

# Architecture of a Tangible Interface for Modeling Plant Cell Cycle





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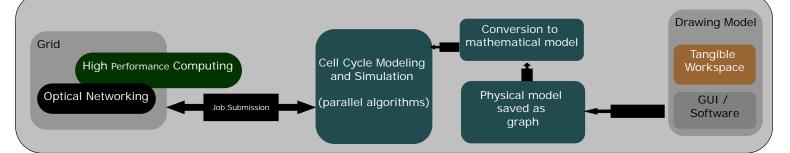
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#### Motivation:

Cell cycle simulation using HPC techniques had been successfully done using supercomputers [2]. techniques used to model cell cycle is not user-friendly for biologists. Also, there is no way in which a schematic diagram of a part of the model can be converted to corresponding mathematical model for simulation purposes. We seek prospects of interfacing supercomputers with tangible interfaces to serve this purpose.

#### **Interfacing Grid with Tangible Workspace:**



# **Tangible Workspace (TW)**

- Includes both visual+physical components
- Easy for human manipulation
- Maintains back-end connectivity with cluster computers, and grid resources for simulation.

## **Back-end Software GUI**

- needs to save the physical model drawn on the tangible workspace
- needs to retrieve a saved model for further editing
- needs to create new models, if necessary, and save it.

#### **Design considerations** for TW

- · the material for the workspace area
- · how the objects will hold on to the workspace
- · how the objects will sense presence of each other

#### Handling simulations, Computational Issues, back-end gridconnectivity

- · comparing results of mathematical model and experimental data
- · very hard to set numerical values of the "parameters" in the mathematical model equations to create more accurate representations of steps in each cell cycle

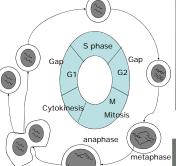








Fig. 1: (Some existing tangible interfaces): (left top) Graphical, actuated workbench for cell tower placement [Patten et al. 2007]; (left bottom) PostIt-based web layout [Klemmer et al. 2001]; (middle top/bottom) collaborative visualization using viz tangibles [Ullmer et al. 2008]; (left top) Microsoft Surface Technology; (left bottom) String-based interface;



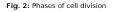




Fig. 3: Existing physical workspace for cell-cycle modeling, without any computer connectivity

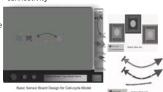


Fig. 4: Preliminary sketches of Tangible kspace, objects, and connectors

## **Basic Workflow**

- (1) Actual physical design of a cell-cycle model on the tangible interface workspace;
- (2) Saving of the physical model as a softcopy for future retrieval, using controls available on the workspace;
- (3) Editing the saved model in a GUI, if needed;
- (4) Choosing a cell-cycle model as input to the parallel algorithm for simulation and verification of the model's
- (5) Submitting the resulting compute job description to the grid;
- (6) Collecting the output from the compute resources for further analysis and archiving

## References

- [1] Thomas Haberichter et. al., A systems biology dynamical model of mammalian G1cell cycle progression, Molecular Systems Biology 3; Article number 84; doi:10.1038/msb4100126, Nature Publishing Group, 2007.
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