



# Colonic Crypts Modeling: An Exercise with SBML Spatial and VCell

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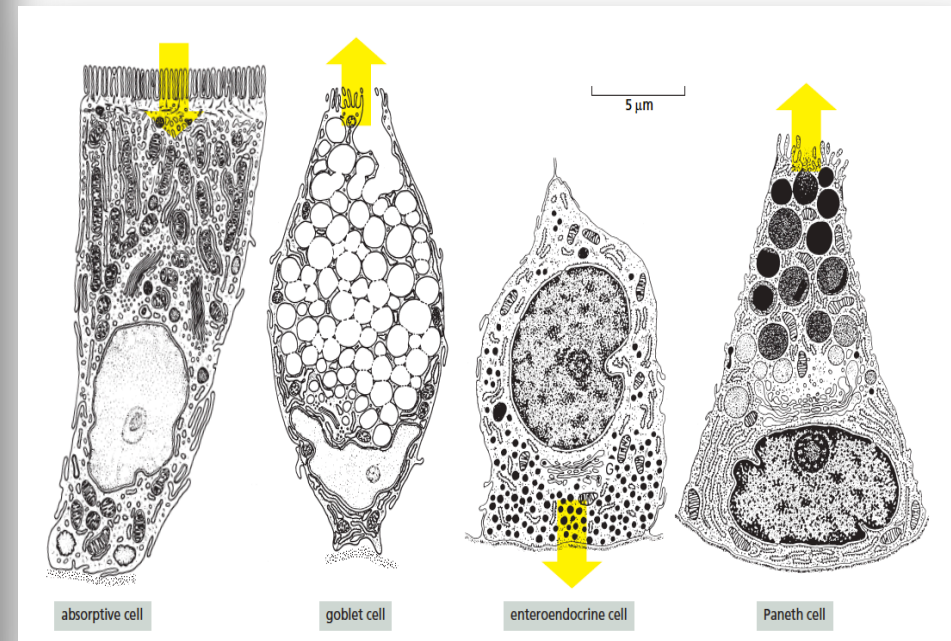
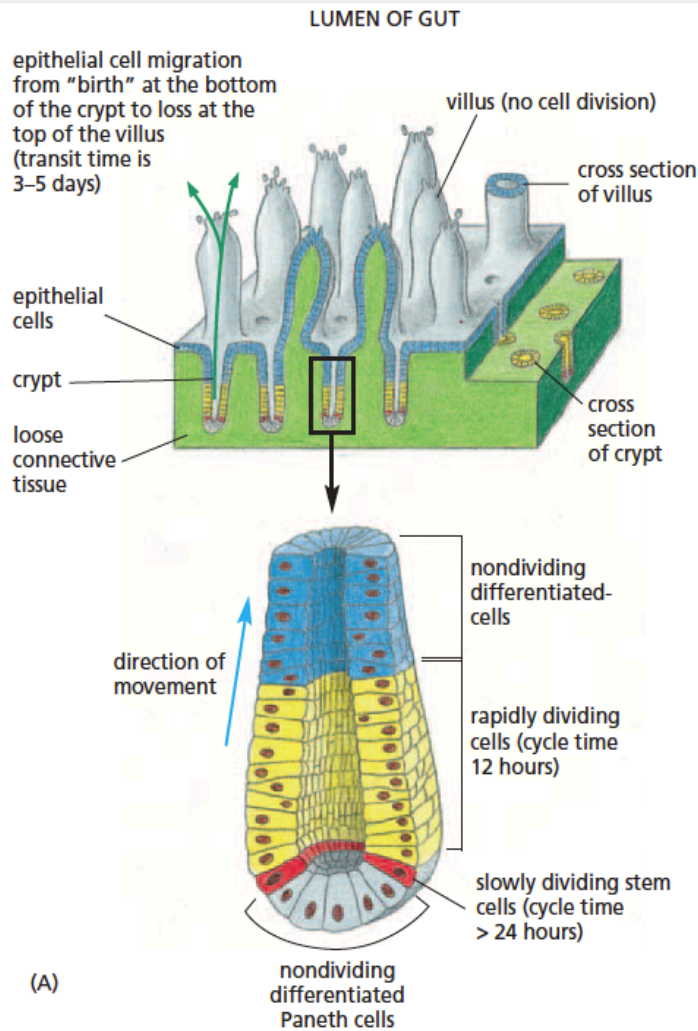


# Outline

- A bit of context: Colorectal crypts, colorectal cancer, simulation schemes (in-lattice, off-lattice), multiscale simulations
- The exercise
  - Rendering a crypt with SBML Spatial
  - Using VCell
  - Issues found

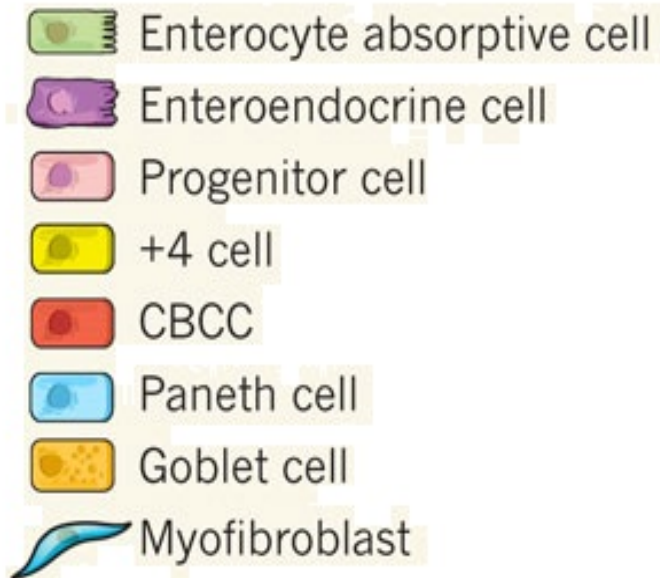


# Intestinal crypts

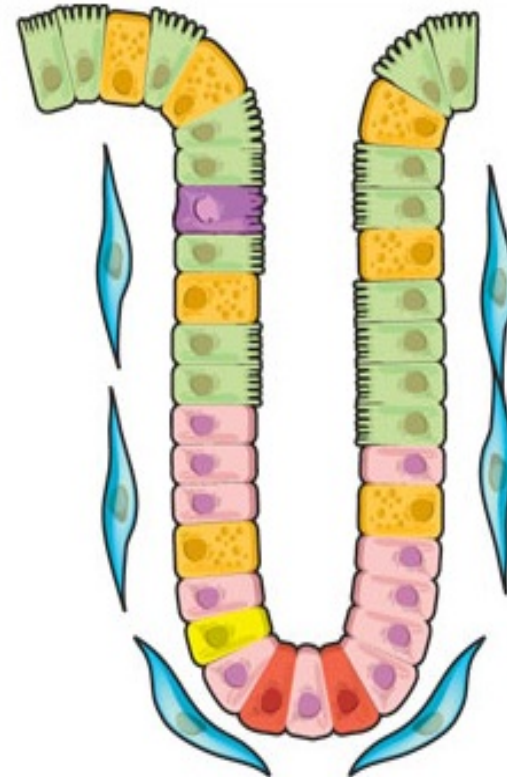


From: Alberts, B. et al. Molecular Biology of the Cell, 5th ed. Garland Science (2007)

# Cellular Types

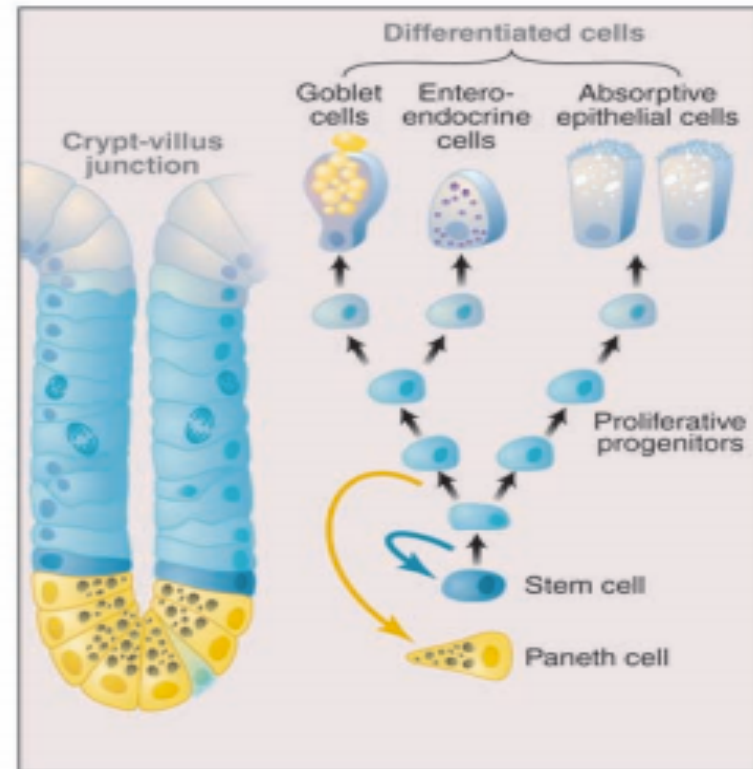
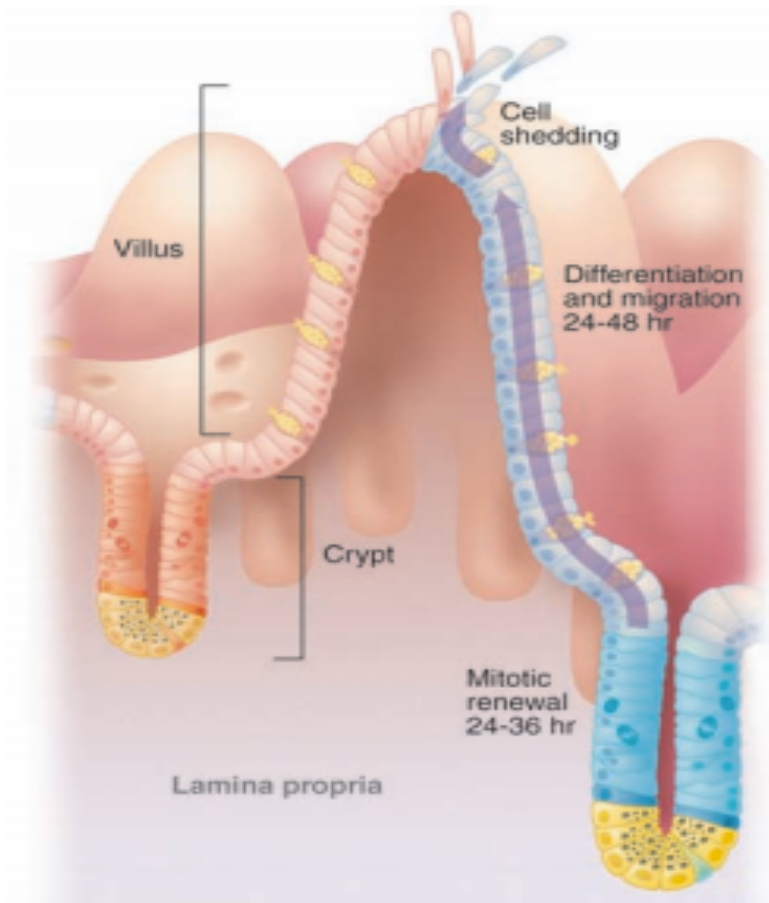


Colon crypt



Cfr. Microenvironmental regulation of stem cells in intestinal homeostasis and cancer  
doi:10.1038/nature10212  
Jan Paul Medema & Louis Vermeulen

# Lineage Commitment Tree







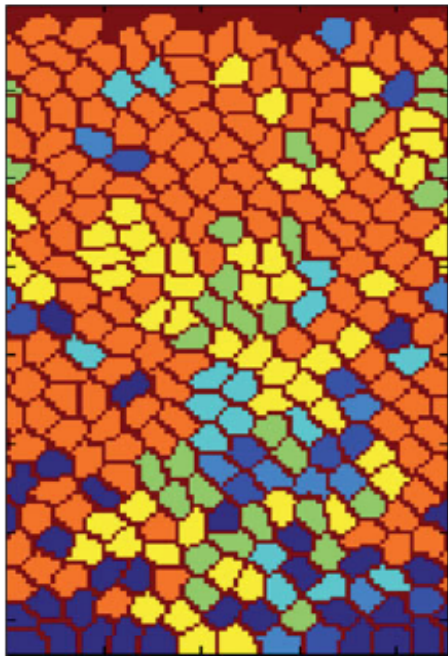
# Potential for in Vitro Experimental Validation

- Recently, several researchers interested in cancer (especially in CSCs) have looked at the potential of growing “mini-guts” in vitro
- As an example:  
**Growing Self-Organizing Mini-Guts from a Single Intestinal Stem Cell: Mechanism and Applications**  
Toshiro Sato and Hans Clevers  
*Science* **340**, 1190 (2013);  
DOI: 10.1126/science.1234852



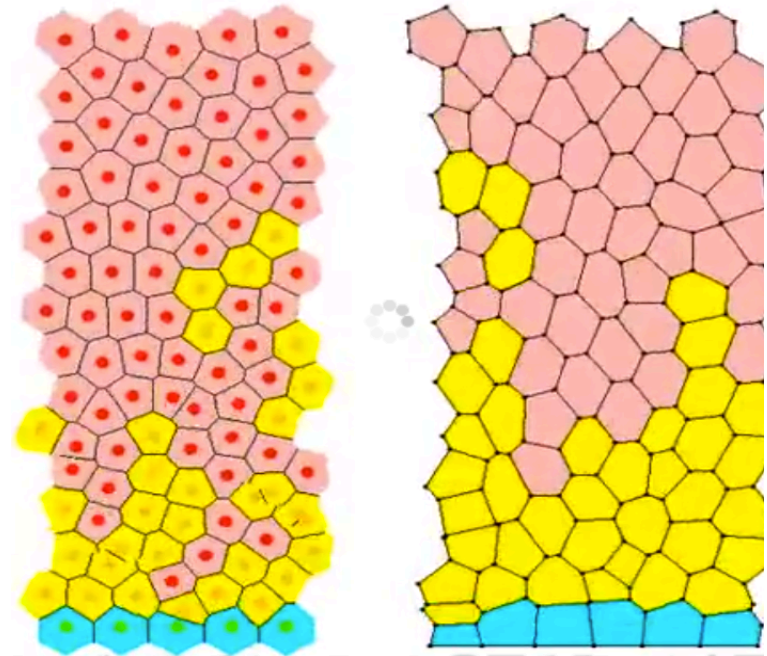
# Spatial Models for Intestinal crypt Simulations

**In lattice – CPM**  
(e.g., CompuCell3D)



From: Wong et al. Journal of The Royal Society Interface  
7(Suppl 3), S351-S363 (2010).

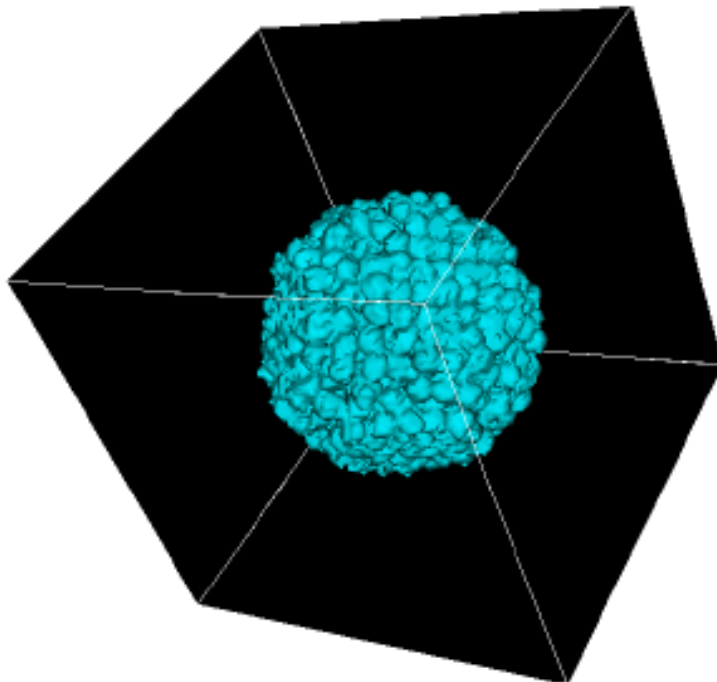
**Off lattice**  
(e.g. CHASTE, CellSys)



From: Pitt-Francis, J. et al. Computer Physics Communications  
180(12), 2452 - 2471 (2009).

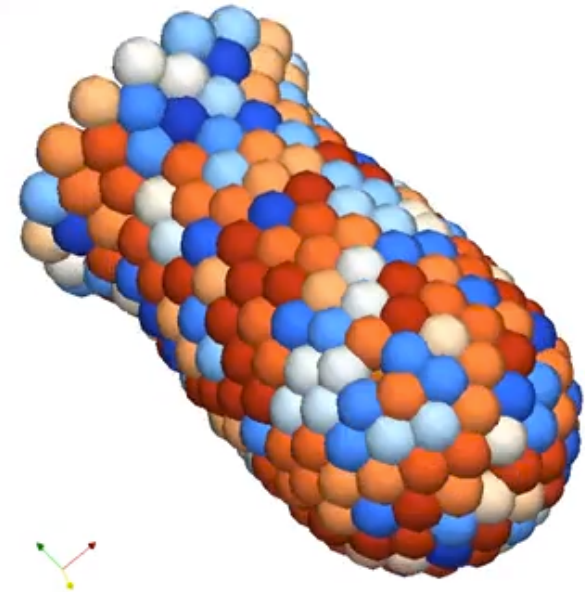
# Extensions in 3D

**In lattice (e.g. CPM)**



<http://www.compuCell3d.org/>

**Off lattice (e.g. CHASTE, CellSys)**



From: Pitt-Francis, J. et al. Computer Physics Communications 180(12), 2452 - 2471 (2009).



# A multiscale model of intestinal crypts

- Intestinal crypts are a clear example of multiscale system:
  - Low-level processes:
    - Gene regulatory networks
    - Signaling pathways
    - Intercellular communication
- High-level “hallmark” phenomena
  - Angiogenesis
  - Enabling replicative immortality
  - Metastasis
  - Evading growth suppressors
  - Sustaining proliferative signaling
  - Resisting cell death
- Tumor microenvironment
  - Stroma interaction



Model of Gene Regulatory Network (GRN):  
**Noisy Random Boolean Networks (RBN)**



Spatial/morphological model of crypt dynamics:  
**Cellular Potts Model (CPM)**



**MULTISCALE MODEL**



# SBML SPATIAL

## SBML 3 Spatial Extension:

- the current version of the extension is the 0.81 release of July 2012\*.
- a new tag, named *geometry*, has been introduced and it enables an explicit definition of a spatial environment for simulation.

\*James Schaff. SBML Spatial Extension proposal – Oct 9, 2010 – COMBINE meeting, Version 0.81 release on 15 August 2012.



# SBML3 Spatial Tags 1/2

- **ListOfCoordinateCompartments:** Definition of the spatial reference frame .
  - In our case it is a 3-dimensional Cartesian System where the x-axis represent the width, the y-axis the height and the z-axis the depth.
- **ListOfDomainTypes:** Definition of homogeneous spatial zones present in the system.
  - In our case the domain type is a cell in the colonic crypt.
- **ListOfDomains:** Definition of contiguous regions identified by the same domain type.
  - In our case the specific cells in the colonic crypt.

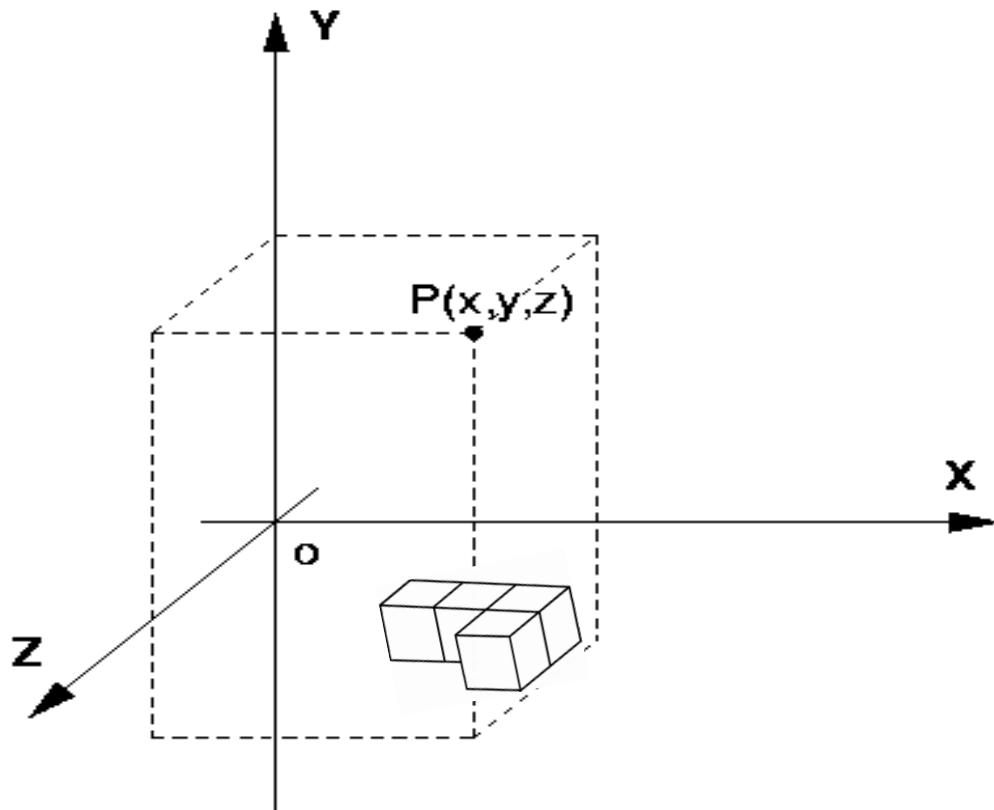


# SBML3 Spatial Tags 2/2

- **ListOfAdjacentDomains:** Definition of adjacent domain types.
  - In our case the adjacent cells in the colonic crypt.
- **ListOfGeometryDefinitions:** Definition of the geometrical structure of each domain type (i.e., analytic, sampledField, constructive solid geometry, and parametric shapes).
  - In our case the `AnalyticalGeometry` tag has been adopted, which, in our case, was a stop-gap measure



# Spatial Structure



- The crypt as a hollow parallelepiped
- The cells as cubes





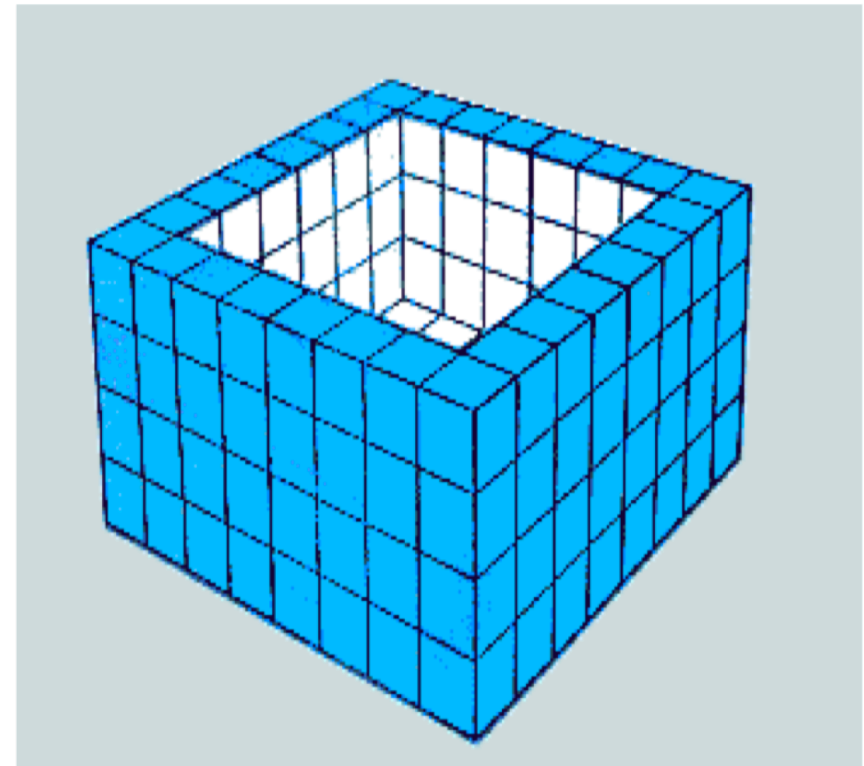
# Spatial Initial Condition

- A set of stem cellular types (no cells of other types).
- Located in the bottom of the colonic crypt.
- The cells above and below the layer of stem cellular types are empty



# Crypt SBML Spatial 1/3

- **Dynamic model:**  
cellular differentiation
- **Spatial model:** the movement is simulated by consuming a cellular type in a given cell and creating a new one in the upper or lower cell





# Crypt SBML Spatial 2/3

The dynamic model based on SBML core and the geometry based on SBML spatial are connected by the **compartmentMapping** tag

```
<listOfCompartments>
  <compartment id="id_comp_cel_1" name="comp_cel_1" size="1" spatialDimensions="3" constant="false">
    <spatial:compartmentMapping compartment="id_comp_cel_1" spatial:domainType="colonic_crypt_cell_1" spatial:unitSize="1" spatial:spatialId="com_dom_mapping_1"/>
  </compartment>
  . . . . .
  . . . . .

<spatial:geometry spatial:coordinateSystem="XYZ">
  <spatial:listOfCoordinateComponents>
    <spatial:coordinateComponent spatial:spatialId="x" spatial:componentType="cartesianX" spatial:index="0" spatial:sbmlUnit="microns">
      <spatial:boundaryMin spatial:spatialId="x_min" spatial:value="0"></spatial:boundaryMin>
      <spatial:boundaryMax spatial:spatialId="x_max" spatial:value="2"></spatial:boundaryMax>
    </spatial:coordinateComponent>
    <spatial:coordinateComponent spatial:spatialId="y" spatial:componentType="cartesianY" spatial:index="1" spatial:sbmlUnit="microns">
      <spatial:boundaryMin spatial:spatialId="y_min" spatial:value="0"></spatial:boundaryMin>
      <spatial:boundaryMax spatial:spatialId="y_max" spatial:value="6"></spatial:boundaryMax>
    </spatial:coordinateComponent>
    <spatial:coordinateComponent spatial:spatialId="z" spatial:componentType="cartesianZ" spatial:index="2" spatial:sbmlUnit="microns">
      <spatial:boundaryMin spatial:spatialId="z_min" spatial:value="0"></spatial:boundaryMin>
      <spatial:boundaryMax spatial:spatialId="z_max" spatial:value="2"></spatial:boundaryMax>
    </spatial:coordinateComponent>
  </spatial:listOfCoordinateComponents>
</spatial:geometry>
```



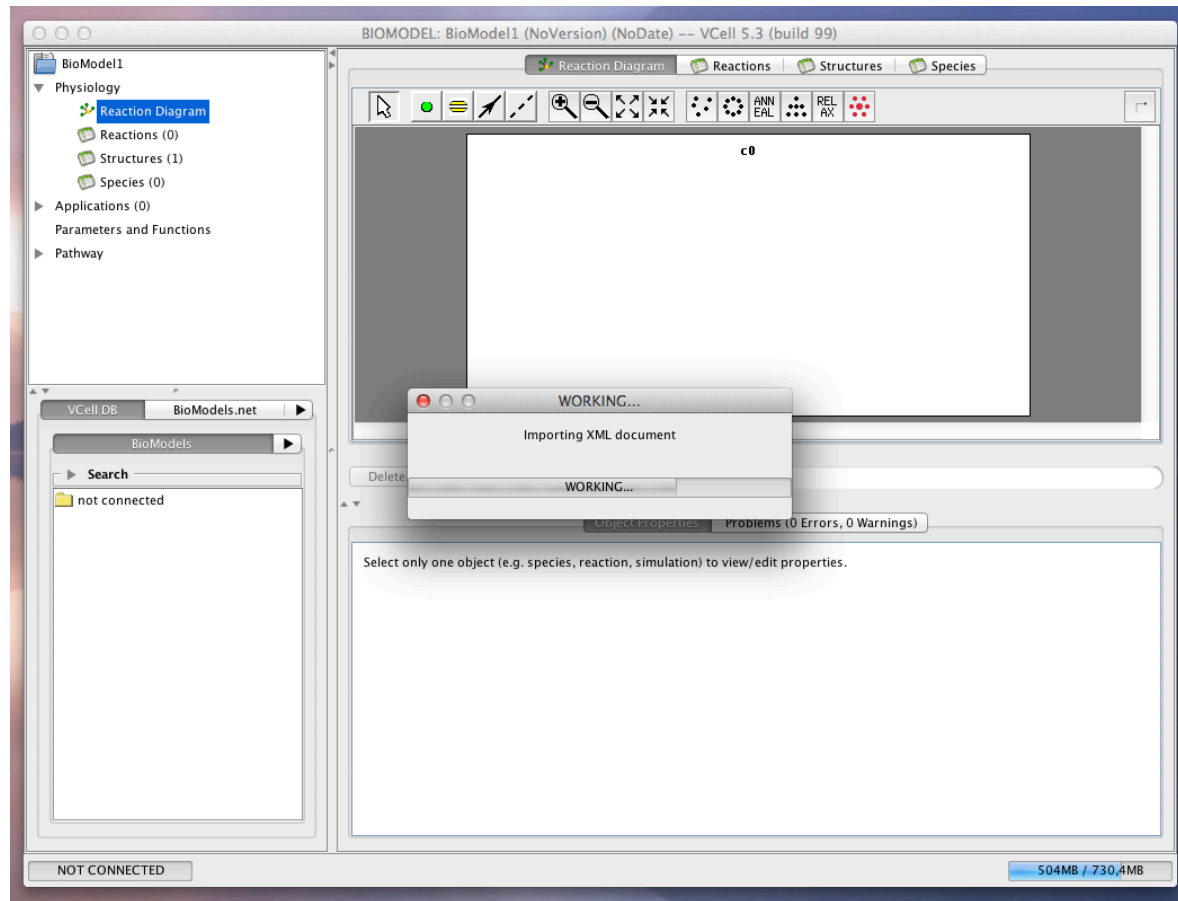
# Crypt SBML Spatial 3/3

In the **domainType** tags homogeneous spatial zones are defined. The specific zone associated with a **domainType** are then instantiated by the **domain** tag

```
<spatial:listOfDomainTypes>
  <spatial:domainType spatial:spatialId="colonic_crypt_cell_1" spatial:spatialDimension="3"></spatial:domainType>
  <spatial:domainType spatial:spatialId="colonic_crypt_cell_2" spatial:spatialDimension="3"></spatial:domainType>
  <spatial:domainType spatial:spatialId="colonic_crypt_cell_3" spatial:spatialDimension="3"></spatial:domainType>
  <spatial:domainType spatial:spatialId="colonic_crypt_cell_4" spatial:spatialDimension="3"></spatial:domainType>
  <spatial:domainType spatial:spatialId="colonic_crypt_cell_5" spatial:spatialDimension="3"></spatial:domainType>
</spatial:listOfDomainTypes>

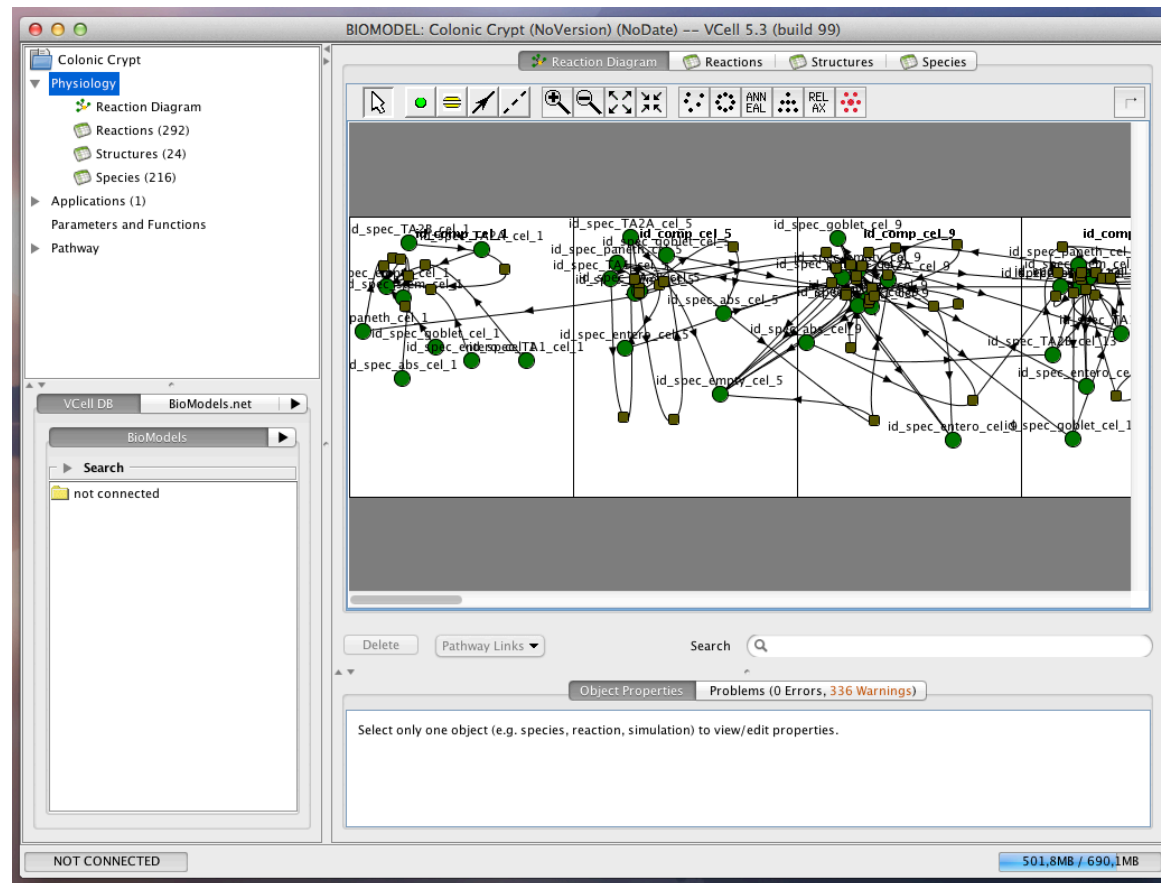
<spatial:listOfDomains>
  <spatial:domain spatial:spatialId="colonic_crypt_cell_domain_1" spatial:domainType="colonic_crypt_cell_1">
    <spatial:listOfInteriorPoints>
      <spatial:interiorPoint spatial:coord1="0" spatial:coord2="0" spatial:coord3="0"></spatial:interiorPoint>
      <spatial:interiorPoint spatial:coord1="1" spatial:coord2="1" spatial:coord3="1"></spatial:interiorPoint>
    </spatial:listOfInteriorPoints>
  </spatial:domain>
  <spatial:domain spatial:spatialId="colonic_crypt_cell_domain_2" spatial:domainType="colonic_crypt_cell_2">
    <spatial:listOfInteriorPoints>
      <spatial:interiorPoint spatial:coord1="0" spatial:coord2="0" spatial:coord3="1"></spatial:interiorPoint>
      <spatial:interiorPoint spatial:coord1="1" spatial:coord2="1" spatial:coord3="2"></spatial:interiorPoint>
    </spatial:listOfInteriorPoints>
  </spatial:domain>
</spatial:listOfDomains>
```

# Using VCell 1/3





# Using VCell 2/3



# Using VCell 3/3

BIOMODEL: Colonic Crypt (NoVersion) (NoDate) -- VCell 5.3 (build 99)

Geometry Specifications Protocols Simulations

Structure Mapping Geometry Definition

All structures and subdomains must be mapped to run a simulation.  
Use line tool or drop down menu in the 'subdomain' column.

Physiology (structures)

Geometry (subdomains)

Structure	Subdomain	Size Ratio	X-	X+	Y-	Y+	Z-	Z+
id_comp_cel_1	colonic_crypt_c...	1 [1]	Flux	Flux	Flux	Flux	Flux	Flux
id_comp_cel_2	colonic_crypt_c...	1 [1]	Flux	Flux	Flux	Flux	Flux	Flux
id_comp_cel_3	colonic_crypt_c...	1 [1]	Flux	Flux	Flux	Flux	Flux	Flux
id_comp_cel_4	colonic_crypt_c...	1 [1]	Flux	Flux	Flux	Flux	Flux	Flux
id_comp_cel_5	colonic_crypt_c...	1 [1]	Flux	Flux	Flux	Flux	Flux	Flux

Object Properties Problems (0 Errors, 336 Warnings)

Select only one object (e.g. species, reaction, simulation) to view/edit properties.

NOT CONNECTED 245,3MB / 748,2MB



# Open Issues

- Link between dynamic and spatial components into an unique model of kinematic evolution of the colonic crypt.
  - Associate to each cellular transformation a spatial movement (i.e., stem to paneth → downward movement)
  - Properly defining cell division (mitosis and cytokinesis)
  - Defining the initial conditions of a given simulation run
  - ... and more.



# Conclusions

- As usual the interplay between static and dynamic representation poses several problems, as revealed, in our case, by the issue of “events”
- The interplay of SBML Spatial with other SBML packages should be clarified: SBML Multi (and SBML Dynamic?) being an example
- Another issue, as far as we could tell, is that the **ListOfGeometryDefinitions** does not take into account the simple “in-lattice/cellular-automata” simulation schemes
  - Could we just have a ‘set-of-voxels’ definition?
  - As usual, how much of the “simulation engine” should be represented in SBML
- Finally, spatial simulations should be replicable, which is another topic for SED-ML



# Acknowledgements

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