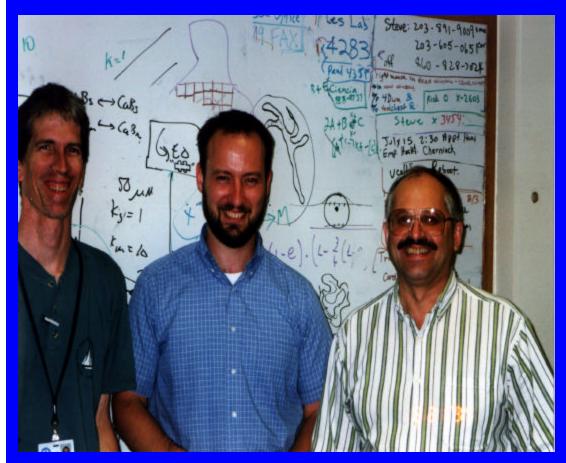
"Every attempt to employ mathematical methods in the study of biological questions must be considered profoundly irrational and contrary to the spirit of biology.

"If mathematical analysis should ever hold a prominent place in biology - an aberration which is happily almost impossible - it would occasion a rapid and widespread degeneration of that science"

-- Auguste Comte, Pilosophie Positive, 1830

The Virtual Cell Project

Jim Schaff Boris Slepchenko



Chuck Fink Yung-Sze Choi **Talitha Washington Diana Resasco** John Carson Frank Morgan John Wagner Ion Moraru **Jim Watras** Mark Terasaki **Dong-Guk Shin** NIGMS, NSF

Features of the VC Modeling Environment

N		
File Tools Connection Help		
Physiology Geometry Simulation Con	text Simulation	2
MathDescription: NE_Full_generated		
VCML Editor C Equation Viewer CODE S	Simulator PDE Simulator	N
Time Step 0.01 End Time	105.0 Save interval 0.5	Run Stop
Valid Spatial Model, PDE solver ready		
Variable	Time C Spatial Plot	Time Plot
RactCa 2.0 Ca_ER 2.5 Fura2_bound 3.0	Refresh List	size = (167,63,1)
CalciumBufferBound 3.5	_	
Calcium . 4.5	<u>-</u>	
min = 0.0	max = 5.319894054213793	
AXIS	slice	C Auto Scale
• XY • YZ • ZX	- 0 +	Manual Scale
		Min 0.0
		0.0
		Max 5.3198940
		5.0
		C Grayscale
		Blue to red
🖆 Unsigned Java Applet Window		

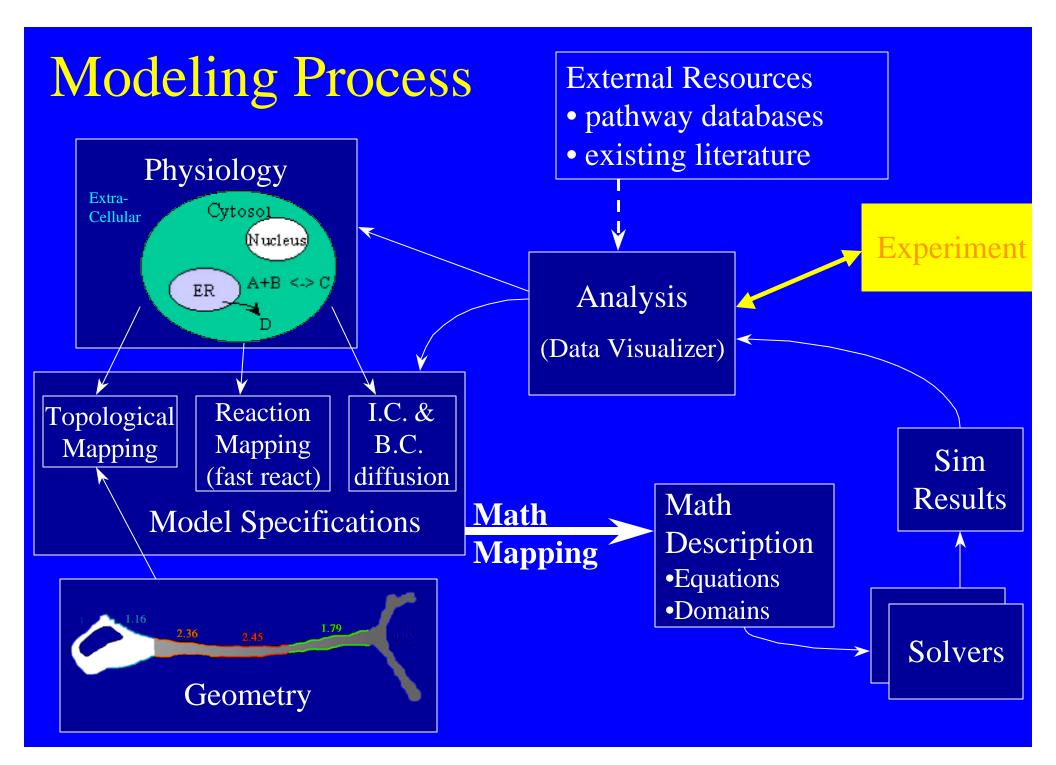
- Designed to be used Interactively with Experiment
- Enables Construction and Testing of Complex Models of Rapid Investigation of Simple Hypotheses
- Geometry from Experimental Images
- Math, Physics, and Numerics are Transparent to an Experimentalist while fully Accessible to a Theorist

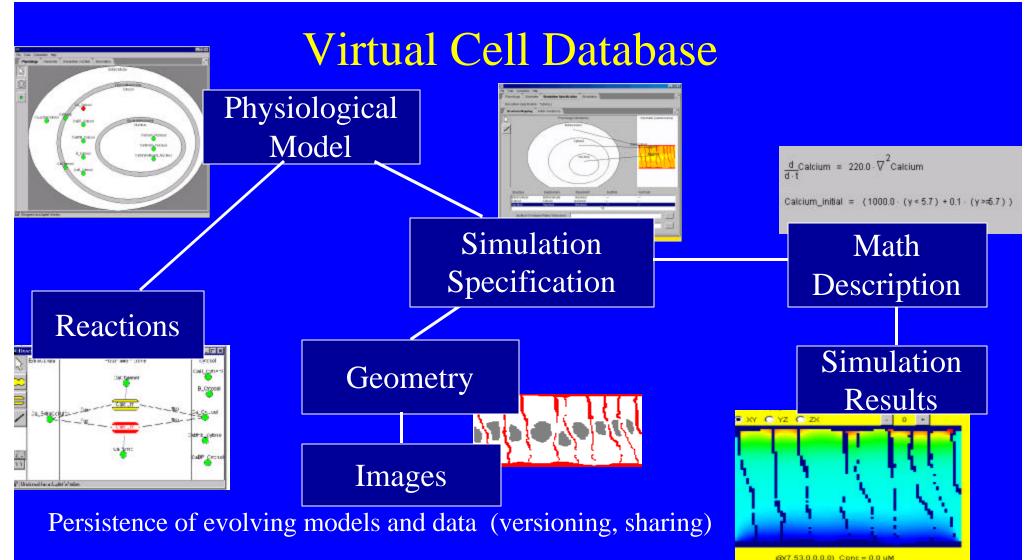
Math and Physics

$$\boldsymbol{F}_{i} = -D_{i}\nabla C_{i} - z_{i}\boldsymbol{m}_{i}C_{i}\nabla\Phi, \qquad \boldsymbol{m}_{i} = \frac{D_{i}F}{RT}$$

$$A + B \quad \stackrel{k_1}{\longleftrightarrow} \quad C, \qquad R_i = k_1 [A] [B] - k_{-1} [C]$$

$$\frac{\P C_i}{\P t} = -\text{div} \boldsymbol{F}_i + R_i$$

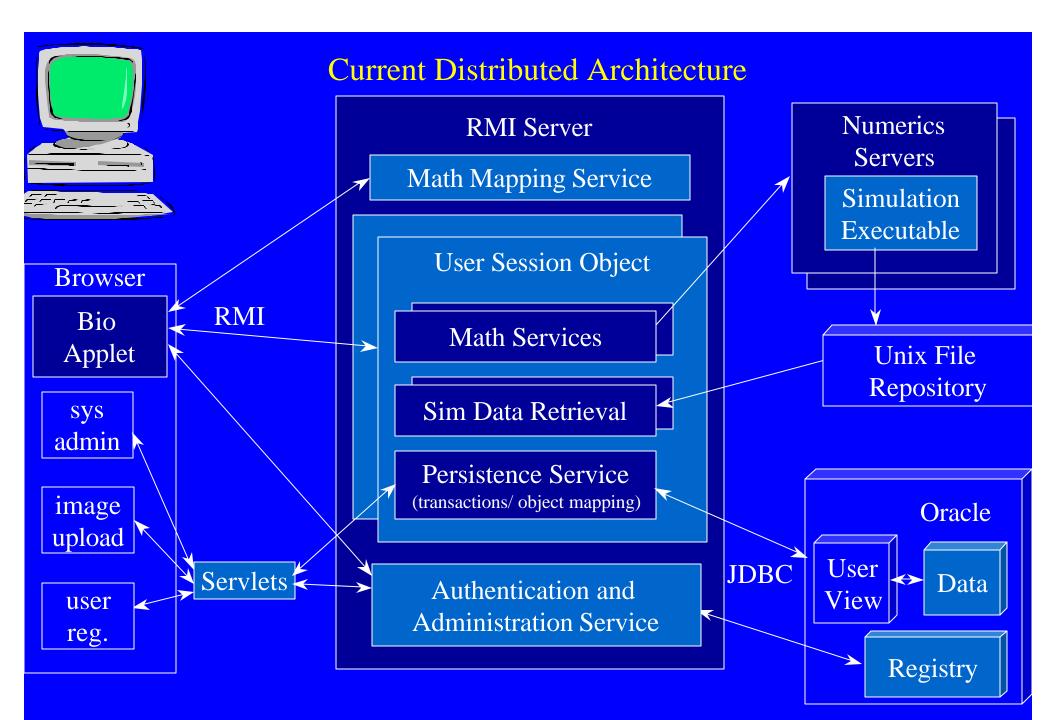




Allows future integration with WWW accessible resources

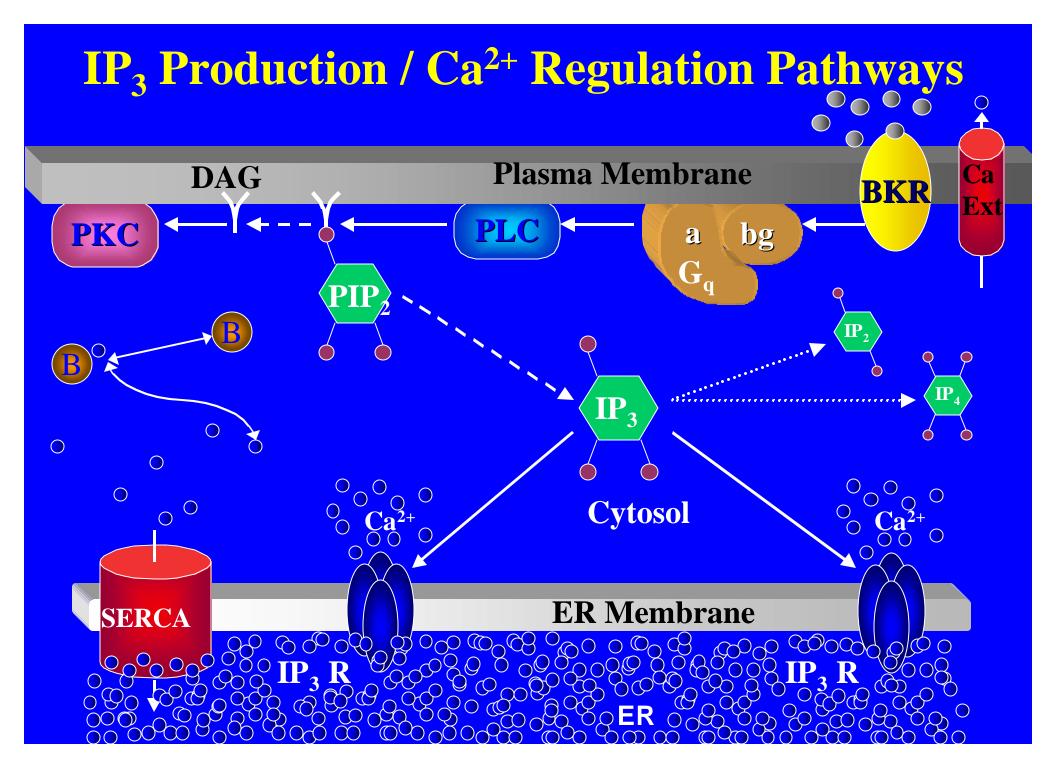
- Pathway databases (EcoCyc, KEGG, EMP)
- Other modeling tools

Requires development of standard vocabularies for modeling entities.



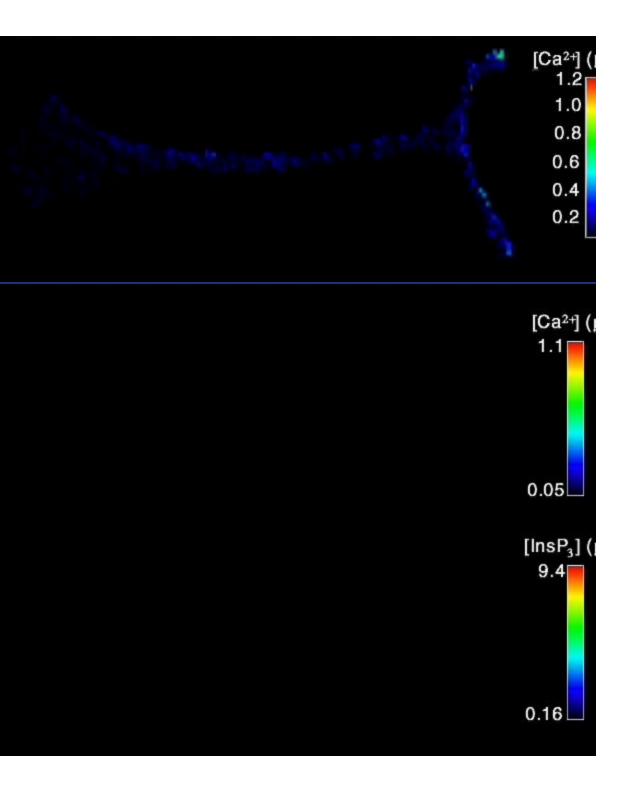
Why?

- Explore the fit of mechanistic hypotheses
- Make new predictions that can be tested experimentally
 - Results could either strengthen a model or force us to modify hypotheses
- Show species that can't be visualized experimentally
- Examples: calcium wave in a neuronal cell, long term depression in cerebellar Purkinje cells, fertilization-wave in frog egg, mitochondrial respiration, RNA granule trafficking, nuclear envelope breakdown, epithelial cell transport.

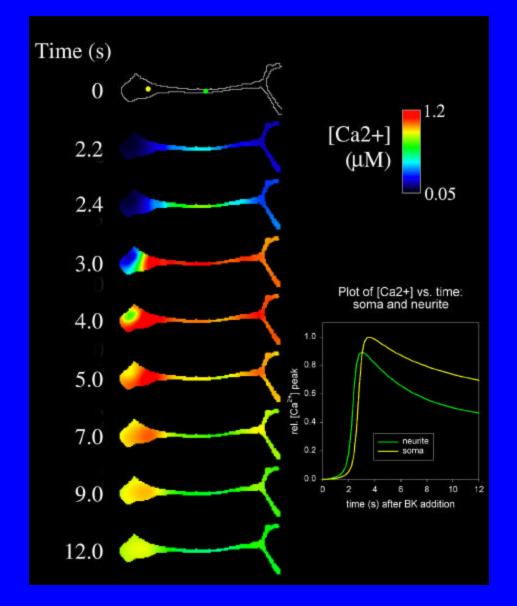


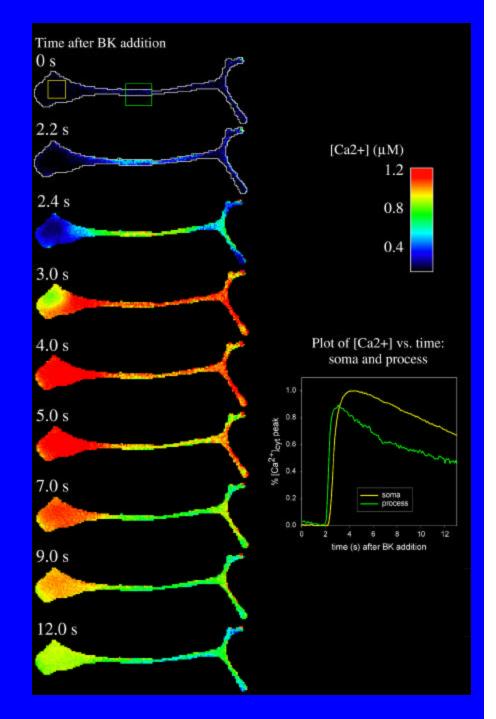
Bradykinin Induced Calcium Wave in a Neuroblastoma Cell

Modeling Result for Ca²⁺ and InsP₃



Comparison of Simulation and Experiment





"What are we to do with the enormous cornucopia of genes and molecules we have found in living cells? How can we see the wood for the trees and understand complex cellular processes? Although we poor mortals have difficulty manipulating seven things in our head at the same time, our silicon protégés do not suffer this limitation. ... The data are accumulating and the computers are humming. What we lack are the words, the grammar and the syntax of the new language." -- Dennis Bray, 1997