

# Cell Simulation Software



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Bioengineering 6003 Cellular Electrophysiology & Biophysics

## Software for Cell Simulation

Results 1 - 10 of about 344,000 for [action potential simulation software](#)

1. [Propagated Action Potential](#) This program simulates the generation and propagation of the **action potential** by solving the Hodgkin and Huxley (HH) model for the space-clamped (membrane ...

2. [nerve.bsd.uchicago.edu/nerve1.html](http://nerve.bsd.uchicago.edu/nerve1.html) - 7k - [Cached](#) - [Similar pages](#) -

### [iROCKET Learning Module: Cardiac Action Potential Simulator](#)

The Cardiac **Action Potential Simulator** aims to conduct a graphical review of cardiac membrane physiology and to demonstrate the effects of common ...

[missinglink.ucsf.edu/lm/cardiac\\_action\\_potential\\_sim/](http://missinglink.ucsf.edu/lm/cardiac_action_potential_sim/) - 10k

### [Simulation of developmental changes in action potentials with ...](#)

Computational Challenges in Cell **Simulation**: A **Software** Engineering Approach · ArticleReference **Simulation** of developmental changes in **action potentials** ...

[www.e-cell.org/ecell/bibliographyfolder/itoh\\_simulation\\_2007](http://www.e-cell.org/ecell/bibliographyfolder/itoh_simulation_2007) - 53k

### [Software for characterizing the ionic basis of the molluscan ...](#)

**simulation software** will be useful for developing hypotheses ... Recordings of **action potentials** in the molluscan cardiac muscle by use of the single- ... [ieeexplore.ieee.org/iel5/6792/18228/00842415.pdf?arnumber=842415](http://ieeexplore.ieee.org/iel5/6792/18228/00842415.pdf?arnumber=842415)



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# Computing Action Potentials

## Compute new ion currents

```
[iNa(i),gNa(i),mNa,hNa,jNa] = INa(V(i),th,mNa,hNa,jNa);
[isi(i),gsi(i),msi,hsi] = Isi(V(i),th,msi,hsi);
[iK(i),gK(i),mK] = IK(V(i),th,mK);
[iK1(i),gK1(i),ht,tauht] = IK1(V(i));
[iKp(i),gKp(i)] = IKp(V(i));
[ib(i),gb(i)] = Ib(V(i));
```

## Compute total current (I<sub>d</sub> = stimulus)

```
I = Id+iNa(i)+isi(i)+iK(i)+iK1(i)+iKp(i)+ib(i);
```

## Compute incremental voltage change

```
dV = -I/Cm;
```

## Update value of voltage and time

```
V(i+1) = V(i) + dV*th;
```

```
t(i+1) = t(i) + th;
```

## Increment counter

```
i = i+1;
```



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# Computing the Currents: I<sub>Kp</sub> (plateau)

$$i_{Kp} = g\bar{K}_p K_p (V_m - E_K)$$

$$K_p = \frac{1}{1 + e^{\frac{7.488 - V_m}{5.98}}}$$

```
function [I,gKp] = IKp(V)
% IKp Plateau potassium current
% I = IKp(V)
% I potassium current
% gKp conductance
% V membrane potential (mV)
% Reference: QN:Luo94a
```

```
global R T F K_o K_i;
GKp_ = 0.0183;
EKp = (R*T/F)*Log(K_o/K_i);
```

```
numTimeStep = length(V);
```

```
for i=1:numTimeStep
Kp = 1/(1+exp((7.488-V(i))/5.98));
gKp(i) = GKp_*Kp;
I(i) = gKp(i)*(V(i)-EKp);
end
```



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## Computing the Currents: IK (time dependent)

$$i_K = \bar{g}_K m h (V_m - E_K)$$

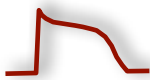
$$\bar{g}_K = g_K \sqrt{\frac{[K^+]_e}{5.4}}$$

$$h = 1.0 \quad h = 2.837 \frac{e^{0.04(V_m + 77)} - 1}{(V_m + 77)e^{0.04(V_m + 35)}}$$

```
function [I,gK,mm] = IK(V,th,m)
GK_ = 0.282*sqrt(K_o/5.4); % millisiemens/uF
EK = (R*T/F)*log((K_o+PNa_K*Na_o)/(K_i+PNa_K*Na_i)); % mV

if V > -100
    h = 2.837*(exp(0.04*(V+77))-1)/((V+77)*exp(0.04*(V+35)));
else
    h = 1;
end

gK = GK_*m*h;
I = gK*(V-EK);
```



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## Computing the Currents: IK activation

$$m = m_\infty(v_o) - (m_\infty(v_o) - m_0)e^{-t/\tau_m(v_o)}$$

$$\tau_m = \frac{1}{\alpha_m(V) + \beta_m(V)} \quad m_\infty = \frac{\alpha_m(V)}{\alpha_m(V) + \beta_m(V)}$$

$$\alpha_m = 0.005 \left( \frac{e^{0.083(V_M + 50)}}{1 + e^{0.057(V_M + 50)}} \right) \quad \beta_m = 0.0013 \left( \frac{e^{-0.06(V_M + 20)}}{1 + e^{-0.04(V_M + 20)}} \right)$$

```
am = 0.0005*exp(0.083*(V+50))/(1+exp(0.057*(V+50)));
bm = 0.0013*exp(-0.06*(V+20))/(1+exp(-0.04*(V+20)));

minf = am/(am+bm);
taum = 1/(am+bm);

mm = minf - (minf-m)*exp(-th/taum);
```



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## Computing the Currents: IK1 (time independent)

$$i_{K1} = g_{\bar{K}1} K_{1\infty} (V_m - E_K)$$

$$g_{\bar{K}1} = g_{K1} \sqrt{\frac{[K^+]_e}{5.4}} \quad K_{1\infty} = \frac{\alpha_{K1}}{\alpha_{K1} + \beta_{K1}}$$

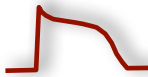
```
function [I,gK1,hinf,tauh] = IK1(V)
GK1_ = 0.6047*sqrt(K_o/5.4);
EK1 = (R*T/F)*log(K_o/K_i);

numTimeStep = length(V);

for i=1:numTimeStep
    ah(i) = 1.02/(1+exp(0.2385*(V(i)-EK1-59.215)));
    bh(i) = (0.49124*exp(0.08032*(V(i)-EK1+5.476)) + ...
        exp(0.06175*(V(i)-EK1-594.31)))/ (1+exp(-0.5143*(V(i)-EK1+4.753)));

    hinf(i) = ah(i)/(ah(i)+bh(i));
    tauh(i) = 1/(ah(i)+bh(i));

    gK1(i) = GK1_*hinf(i);
    I(i) = gK1(i)*(V(i)-EK1);
end
```



## Optimization of Code

- Time step

```
I = Id+iNa(i)+isi(i)+iK(i)+iK1(i)+iKp(i)+ib(i);
dV = -I/Cm;
th = TSmín*dVmax/abs(dV);
if th > TSmáx
    th = TSmáx;
elseif th < TSmín
    th = TSmín;
end
```



# DEMO

<http://thevirtualheart.org/>

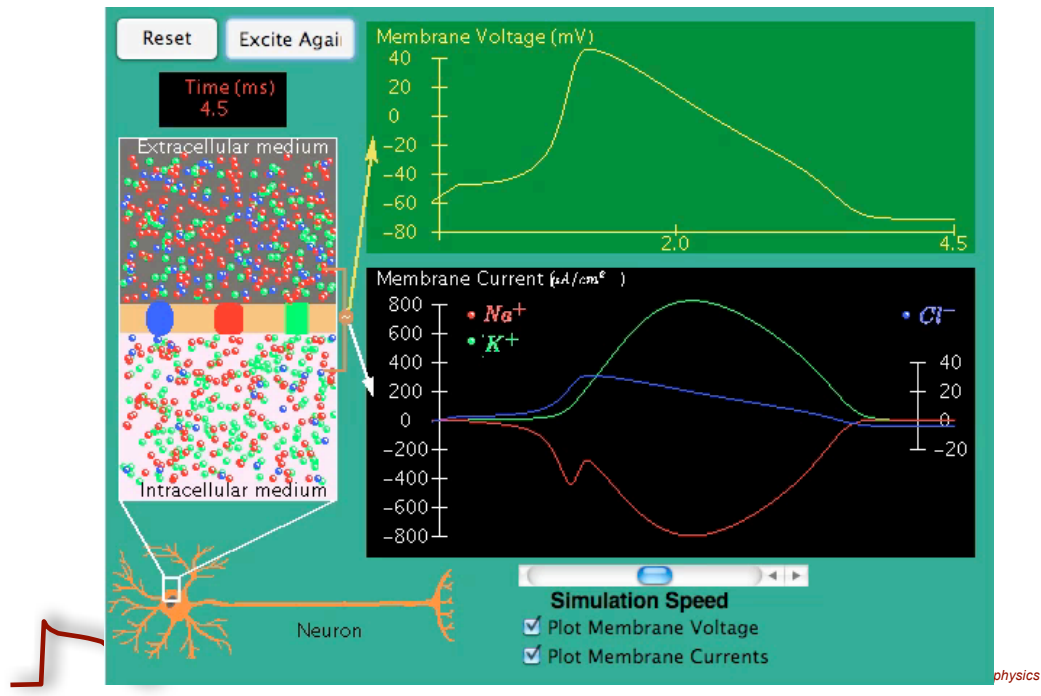


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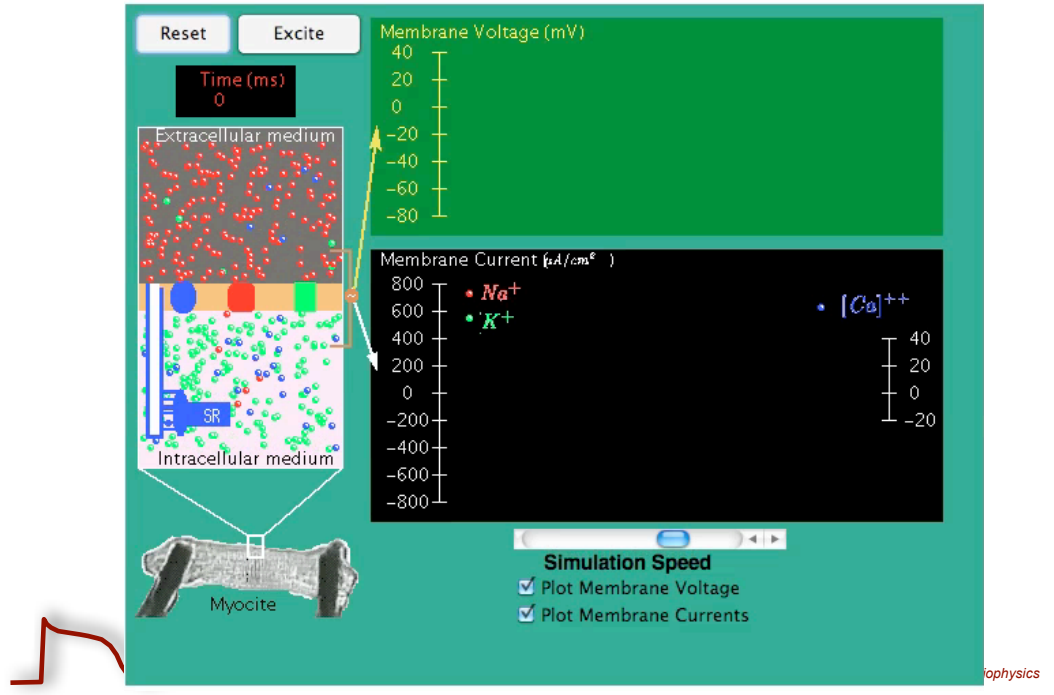
## Virtual Heart Examples

### Nerve Action Potential



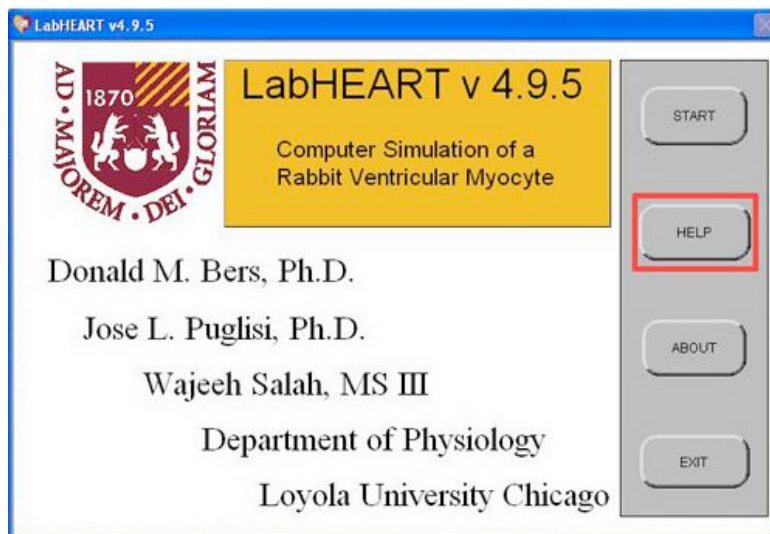
# Virtual Heart Examples

## Cardiac Action Potential



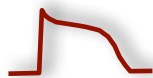
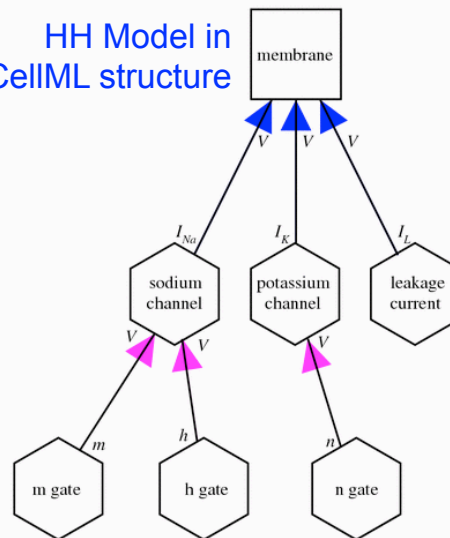
# LabHeart

<http://www.labheart.org>



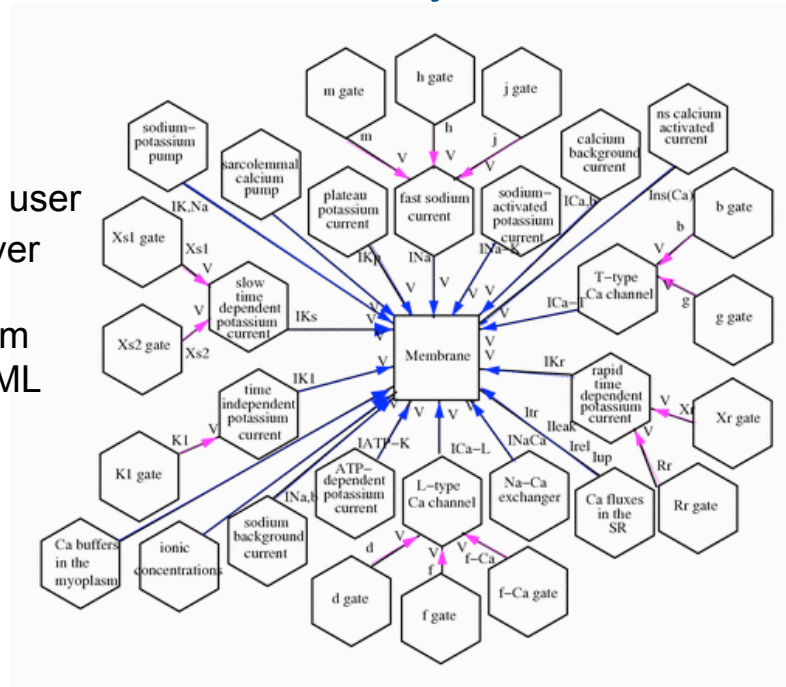
- [www.cellml.org](http://www.cellml.org)
- Standardized interface to cell simulations
- Includes geometry, electrophysiology and signal transduction
- Based on markup language (ML) standards
- Part of larger set of ML's for simulation

HH Model in CellML structure



## CellML Luo-Rudy II

- Additional complexity hidden from user
- Software layer provides interface from user to CellML functions



# CellML + JSim

- Tcl script based interface to CellML functions

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**DRGL J-Sim**

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