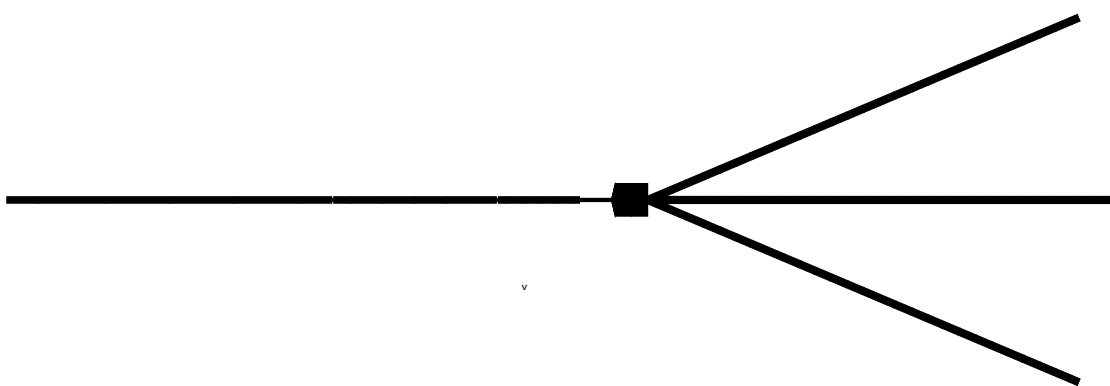


The NEURON Simulation Environment

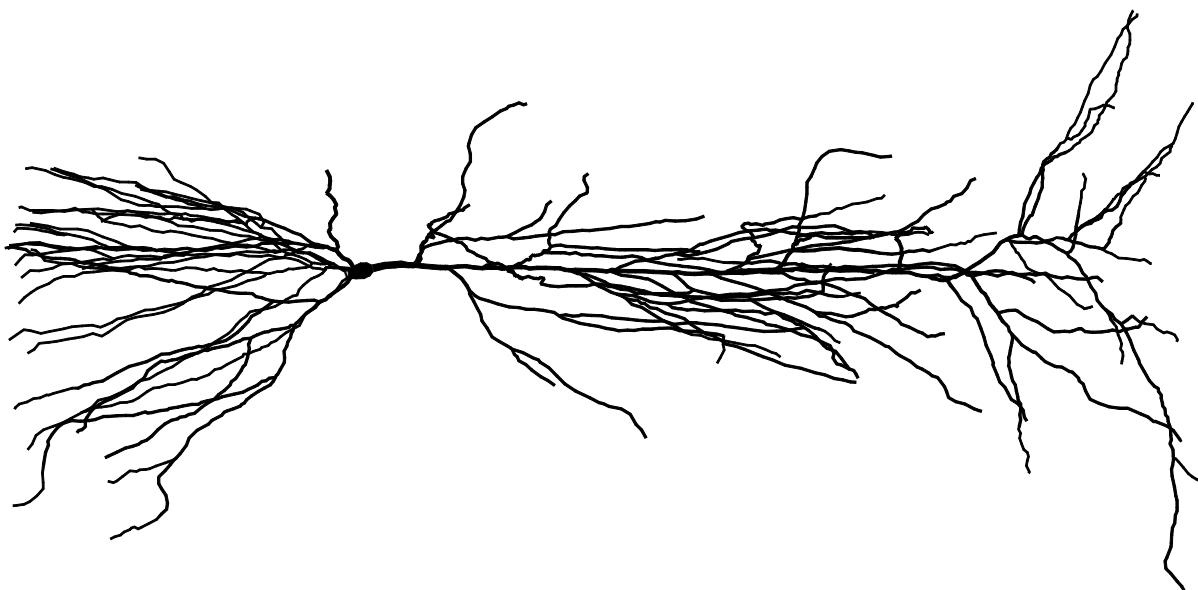
**Ted Carnevale
Michael Hines
John Moore
Gordon Shepherd**

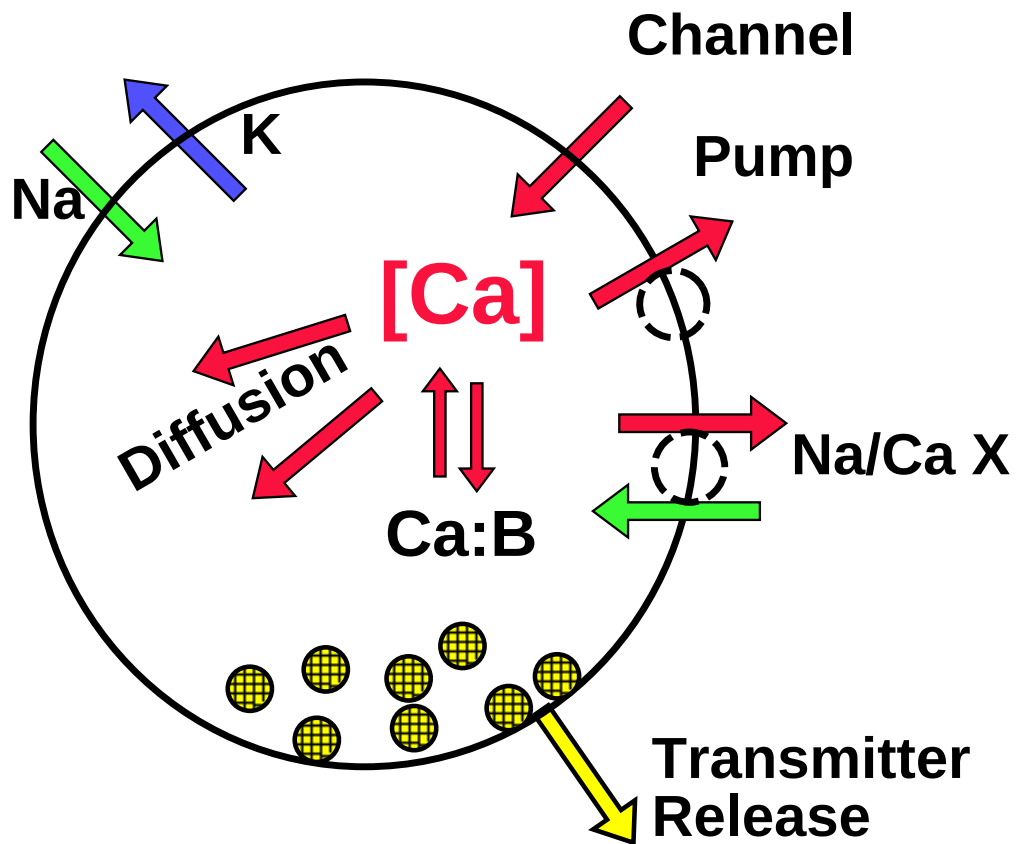
<http://www.neuron.yale.edu>

Supported by NINDS



v





Extracellular fields
Linear circuits
Synapses
Networks

Physical System



Model



Representation in NEURON

Create Representation

**Investigate/Explore/Control/Use
Representation**

Hodgkin - Huxley axon

Physical System



From <http://www.mbl.edu>

Model

Hodgkin-Huxley cable equations

$$\frac{D}{4R_a} \cdot \frac{\partial^2 V}{\partial x^2} = C_m \frac{\partial V}{\partial t} + \bar{g}_{na} m^3 h \cdot (V - E_{na}) + \bar{g}_k n^4 \cdot (V - E_k) + g_l \cdot (V - E_l)$$

$$\begin{aligned} \frac{dm}{dt} &= -\alpha_m m + \beta_m \cdot (1 - m) & \alpha_m &= \frac{.1(V+40)}{1 - e^{-.1(V+40)}} & \beta_m &= 4e^{-(V+65)/18} \\ \frac{dh}{dt} &= -\alpha_h h + \beta_h \cdot (1 - h) & \alpha_h &= .07e^{-.05(V+65)} & \beta_h &= \frac{1}{1 + e^{-.1(V+35)}} \\ \frac{dn}{dt} &= -\alpha_n n + \beta_n \cdot (1 - n) & \alpha_n &= \frac{.01(V+55)}{1 - e^{-.1(V+55)}} & \beta_n &= .125e^{-(V+65)/80} \end{aligned}$$

Simulation

Representation

```
create axon
axon {
    nseg = 50
    diam = 100
    L = 20000
    insert hh
}
```

Interpreted

Setup

Control

Presentation



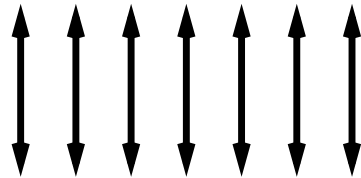
Functions
Variables

Compiled

Parameterized
Equations

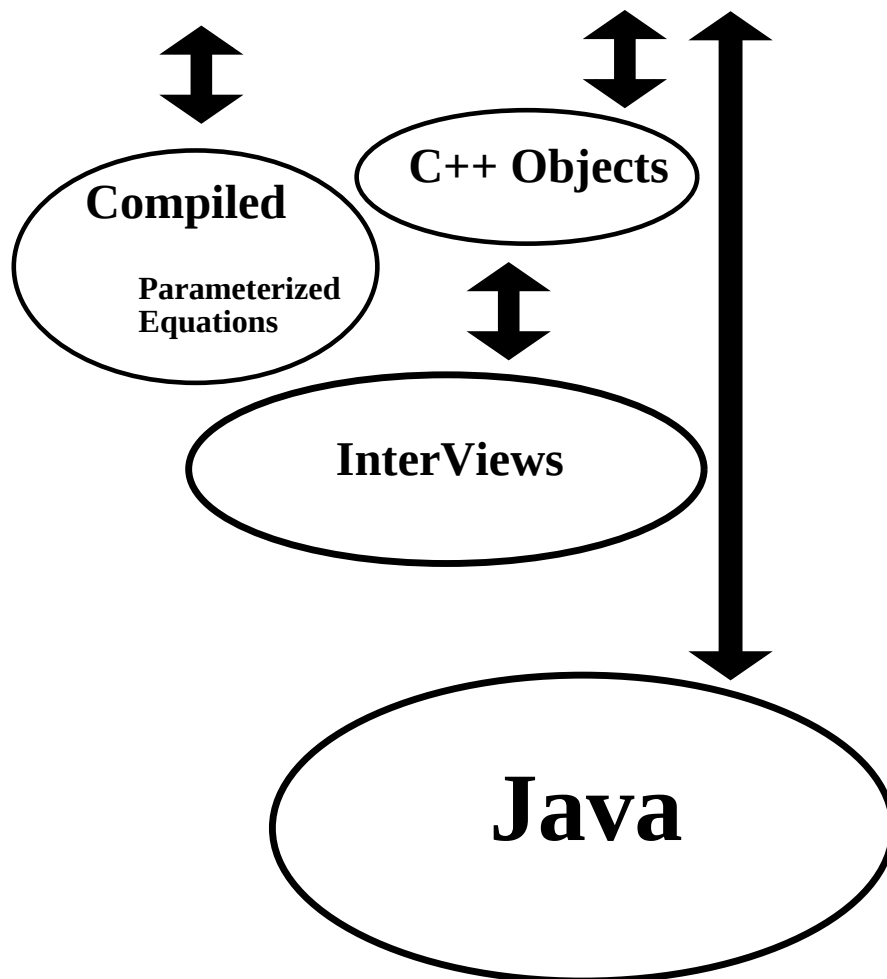
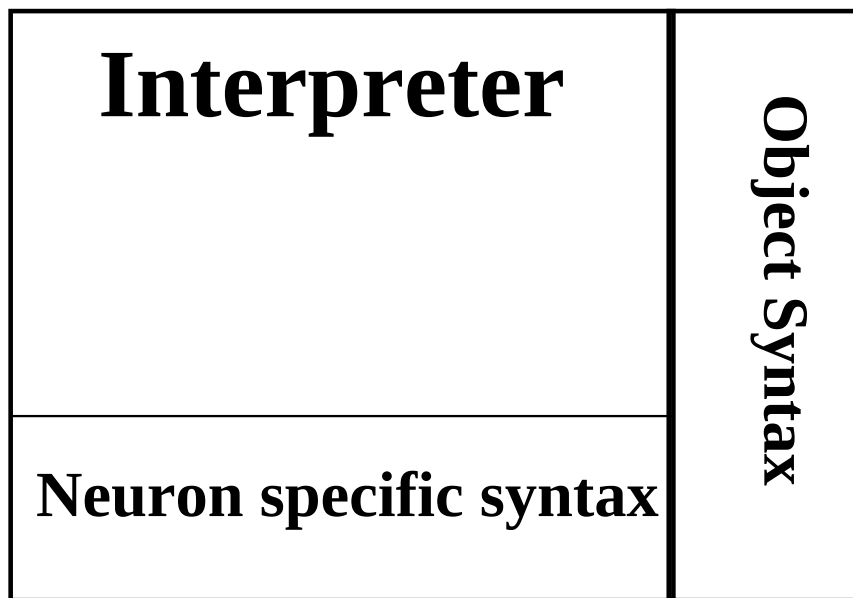
Interpreter

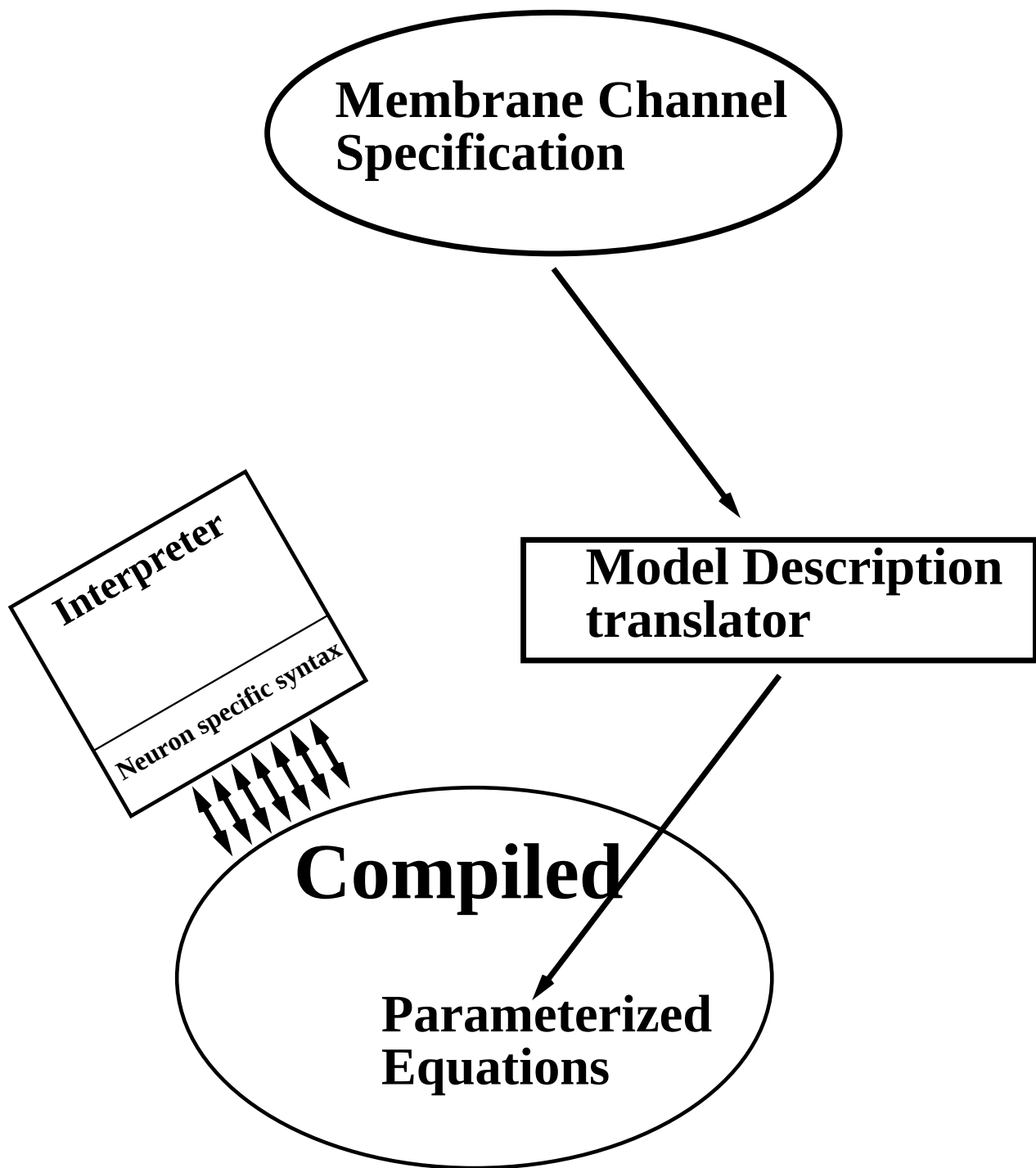
Neuron specific syntax



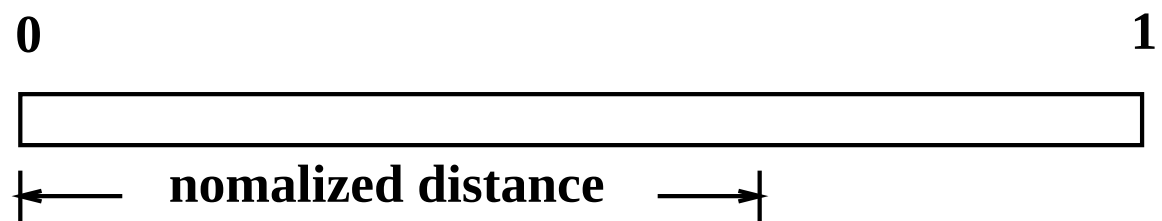
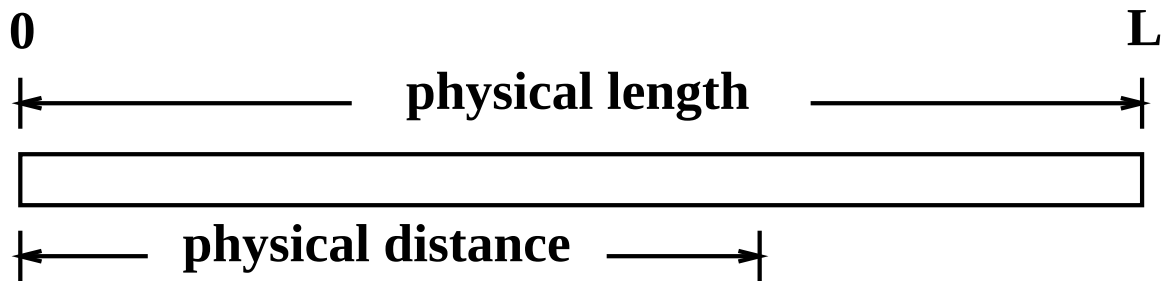
Compiled

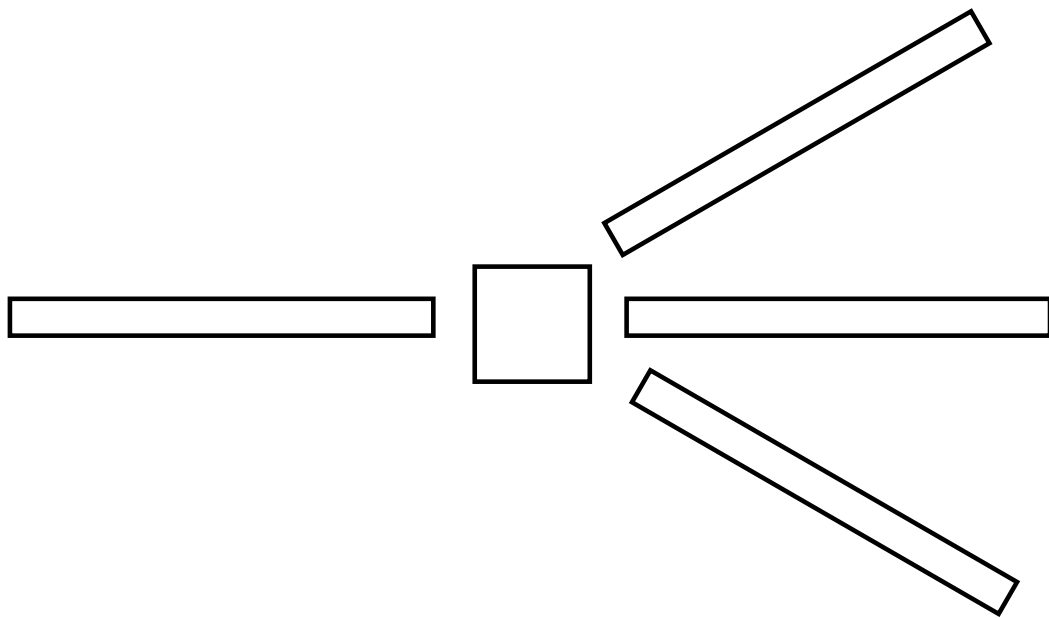
**Parameterized
Equations**





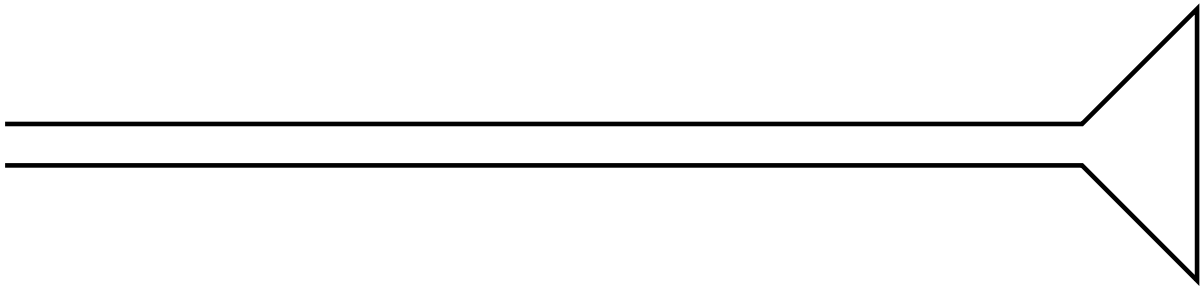
Cable Section



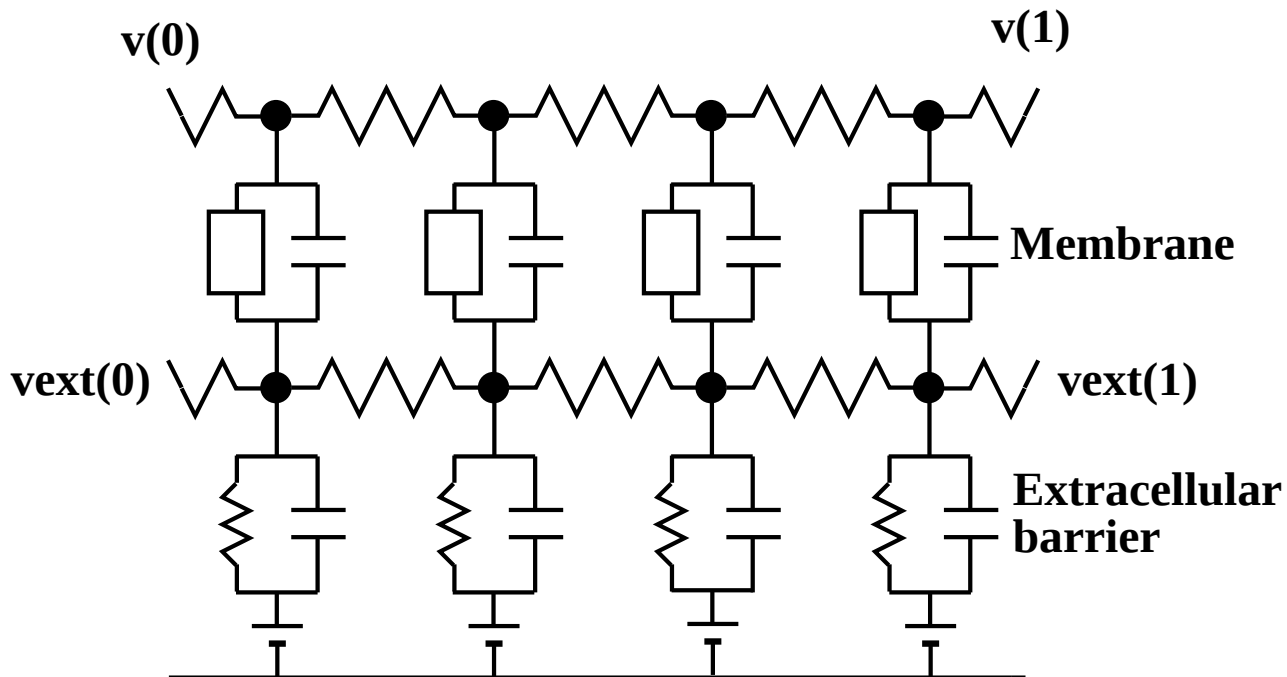
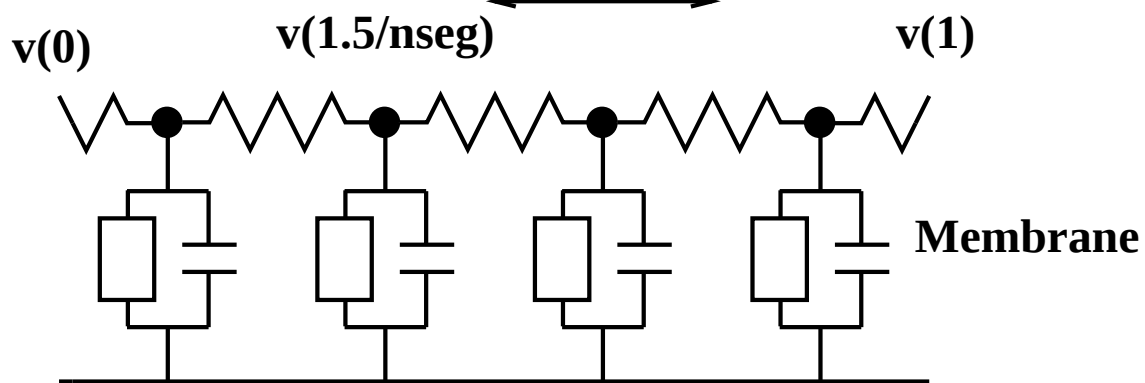
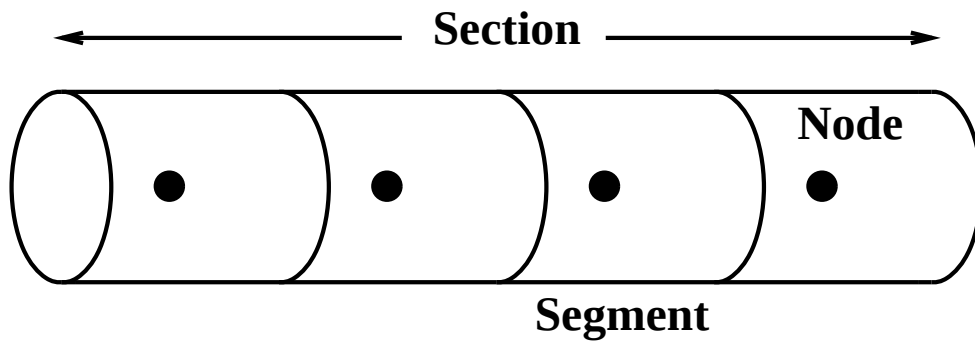


```
create axon, soma, dendrite[3]
connect axon(0), soma(0)
for i=0,2 connect soma(1), dendrite[i](0)
```

```
axon.L = 1000
axon {
    insert hh
    gnabar_hh = .120
}
```

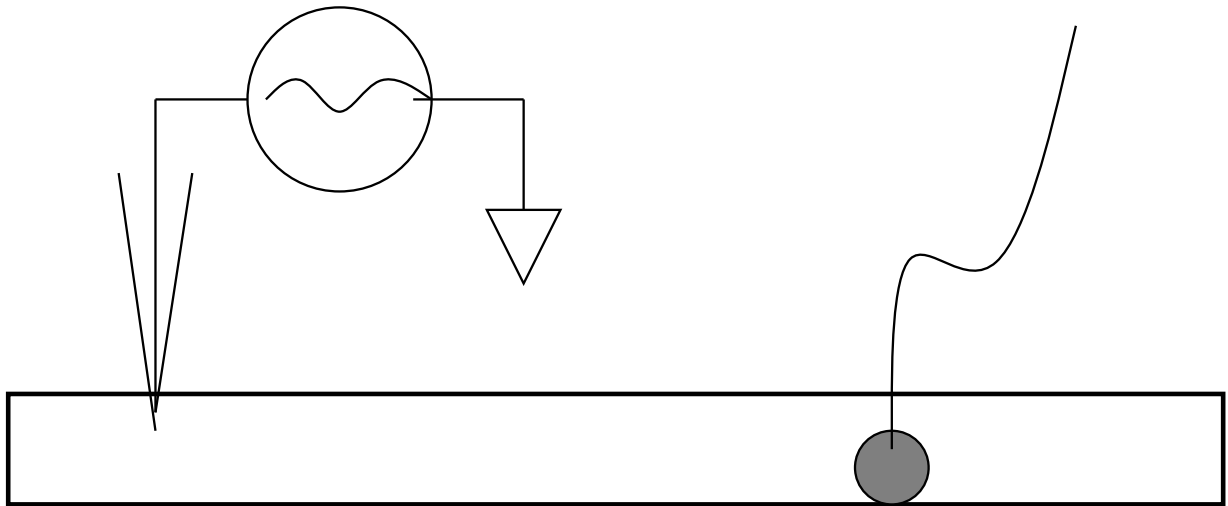


```
nseg = 20
diam(0:.1) = 20:2
diam(.1:1) = 2:2
```





forall nseg *= 3

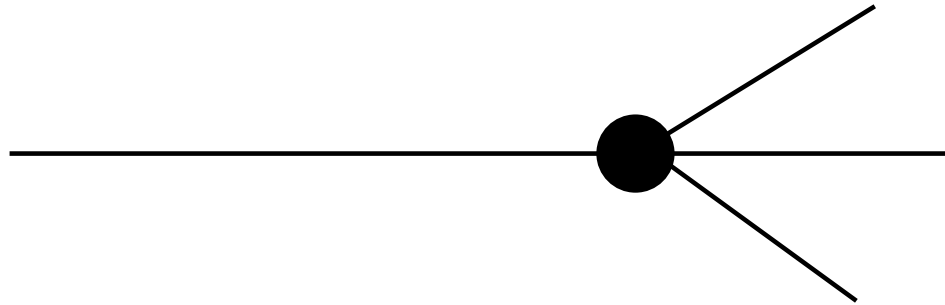


```
objref stim, syn  
stim = new IClamp(.1)  
syn = new Synapse(.7)
```

```
stim.dur = .1  
stim.amp = 5
```

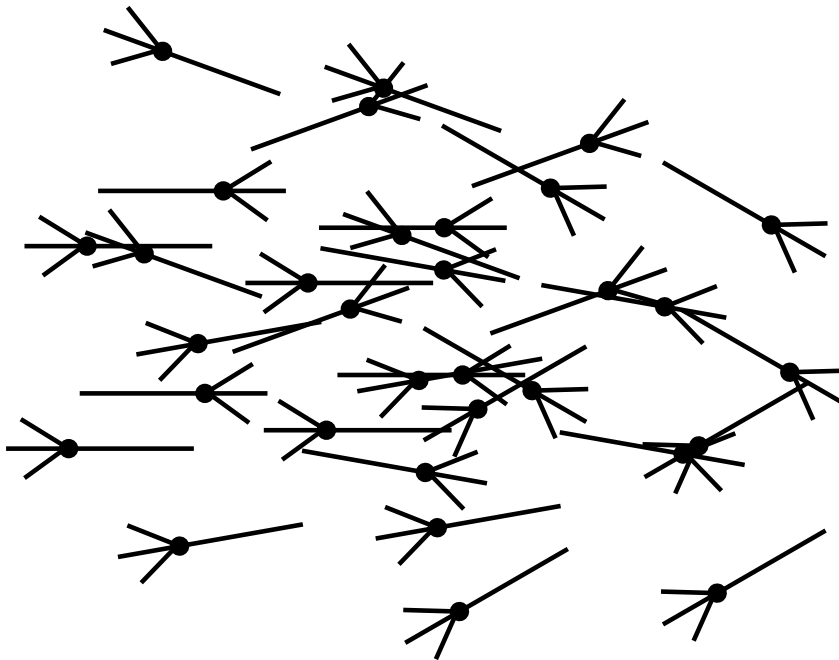
```
setpointer syn.vpre, cell[15].axon.v(1)
```

```
begintemplate Cell  
  public axon, soma, dendrite[3]
```



```
endtemplate Cell
```

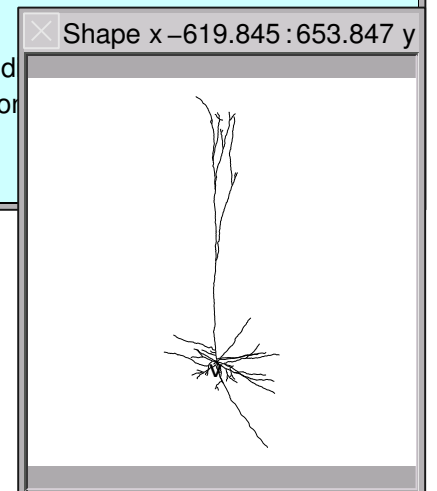
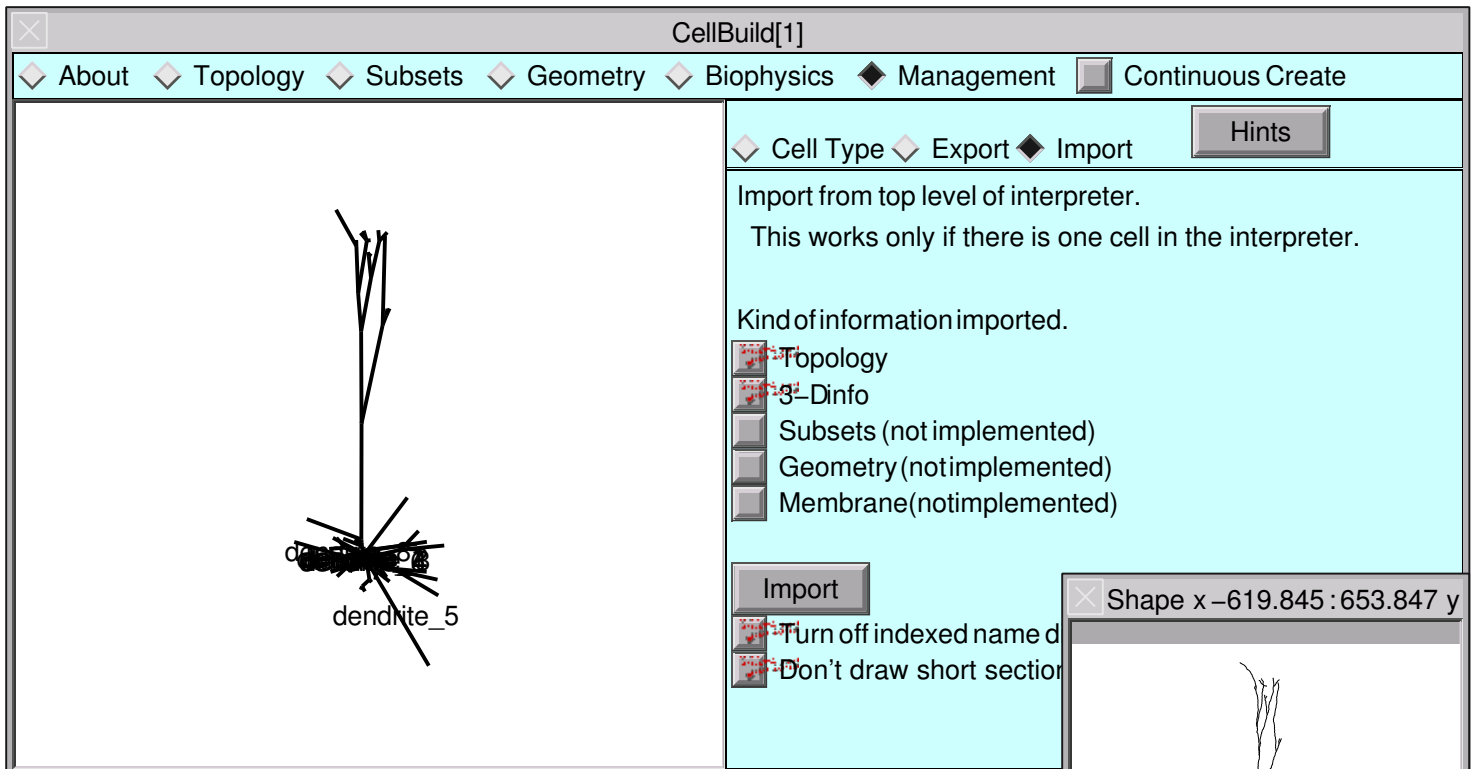
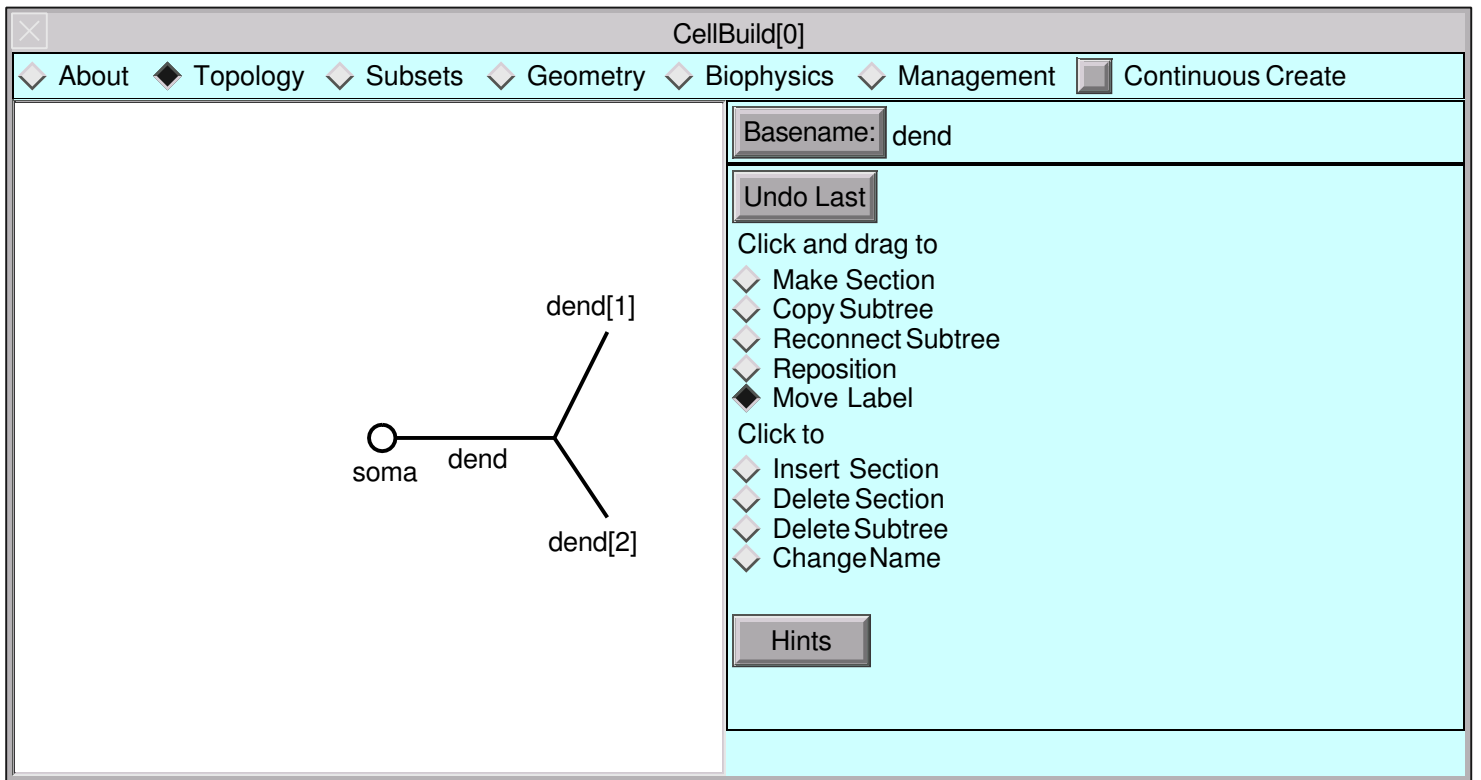
```
objectvar cell[32]  
for i=0,31  cell[i] = new Cell()
```

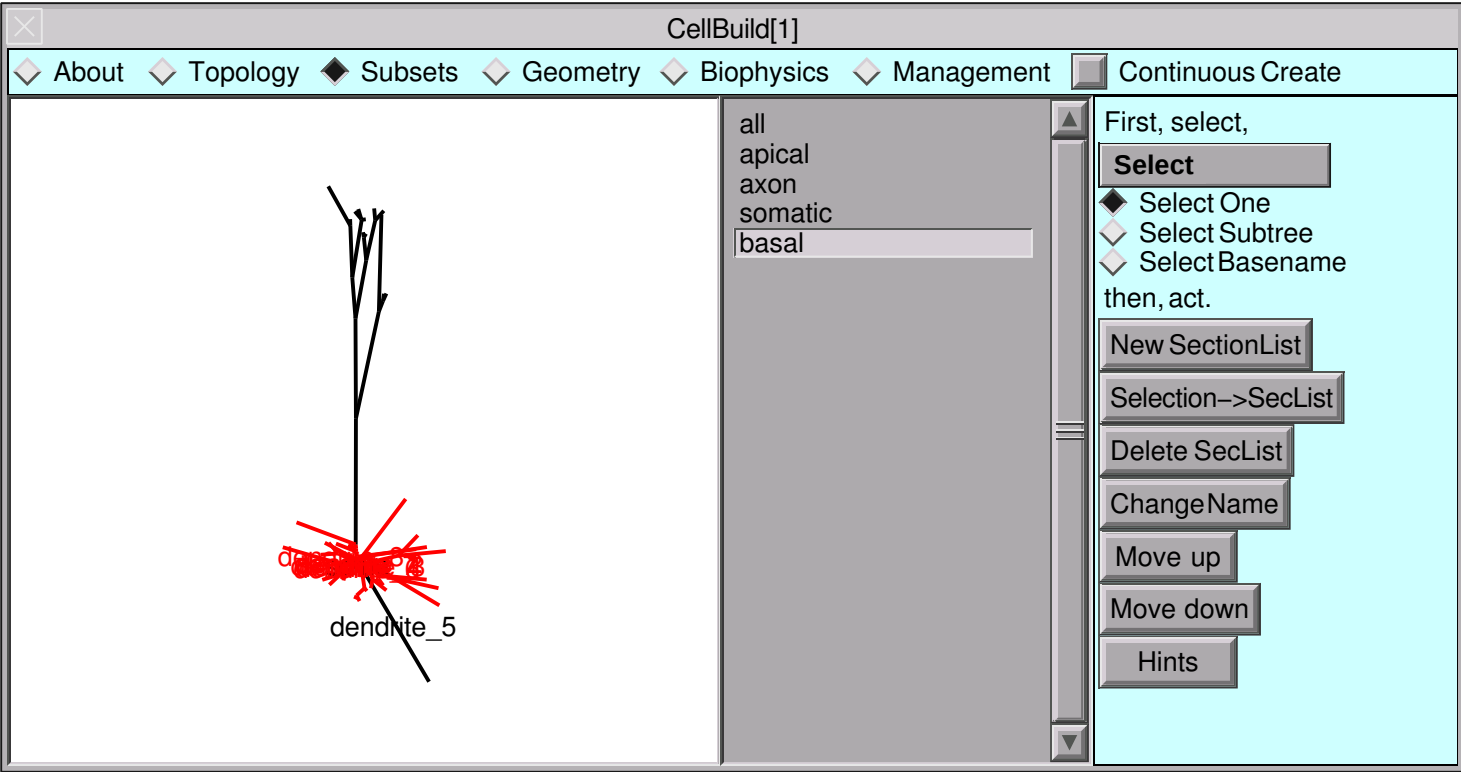


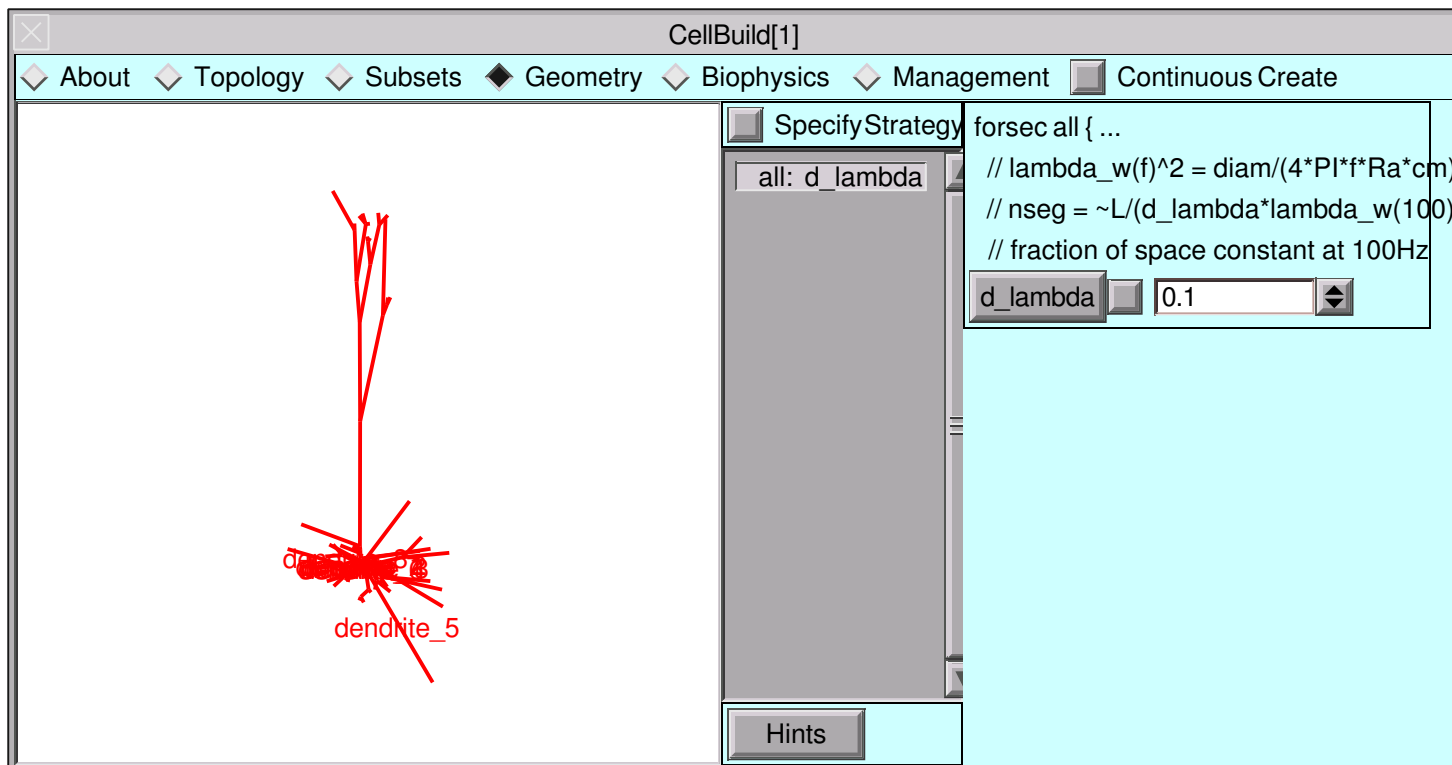
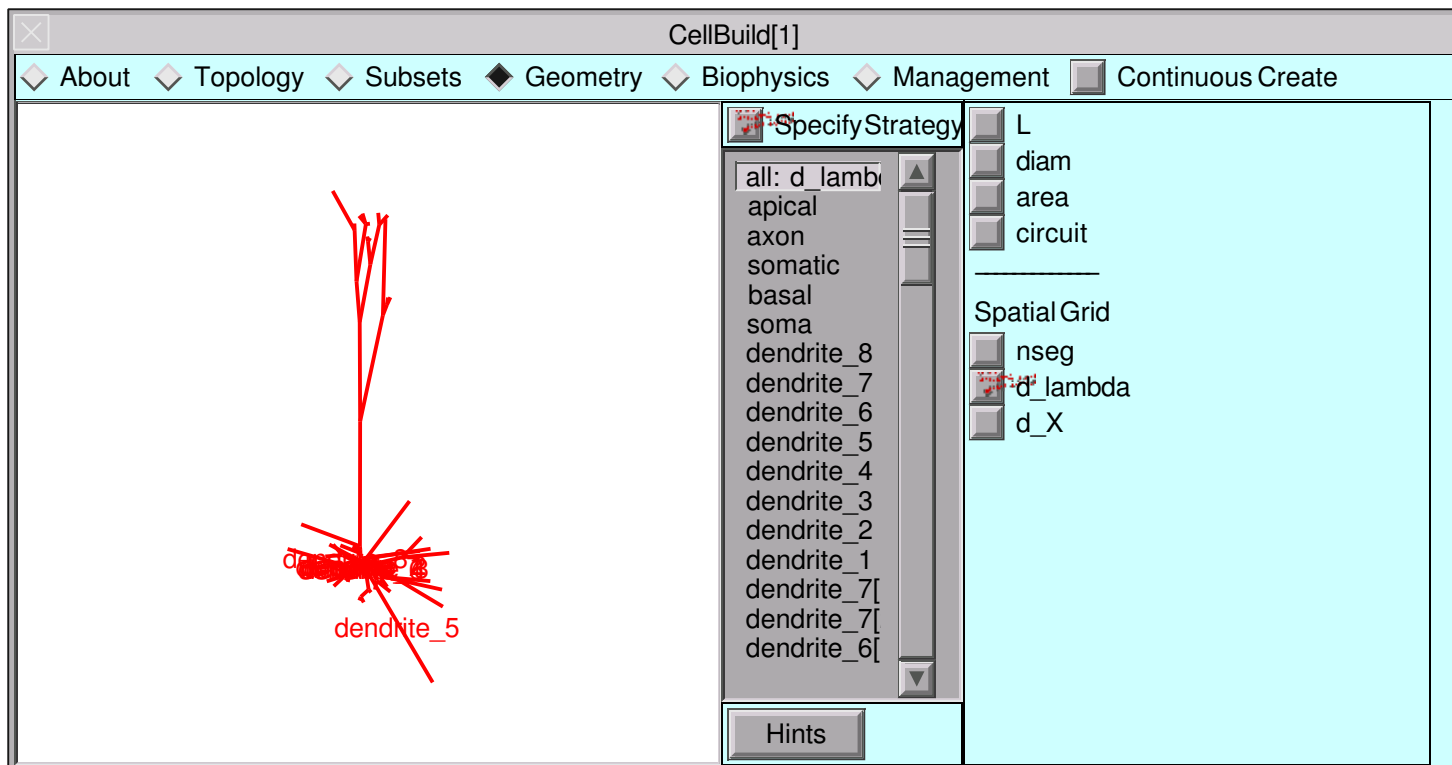
```
cell[15].axon.v(1)
```

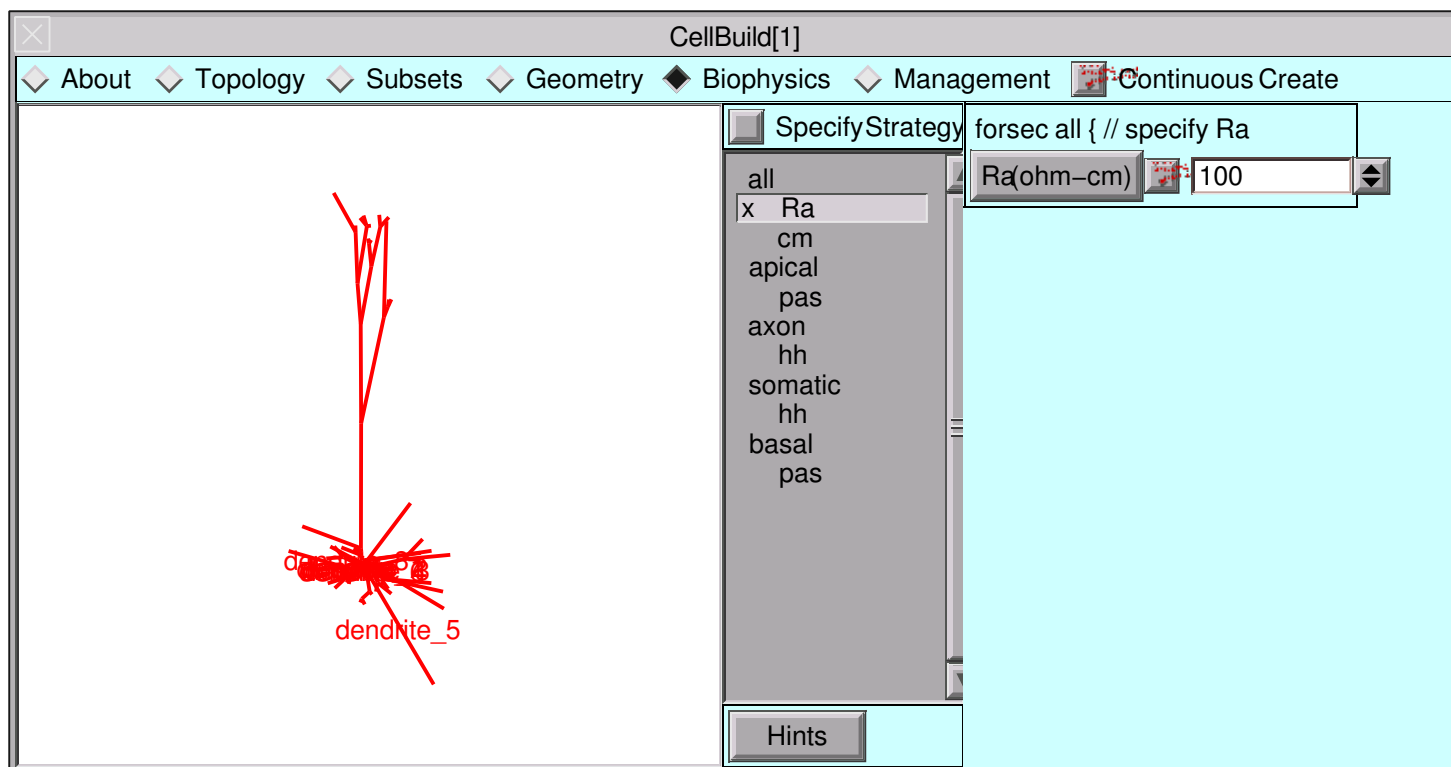
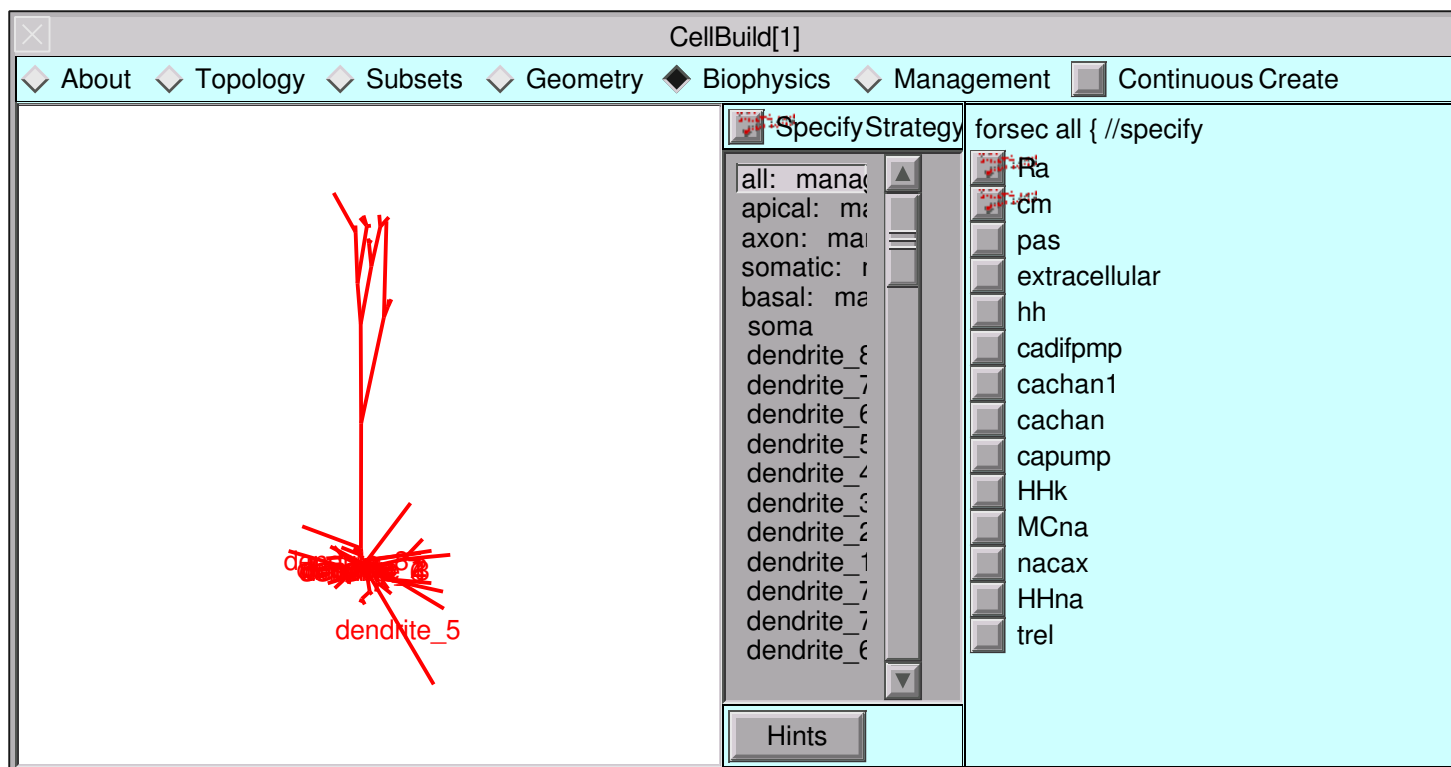
```
proc run() {  
    finitalize(-65)  
    while (t < tstop) {  
        fadvance()  
    }  
}
```

t  **t + dt**









✕

NEURON Main Menu

Iconify

File

Edit

Build

Tools

Graph

Vector

Window

- RunControl
- RunButton
- VariableStepControl
- Point Processes
- Distributed Mechanisms
- Fitting
- Impedance
- Miscellaneous

- Managers
- Viewers

- Point Manager
- Point Group
- Electrode

✕

PointProcessManager

Close

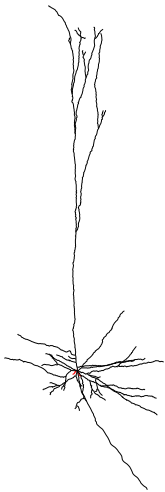
Hide

SelectPointProcess

Show

None

at:soma(0.5)



×

PointProcessManager

CloseHide

SelectPointProcess

none

IClamp

AlphaSynapse

ExpSyn

Exp2Syn

SEClamp

VClamp

OClamp

APCount

NetStim

IntFire1

IntFire2

IntFire4

PointProcessMark

×

PointProcessManager

CloseHide

SelectPointProcess

Show

IClamp[1]
at:soma(0.5)

IClamp[1]

del (ms)

1

dur (ms)

2

amp (nA)

25

i (nA)

0

×

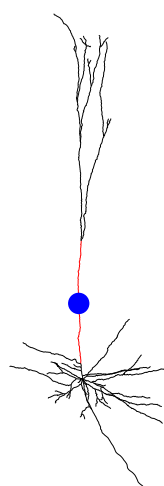
PointProcessManager

CloseHide

SelectPointProcess

Show

IClamp[0]
at:dendrite_1[7](0.5)



RunControl

Init (mV)

Init & Run

Stop

Continue til (ms)

Continue for (ms)

Single Step

t (ms)

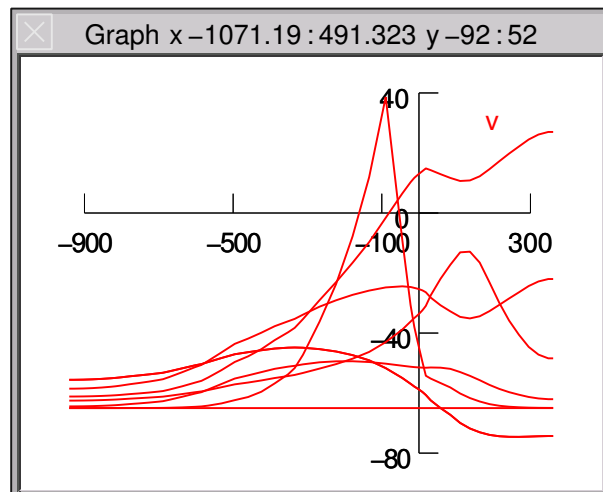
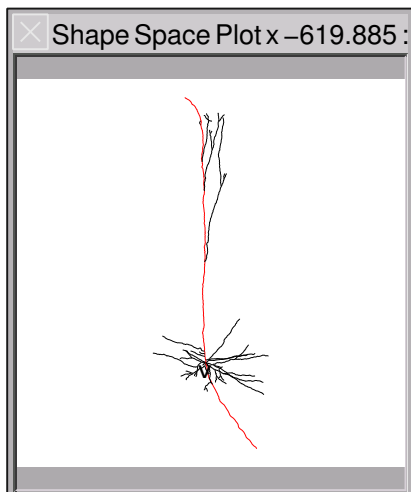
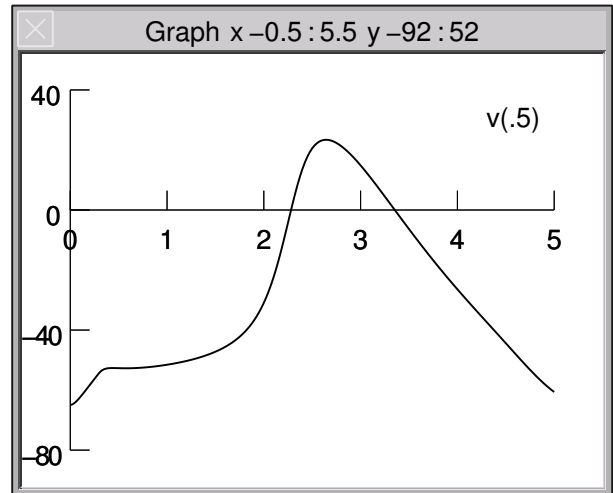
Tstop (ms)

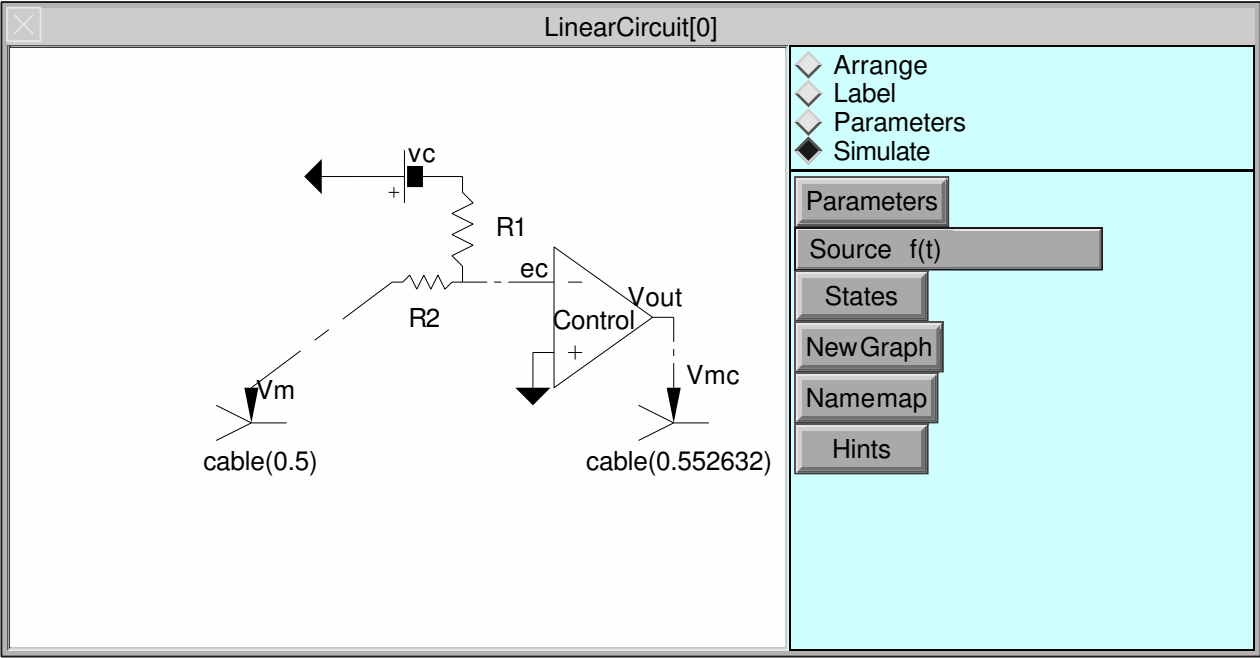
dt (ms)

Points plotted/ms

Quiet

Real Time (s)





Cable

$$C y' + G y = b$$

d a
b d a
b d a
b d a
b d

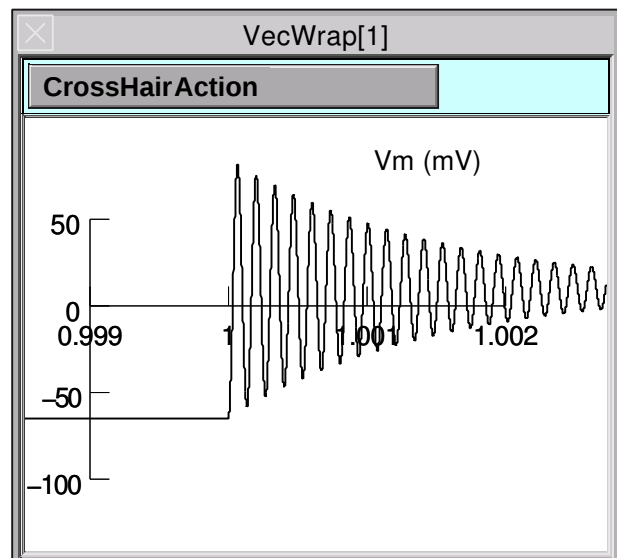
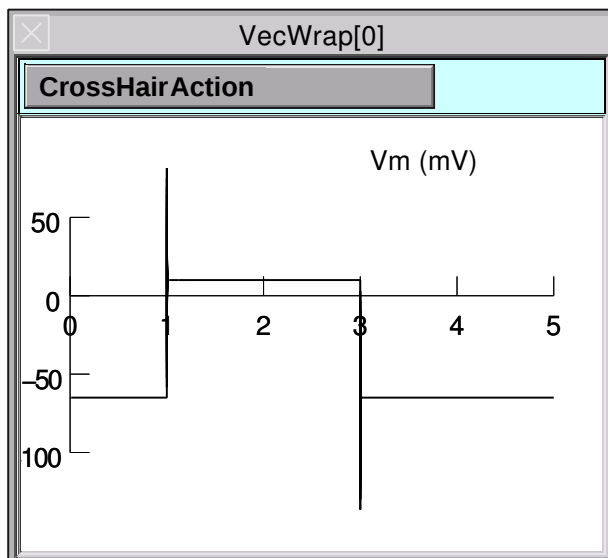
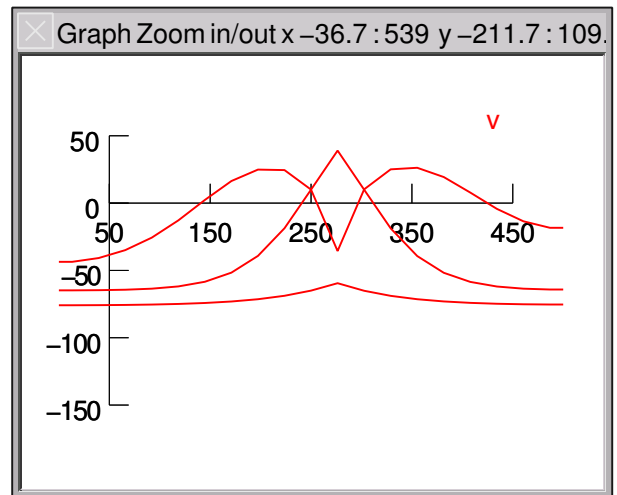
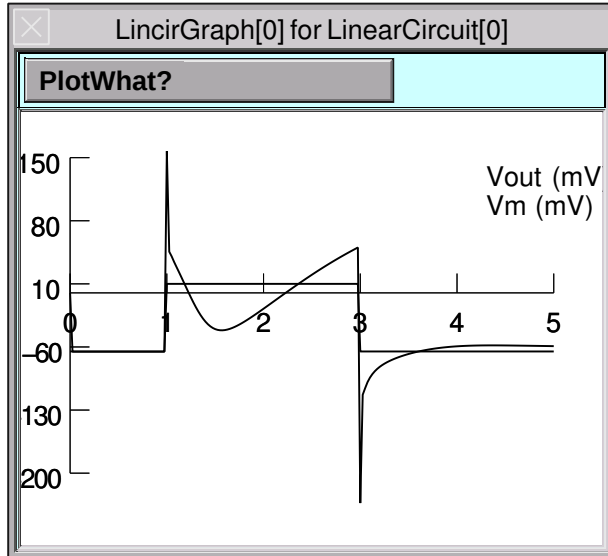
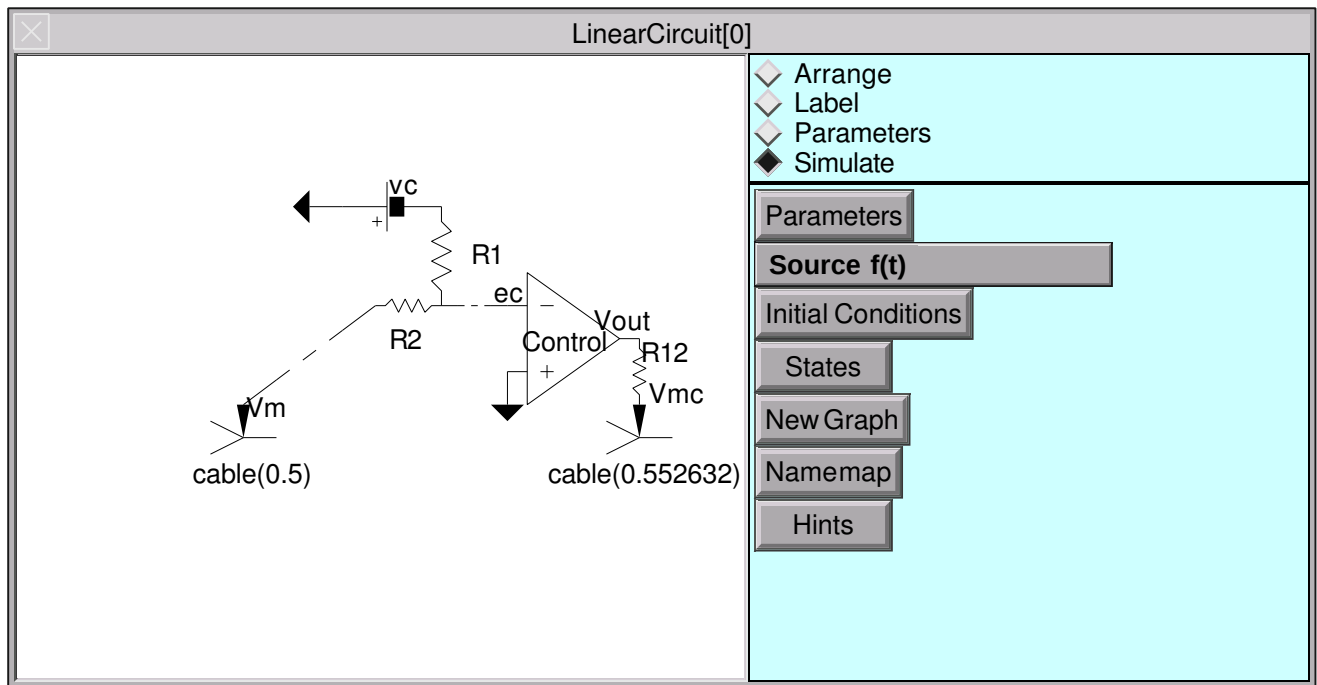
v1
vm
v3
vmc
v5

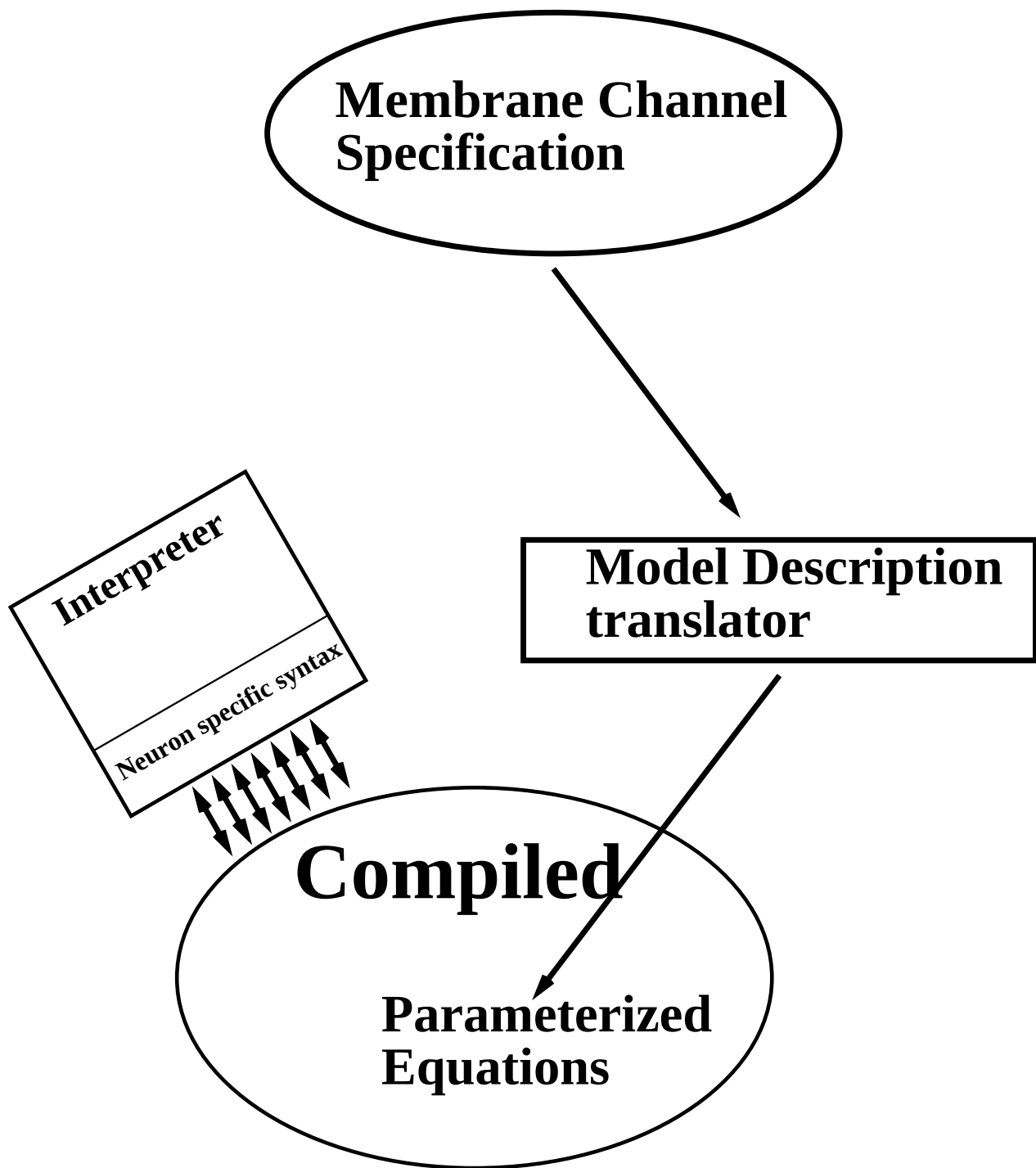
x
x x x
x x x x
x x
x
x

vm
vmc
ec
vc
IC
Ivc

d a
b d a
b d a
b d a
b d
x x x x
x x x
x
x

v1
vm
v3
vmc
v5
ec
vc
IC
Ivc





NMODL

NEURON Model Description Language

Add new membrane mechanisms to NEURON

Density mechanisms

- Distributed Channels
- Ion accumulation

Point Processes

- Electrodes
- Synapses

Described by

- Differential equations
- Kinetic schemes
- Algebraic equations

Benefits

- Specification only -- independent of solution method.
- Efficient -- translated into C.
- Compact
 - One NMODL statement -> many C statements.
 - Interface code automatically generated.
- Consistent ion current/concentration interactions.
- Consistent Units

NMODL general block structure

What the model looks like from outside

```
NEURON {  
    SUFFIX kchan  
    USEION k READ ek WRITE ik  
    RANGE gbar, ...  
}
```

What names are manipulated by this model

```
UNITS { mV = (millivolt) ... }  
PARAMETER { gbar = .036 (mho/cm2) <0, 1e9>... }  
STATE { n ... }  
ASSIGNED { ik (mA/cm2) ... }
```

Initial default values for states

```
INITIAL {  
    rates(v)  
    n = ninf  
}
```

Calculate currents (if any) as function of v, t, states

(and specify how states are to be integrated)

```
BREAKPOINT {  
    SOLVE deriv METHOD cnexp  
    ik = gbar * n^4 * (v - ek)  
}
```

State equations

```
DERIVATIVE deriv {  
    rates(v)  
    n' = (ninf - n)/ntau  
}
```

Functions and procedures

```
PROCEDURE rates(v(mV)) {  
    ...  
}
```

Ion Channel

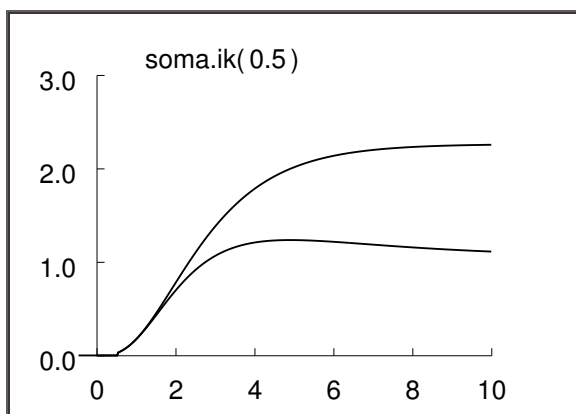
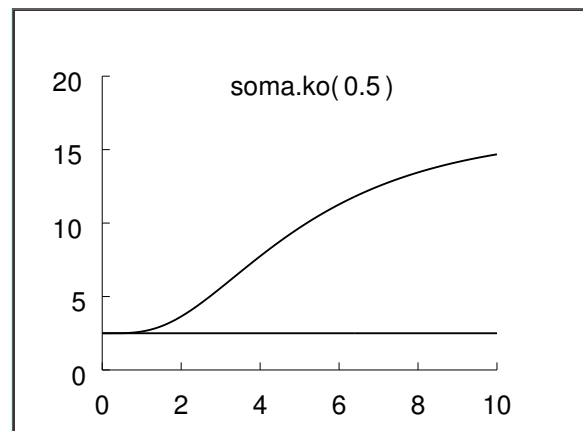
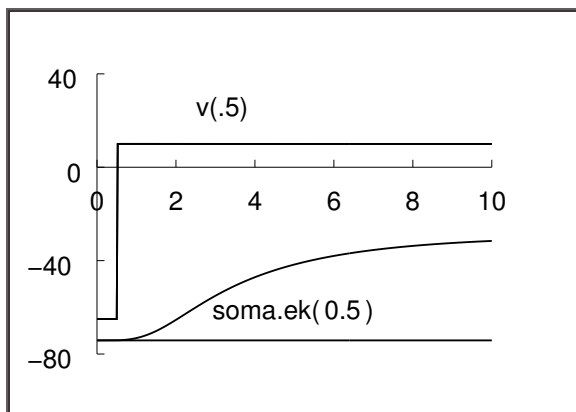
```

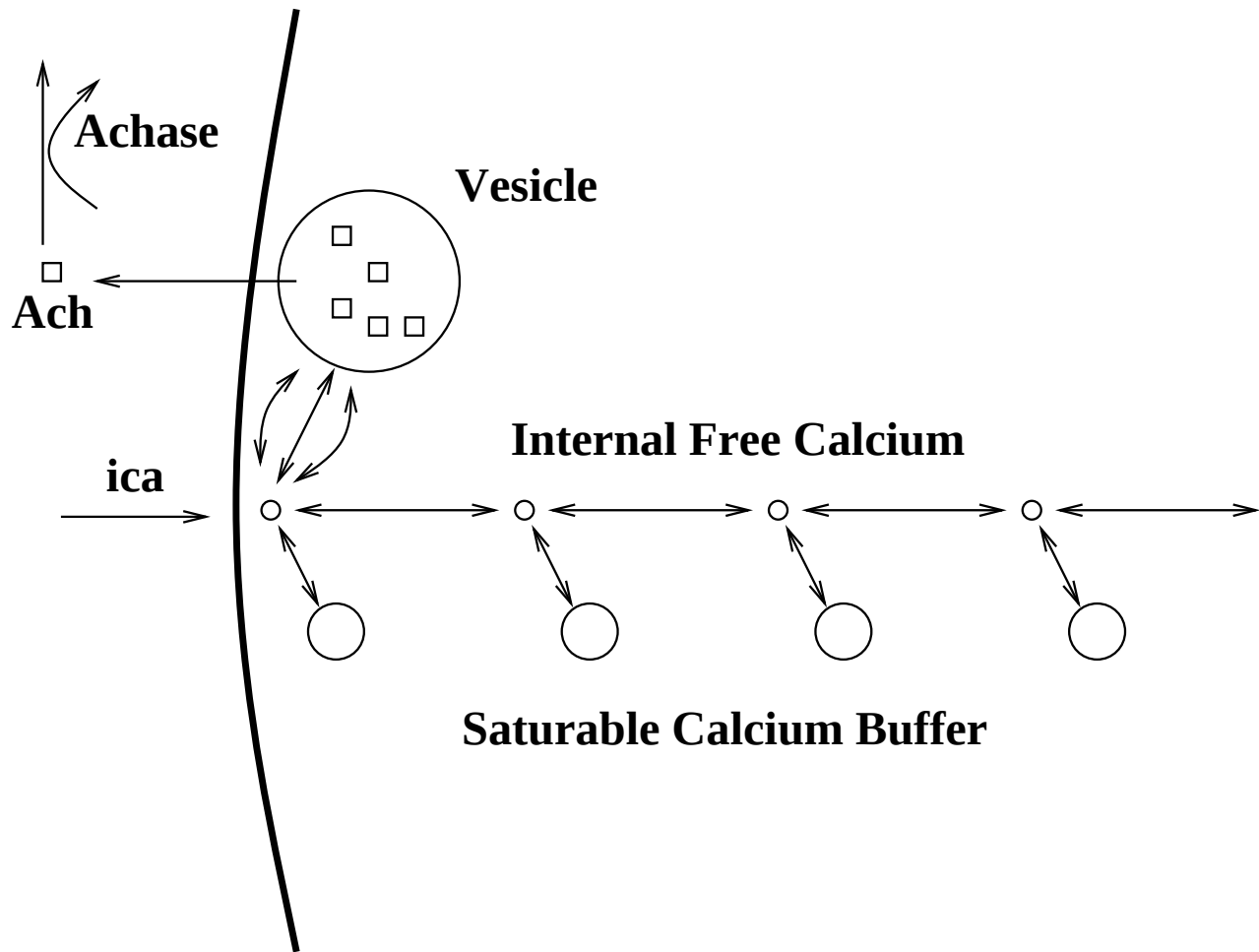
NEURON {
  USEION k READ ek WRITE ik
}
BREAKPOINT {
  SOLVE states METHOD cnexp
  ik = gbar*n*n*n*n*(v - ek)
}
DERIVATIVE states {
  rate(v*1(/mV))
  n' = (inf - n)/tau
}
  
```

Ion Accumulation

```

NEURON {
  USEION k READ ik WRITE ko
}
BREAKPOINT {
  SOLVE state METHOD cnexp
}
DERIVATIVE state {
  ko' = ik/fhspace/F*(1e8)
  + k*(kbath - ko)
}
  
```





```

STATE {
Vesicle Ach Achase Ach2ase X Buffer[N] CaBuffer[N] Ca[N]
}

KINETIC calcium_evoked_release {
  : release
  ~ Vesicle + 3Ca[0] <=> Ach      (Agen, Arev)
  ~ Ach + Achase <=> Ach2ase      (Aase2, 0)  :idiom for enzyme reaction
  ~ Ach2ase <=> X + Achase         (Aase2, 0)  : requires two reactions

  : Buffering
  FROM i = 0 TO N-1 {
    ~ Ca[i] + Buffer[i] <=> CaBuffer[i]    (kCaBuffer, kmCaBuffer)
  }

  :Diffusion
  FROM i = 1 TO N-1 {
    ~ Ca[i-1] <=> Ca[i]                  (Dca*a[i-1], Dca*b[i])
  }

  : inward flux
  ~ Ca[0] << (ica)
}

```

UNITS Checking

```
NEURON { POINT_PROCESS Shunt ... }

PARAMETER {
    e = 0 (millivolt)
    r = 1 (gigaohm) <1e-9,1e9>
}

ASSIGNED {
    i (nanoamp)
    v (millivolt)
}

BREAKPOINT {
    i = (v - e)/r
}
```

Units are incorrect in the "i = ..." current assignment.

The output from

```
modlunit shunt
```

is:

```
Checking units of shunt.mod
The previous primary expression with units: 1-12 coul/sec
is missing a conversion factor and should read:
    (0.001)*()
at line 14 in file shunt.mod
    i = (v - e)/r<>
```

To fix the problem replace the line with:

```
i = (.001)*(v - e)/r
```

What conversion factor will make the following consistent?

$$\frac{\text{nai}'}{(\text{uM/ms})} = \frac{\text{ina}}{(\text{mA/cm}^2)} \div \frac{\text{FARADAY} * (\text{c/radius})}{(\text{coulomb/mole}) \div (\text{um})}$$