CNS Tutorial T3

Modelling of spiking neural networks with the Brian simulator

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Schedule for today

Morning

Time	Торіс			
9.00 - 9.30	Introduction & installation			
9.30 - 10.10	Core concepts of Brian2			
10.10 - 10.30	COFFEE BREAK			
10.40 - 12.00	Hands-on tutorial			

Afternoon

13.30 - 13.45	Going from Brian 1 to Brian 2			
13.45 – 15.10	Advanced Brian 2 + Extending Brian 2			
15.10 - 15.40	COFFEE BREAK			
15.40 - 16.30	Question & answers			

The spirit of

"A simulator should not only save the time of processors, but also the time of scientists"



scientist



computer

Writing code often takes more time than running it

Goals:

- Quick model coding
- Flexible

models are defined by equations (rather than pre-defined)

Goodman, D. and R. Brette (2009). The Brian simulator. Front Neurosci, doi:10.3389/neuro.01.026.2009.

An example

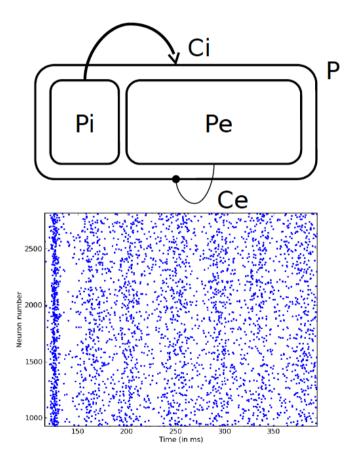
```
from brian2 import *
```

```
tau m = 20*ms
                      Stimberg M, Goodman DFM, Benichoux V, Brette R
tau e = 5*ms
tau i = 10*ms
                      (2014). Equation-oriented specification of neural models for
V_th = -50*mV
                      simulations. Frontiers Neuroinf. doi: 10.3389/fninf.2014.00006.
E L = -60*mV
I_const = 11*mV
egs = '''
dv/dt = (-(v-E_L) + I_{const} + g_e + g_i)/tau_m : volt
dg e/dt = -g e/tau e : volt
dg i/dt = -g i/tau i : volt
P = NeuronGroup(4000, model=eqs,
                threshold='v>V th', reset='v=E L')
P.v = 'E L+10*mV*rand()'
Pe = P[:3200]
Pi = P[3200:]
w e = 1.62*mV
w i = 9*mV
Ce = Synapses(Pe, P, pre='g_e+=w_e')
Ci = Synapses(Pi, P, pre='g i-=w i')
Ce.connect(True, p=0.02)
Ci.connect(True, p=0.02)
M = SpikeMonitor(P)
run(1*second)
plot(M.t/ms, M.i, '.')
xlabel('Time (in ms)'); ylabel('Neuron number')
show()
```

$$\tau_m \frac{\mathrm{d}V}{\mathrm{d}t} = -(V - E_L) + g_e + g_i$$

$$\tau_e \frac{\mathrm{d}g_e}{\mathrm{d}t} = -g_e$$

$$\tau_i \frac{\mathrm{d}g_i}{\mathrm{d}t} = -g_i$$



Standardization issues

Key issue: each simulator has its own language, how to communicate models?

Example 1: PyNEST (Python interface to NEST)

Component-based approach:

you need to know what the components are exactly you need to know parameter names you need to know implicit units

Standardization issues

Solution: equation-oriented approach

Example 2: NineML (Izhikevich model)

Issues:

- can't specify a full simulation (including initialization & stimulus)
- heavy!

```
<?xml version='1.0' encoding='UTF-8'?>
<NineML xmlns="http://nineml.org/9ML/0.1">
   xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
   xsi:schemaLocation="http://nineml.org/9ML/0.1 NineML\_v0.2.xsd">
  <ComponentClass name="izhikevichCellNew">
    <Parameter name="a" dimension="none"/>
    <Parameter name="c" dimension="voltage"/>
    <Parameter name="b" dimension="per_time"/>
    <Parameter name="d" dimension="voltage per time"/>
    <Parameter name="theta" dimension="voltage"/>
    <AnalogPort name="iSvn" mode="reduce" reduce op="+" dimension="current"/>
    <AnalogPort name="U" mode="send" dimension="none"/>
    <AnalogPort name="V" mode="send" dimension="voltage"/>
    <EventPort name="spikeOutput" mode="send"/>
    <Dynamics>
        <StateVariable name="V" dimension="voltage"/>
        <StateVariable name="U" dimension="voltage_per_time"/>
        <Alias name="rv" dimension="none">
            <MathInline>V*U</MathInline>
        </Alias>
        <Regime name="subthresholdRegime">
          <TimeDerivative variable="U">
           <MathInline>a*(b*V - U)</MathInline>
          </TimeDerivative>
          <TimeDerivative variable="V">
            <MathInline>0.04*V*V + 5*V + 140.0 - U + iSyn</mathInline>
          </TimeDerivative>
          <OnCondition>
             <MathInline>V \> theta </MathInline>
            </Trigger>
            <StateAssignment variable="V" >
             <MathInline>c</MathInline>
            </StateAssignment>
            <StateAssignment variable="U" >
             <MathInline>U+d</MathInline>
            </StateAssignment>
           <EventOut port="spikeOutput" />
          </OnCondition>
        </Regime>
    </Dynamics>
  </ComponentClass>
</NineML>
```

Standardization issues: the Richard way



Goal: to minimize «language entropy» = uncertainty about syntax and names, given the meaning

Brette R (2012). On the design of script languages for neural simulation. Network 23(4), 150-156

Key points:

- 1) There is already an accepted standard for models: math!
- 2) If you specify names, units and equations yourself, you don't need to know them in advance
- 3) Full expressivity requires a programming language

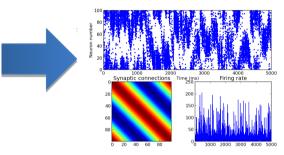
The future of



37772

relies on code generation to run on multiple targets

Simulation on PC, clusters, GPU







Interface with robots



Embedded simulation on Android smartphones





Website

briansimulator.org



The Brian spiking neural network simulator

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Brian tutorial at CNS 2015

17 Jul 2015

Google Summer of Code 2015

5 Mar 2015

Brian tutorial at CNS 2014

8 Jul 2014

New Brian paper: Equation-oriented specification of neural models for simulations

About

* Brian 2.0 fourth beta release: release notes *

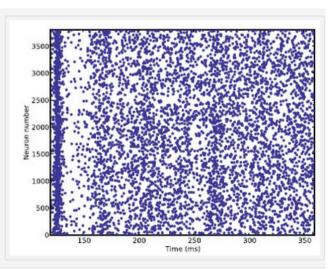
* Brian tutorial at the CNS 2015 meeting in Prague: more information *

Brian is a simulator for spiking neural networks available on almost all platforms. The motivation for this project is that a simulator should not only save the time of processors, but also the time of scientists.

Brian is easy to learn and use, highly flexible and easily extensible. The Brian package itself and simulations using it are all written in the Python programming language, which is an easy, concise and highly developed language with many advanced features and development tools, excellent documentation and a large community of users providing support and extension packages.

The following code defines a randomly connected network of integrate and fire neurons with exponential inhibitory and excitatory currents, runs the simulation and makes the raster plot on the right.

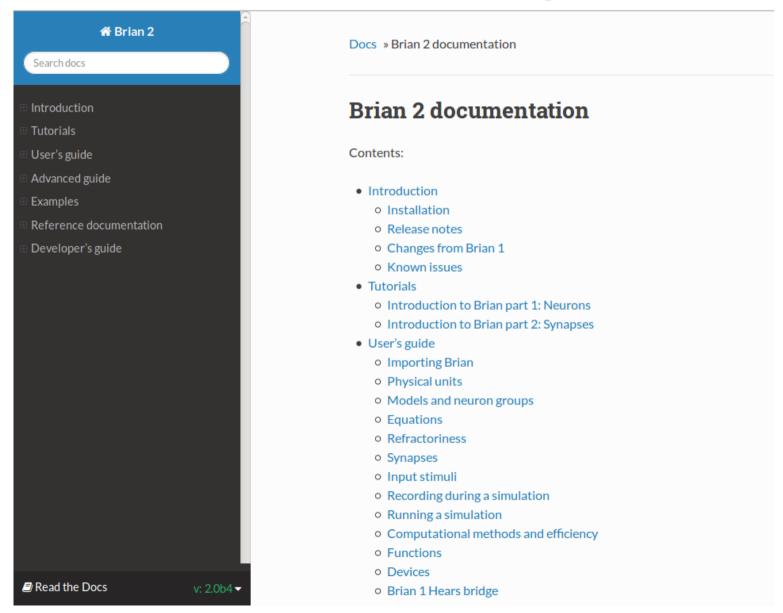
```
from brian2 import *
eas = '''
dv/dt = (qe+qi-(v+49*mV))/(20*ms) : volt
dqe/dt = -qe/(5*ms) : volt
dqi/dt = -qi/(10*ms) : volt
P = NeuronGroup(4000, eqs, threshold='v>-50*mV', reset='v=-60*mV')
P.v = -60*mV
Pe = P[:3200]
Pi = P[3200:1]
Ce = Synapses(Pe, P, pre='ge+=1.62*mV')
Ce.connect(True, p=0.02)
Ci = Synapses(Pi, P, pre='qi-=9*mV')
Ci.connect(True, p=0.02)
M = SpikeMonitor(P)
run(1*second)
plot(M.t/ms, M.i, '.')
show()
```



See the manuals for more examples.

Documentation

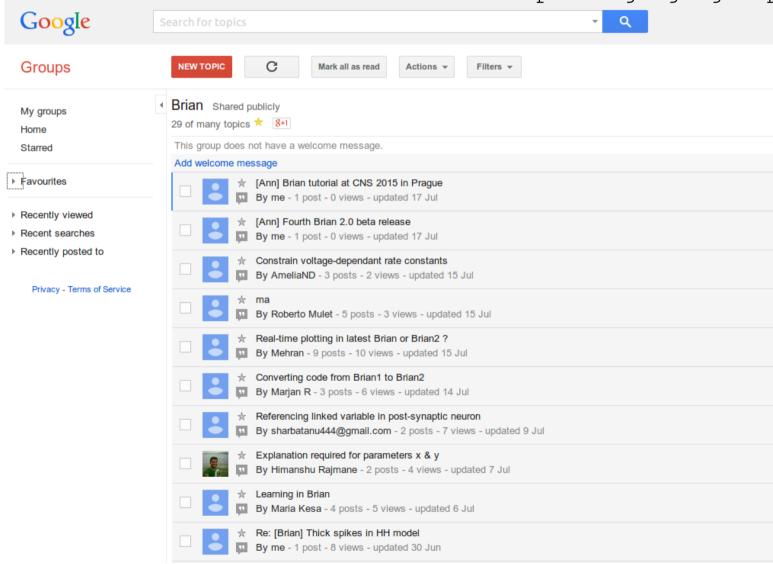
brian2.readthedocs.org



Mailing lists

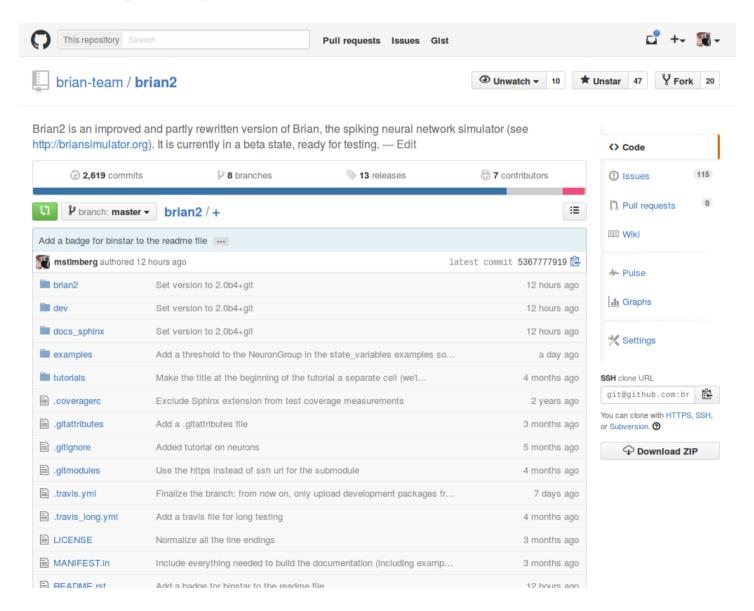
briansupport@googlegroups.com

brian-development@googlegroups.com



Code repository

https://github.com/brian-team/brian2



Installation

Recommended way:

- Use Anaconda distribution
- Add the "brian-team" channel
 conda config --add channels brian-team
- Install brian2conda install brian2

More infos and alternative installation: **brian2.readthedocs.org**

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Core concepts of



Anatomy of a Brian script

```
from brian2 import *
                                                                   Import Brian library
tau m = 20*ms
                                                                   Set constants
tau e = 5*ms
tau i = 10*ms
V th = -50*mV
E L = -60*mV
I const = 11*mV
egs = '''
dv/dt = (-(v-E_L) + I_{const} + g_e + g_i)/tau_m : volt
dq e/dt = -g e/tau_e : volt
                                                                   Specify the neuronal model
dg_i/dt = -g_i/tau_i : volt
P = NeuronGroup(4000, model=eqs,
               threshold='v>V th', reset='v=E L')
                                                                   Initialise state variables
P.v = 'E L+10*mV*rand()'
Pe = P[:3200]
Pi = P[3200:]
                                                                   Specify subgroups of
                                                                   neurons
w e = 1.62*mV
w i = 9*mV
                                                                   Specify synaptic model
Ce = Synapses(Pe, P, pre='g e+=w e')
Ci = Synapses(Pi, P, pre='g i-=w i')
                                                                   Connect neurons
Ce.connect(True, p=0.02)
Ci.connect(True, p=0.02)
                                                                   Record activity
M = SpikeMonitor(P)
                                                                   Run the simulation
run(1*second)
plot(M.t/ms, M.i, '.')
                                                                   Plot the activity
xlabel('Time (in ms)'); ylabel('Neuron number') 
show()
```

Constants

 Constants defined outside of strings can be used inside strings (only scalar values):

```
tau = 10*ms
G = NeuronGroup(1, 'dv/dt = -v / tau : volt')
```

Values of constants are resolved at the run call:

Variables and equations

Equations define state variables of an object:

Additional variables are defined automatically

Variables and equations

A Hodgkin-Huxley equations

B Noisy membrane

```
G = NeuronGroup(number_of_neurons,

'dv/dt = -(v-v_0)/tau_m +

tau_m**-0.5*3*mV*xi : volt # membrane potential')
```

C Leaky integrate-and-fire neuron

D Leaky integrate-and-fire neuron with adaptive threshold

special variable provided for noise

Expressions

- Can be used to
 - Specify conditions (threshold, refractoriness, synaptic connection)

Index and set state variables

```
min_freq = 100*Hz
print G.v['freq > min_freq']
G.v = '0*mV + rand()*10*mV'
```

• Can refer to state variables, constants, units

Defining synaptic connections with string expressions

Full connectivity:

S.connect('True') Condition for a connectivity:

S.connect('i == j')

Convergent connectivity:

S.connect('(i/N) == j')

Ring structure, connecting to the immediate neighbours:

S.connect('abs((i - j + N/2)%N - N/2) == 1')

Connections to 2d neighbourhood:

S.connect('sqrt((x_pre-x_post)**2+(y_pre-y_post)**2) < 250*umeter')

Sparse random connectivity without self-connections:

Probability for a connection

S.connect('i !=j', p=0.1)

Random connectivity to a 2d neighbourhood without self-connections:

S.connect('i != j', p='p_max*exp(-(x_pre-x_post)**2+(y_pre-y_post)**2) / (2*(125*umeter)**2)'

One-to-one connectivity with two synapses per connection:

S.connect('i == j', n=2)

Number of synapses per connection

Abstract code statements

• Event-triggered operations (reset, synaptic event) are specified as *abstract code*:

```
G = NeuronGroup(..., reset='v = E_L')
S = Synapses(..., pre='v+=w')
G.run_regularly('stim = rand()')
```

- Again: can refer to state variables, constant, units
- Only assignments (and "+=" etc.) allowed
- Automatically interpreted as referring to all "relevant" elements of a group (neurons that spiked for reset, synapses that received a pre-synaptic spike for "pre", all neurons/synapses for custom operations)

Functions

- Abstract code ≠ Python code
- Functions have to be explicitly supported
- Built-in functions:
 - Random numbers: rand(), randn()
 - Elementary functions: sqrt, exp, log, log10, abs
 - Trigonometric functions: sin, cos, tan, sinh, cosh, tanh, arcsin, arccos, arctan
 - General utility functions: clip, floor, ceil, sign
 - Boolean → integer: int
- Support for other functions can be added (afternoon session)

Brian's unit system

Brian allows to use units for scalars and vectors

```
>>> E_L = -70*mV

>>> print E_L

-70.0 mV

>>> freqs = [100, 200, 300] * Hz

>>> print freqs

[ 100. 200. 300.] Hz
```

 Most numpy functions work correctly with units (make sure to not import from numpy directly)

```
>>> mean(freqs)
200.0 * hertz
>>> diff(freqs)
array([ 100., 100.]) * hertz
```

Brian's unit system

 To remove units, use numpy.asarray or divide by the unit

```
>>> print freqs/Hz
[ 100. 200. 300.]
>>> print asarray(freqs)
[ 100. 200. 300.]
```

For state variables: adding an underscore returns unitless value

```
>>> print G.v
<neurongroup.v: array([-70., -70., -70., -70., -70.]) * mvolt>
>>> print G.v_
<neurongroup_1.v: array([-0.07, -0.07, -0.07, -0.07, -0.07])>
```

 Unit consistency is also checked in equations, expressions and abstract code statements

Running simulations: "magic"

 "Magic" network – Brian collects all the object it "sees":

```
G = NeuronGroup(...)
S = Synapses(...)
mon = SpikeMonitor(...)
run(runtime) # G, S and mon
```

Running simulations: explicit

 Explicitly constructed network, recommended for complicated setups:

```
G = NeuronGroup(...)
S = Synapses(...)
monitors = [SpikeMonitor(...), StateMonitor(...)]
net = Network(G, S, monitors)
net.run(runtime)
```

Coffee break!



Topographical connections in Brian

Topographical connections in Brian

