

# Tutorial: Parallel Network Simulations with NEURON

## Installation

MPI

NEURON

Launch

## Network specification

GID distribution

Create cells

Connections

Random

cell parameters, connections, weight, delay

spike input stimulus

## Running

performance measurement

efficiency

## An example

Vogels and Abbott (2005)

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## An example

Vogels and Abbott (2005)

Same result, no matter

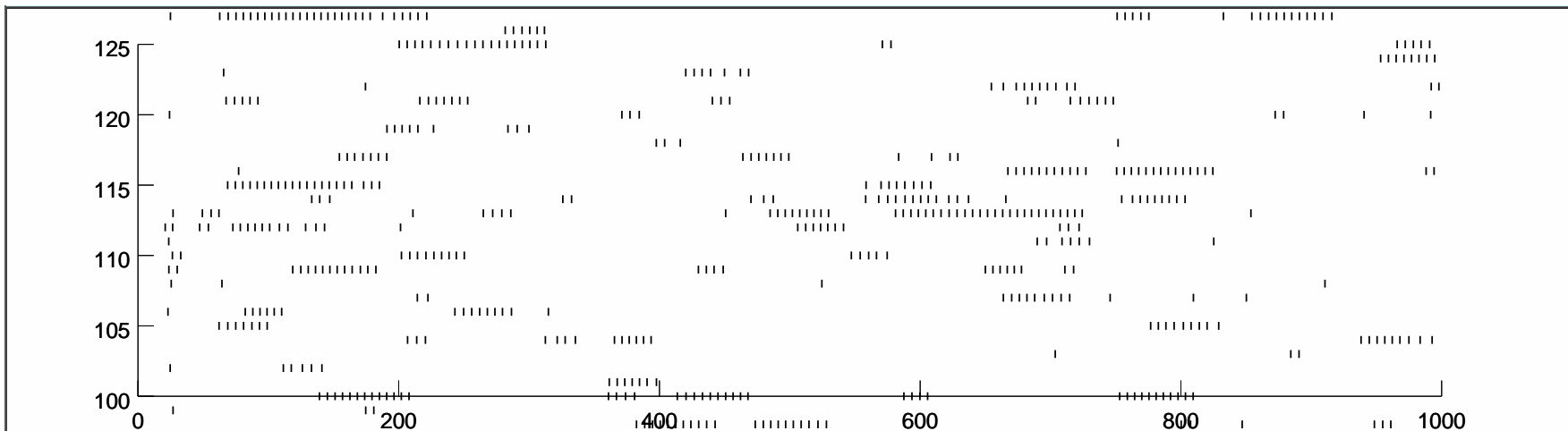
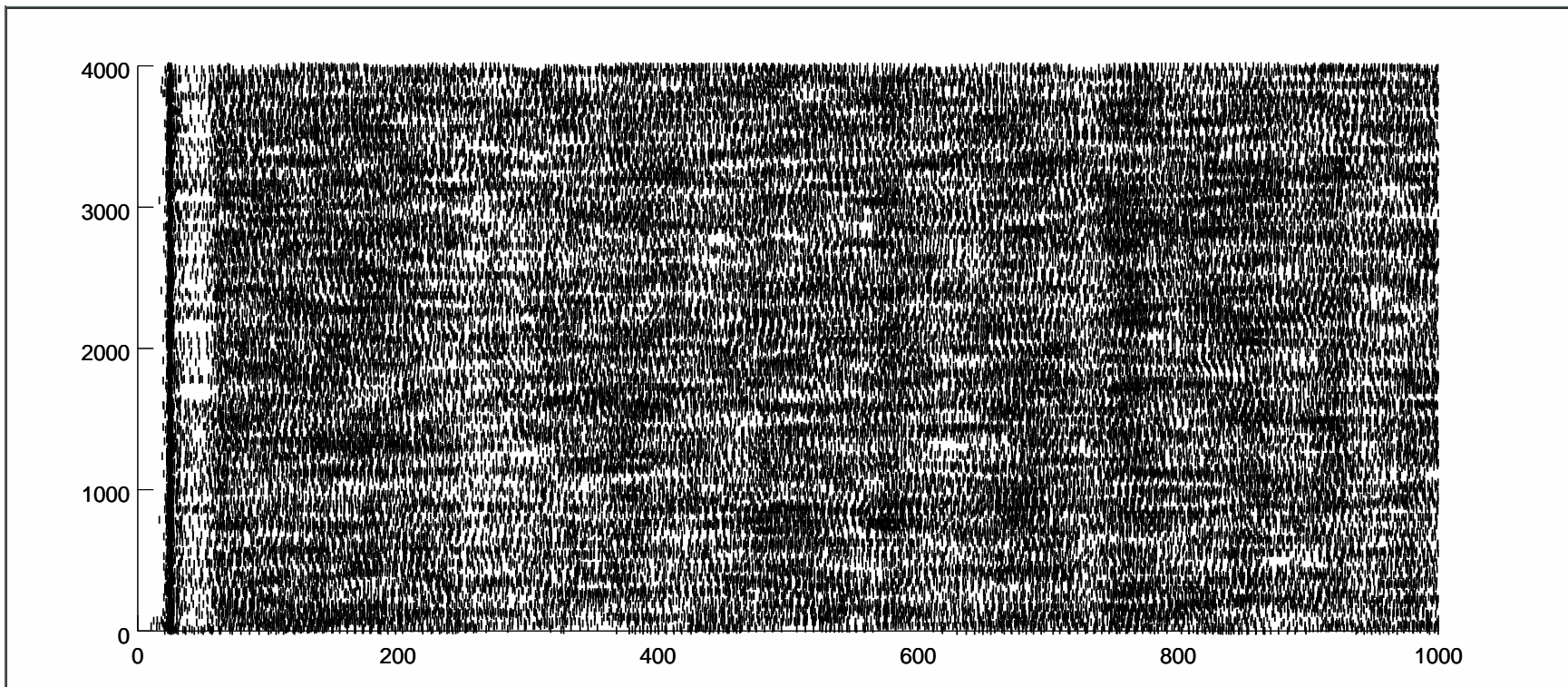
how many machines

how cells distributed

reproducible

statistically

independent



```
$ mpirun ... -np 10 ...
```

```
$ cat out.dat
```

```
18.1  2410          999.9  1190
20.1  530          999.9  1810
21.2  3880         18.2    75
21.4  2670         19.6   1915
...
999.8  550         19.6   3805
...
```

```
$ cat ~/bin/sortspike
```

```
#!/bin/sh
```

```
sort -k 1n,1n -k 2n,2n $1 > $2
```

```
$ sortspike out.dat out.10
```

```
$ mpirun ... -np 5 ...
```

```
$ sortspike out.dat out.5
```

```
$ diff out.5 out.10
```

Beowulf cluster or multiprocessor personal machine. (Linux)

See MPI under ParallelContext

Can you login to any node without a password?

```
ssh `hostname`
```

Is MPI installed?

```
which mpicc
```

```
which mpic++ # or mpicxx, mpiCC, ...
```

Beowulf cluster or multiprocessor personal machine. (Linux)

See MPI under ParallelContext

Can you login to any node without a password? If not...

```
ssh-keygen -t rsa
cd $HOME/.ssh; cat id_rsa.pub >> authorized_keys
```

Is MPI installed? If not...

Get the latest stable version from:

```
http://www.mcs.anl.gov/research/projects/mpich2/
```

```
tar xzf mpich2-1.2.1p1.tar.gz
```

```
cd mpich2-1.2.1p1
```

```
./configure --prefix=$HOME/mpich2
```

```
make
```

```
make install
```

```
export PATH=$HOME/mpich2/bin:$PATH
```

```
cd $HOME;echo 'secretword=nrnmpi'>.mpd.conf;chmod 600 .mpd.conf
```

Test it.

```
mpd&
```

```
mpiexec -n 2 echo 'hello' # should print twice
```

```
mpdallexit
```

# Beowulf cluster or multiprocessor personal machine. (Linux)

## Installing NEURON (without InterViews)

Assume sources are in \$HOME/neuron/nrn

```
cd $HOME/neuron
```

```
mkdir nrnmpi ; cd nrnmpi
```

```
../nrn/configure --prefix=`pwd` --with-paranrn --with-nrnpython \  
--without-x
```

```
make install
```

# Supercomputers need special configuration...

## SDSC IBM DataStar

[http://www.sdsc.edu/user\\_services/datastar/](http://www.sdsc.edu/user_services/datastar/)

272 (8-way) P655+ and 7 (32-way) P690 compute nodes.

could not get autoconf-5.9 installed so extracted a nrn-x.tar.gz file over the cvs sources. /usr/bin/oslevel does not work so, in nrn, run

```
grerep /usr/bin/oslevel /usr/local/bin/oslevel config.guess configure
```

```
sh ../nrn/configure --prefix='pwd' \  
  '--srcdir=../nrn' --with-nmodl-only --without-x CC=gcc CXX=g++  
make; make install
```

```
sh ../nrn/configure --prefix='pwd' '--srcdir=../nrn' \  
  '--without-x' '--without-memacs' '--without-readline' \  
  '--disable-shared' '--with-paranrn' '--without-nmodl' \  
  CC=mpcc_r CXX=mpCC_r MPICC=mpcc_r MPICXX=mpCC_r \  
  CFLAGS=-q64 CXXFLAGS=-q64 linux_nrnmech=no AR="ar -X64" \  
  always_call_mpi_init=yes java_dlopen=no \  
  --host=powerpc-ibm-aix5.2.0.0  
make; make install
```

## **Install: MSWindows**

Vista or Windows 7 install

<http://www.neuron.yale.edu/ftp/neuron/versions/alpha/nrn-7.2.alpha-426-setup.exe>

Test by opening an rxvt window and typing

```
mpd&
```

```
mpiexec -n 2 /cygdrive/c/nrn72/bin/echo 'hello'
```

```
mpdallexit
```

Test that NEURON works with mpi by creating a test0.hoc program:

```
objref pc
```

```
pc = new ParallelContext()
```

```
{printf("I am %d of %d\n", pc.id, pc.nhost)}
```

```
{pc.runworker() pc.done() quit() }
```

and launch with

```
mpiexec -n 4 /cygdrive/c/nrn72/bin/nrniv -mpi test0.hoc
```

(try leaving out the -mpi arg)

# Launch (Linux)

```
cd neuron/nrn/src/parallel  
mpiexec -n 3 nrniv -mpi test0.hoc
```

```
numprocs=3
```

```
NEURON -- Version 7.2 (428:986821b56b98) 2010-03-07
```

```
Duke, Yale, and the BlueBrain Project -- Copyright 1984-2008
```

```
See http://www.neuron.yale.edu/credits.html
```

```
I am 0 of 3
```

```
I am 1 of 3
```

```
I am 2 of 3
```

## **nrn/src/parallel/test0.hoc**

```
objref pc
pc = new ParallelContext()
{printf("I am %d of %d\n", pc.id, pc.nhost)}
{pc.runworker()}
{pc.done()}
quit()
```

## common/net.hoc

```
proc create_cells() { local i, gid localobj cell, nc
  for pcitr(&i, &gid) {
    cell = newcell(gid)
    pnm.register_cell(gid, cell)
  }
}
```

## coba/init.hoc

```
obfunc newcell() {
  return new CobaCell()
}
```

## coba/cobacell.hoc

```
begintemplate CobaCell
public ...connect2target...
proc init() {
  soma spkout = new SpikeOut(.5)
}
proc connect2target() { //$o1 target point process, $o2 returned NetCon
  $o2 = new NetCon(spkout, $o1)
}
endtemplate CobaCell
```

# Create

## cuba/cubacell.hoc

```
obfunc newcell() {localobj cell
  cell = new IF3()
  cell.taum = 20
  cell.taue = 10
  cell.taui = 20
  return cell
}
```

PointProcessManager

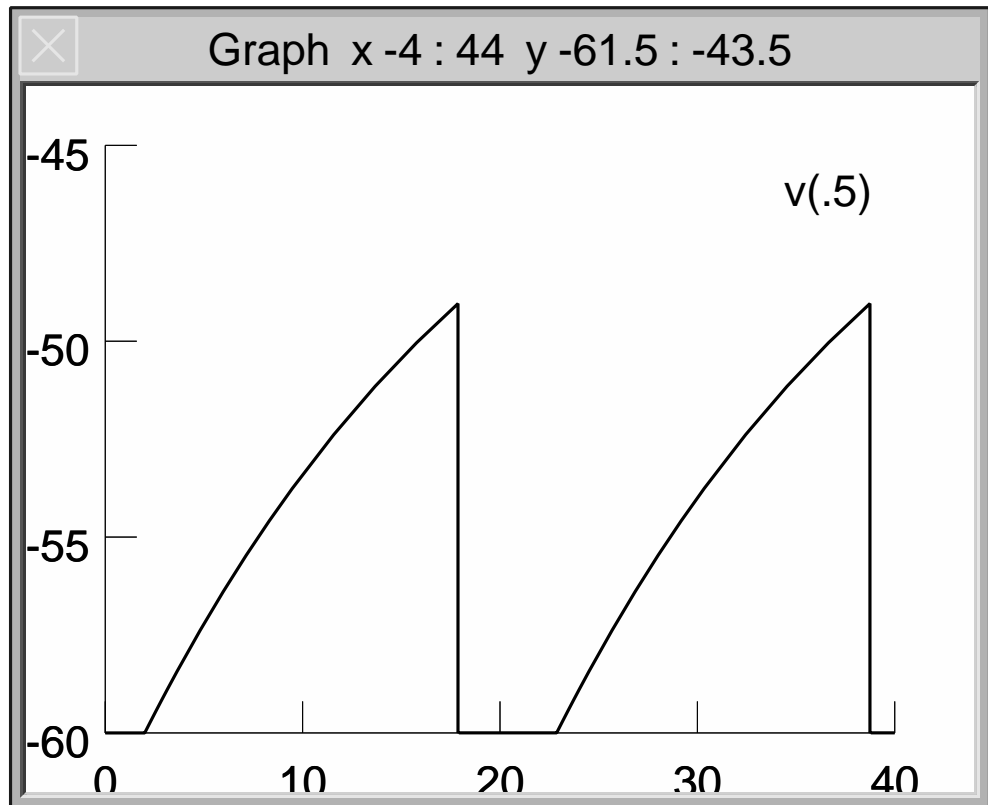
SelectPointProcess

Show

IClamp[0]  
at:CobaCell[0].soma(0.5)

IClamp[0]

del (ms)	<input checked="" type="checkbox"/>	2
dur (ms)	<input checked="" type="checkbox"/>	100
amp (nA)	<input checked="" type="checkbox"/>	0.2
i (nA)		0.2



# nrn/lib/hoc/netparmpi.hoc

begintemplate ParallelNetManager

...

```
proc register_cell() { localobj nc
    if (!pc.gid_exists($1)) { pc.set_gid2node($1, pc.id) }
    // all existing cells must have an associated gid which
    // is stored in the cell's PreSyn. The nc below will be
    // unreffed but the PreSyn will continue
    // in existence and from the gid we will quickly be able
    // to find the PreSyn and from that the Cell
    // we force the cell to be an outputcell due to the danger of
    // user error
    cells.append($o2)
    if (hoc_sf_.is_artificial($o2)) {
        nc = new NetCon($o2, nil)
    }else{
        $o2.connect2target(nil, nc)
    }
    pc.cell($1, nc, 1)
}
```

```

proc connect_cells() { local i, j, gid, r, d localobj cell, u, rs
  d = DELAY // in the paper it is 0
  mcell_ran4_init(connect_random_low_start_)
  u = new Vector(ncell) // for sampling without replacement
  for pcitr(&i, &gid) { // target cell
    u.fill(0)
    cell = pnm.cells.object(i)
    rs = ranlist.object(i) rs.start()
    rs.r.discunif(0, N_E-1)
    j=0 while(j < C_E) { // Excitatory sources
      r = rs.repick()
      if (r != gid) if (u.x[r] == 0) {
        pnm.nc_append(r, gid, AMPA_INDEX, AMPA_GMAX, d)
        u.x[r] = 1    j += 1
      }
    }
    rs.r.discunif(N_E, ncell-1)
    j=0 while(j < C_I) { // Inhibitory sources
      r = rs.repick()
      if (r != gid) if (u.x[r] == 0) {
        pnm.nc_append(r, gid, GABA_INDEX, GABA_GMAX, d)
        u.x[r] = 1    j += 1
      }
    }
  }
}

```

pick exactly C\_E and C\_I unique non-self random connections for each target. Assume many more sources than target connections.

# Connect

```
proc create_stim() {local i, gid localobj stim, cell, nc, rs
```

```
  mcell_ran4_init($1)
```

```
  stimlist = new List()
```

```
  ncstimlist = new List()
```

```
  // first N_STIM (excitatory) cells stimulated
```

```
  for pcitr(&i, &gid) {
```

```
    if (gid >= N_STIM) { break }
```

```
    cell = pc.gid2cell(gid)
```

```
    rs = ranlist.object(i)
```

```
    stim = new NetStim()
```

```
    stim.interval = STIM_INTERVAL
```

```
    stim.number = 1000 // but will shut off after STOPSTIM
```

```
    stim.noise = 1      stim.start = 0
```

```
    // use the gid specific random generator so random streams
```

```
    // independent of where and how many stims there are
```

```
    stim.noiseFromRandom(rs.r)
```

```
    rs.r.negexp(1)      rs.start()
```

```
    nc = new NetCon(stim, cell.synlist.object(0))
```

```
    nc.delay = 1  nc.weight = $2
```

```
    ncstimlist.append(nc)
```

```
    stimlist.append(stim)
```

```
  }
```

# Stimulate

```
  stim = new NetStim() // will turn off all the others
```

```
  stim.number = 1
```

```
  stim.start = STOPSTIM
```

```
  for i=0, stimlist.count-1 {
```

```
    nc = new NetCon(stim, stimlist.object(i))
```

```
    nc.delay = 1  nc.weight = -1
```

```
    ncstimlist.append(nc)
```

```
  }
```

```
  stimlist.append(stim)
```

```
}
```