

Simulator-independent model specification ("Meta-simulators")

Simulator-independent environments for developing neuroscience models:

- keep the advantages of having multiple simulators
- but remove the translation barrier.

Three (complementary) approaches:

- GUI (e.g. neuroConstruct)
- XML-based language (e.g. NeuroML)
- interpreted language (e.g. Python)





A common scripting language for neuroscience simulators

Goal

Write the code for a model simulation *once*, run it on any supported simulator* without modification.

* or hardware device



A common scripting language for neuroscience simulators

Simulator Language

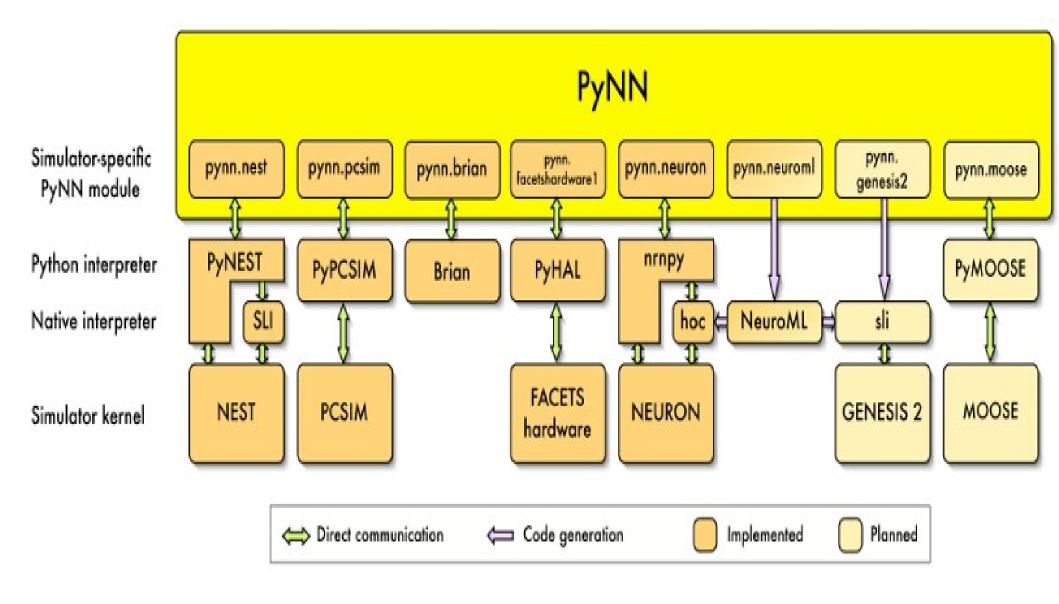
PCSIM C++ or Python
MOOSE SLI or Python
MVASpike C++ or Python
NEST sli or Python
NEURON hoc or Python

SPLIT C++ (Python interface planned)

Brian Python FACETS hardware Python



PyNN Approach: A standardized API



Selecting the simulator

```
from pyNN.neuron import *
from pyNN.nest1 import *
from pyNN.nest2 import *
from pyNN.pcsim import *
from pyNN.moose import *
from pyNN.brian import *
import pyNN.neuron as sim
```



Overview: PyNN low-level API

setup() and end()



create()

```
create(IF_curr_alpha)
create(IF_curr_alpha, n=10)
create(IF_curr_alpha, {'tau_m': 15.0, 'cm': 0.9}, n=10)
>>> IF_curr_alpha.default_parameters
{'tau_refrac': 0.0, 'tau_m': 20.0, 'i_offset': 0.0,
 'cm': 1.0, 'v_init': -65.0, 'v_thresh': -50.0,
 'tau_syn_E': 0.5, 'v_rest': -65.0, 'tau_syn_I': 0.5,
 'v_reset': -65.0}
```



create()

```
>>> create(IF_curr_alpha, param_dict='foo': 15.0)
    Traceback (most recent call last):
    NonExistentParameterError: foo
    >>> create(IF_curr_alpha, param_dict='tau_m': 'bar')
    Traceback (most recent call last):
    InvalidParameterValueError:
        (<type 'str'>, should be <type 'float'>)
```



create()

```
create(IF_curr_alpha, 'v_thresh': -50, 'cm': 0.9)
create('iaf_neuron', 'V_th': -50, 'C_m': 900.0)
```



Standard cell models

```
IF_curr_alpha
IF_curr_exp
IF_cond_alpha
IF_cond_exp,
IF_cond_exp_gsfa_grr
IF_facets_hardware1
HH_cond_exp,
EIF_cond_alpha_isfa_ista
SpikeSourcePoisson
SpikeSourceInhGamma
SpikeSourceArray
```



Standard cell models

Example: Leaky integrate-and-fire model with fixed firing threshold, and current-based, alpha-function synapses.

Name	Units	NEST	NEURON
v_rest	mV	UO	v_rest
v_reset	mV	Vreset	v_reset
cm	nF	C†	CM
tau_m	ms	Tau	tau_m
tau_refrac	ms	TauR	t_refrac
tau_syn	ms	TauSyn	tau_syn
$v_{\mathtt{-}}thresh$	mV	Theta	v_{-} thresh
i_offset	nA	IO†	i_offset

 $\dagger Unit \ differences: \ C \ is \ in \ pF, \ IO \ in \ pA.$



ID objects

```
>>> my_cell = create(IF_cond_exp)
>>> print my_cell
1
>>> type(my_cell)
<class 'pyNN.nest2.ID'>
>>> my_cell.tau_m
20.0
>>> my_cell.position
(1.0, 0.0, 0.0)
>>> my_cell.position = (0.76, 0.54, 0.32)
```



connect()

p=0.2, synapse_type='inhibitory')



record()

```
record(cell, "spikes.dat")
record_v(cell_list, "Vm.dat")
```

Writing occurs on end()



run()

run(100.0)



Simulation status

```
get_current_time()
get_time_step()
get_min_delay()
num_processes()
rank()
```



Random numbers

```
>>> from pyNN.random import NumpyRNG, GSLRNG, NativeRNG
>>> rng = NumpyRNG(seed=12345)
>>> rng.next()
0.6754034
>>> rng.next(3, 'uniform', (-70,-65))
[-67.4326, -69.9223, -65.4566]
```

- Use NativeRNG or GSLRNG to ensure different simulators get the same random numbers
- Use NativeRNG to use a simulator's built-in RNG





Random numbers

```
>>> from pyNN.random import RandomDistribution
>>> distr = RandomDistribution('uniform', (-70, -65),
... rng=rng)
>>> distr.next(3)
[-67.4326, -69.9223, -65.4566]
```



Overview: PyNN high-level API

Populations

```
p1 = Population((10,10), IF_curr_exp)
p2 = Population(100, SpikeSourceArray,
                label="Input Population")
p3 = Population(dims=(3,4,5), cellclass=IF_cond_alpha,
                cellparams={'v_thresh': -55.0},
                label="Column 1")
p4 = Population(20, 'iaf_neuron', {'Tau': 15.0,
                                    'C': 100.0})
```



Populations

Accessing individual members

```
>>> p1[0,0]
    >>> p1[9,9]
    100
    >>> p3[2,1,0]
    246
    >>> p3.locate(246)
    (2, 1, 0)
>>> p1.index(99)
100
>>> p1[0,0].tau_m = 12.3
```



Populations

Iterators

```
>>> for id in p1:
    ... print id, id.tau_m
    0 12.3
    1 20.0
    2 20.0
    >>> for addr in p1.addresses():
    ... print addr
    (0, 0)
    (0, 1)
    (0, 2)
    (0, 9)
```



set(), tset(), rset()

```
>>> p1.set("tau_m", 20.0)
>>> p1.set('tau_m':20, 'v_rest':-65)
>>> distr = RandomDistribution('uniform', [-70,-55])
>>> p1.rset('v_init', distr)
>>> import numpy
>>> current_input = numpy.zeros(p1.dim)
>>> current_input[:,0] = 0.1
>>> p1.tset('i_offset', current_input)
```



Recording

```
# record from all neurons in the population
>>> p1.record()
# record from 10 neurons chosen at random
>>> p1.record(10)
# record from specific neurons
>>> p1.record([p1[0,0], p1[0,1], p1[0,2]])
>>> p1.printSpikes("spikefile.dat")
>>> p1.getSpikes()
array([])
```



Position in space

```
\Rightarrow p1[1,0].position = (0.0, 0.1, 0.2)
>>> p1[1,0].position
array([ 0. , 0.1, 0.2])
>>> p1.positions
array([[...]])
>>> p1.nearest((4.5, 7.8, 3.3))
48
>>> p1[p1.locate(48)].position
array([ 4., 8., 0.])
```



Projections



Connectors

AllToAllConnector

OneToOneConnector

FixedProbabilityConnector

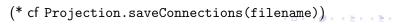
 ${\tt Distance Dependent Probability Connector}$

 ${\tt Fixed Number Post Connector}$

 ${\tt Fixed Number Post Connector}$

FromFileConnector*

FromListConnector





Connectors

```
c = DistanceDependentProbabilityConnector(
    "exp(-abs(d))",
    axes='xy',
    periodic_boundaries=(500, 500, 0),
    weights=0.7,
    delays=RandomDistribution('gamma', [1,0.1])
)
```



Weights and delays

```
>>> prj1_1.setWeights(0.2)
>>> weight_list = 0.1*numpy.ones(len(prj2_1))
>>> weight_list[0:5] = 0.2
>>> prj2_1.setWeights(weight_list)
>>> prj1_1.randomizeWeights(weight_distr)
>>> prj1_2.setDelays('exp(-d/50.0)+0.1')
```

[Note: synaptic weights are in nA for current-based synapses and μS for conductance-based synapses]



Weights and delays

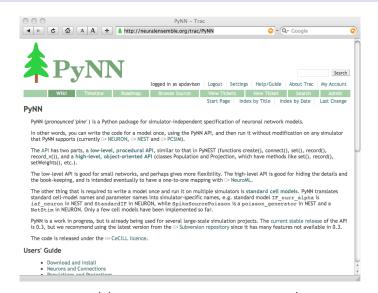
```
w_array = prj.getWeights()
prj.printWeights(filename)
```

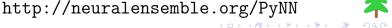


Synaptic plasticity

```
# Facilitating/depressing synapses
depressing_syn = SynapseDynamics(
                     fast=TsodyksMarkramMechanism(**params))
prj = Projection(pre, post, AllToAllConnector(),
                 synapse_dynamics=depressing_syn)
# STDP
stdp_model = STDPMechanism(
               timing_dependence=SpikePairRule(
                                   tau_plus=20.0,
                                    tau minus=20.0).
               weight_dependence=AdditiveWeightDependence(
                                    w_{min=0}, w_{max=0.02},
                                    A_plus=0.01, A_minus=0.012)
prj2 = Projection(pre, post, FixedProbabilityConnector(p=0.1),
                  synapse_dynamics=SynapseDynamics(slow=stdp_model))
```

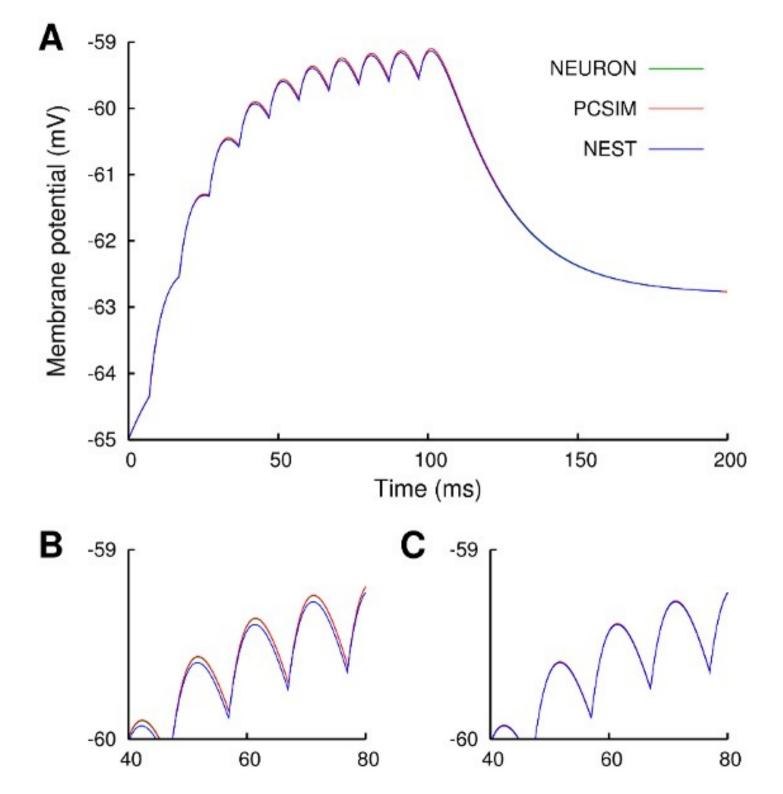
How to participate in PyNN development

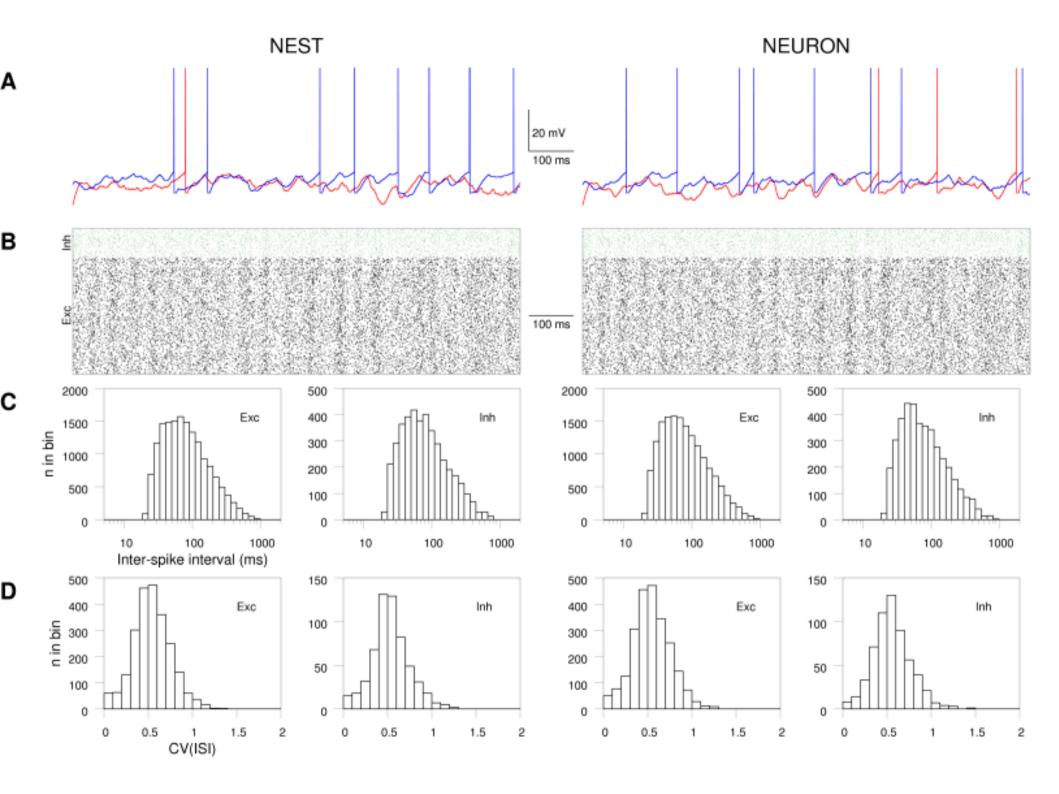




Why might PyNN be important for this INCF initiative?

- Declarative XML
 - Not generally human generatable or even readable
 - PyNN is human generatable and can generate NeuroML
- Standards compliance verification
 - A test-suite for compliance
 - Write once in PyNN for all
 - No testing of the various test suites, no bash|perl





- Implement XML importer/parser for PyNN
 - Writing and maintaining Python+PyNN support for a given simulator is orders of magnitude more appealing than writing and maintaining a compliant XML parser. PyNN can provide this.
- PyNN API user extensions in terms of the API
 - Such extensions supported by all simulators implementing the API
 - Maintain existing isomorphism between PyNN API Object Model and NeuroML for I&F networks
 - Avoid re-inventing a general modeling language in XML with associated difficulties to parse and support downstream.

- 1 API (PyNN low-level)
 - create, connect
- 2 Object model (PyNN high-level)
 - Population, Projection
- 3 Object Declarative Mapping (à la ORM for DBs)
 - PyNN <=> NeuroML

Main message:

- Unification of simulator object models (hard)
- Declarative model format through ODM (easy)