Sub-cellular and Multicellular Modeling Workshop at The Hamner Institutes for Health Sciences Research Triangle Park, NC USA

Instructors: Herbert M. Sauro (Univ. Washington),
Maciej Swat, James Glazier, Julio Belmonte, Jim Sluka (Indiana Univ.)

The following materials will be provided to Participants during the Course:

- 1. Discounted copy of Text Book on Enzyme and Gene Regulatory Kinetics (Paper and ebook)
- 2. CompuCell3D Introduction (all other manuals including CC3D reference, Python scripting etc... will be available in the electronic form)
- 3. PowerPoint/PDF copies of lecture slides
- 4. Folder with simulation notes and other materials
- 5. Model Files from the course web site
- 6. Selected readings and background material

Aug 10th, 2014 (Sunday)

Introduction to Python Programming (optional):

Day 0: Course: 10AM - 5.00PM:

10.00 – 11.00: Basic Python Programming (if, for, list, dictionaries etc...)

11.00 – 12.00: Basic Python Programming - Tutorials

12.00 – 1.00: Lunch Break

1.00 – 2.00: Slightly More Advanced Python Programming (classes and objects)

2.00 – 3.00: Slightly More Advanced Python Programming Tutorials

3.00 – 3.15: *Break*

3.15 – 4.15: Python Standard Library (file I/O, file management - os, path modules, sys module, search paths etc...)

4.15 – 5.00: Python Standard Library Tutorials

Aug 11th, 2014 (Monday)

Day 1: Course: 9AM – 6.00PM:

9.00 - 9.10: Welcome note (Jim Sluka, Indiana Univ.)

9.10 – 9.20: Welcome note (Sudin Bhattacharya, The Hamner Institute)

9.20 – 10.00: *Lecture:*

a) Introduction to Modeling with JDesigner and Jarnac

b) Definitions of: Stoichiometry, Rate Laws, Boundary Species, Steady States and Transients

10.05-10.30: Hands on exercises

a) Getting Help: Documentation and tutorials

b) Simple Closed Systems

c) Rate Law Selection

10.30 – 11.00: *Break*

11:00 – 12:30: Hands on exercises

a) Simple Open Systems

b) Applying Perturbations

c) Plotting Simulations

d) SBW Simulation Tool

12.30 – 1.30: *Lunch* **1.30 – 2.30:** *Lecture:*

SBML, Sensitivity Analysis, Parameter Scans

3.30 – 4.00: *Break*

4:00 – 5:40: *Hands on exercises*

a) Exchanging models, SBML, Matlab

b) Using other models, obtaining model components

c) Sensitivity Analysis, Interactive Modeling and Parameter Scans

5.40 - 6.00: Road map for 2012-2014 and developer information

Aug 12th, 2014 (Tuesday)

Day 2: Course: 9AM - 5.30PM:

9.00 – 10.30: Generalized enzyme kinetic rate laws 9.35 – 10.30: Mini Project: Build a Signaling Pathway

10.30 – 11.00: *Break*

11:00 – 12:30: Mini Project: Build a Signaling Pathway

12.30 – 1.30: *Lunch*

1.30 – 5.30: Mini Project: Build a Signaling Pathway, and Q&A session on Reaction Kinetics Modeling

August 13th, 2014 (Wednesday)

Day3: Course: 9AM - 5.30PM:

9.00 – 10.30: Overview of multi-cell modeling (James Glazier, Indiana Univ.)

10.30 – 11.00: *Break*

11.00 – 12.30: Introduction to CompuCell3D (Maciej Swat, Indiana Univ.)

12.30 - 1.30: Lunch

1:30 – 3.00: *CompuCell3D 101 tutorials*

3:00 - 3.30: Break

3:30 – 5.30: CompuCell3D (Introduction to Python-based Simulations-Mastering Twedit++, Python Mini-Tutorial, CC3D-Hands-on Exercises - Maciej Swat, Indiana Univ.)

August 14th, 2014 (Thursday)

Day 4: Course: 9AM - 5.30PM:

9.00 - 10.30: CC3D-Hands-on Exercises - (Maciej Swat, Indiana Univ.)

10.30 – 11.00: *Break*

11.00 – 12.30: CompuCell3D (Fields, Basic Diffusion-based PDEs, Chemotaxis, Hands-on Tutorials - Maciej Swat, Indiana Univ.)

12.30 - 1.30: Lunch

1:30 - 2:15: Invited talk- Tom Knudsen EPA

2:15 – 3:30: CompuCell3D (continue Hands-on Tutorial - Maciej Swat, Indiana Univ.)

3:30 – 4.00: *Break*

4.00 – 5.30: CompuCell3D (Advanced Python Scripting in CC3D –Attaching Extra Cell Attributes, Mitosis-Based Simulations, Cell Shape Constraints etc..., Hands-on Tutorials - Maciej Swat, Indiana Univ.)

August 15th, 2014 (Friday)

9.00 – 10.30: CompuCell3D (continue Advanced Python Scripting in CC3D –Attaching Extra Cell Attributes, Mitosis-Based Simulations, Cell Shape Constraints etc..., Hands-on Tutorials - Maciej Swat, Indiana Univ.)

10.30 – 11.00: *Break*

11:00 – 12:30: CompuCell3D (continue Hands-on Tutorial - Maciej Swat, Indiana Univ.)

12.30 - 1.30: Lunch

1:30 - 2:15: Invited talk- Imran Shah EPA

2.15 – 3.30: CompuCell3D (Advanced Python Scripting in CC3D Part 2 – using third party modules in Python, File operations, Post Processing, Simulation Steering – changing CC3DML on-the-fly - Maciej Swat, Indiana Univ.)

3.30 - 4.00: Break

4.00 – 5.30: CompuCell3D (Plots and Graphs, configuring CC3D GUI from Python/XML, continue Handson Tutorial - Maciej Swat, Indiana Univ.)

7.00 – 9.30: Hackathon: Combining a PBPK (as SBML) whole body, CompuCell3D multi-cellular and sub-cellular metabolic modeling (SBML) to create a multi-scale model of Acetaminophen (Paracetamol) liver toxicity and therapeutic effects. (CC3D group and The Hamner group but everyone is invited.)

August 16th, 2014 (Saturday)

Day 5: Course: 9AM – 3.30PM:

9.00 – 10.30: CompuCell3D (SBML-Based Models in CC3D – Reaction Kinetics, ODE, PBPK models, Hands On Tutorials – Julio Belmonte, Indiana Univ.)

10.30 - 11.00: Break

11:00 - 12:30: CompuCell3D (continue Hands-on Tutorial - Julio Belmonte, Indiana Univ.)

12.30 – 1.30: *Lunch*

1:30 – 2:15: Invited talk- Dr. Jeff Woodhead, The Hamner.

Title: "The DILIsym Model and its Application to Hepatotoxicity Testing in Drug Development." (It will include their work on APAP toxicity modeling.)

2.15 – 3.30: CompuCell3D (Developing CC3D extension modules in C++ using Twedit++ - demo, Questions-And-Answers session – Suggestions For Future CC3D improvements - Maciej Swat, James Glazier, Jim Sluka, Julio Belmonte, Indiana Univ.)