

**„Modeling Multiscale Cardiovascular and Respiratory System Dynamics**

**August 22-26, 2011 N140 William H. Foege Building**

**University of Washington, Seattle, WA 98195**

**(EarlyDraft Syllabus 9aug11)**

	<b>830-1015 AM</b>	<b>1030-1215 AM</b>	<b>1315-1500 PM</b>	<b>1515-1730 PM</b>
Monday 8:30 AM 22 August	<p>Introductions: Bassingthwaighte Modeling overview</p> <p>Strategies, Objectives in Multi-scale Modeling</p> <p>Hands on: Login compartmental models, 2-comp exchanges M-M reaction</p>	<p><u>Gary Raymond:</u> Dan Beard's Aquarium study</p> <p><u>Butterworth:</u> JSim structure and function ----Download JSim-----</p>	<p><u>Gary Raymond:</u> Aspirin: discussion of hypotheses, modeling clearance by reaction</p> <p>Hands on: fit 3 data sets with one model <u>Bassingthwaighte</u> : Prog Curves: Xanthine Oxidase (Optimize)</p> <p>ODEs--Hands on:</p>	<p>Baasingthwaighte Lecture: Blood Tissue Exchange</p> <p>Multiple Indicator Dilution Expt</p> <p>Crone Extraction Data</p> <p><b>Mass Balance</b></p> <p>-work Time on fitting with PDEs -</p>
Tuesday 8:30 AM 23 August	<p><u>Hong Qian:</u> Stochastic and Deterministic Modeling. Where do they intersect? 1. <b>Stochastic chem</b> 2. Gillespie algorithm</p>	<p><u>Hong Qian:</u> Biochem Network Models and their analysis. Building biochem <b>networks</b>. Critical paths.</p>	<p><u>Adam Alessio:</u> PET analysis, using 13N-NH3 to estimate regional blood flow distributions in dog and human hearts. Distributed and compartmental modeling. Data acquisition, ROI selection, and automating analysis w/ QPP</p>	<p><u>Butterworth:</u> 1. Computing <b>Platforms:</b> Matlab, JSim, Modelica, PEnv, etc</p> <p><u>Lucian Smith:</u> 2. <b>SBML, CellML, JSim</b> and others, incl. platform dependent ones. --Importing from CellML and SBML-</p>
Wednesday 8:30 AM 24 August	<p><u>Joseph Anderson:</u> Modeling Pulmonary Mechanics and Gas Exchange I: modeling with electrical analog</p> <p><b>Model Verification</b></p>	<p><u>Joseph Anderson:</u> Modeling Pulmonary Mechanics and Gas Exchange II. spatially distributed permeation. Spatial profiles in capillary. Diffusion. Pendelluft (Lutchen)</p>	<p><u>Max Neal:</u> CV-Resp System Model: Integrative Control.</p> <p>Built-in interventions</p>	<p><u>Max Neal:</u> Extending the Model.</p> <p>Modularizing a model for repeated automated reconstructions.</p>

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Thursday 8:30 AM 25 August	<u>Herbert Sauro</u> : Quantitative Synthesis of real Genetic Networks Working with Jarnac and <b>SBML</b> and <b>Biomodels</b> and <b>CellML</b> databases	<u>Herbert Sauro</u> : Quantitative Simulation Analysis of Metabolic Nets  <b>Validation:</b>	<u>Bart Jardine</u> : <u>Matlab Intro</u> <u>CircAdapt</u>  --diffusion model <b>Model</b> <b>Validation:</b>	<u>Bassingthwaighte</u> Cardiac Mechanics. Varying elastance modeling  5:30 PM: drive to Pier 55 ferry to Blake Island, Tillicum Village salmon dinner
Friday 8:30 AM 26 August	<u>Bassingthwaighte</u> Model stages for rMBF estimation from PET images  <b>??</b> : Effect of Noise in the data <b>Parameter</b> <b>Evaluation</b>	<u>James Caldwell</u> : rMBF: Regional Myocardial Blood Flow using PET Imaging and QPP <b>Test of</b> <b>Model Validity</b> Optimization for rMBF ====Hands on ===	<u>Gary Raymond</u> : Modular Modeling Example constructions with FORTRAN  <b>Standards for</b> <b>Modules</b> Mono- versus multi- scale. Reproducible science	<u>Wrap-up</u> : Key strategies? Review Platt 1964. Integration Biology More sets of standards? <b>Code Sharing</b>  5:30 PM

~jbb/COURSE2011/Sched2011.15aug11

BassingthwaighteElectrophys, cell models: Boltzmann chan Boltz => HHActionPot  
Beeler-Reuter Action Pot.

Real Multi-scale: Rudy2010