$egin{aligned} extbf{Circuit-Tool}: & A & Tool for generating \\ Neural & Micro Circuits \end{aligned}$

Version 1.0

User Manual

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1 Preliminaries

1.1 What is Circuit-Tool?

Circuit-Tool is a set of Matlab objects and scripts that allow the construction of multi-"column" neural microcircuits with a distribution of parameters that match those reported in the literature. This neural microcircuit models can then be simulated efficiently using CSIM .

1.2 About this Manual

This manual is intended to describe how to use Circuit-Tool from the (Matlab) users point of view. It does not try to explain (or give an introduction to) the type of models which can be constructed an simulated with Circuit-Tool . Regarding neural modeling we refer the

reader to [Dayan and Abbott, 2001] and [Gerstner and Kistler, 2002]. Furthermore Matlab programming knowledge is assumed.

1.3 Features of the current version

Customizeable intra and inter "column" connectivity

Runs under Unix (Linux) and Windows

Object oriented design

Parallel simulation for large set of stimuli

1.4 Getting and Installing Circuit-Tool

Circuit-Tool is distributed under the GNU General Public License and can be downloaded from http://www.igi.tugraz.at/circuits.

To install Circuit-Tool perform the following steps:

- 1. Donwload Circuit-Tool from www.igi.tugraz.at/circuits
- 2. Unzip the file circuits-VER.zip where VER stands for the version you have downloaded. This will create a subdirectory 1sm and 1sm/circuits
- 3. Start Matlab and change into the directory 1sm
- 4. Run the Matlab script install.m.
- 5. Add the path 1sm to the Matlab search path; e. g.
 - addpath('/home/jack/lsm')) or
 - addpath('C:\Work\Neuroscience\lsm').
- 6. Change into the directory lsm/circuits/demos and play around with them.
- 7. Have fun using Circuit-Tool!

2 A short Tutorial

In this section we will introduce Circuit-Tool by means of an example. We will use Circuit-Tool to construct a model which consists of three "columns" or *pools* how they are called in Circuit-Tool. The first pool consists of leaky integrate an fire neurons, the second is made up of sigmoidal neurons and the third uses more detailed conductance based model neurons. The pools will be connected internally and to each other (for details see below). The circuit will be driven by two input spike trains and an analog input current. The input projects in some kind of topographic map into the circuit.

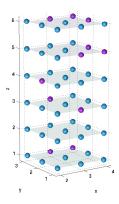


Figure 1: A $3 \times 3 \times 10$ pool of LifNeurons with origin (2,1,1)

2.1 Initializing the model

The Circuit-Tool -implementation of this model employs the Matlab class neural_microcircuit (Sec. 6). To start our construction of the model we will instantiate an "empty" microcircuit:

```
>> nmc = neural_microcircuit;
```

2.2 Creating the Pools

As the next step we will create the individual pools. First we will create a pool of integrateand-fire neurons:

```
>>[nmc,p_lif]=add(nmc,'pool','type','LifNeuron','size',[3 3 6],'origin',[2 1 1]);
```

The above command creates 54 neurons of the class LifNeuron (which is a neuron type available in CSIM) and adds them to nmc. Note that due to the object oriented paradigm used in Matlab the nmc object must appear also on the left hand side of the command. The variable p_lif is a handle/index to refer to that particular pool later in the programm.

The neurons are located on a three dimensional $3 \times 3 \times 6$ integer grid with origin (2,1,1). You can visualize this by issuing the command

```
>> plot(nmc);
```

The plot command should produce a figure which looks very much like Figure ??. As you can see in Figure ?? some neurons are marked by magenta balls. These are inhibitory neurons while the other are excitatory neurons. By default a neuron is choosen to be a excitatory with a probability of 80% (this can be controlled with the frac_EXC parameter).

Now we add the other two pools of neurons where we set some off-default parameters:

```
>> [nmc,p_sig] = add(nmc,'pool','type','SigmoidalNeuron',...
```

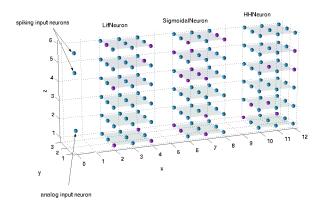


Figure 2: Three pools of model neurons consisting of neurons of the classes LifNeuron, SigmoidalNeuron, and HHNeuron (from left to right) and two input pools.

```
'size',[3 3 6],'origin',[6 1 1],...
'Neuron.thresh',1,'Neuron.beta',2,'Neuron.tau_m',3,...
'Neuron.A_max',4,'Neuron.I_inject',1,'Neuron.Vm_init',0);

>> [nmc,p_hh] = add(nmc,'pool','type','HHNeuron',...
'size',[3 3 6],'origin',[10 1 1],...
'Neuron.Inoise',???,'Neuron.Iinject',[??? ???]);
```

The code fragment above shows how to set off-default values for some parameters of the Neurons generated. One has to add a pair of 'Neuron.<field>',<value> arguments to the function call. Which fields are valid is determined by the class of the neuron (see the CSIM Class Reference for details).

As the next step we create the input neurons. A pool of 2 excitatory spiking input neurons

```
>> [nmc,p_sin] = add(nmc,'pool','type','SpikingInputNeuron',...
'size',[1 1 2],'origin',[0 1 5],'frac_EXC',1.0);
```

and a pool of a single excitatory analog input neuron:

```
>> [nmc,p_ain] = add(nmc,'pool','type','AnalogInputNeuron',...
'size',[1 1 1],'origin',[0 1 2],'frac_EXC',1.0);
```

A visualization of the current model is shown in Figure ?? which was produce by the command plot(nmc);.

2.3 Making synaptic connections

Now we want to set up synaptic connections between the neurons in the individual pools. This is done by commands of the form

```
>> [nmc,c_idx] = add(nmc,'conn','dest',<destination>,'src',<source>,...);
```

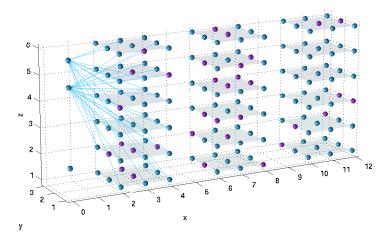


Figure 3: Synaptic connections of the spiking input neurons to the LifNeuron pool.

where **destination** and **source** specify a set of neurons either by the handle/index of a pool or by specifying a volume.

2.3.1 Connecting the input

To see how it works lets start by connecting the spiking inputs to the pool of LifNeurons:

```
>> [nmc,c(1)] = add(nmc,'conn','dest',p_lif,'src',p_sin,...
'Cscale',1,'Wscale',5);
```

Here we used the pool handles/indices as source and destination specification. The additional parameters given specify how to scale the overall connection probability (Cscale) and the synaptic strength (Wscale). To see the actual connectivity pattern generated you can again use the command plot(nmc); and interaktively explore the network structure. By clicking on a neuron you can look at presynaptic as well as postsynaptic connections. Figure ?? shows how the input neurons are connected at the moment. As the next step we connect the spiking input via static synapses (Sec. ??) (default are dynamic synapses (Sec. ??) to some subset of the pool of sigmoidal neurons by specifying a certain volume

```
>> [nmc,c(2)]=add(nmc,'conn','dest',[6 1 1; 6 3 6],'src',p_sin,...
'type','StaticSpikingSynapse','Cscale',Inf);
```

The synaptic connection created by this command are shown in Figure ??. Note that the setting 'Cscale', Inf ensures that there will be a synaptic connection between each pair of neurons in the source region and the destination region. And finally we connect the analog input to the pool of HHNeurons by means of a StaticAnalogSynapse. This is necessary since StaticSpikingSynapses can not transmit analog signals.

```
>> [nmc,c(3)]=add(nmc,'conn','dest',p_hh,'src',p_ain,...
'type','StaticAnalogSynapse','Wscale',0.2);
```

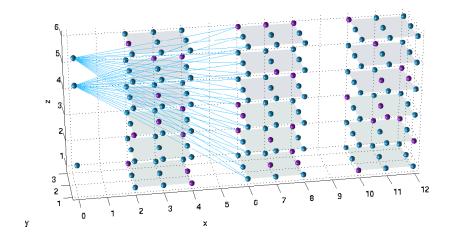


Figure 4: Synaptic connections of the spiking input neurons to the Sigmoidal Neuron pool.

2.3.2 Making recurrent connections

Now we want to create recurrent connections with the pools themselves. The only differene to the previous section is that now the source and destination is the same pool:

Again one can specify in addition to the destination and the source other parameters which determine how individual synaptic connections are created. For example the parameter lambda determines the "average distance" of synaptic connections and SH_W determines the standard deviation SD of the Gamma distribution used to generate the synaptic weights: $SD = SH_W \cdot mean$. Figure ?? shows a typical postsynaptic connectivity pattern for one neuron generated by the code above.

2.3.3 Connecting pools

In this example we connect the pool of HHneurons to the LifNeuron pool by static synapses.

Note that using lambda=Inf has the effect that the distance between the neurons dose not matter in determining wheter a synaptic connection will by generated or not.

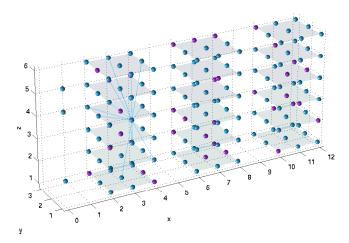


Figure 5: Postsynaptic connections of a typical excitatory LifNeuron.

2.4 Simulating the model

Now that the model is set up we turn to the issue how to define the input and simulate the network with these inputs.

2.4.1 Setting up the input

As we have three input neurons (2 spiking one analog) we have to define a *stimulus* which consists of three channels (2 spiking one analog). The stimulus can be defined as follows:

```
% create empty structure
S = empty_stimulus('nChannels',3,'Tstim',1);
% fill channel 1 with some spikes
S.channel(1).data
                    = 1*rand(1,10);
S.channel(1).spiking = 1;
% fill channel 2 with some spikes
S.channel(2).data
                    = 1*rand(1,20);
S.channel(2).spiking = 1;
S.channel(3).dt
                    = 0.005;
                    = 1+sin(2*pi*10*[0:S.channel(3).dt:1]);
S.channel(3).data
S.channel(3).spiking = 0;
```

Note that each signal/channel can be either spiking (S.channel(i).spiking=1) or analog (S.channel(i).spiking=0). In the later case one has to specify the temporal resolution (S.channel(i).dt) of the signal.

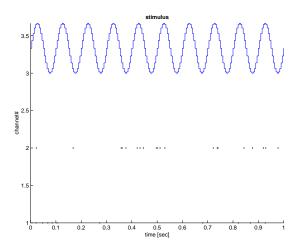


Figure 6: A stimulus with 2 spiking and one analog channel.

We adopt the convention that the first channel of the stimulus is assigned to the first input neuron created by means of [nmc,p]=add(nmc,'pool',... statements. Keep this convention in mind when setting up the stimulus otherwise you may be surprised by error messages or strange results (if the input neurons have different connections to the rest of the network).

You can use the command

```
>> plot_stimulus(S);
```

to plot the stimulus defined above. This results in the plot shown in Figure ??.

2.4.2 Defining the respones

Obviously we want to see how the network responses to the given stimulus. Therefor we must specify what we want to record during the stimulation. The following code fragment shows how to record the spikes of the pool of LifNeurons and the membrane voltage of a certain subset (defined by specifying the appropriate volume) of the SigmoidalNeuron and HHNeuron pool.

```
>> nmc = record(nmc,'Pool',p_lif,'Field','spikes');
>> nmc = record(nmc,'Volume',[6 1 1; 8 3 1],'Field','Vm');
>> nmc = record(nmc,'Volume',[10 1 1; 12 3 1],'Field','Vm');
```

2.4.3 Running the simulation

Now we are ready to run the simulation; lets say for 1 sec.

```
>> R=simulate(nmc,1,S);
```

This returns the cell array R which contains the response of the network. See the section about input and output (Sec. 3) for more details about the structure of R.

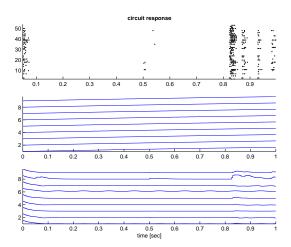


Figure 7: Response.

However, one can use the command

```
>> plot_response(R);
```

to plot the response R. This results in the plot shown in Figure ??.

3 Stimulus and Response

3.1 The Stimuls or Input Signals

When runing a network simulation via simulate(nmc, Tstim, S); one can specify the *stimulus* S.

S has to be a struct array with the following fields:

- S.channel(i).spiking: binary flag (0/1) which determines if S.channel(i).data should be interpreted as spike times or as an analog signal
- S.channel(i).dt: time discretization; for analog signals (S.channel(i).spiking=0) only
- S.channel(i).data : signal date: vektor of the analog values (S.channel(i).spiking=0) or spike times (S.channel(i).spiking=1)
- S.info(i).Tstim: the length of the stimulus (usally used in plotting routines.

3.2 The response or the network output

After a network simulation via a command like

```
>> R=simulate(nmc,Tstim,S);
```

the *response* is stored in the cell array R. $R\{i\}$ contains the traces specified by the *i*-th nmc = record(nmc,...) statement during the setup of the simulation.

R{i} by itself is a struct array with the only field channel which is in turn a struct array with a similar structure as an input signal (Sec. 3.1). That is

- $R\{i\}$.channel(j).data: signal data: vektor of the analog values or spike times. Note that the data always starts at time t = 0.
- R{i}.channel(j).spiking: binary flag (0/1) which determines if data should be interpreted as spike times or as an analog signal
- R{i}.channel(j).dt: time discretization; for analog signals only
- R{i}.channel(j).fieldName: name of the recorded field
- R{i}.channel(j).idx: handle of the object from which field the data was recorded

4 Input Distributions

- 5 Distributed simulations
- 6 neural_microcircuit class reference
- 7 delay_lines class reference
- 8 small_circuits class reference

References

[Dayan and Abbott, 2001] Dayan, P. and Abbott, L. (2001). Theoretical Neuroscience: Computational and Mathematical Modeling of Neural Systems. MIT Press. See also http://people.brandeis.edu/~abbott/book/.

[Gerstner and Kistler, 2002] Gerstner, W. and Kistler, W. (2002). Spiking Neuron Models. Cambridge University Press. See also http://diwww.epfl.ch/~gerstner/BUCH.html.