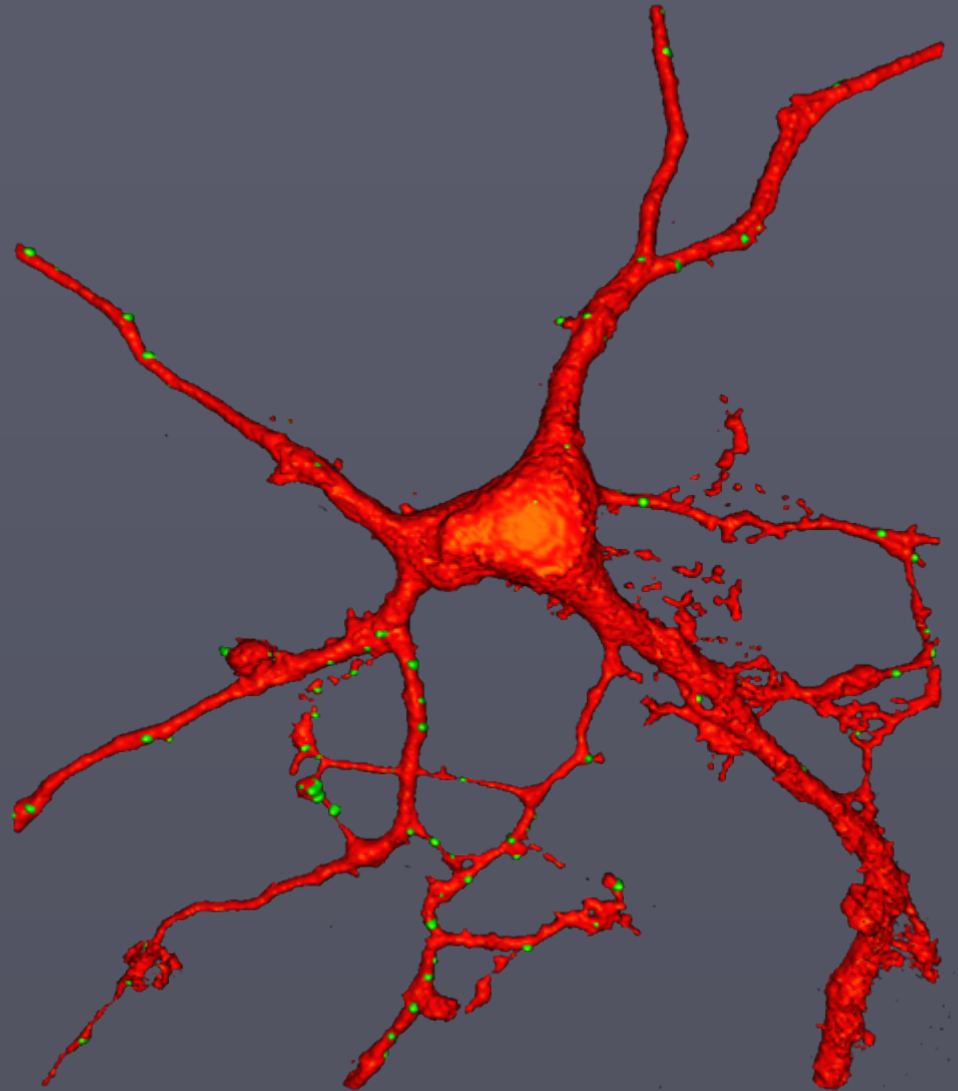
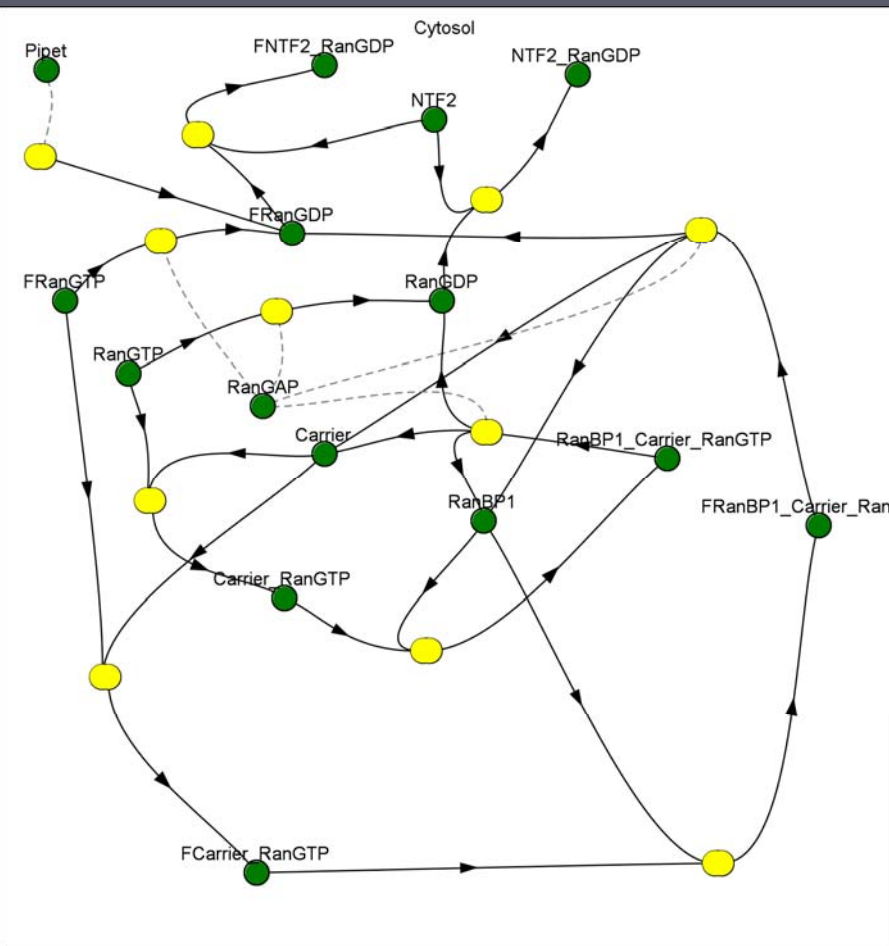


# Space in VCell (“Virtual Cell”)

– Mapping Structures to Geometries –

<http://vcell.org>

# *What is a Model?*

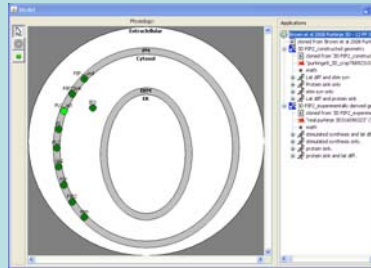


# VCell Top-Level Documents

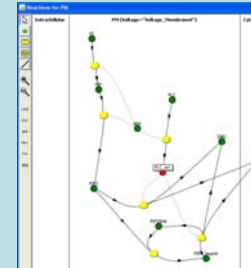
- Database object containers
  - BioModel
  - MathModel
  - Geometry
- Referential objects
  - ResultSet
  - FieldData

# Physiology

Molecules Structures (topology)

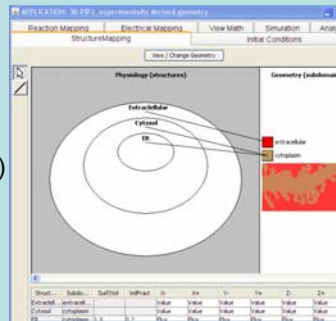


Reactions Fluxes



## Applications

- Structure mapping (topology to geometry)
- Initial Conditions
- Boundary conditions
- Diffusion constants (if spatial)
- Electrophysiology protocols
- Enable/disable reactions
- Fast reactions
- Model analysis
- Stochastic rate conversion



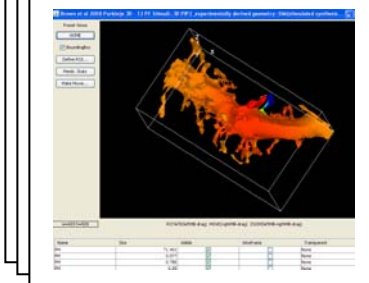
Species	Species Context	Structure	Clamped	Initial Conditions
IP3	IP3_Cyt	Cytosol	<input type="checkbox"/>	0.18
IP3	IP3_In	IP3	<input type="checkbox"/>	([x > 4.024] && [x < 4.32])
PIP2	PIP2_IP3	IP3	<input type="checkbox"/>	4000.0
PIP	PIP_IP3	IP3	<input type="checkbox"/>	200.0
DAG	DAG_IP3	IP3	<input type="checkbox"/>	2000.0
PLC	PLC_IP3	IP3	<input type="checkbox"/>	100.0
PI	PI_IP3	IP3	<input type="checkbox"/>	1.420E+0
PLC_act	PLC_act_IP3	IP3	<input type="checkbox"/>	0.0
PIP2Sink	PIP2Sink_IP3	IP3	<input type="checkbox"/>	2000.0
PIP2_bound	PIP2_bound_IP3	IP3	<input type="checkbox"/>	3000.0

description	Parameter	Expression	Units
Initial concentration	IP3	([x > 4.024] && [x < 4.32]) && 0.18	molecules.um-2
diffusion constant	IP3	0.0	um2.s-1
Boundary Condition X	PLC_IP3		molecules.um-2
Boundary Condition Y	PLC_IP3		molecules.um-2
Boundary Condition Z	PLC_IP3		molecules.um-2



## Results

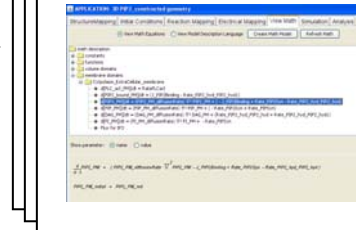


## Simulations

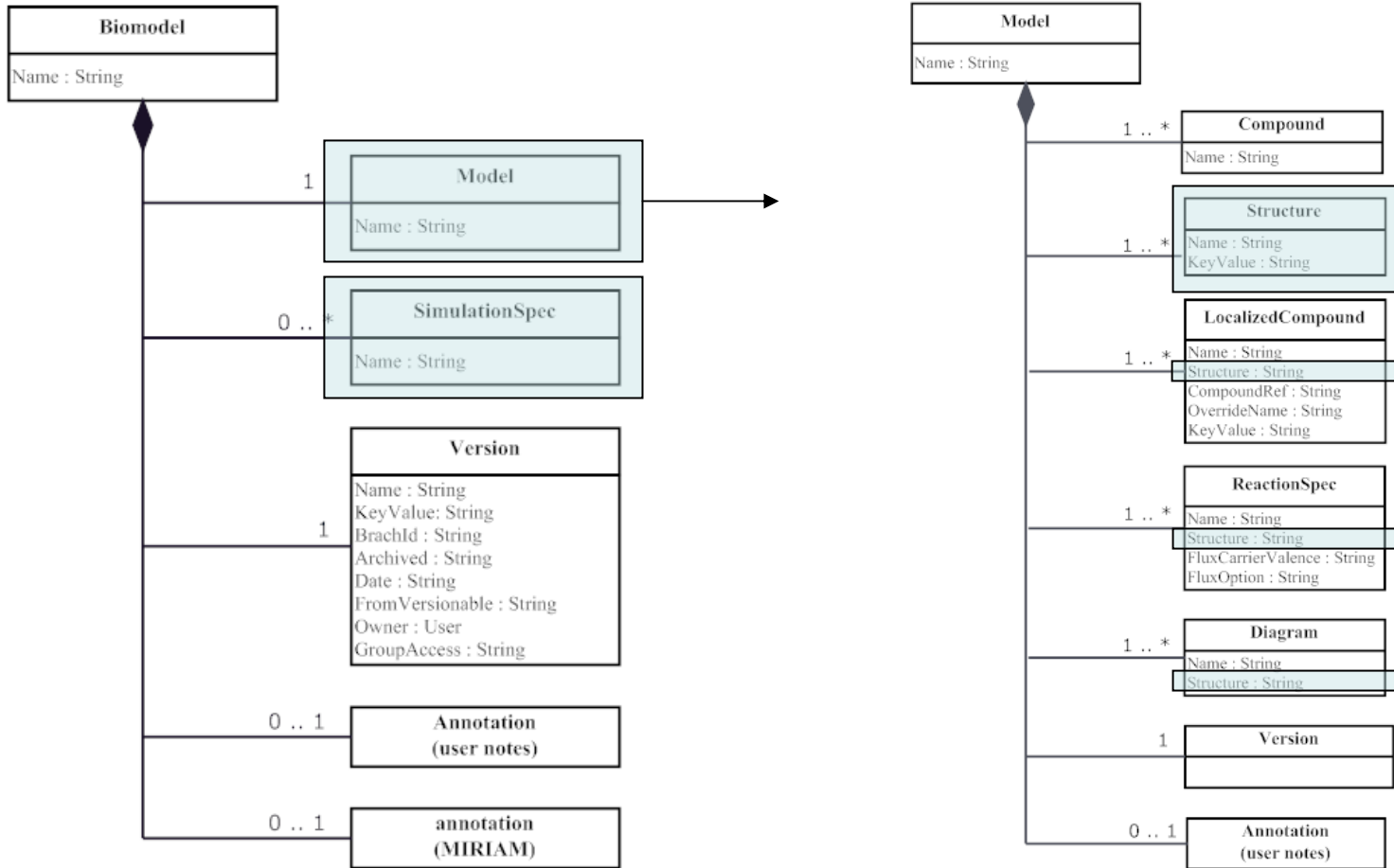
- Timecourse
- Timestep
- Mesh size
- Solver type
- Solver settings
- Parameter changes
- Parameter scans
- Parameter sensitivity

Parameter Name	Default Value	Change Value	Units
IP3	0.18		
PIP2	4000.0		
PIP	200.0		
DAG	2000.0		
PLC	100.0		
PI	1.420E+0		
PLC_act	0.0		
PIP2Sink	2000.0		
PIP2_bound	3000.0		

## Mathematical Description (view-only, automatically generated)

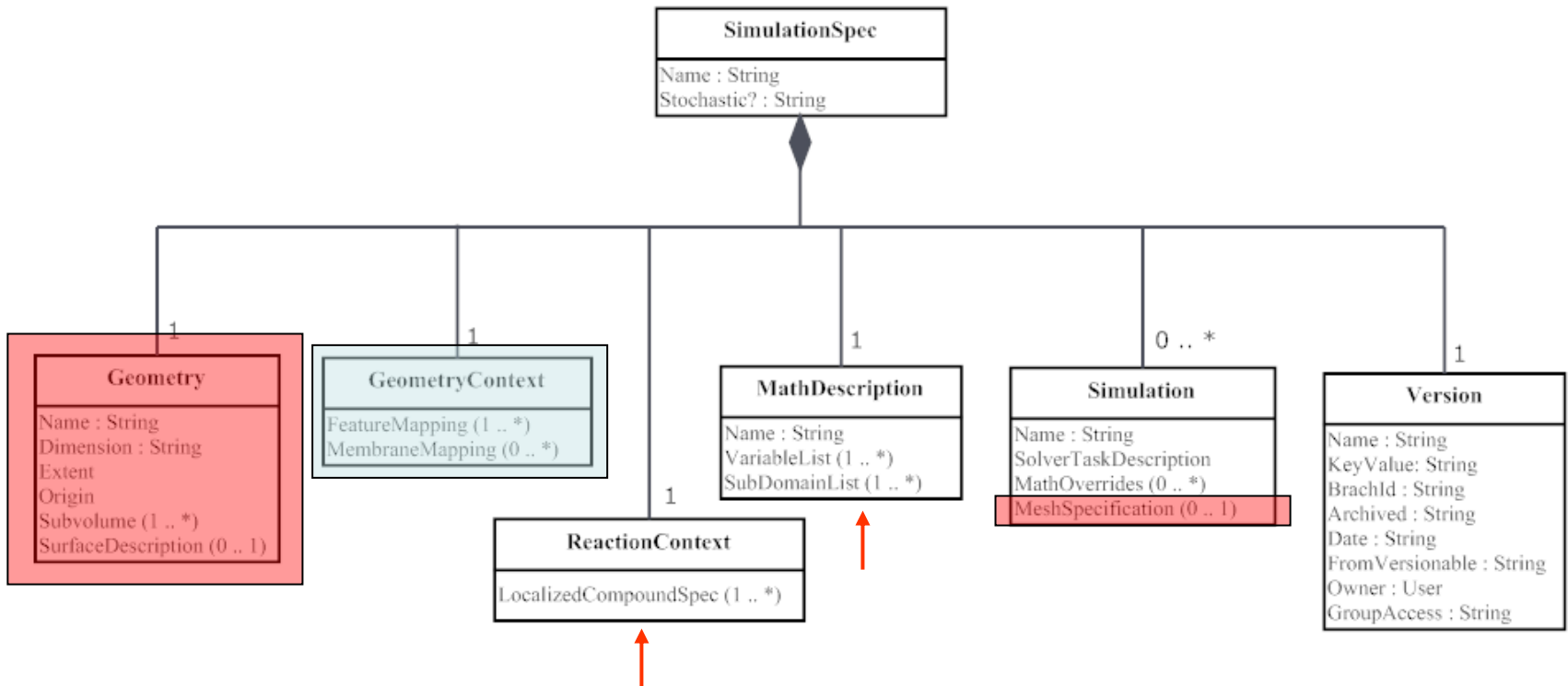


# BioModel Object Hierarchy



# SimulationSpec Element

– a.k.a. “Application” wizard, a.k.a. SimulationContext object –



# Spatially Resolved Species

- Initial Concentrations:
  - PDE:
    - Concentration FIELD
  - Stochastic:
    - Spatial distribution function FIELD
- Boundary Conditions:
  - PDE:
    - specified concentration FIELD (Dirichlet)
    - specified concentration FIELD (Neumann)
    - Periodic
  - Stochastic:
    - specified flux density probability FIELD
    - Periodic
- Diffusion Coefficient:
  - PDE & Stochastic:
    - (An)isotropic (optionally) spatially varying FIELD

# VCell Geometry

## GeometrySpec

(image-based or analytic)

Provides volume sampling  
(surfaces are implicit)

## GeometrySurfaceDesc

- Region Image

(discriminates disjoint regions of same type)

-Volume/Membrane Regions

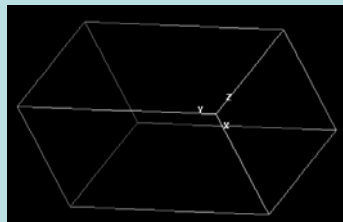
(graph of all volume/surface instances  
with adjacency and size metrics)

-Surface sampling – visualization params

Provides  
Volume  
Sampling  
Function

OR

bounding box  
extent/origin



$$x^2 + y^2 + z^2 < R^2$$

1.0

Analytic  
SubVolume

■ cytosol

■ ec

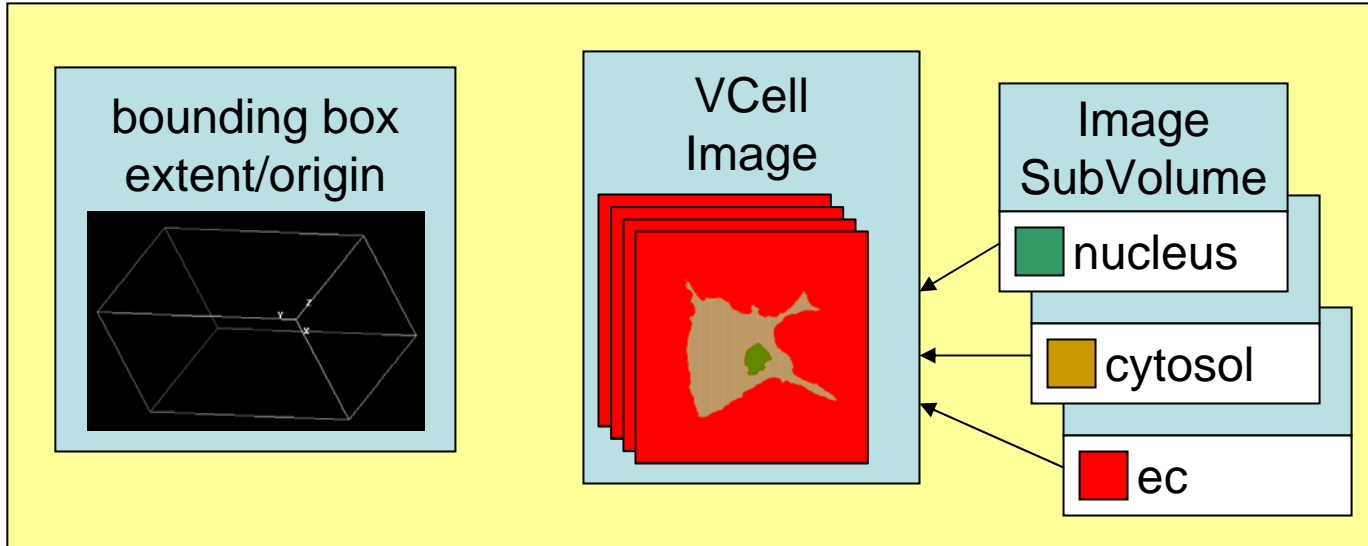
x, y, z



SubVolume

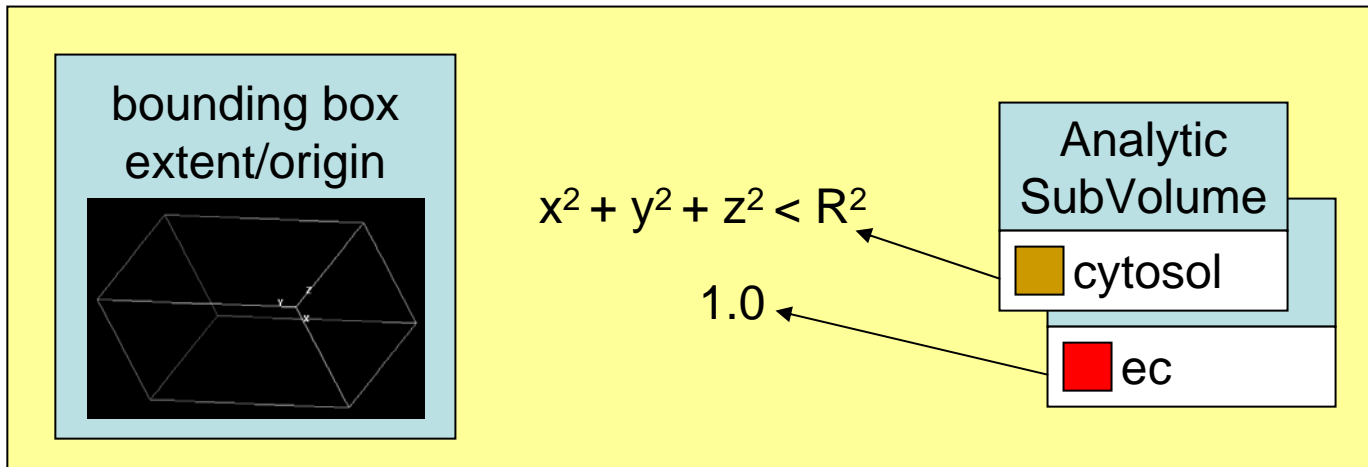


# Geometry Specification



Provides  
Volume  
Sampling  
Function

OR

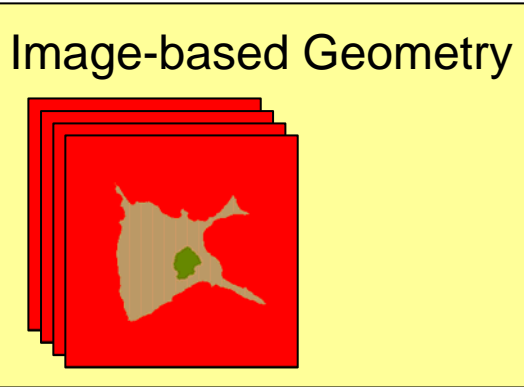


x, y, z

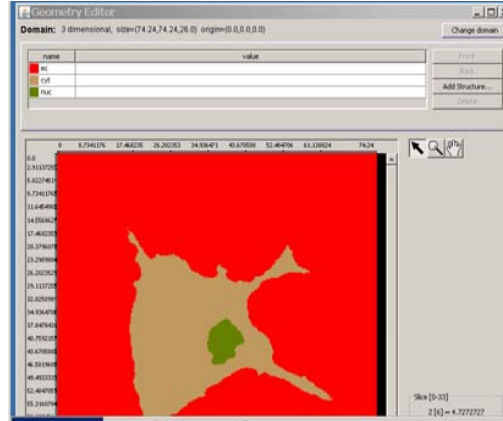


SubVolume

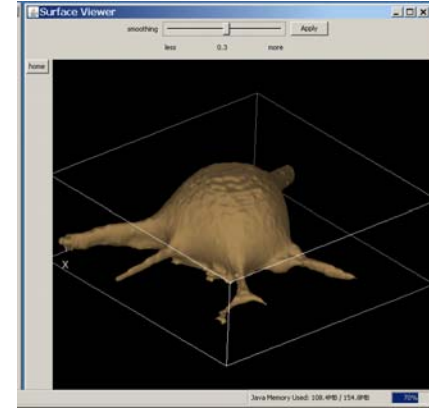
# Geometry Visualization



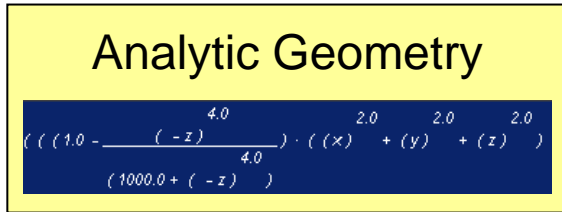
Sampled at original image resolution



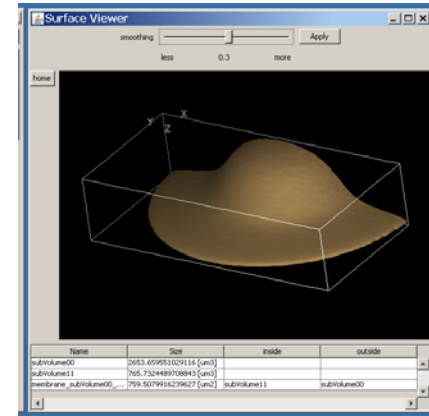
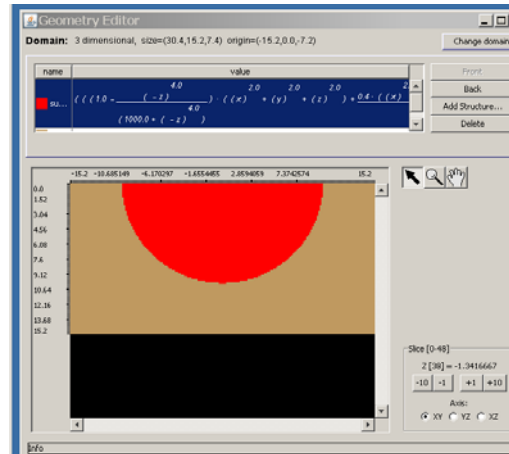
Sampled Volume



Generated Surfaces (smoothed staircase "quads")



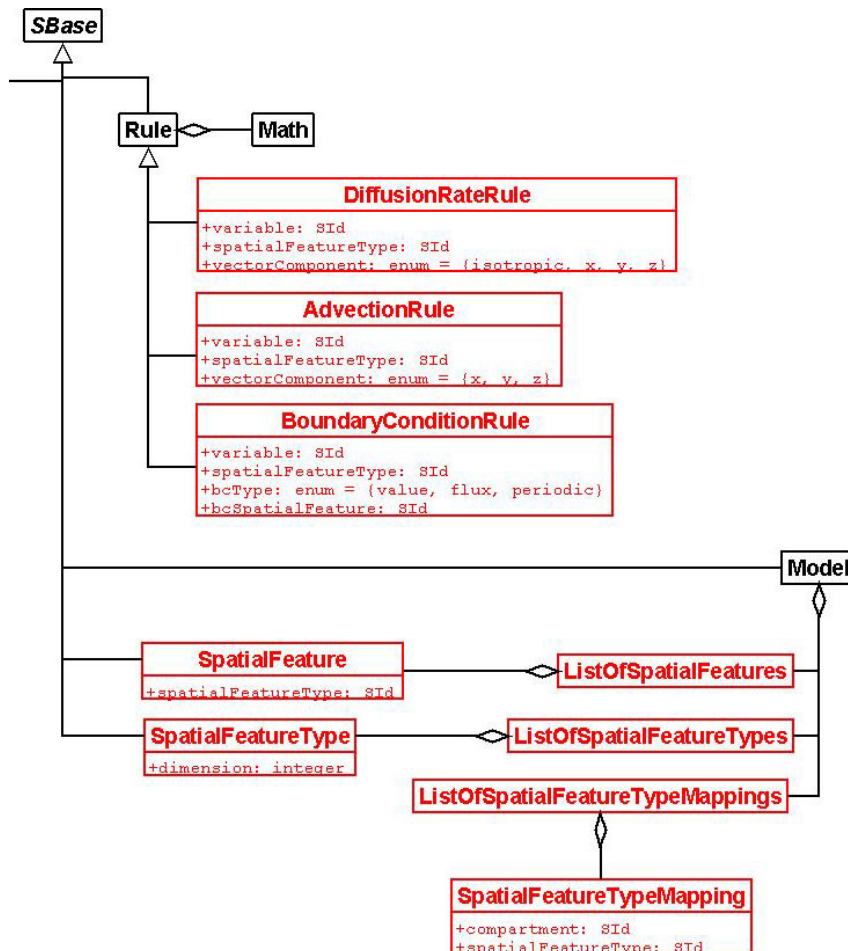
Sampled at "arbitrary" resolution



# Summary – 2 separate domains

...spatial model elements... (~“L3 spatial”)

...geometry descriptions... (~“L3 geom”)



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