the steps below to compile a new pyNN interface (carlsim.i and carlsim_wrap.cxx) with the static library libcarsim.a (generated during a CARLsim build).

```
$gcc -fPIC -c carlsim_wrap.cxx -I/usr/include/python<version> -I<CARLsim4 include dir> -  
I/usr/local/cuda/include -I/usr/local/cuda/samples/common/inc -D__NO_CUDA__
```

Update the above command with the python version of choice and the CARLsim4 installation directory.

3.1 Link the carlsim_wrap.o file to the libcarsim.a file.

```
$ g++ -shared carlsim_wrap.o -o _carsim.so
```

```
$ g++ -shared ~/CARL/lib/libcarsim.a carlsim_wrap.o -o _carsim.so
```

4. Copy the compiled sources and the carlsim/ folder in the pyCARL repo to pyNN.

```
$ cp -r CARLsim4/pyCARL/carlsim <root of pyNN Installation>
```

4.1 Copy the generated _carsim.so and carlsim.py file to <root of pyNN Installation>/carlsim

PyCARL is now integrated with pyNN.

#####

List of APIs implemented.

#####

1. Create spike generator group.

```
#####
```

```
# Define the cell type for all the input groups
```

```
inputCellType = sim.SpikeSourceArray("input", <numNeurons>, "EXCITATORY_NEURON", "CUBA")
```

```
# create a population of <numNeurons> neurons of cellType inputCellType.
```

```
spike_source = sim.Population(<numNeurons>, inputCellType, label='input')
```

```
#####
```

```
#2. Create neuron groups.
```

```
pyCARL only supports Izhikevich neurons
```

```
#####
```

```
#
```

```
# Define the inzhikevich cell type for the group
```

```
izhikevichCellType = sim.Izhikevich("EXCITATORY_NEURON", a=0.02, b=0.2, c=-65, d=8)
```

```
# create a pyNN population of <numNeurons> neurons of cellType izhikevichCellType
```

```
gExc = sim.Population(<numNeurons>, izhikevichCellType, label='Exc')
```

```
#####
```

3. Setup Network

An equivalent for this function does not exist for pyNN.

Therefore the user will have to call this CARLsim function directly in pyNN.

```
#####
```

```
#sim.state.setupNetwork()
```

```
#####
```

4. Run Network

Input - runtime in Ms

```
#####
```

```
sim.run(<time in Ms>)
```

```
#####
```

5. STDP Mechanism for a group

Input - runtime in Ms

```
#####
```

define the properties of the STDP synapse.

```
stdp_model = sim.STDPMechanism(
```

```
    timing_dependence=sim.SpikePairRule(tau_plus=20.0, tau_minus=20.0, A_plus=0.01,  
    A_minus=0.012),
```

```
    weight_dependence=sim.AdditiveWeightDependence(w_min=0, w_max=0.0000001),
```

```
    weight=0.00000005,
```

```
    delay=delay,
```

```
    dendritic_delay_fraction=float(options.dendritic_delay_fraction))
```

connect the groups G1 and G2 using an all-to-all connector, with stdp enabled synapses.

```
connections = sim.Projection(G1, G2, sim.AllToAllConnector(), stdp_model)
```

#####

Connect function

```
{'OneToOneConnector': 'one-to-one', 'AllToAllConnector': "full", "FixedProbabilityConnector": "random",  
"FromListConnector": "FromListConnector"}
```

PyCARL only supports 'full', 'one-to-one' and 'random' connectors

The corresponding

#####

#

```
connectionID = sim.Projection(spike_source, neuron_group1, sim.OneToOneConnector(), synapse_type,  
receptor_type='excitatory')
```