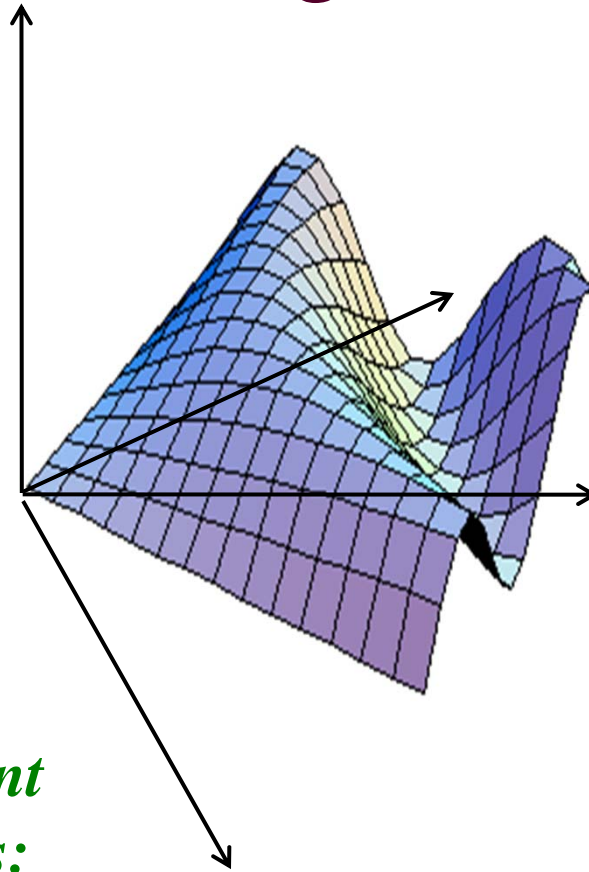


NCI ICBP
Integrative Cancer Biology Program --
MIT Grant, R Hynes PI

- **DNA damage signaling network (Michael Yaffe [Biol/BE], Leona Samson [BE/Biol], DL [BE/Biol/ChE])**
- **ErbB signaling network (Tyler Jacks [Biol], Peter Sorger [Biol/BE], Forest White [BE], DL [BE/Biol/ChE])**
- **Cell migration and invasion processes (Richard Hynes [Biol], Frank Gertler [Biol], DL [BE/Biol/ChE])**
- **Computational modeling (Bruce Tidor [BE/EECS], Jacob White [EECS])**
- **RNAi manipulation (Phil Sharp [Biol], Jianzhu Chen [Biol])**

Integrative Systems Modeling

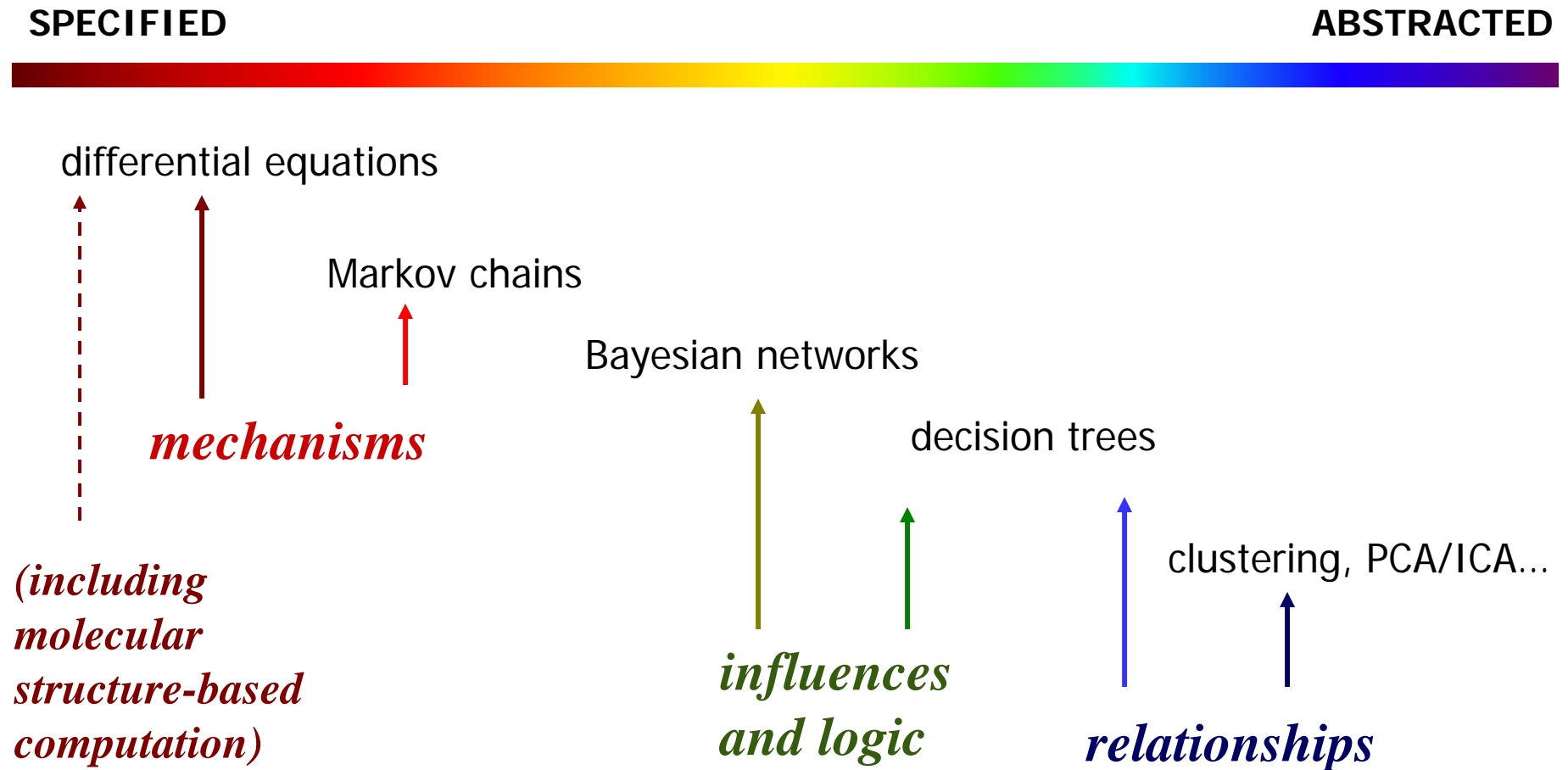
*System
Operation
(‘phenotype’):*
cell,
tissue,
organism,
...



*Component
Properties:*
molecular levels / states /
locations / interactions /
activities... (arising from
sequence & structure)

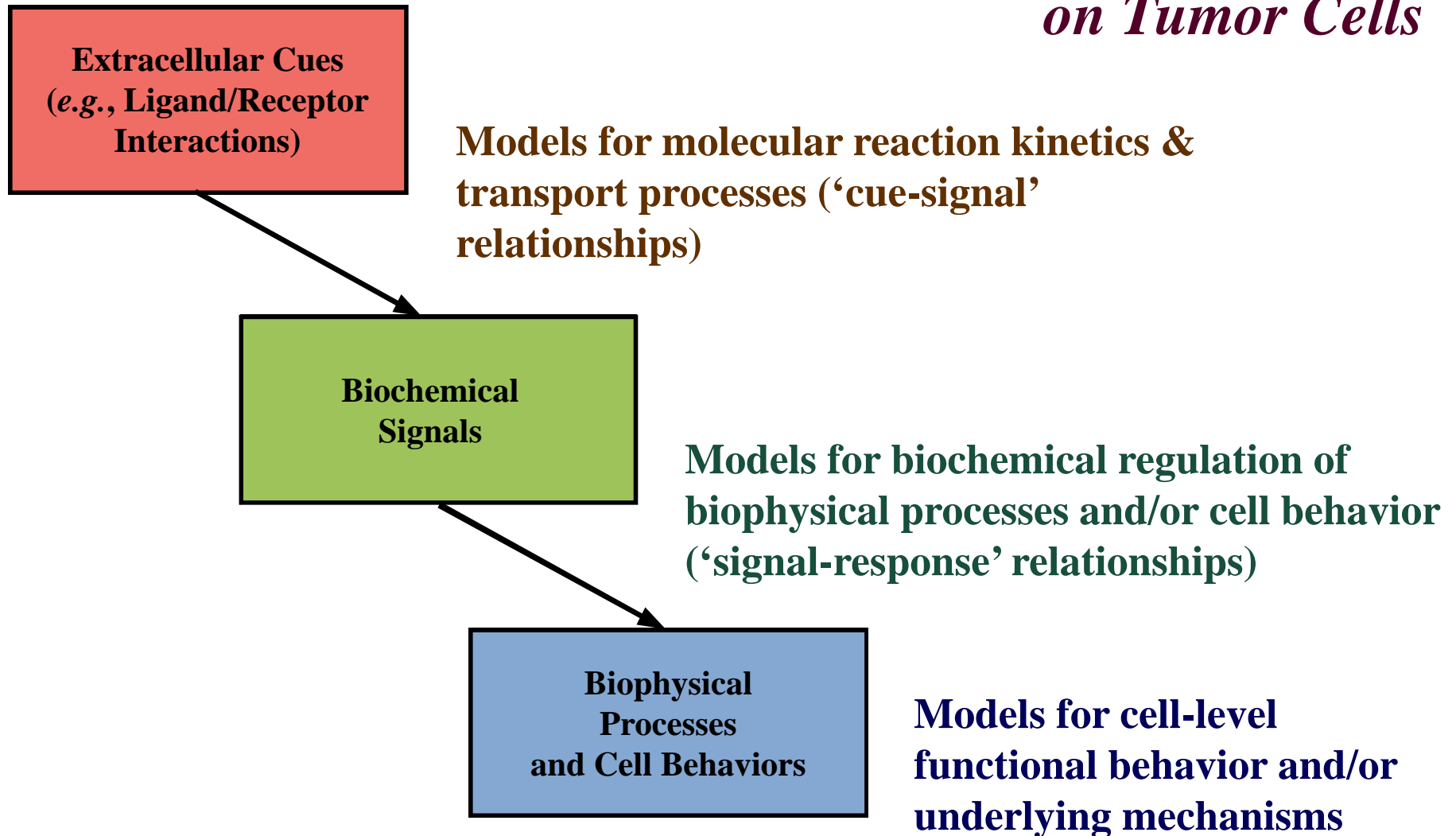
Objective:
**Predictive
understanding
for effect of
component properties**
-- in quantitative,
dynamic, multi-variate
manner

Spectrum of Computational Modeling Methods



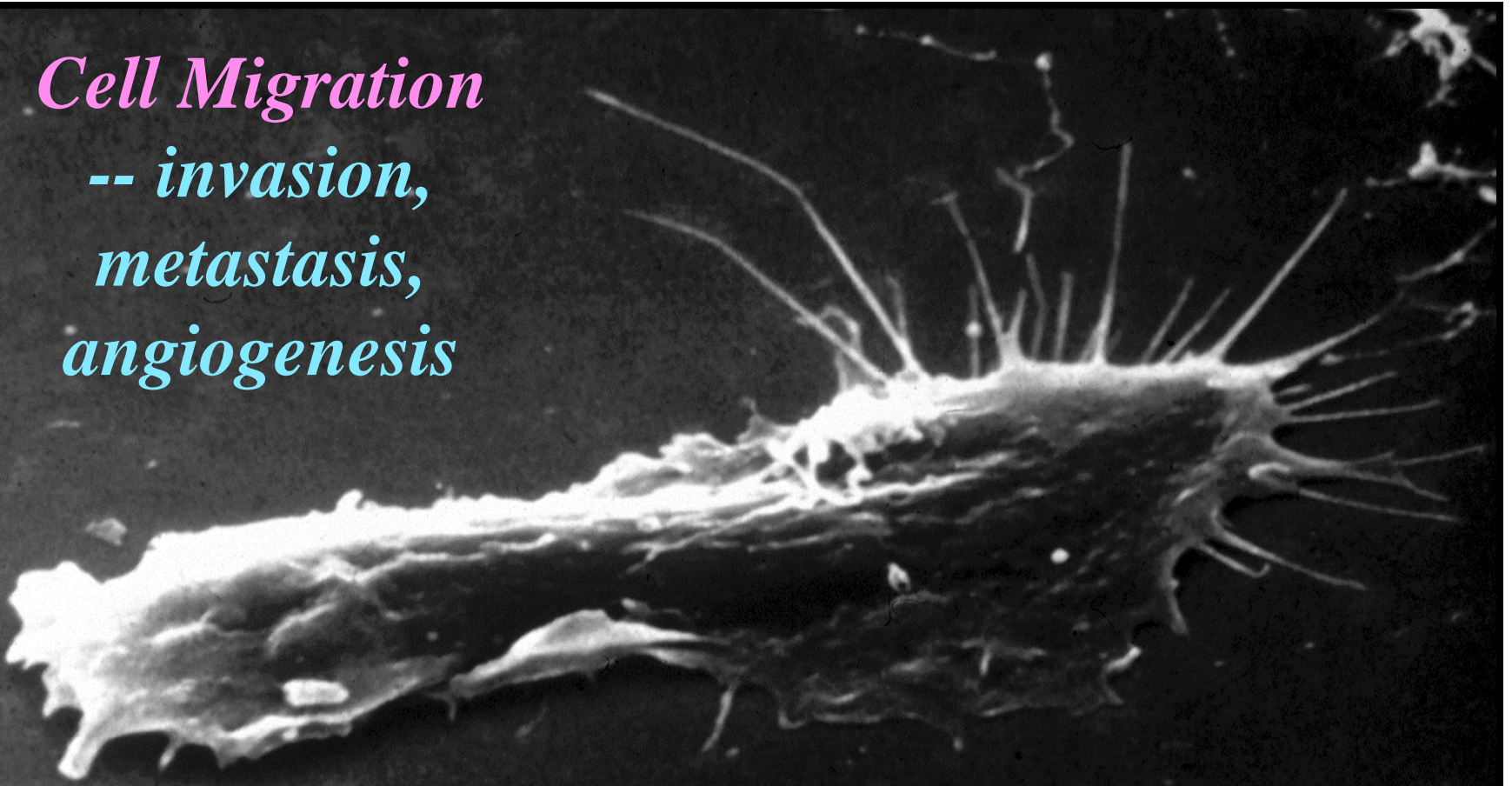
Appropriate approach depends on question and data

Modeling Chain Needed for Prediction of Effects of Gene Mutations and Drug Effects on Tumor Cells

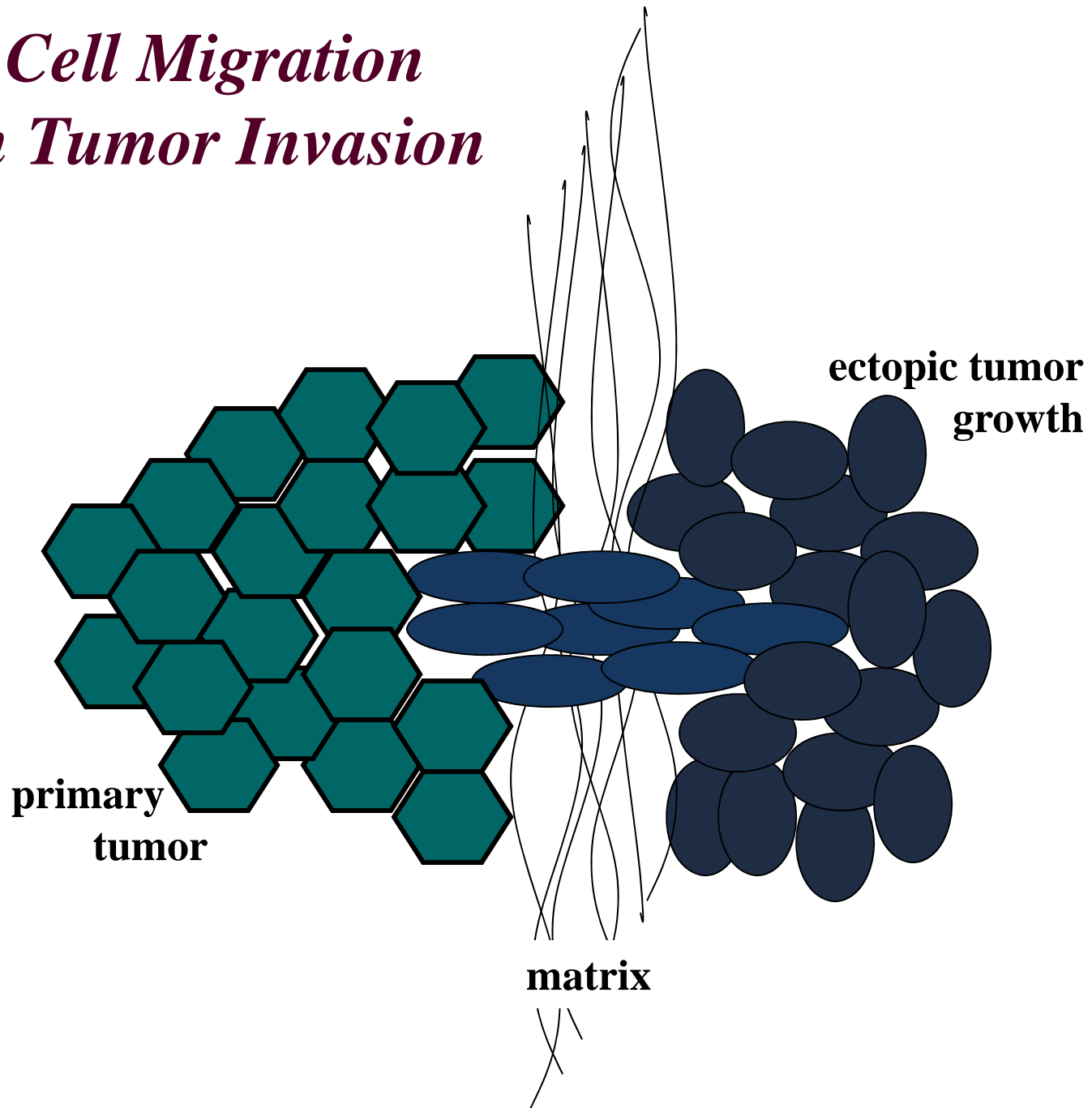


Cell Migration

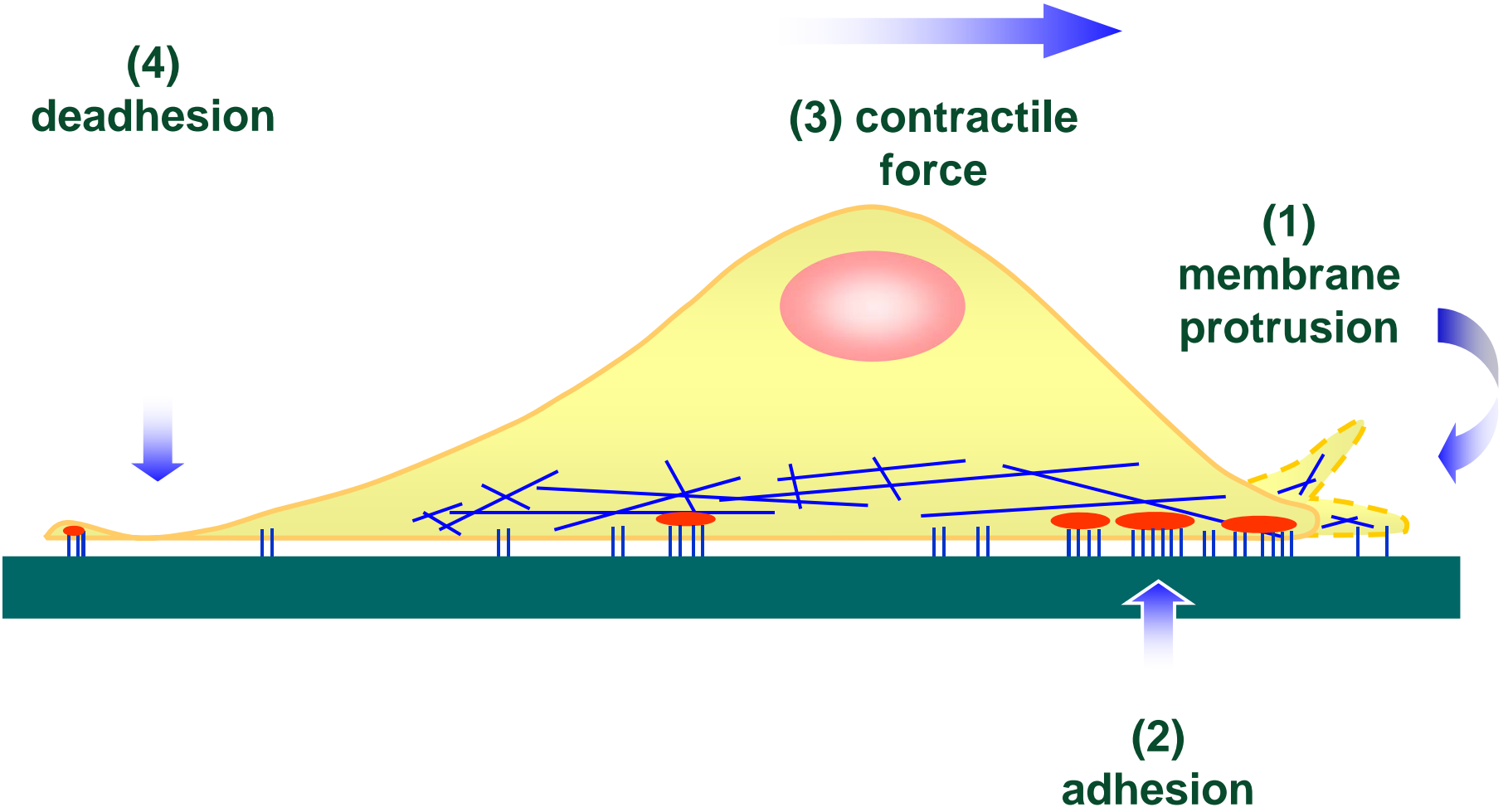
-- *invasion,*
metastasis,
angiogenesis



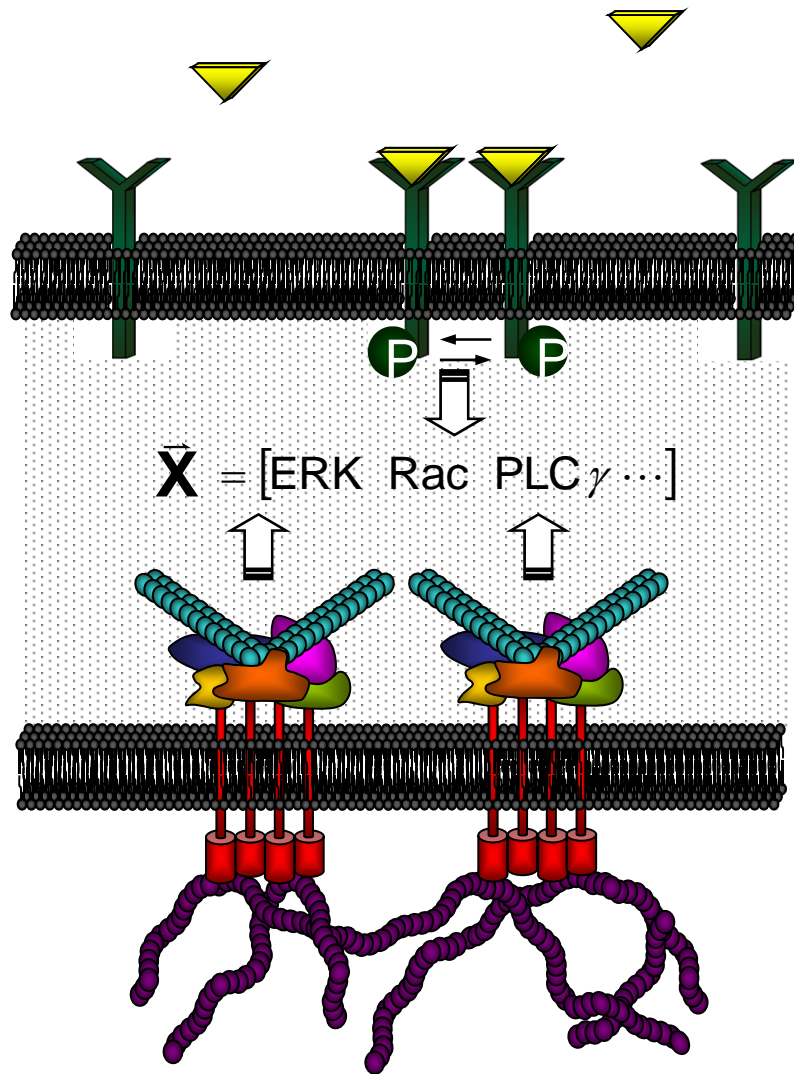
Cell Migration In Tumor Invasion



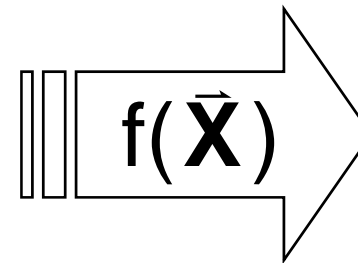
Cell Migration “Cycle” of Biophysical Processes



Regulation of Biophysical Processes by Biochemical Signals



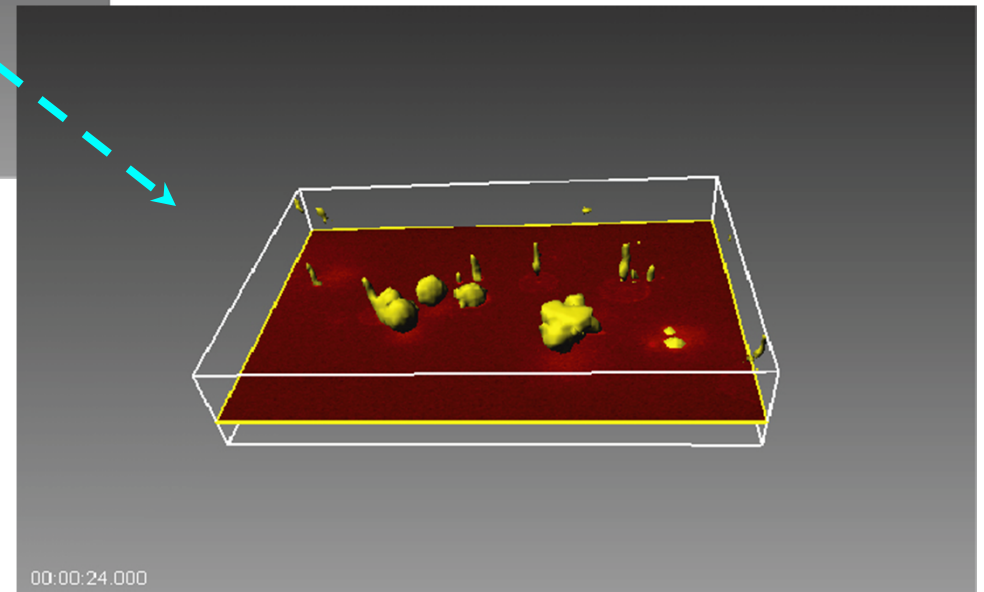
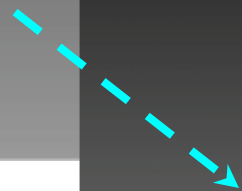
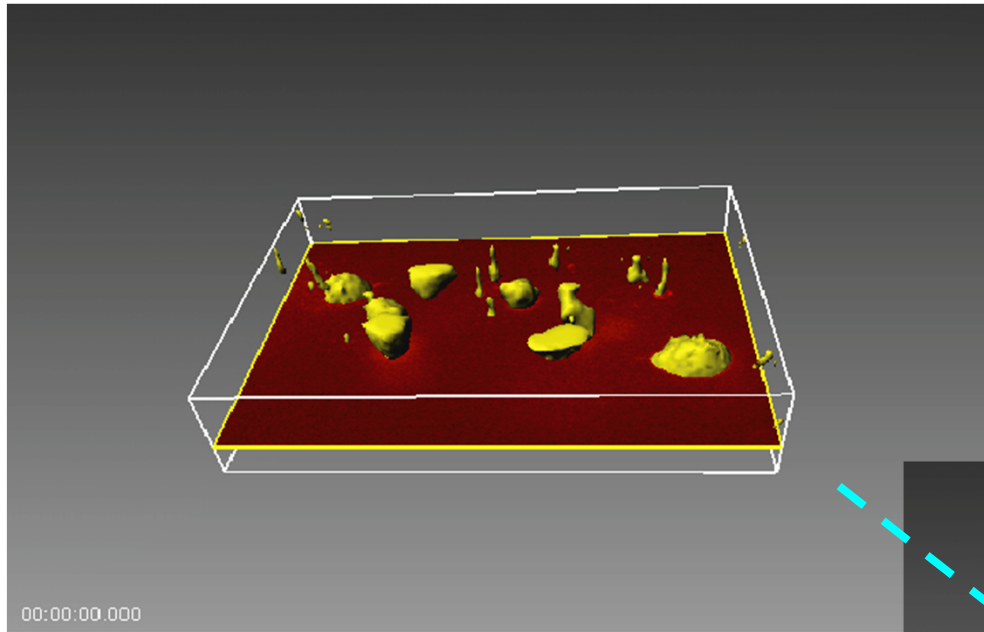
soluble ligands
(e.g., EGF)



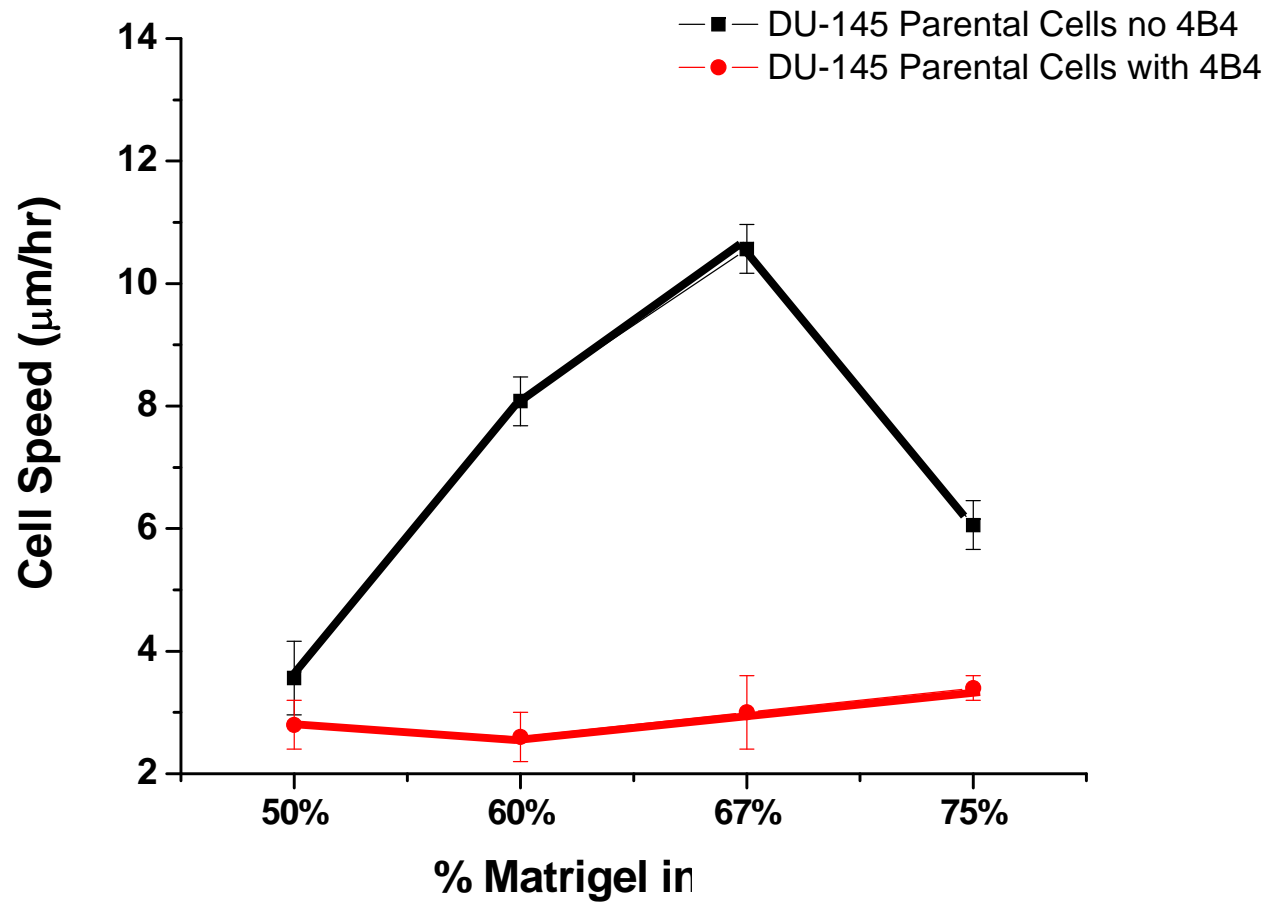
Migration
(speed,
persistence,
orientation)

matrix ligands
(e.g., Fibronectin)

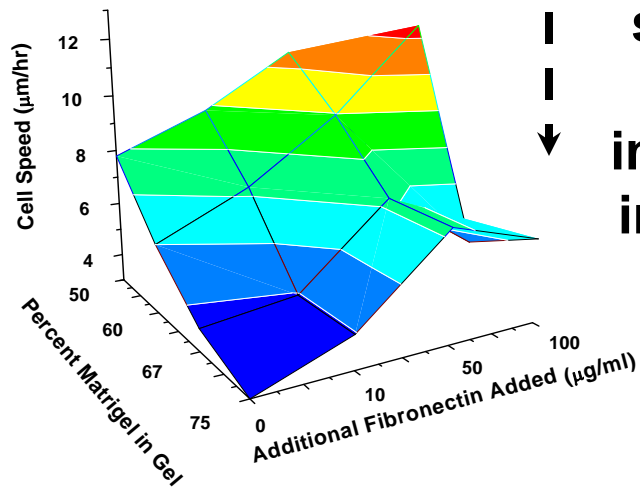
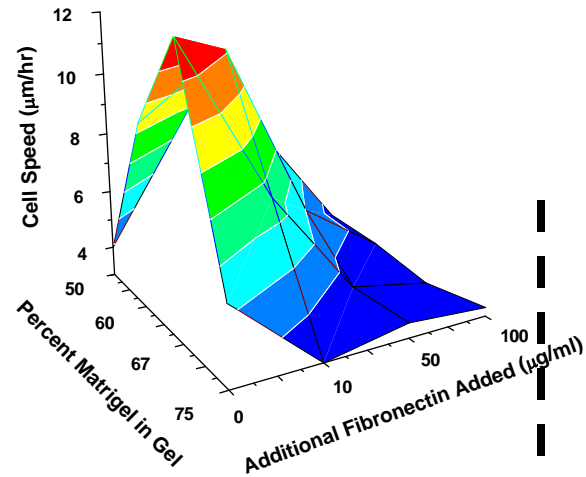
Tumor Cell Migration in 3D Environments
-- DU145 prostate tumor cells (single-cell tracking);
Matrigel / Collagen-I / Fibronectin



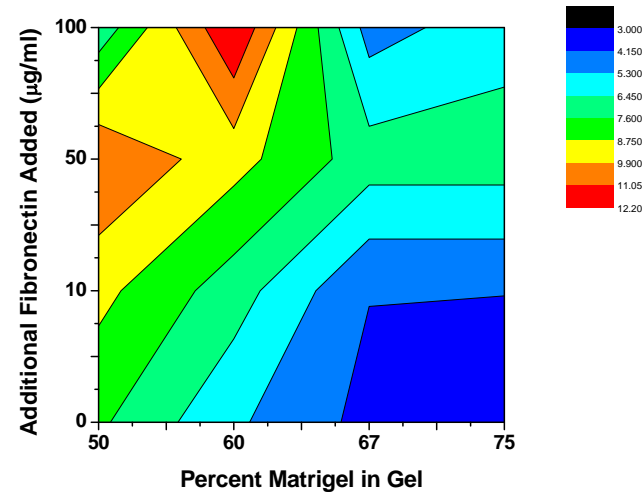
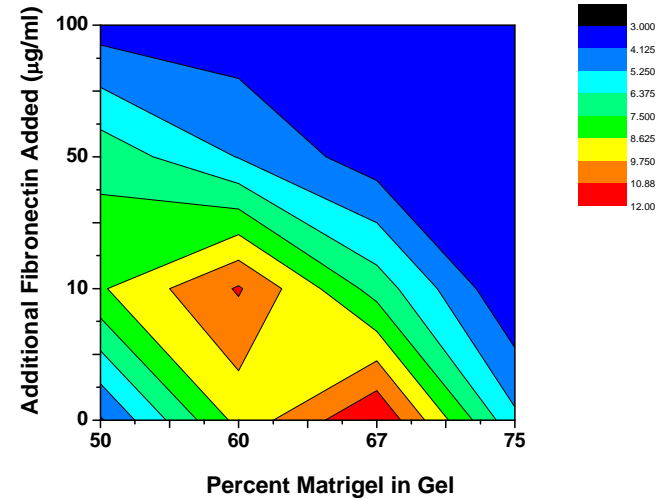
*Biphasic Relationship is Found for
3D Tumor Cell Migration Speed vs Matrix Density
-- Integrin-mediated adhesion?
Matrix sterics and mechanics?*



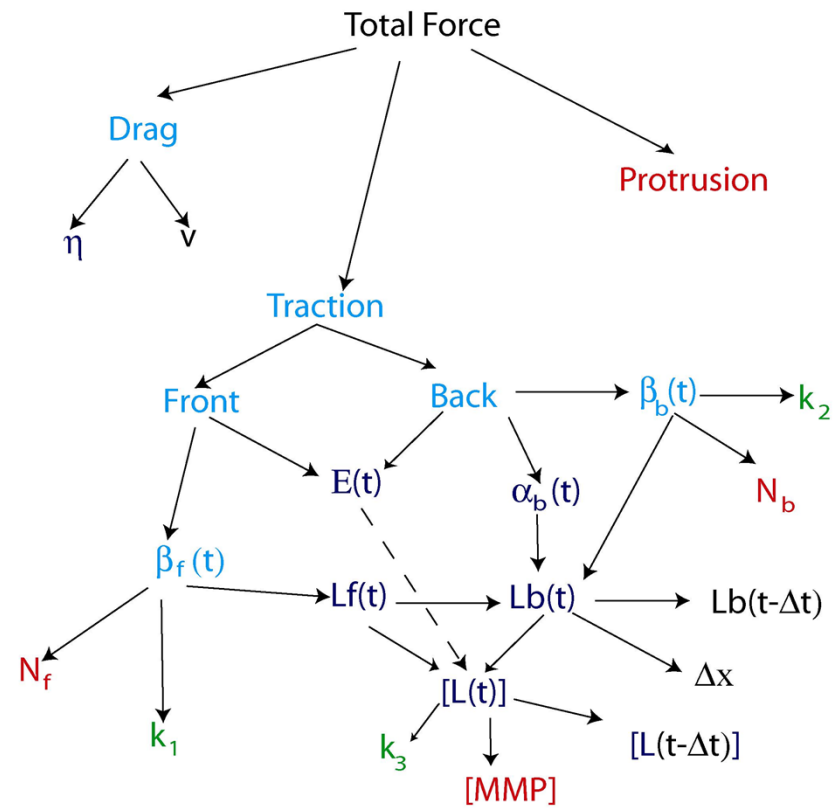
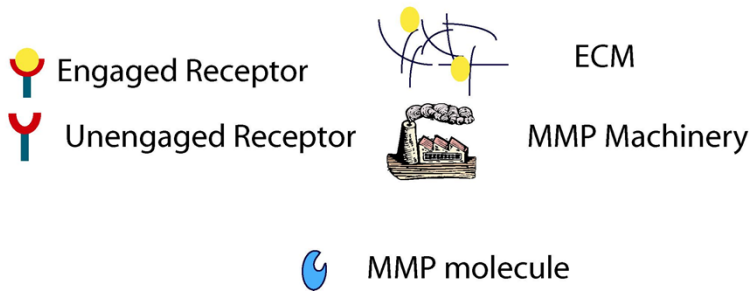
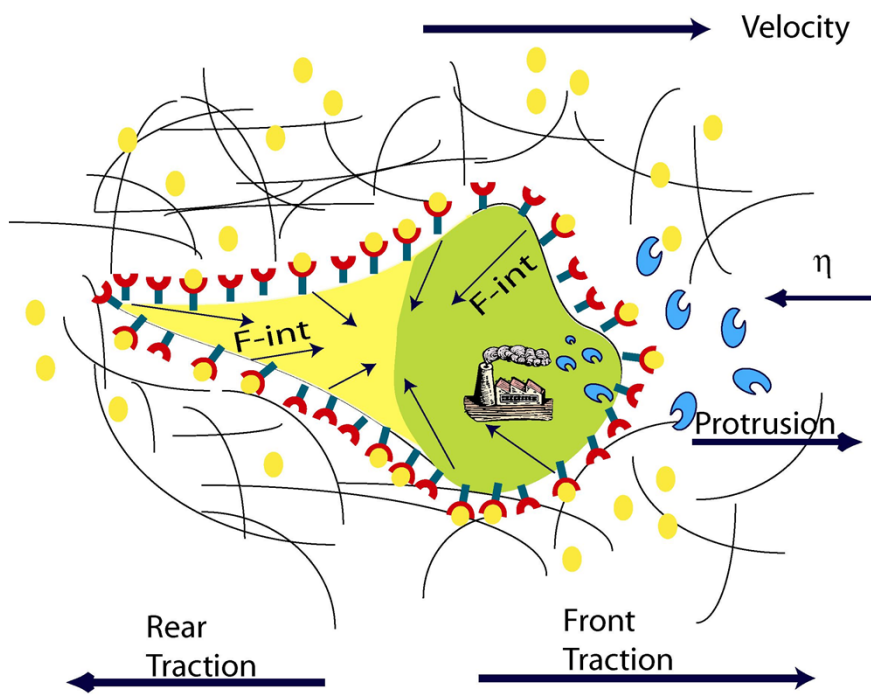
“All of the Above” -- Complex Landscape of Cell/Matrix Adhesion and Matrix Steric & Mechanical Properties Governs 3D Migration



Adhesion-inhibitory antibody decreases migration in some situations but increases it in others...

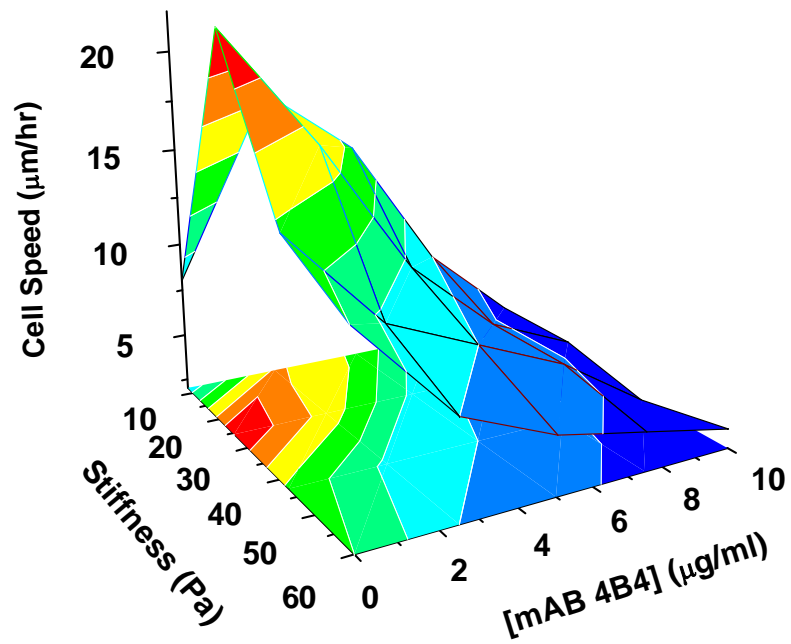


3D Cell Migration Computational Model, Single-Cell Biophysical Simulation (Biophys. J. [2005])

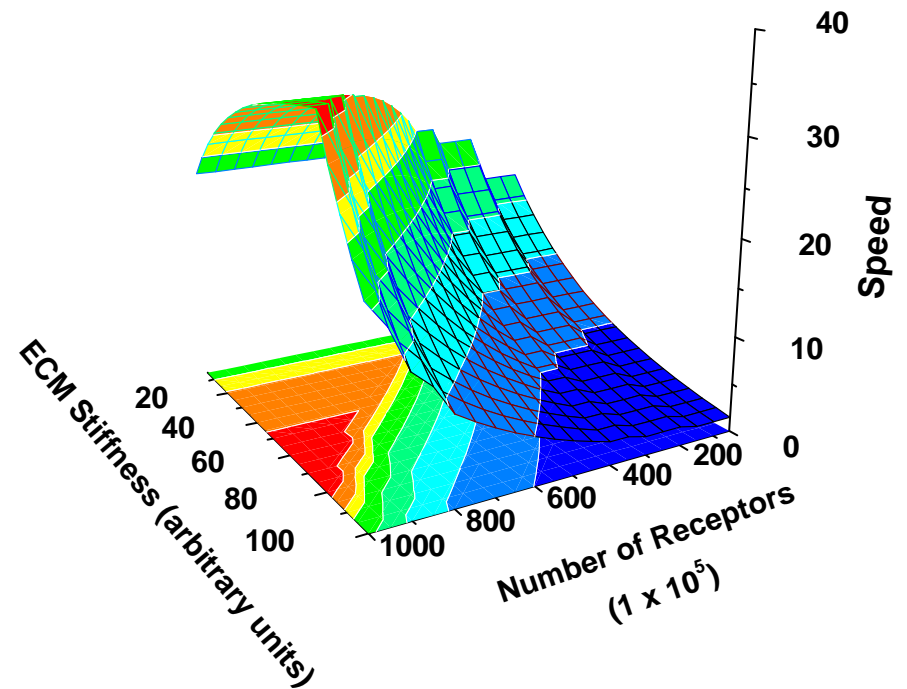


- Cell Dependent parameters
- Cell and Matrix Dependent parameters
- Matrix Dependent parameters
- Binding Constants

Model - Experiment Comparison

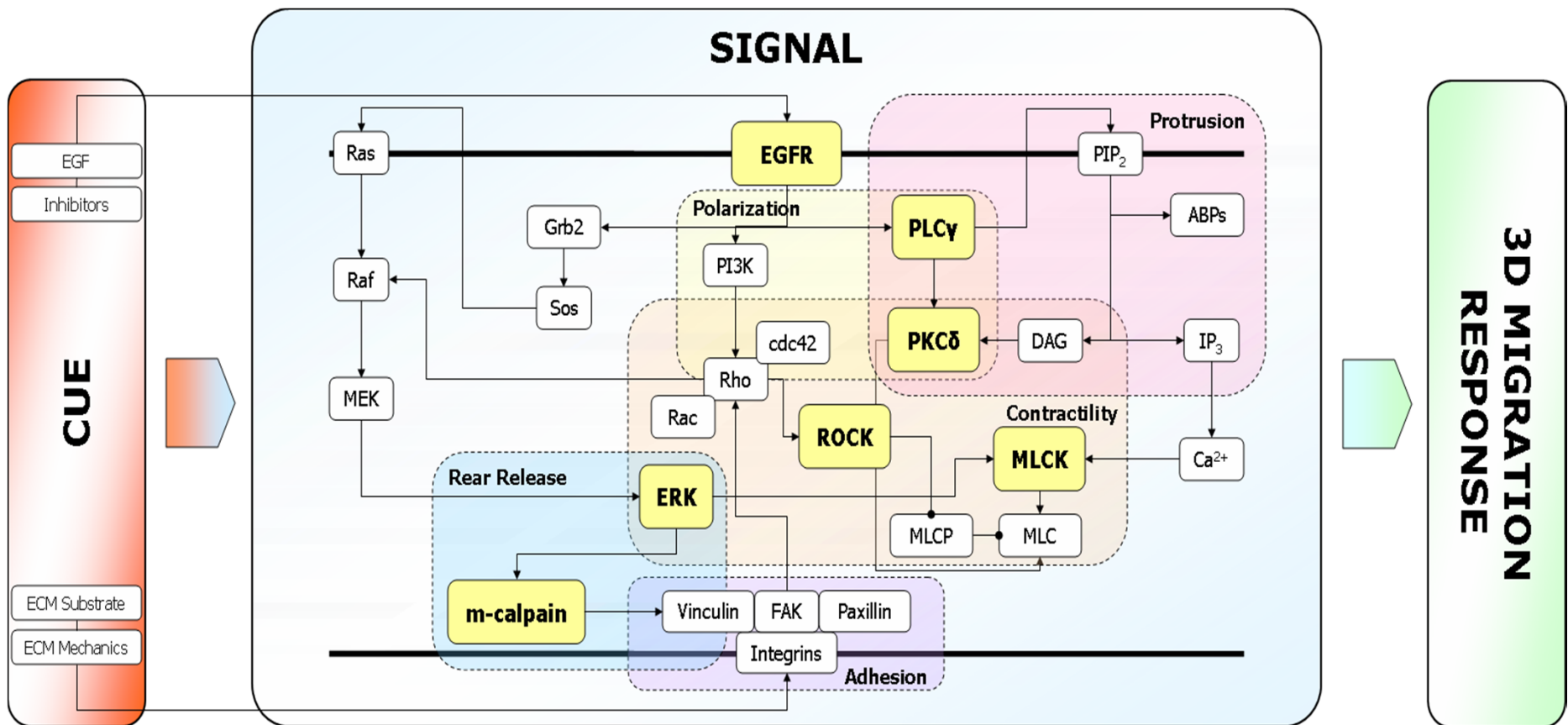


Experimental Data

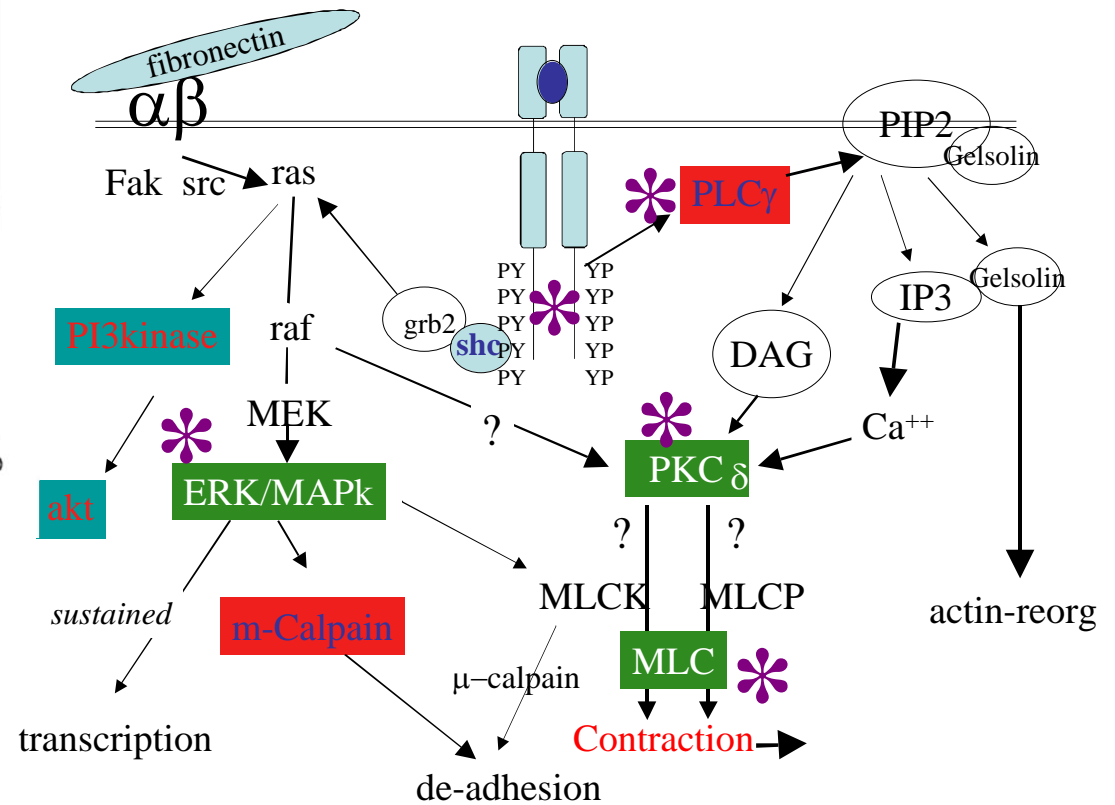
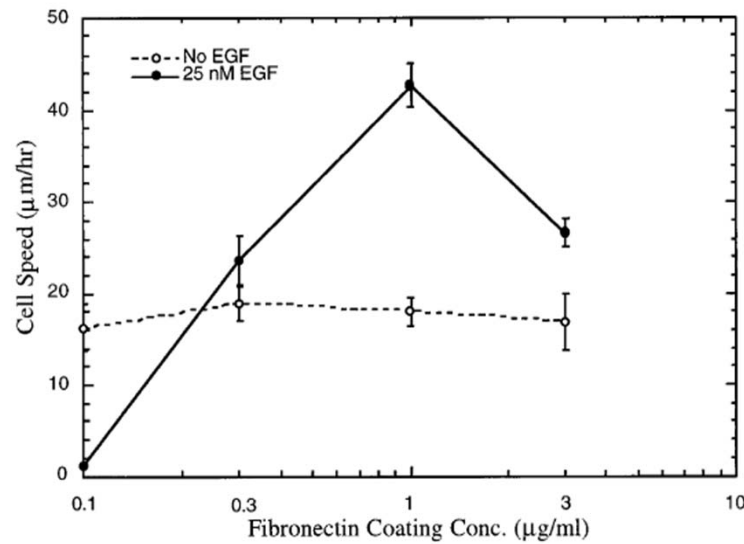


Model Predictions

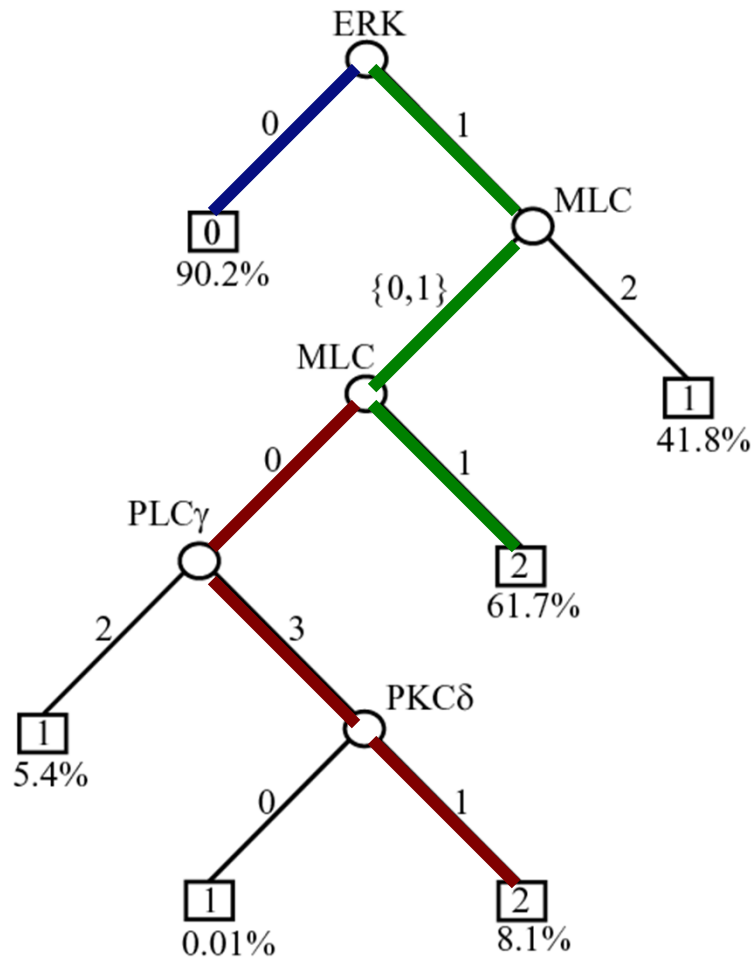
*Then, how are these biophysical motility processes and resulting migration controlled by biochemical signals?
 -- for prospective drug effects*



Example:
How do signals downstream of EGF and Fibronectin integrate to influence cell migration speed?



Decision Tree Model for 5-Minute Signaling Data



(70% overall accuracy)

IF p-ERK is low: cell migration is slow; ~90% of the slowly-migrating cohort observations can be explained with this rule.

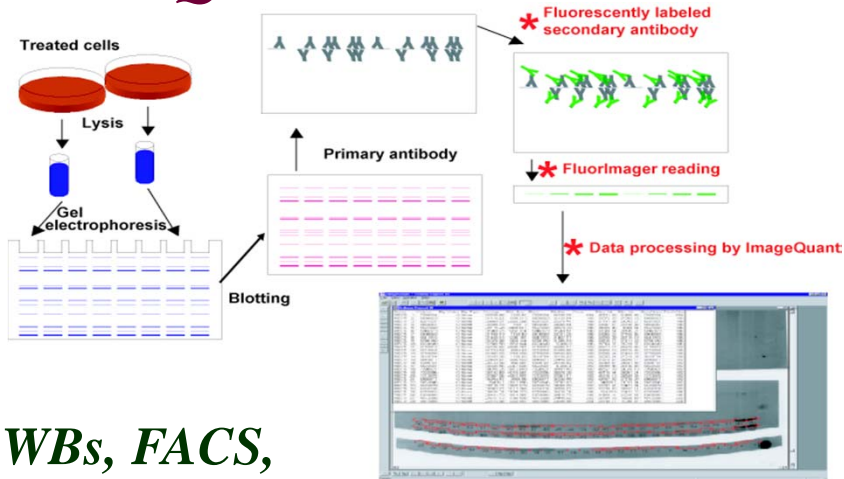
IF p-ERK is high AND p-MLC is intermediate: cell migration is fast; ~60% of the swiftly-moving cohort observations can be explained with this rule.

Another ~10% of the swiftly-moving cohort observations can be explained with the rule IF p-ERK is high AND p-MLC is low AND p-PLCγ is high AND p-PKCδ is high.

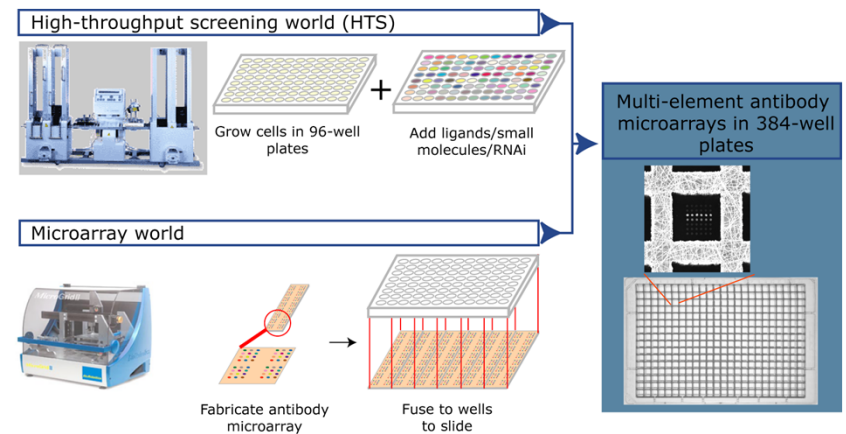
“Network Logic” indicates predominant combined roles of MLC and ERK in regulating the critical balance between cell contractile force and cell/substratum adhesion for governing migration speed.

(Bioinformatics [2005])

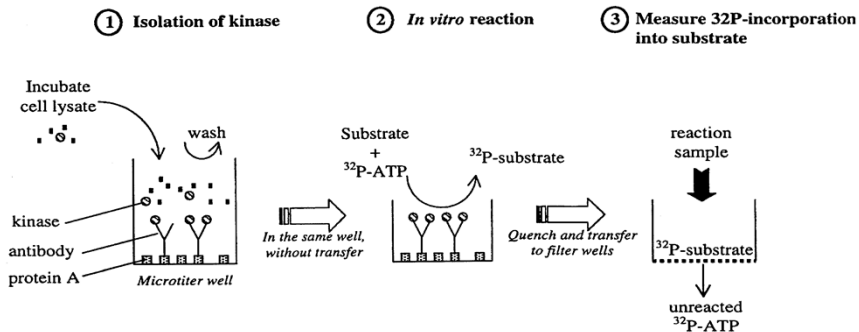
Moving Forward -- "High-Throughput" Protein-Centric Quantitative Measurement -- Protein Levels, States, Activities, Locations, Interactions...



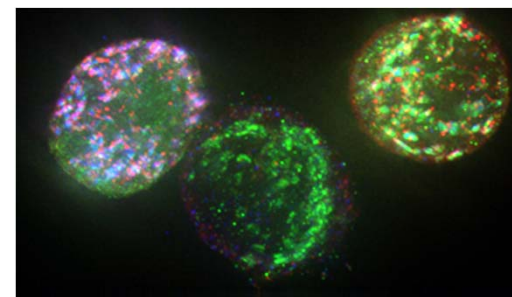
WBs, FACS, mass spectrometry



Multi-well kinase activity assays

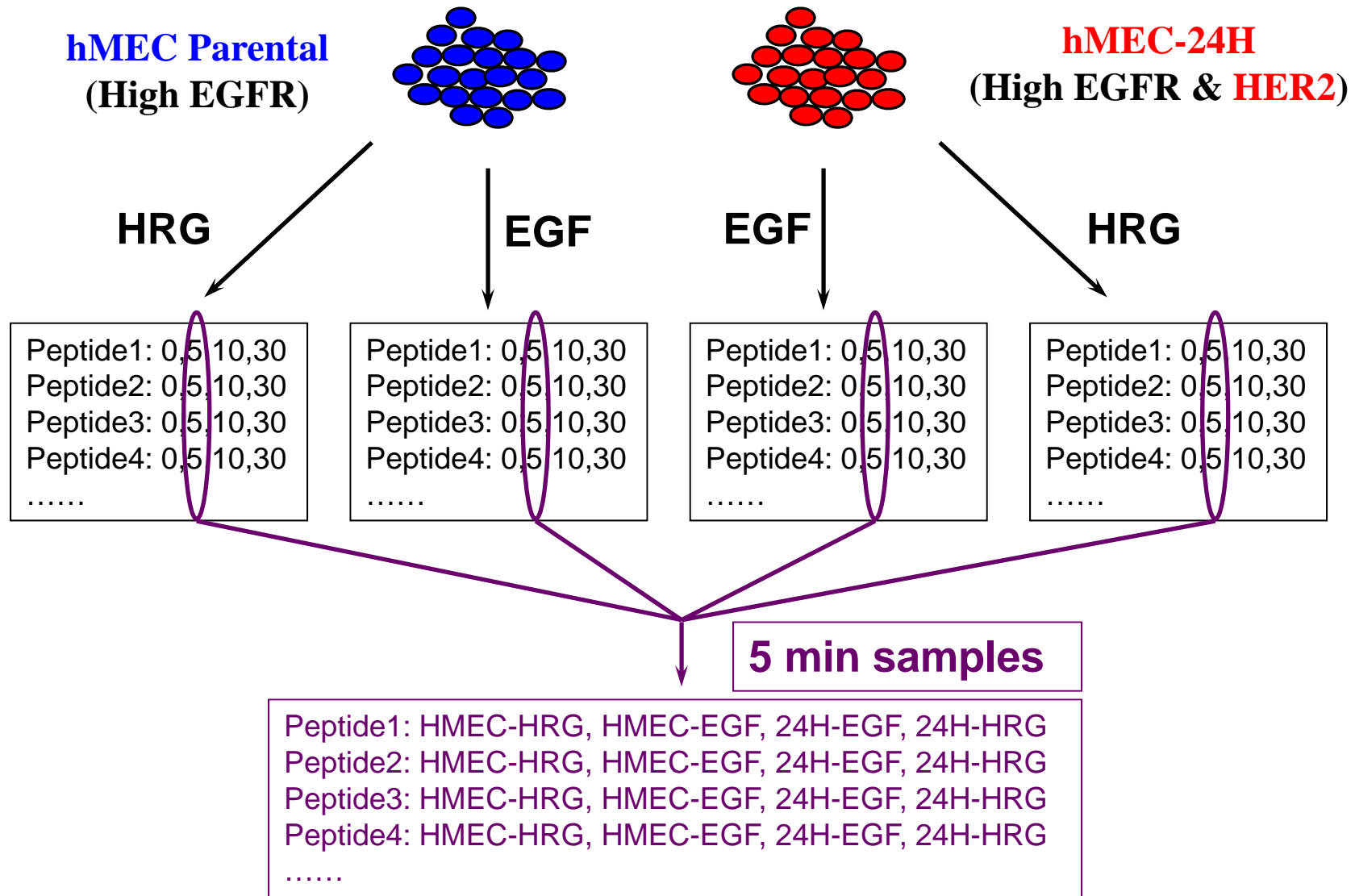


Protein microarrays

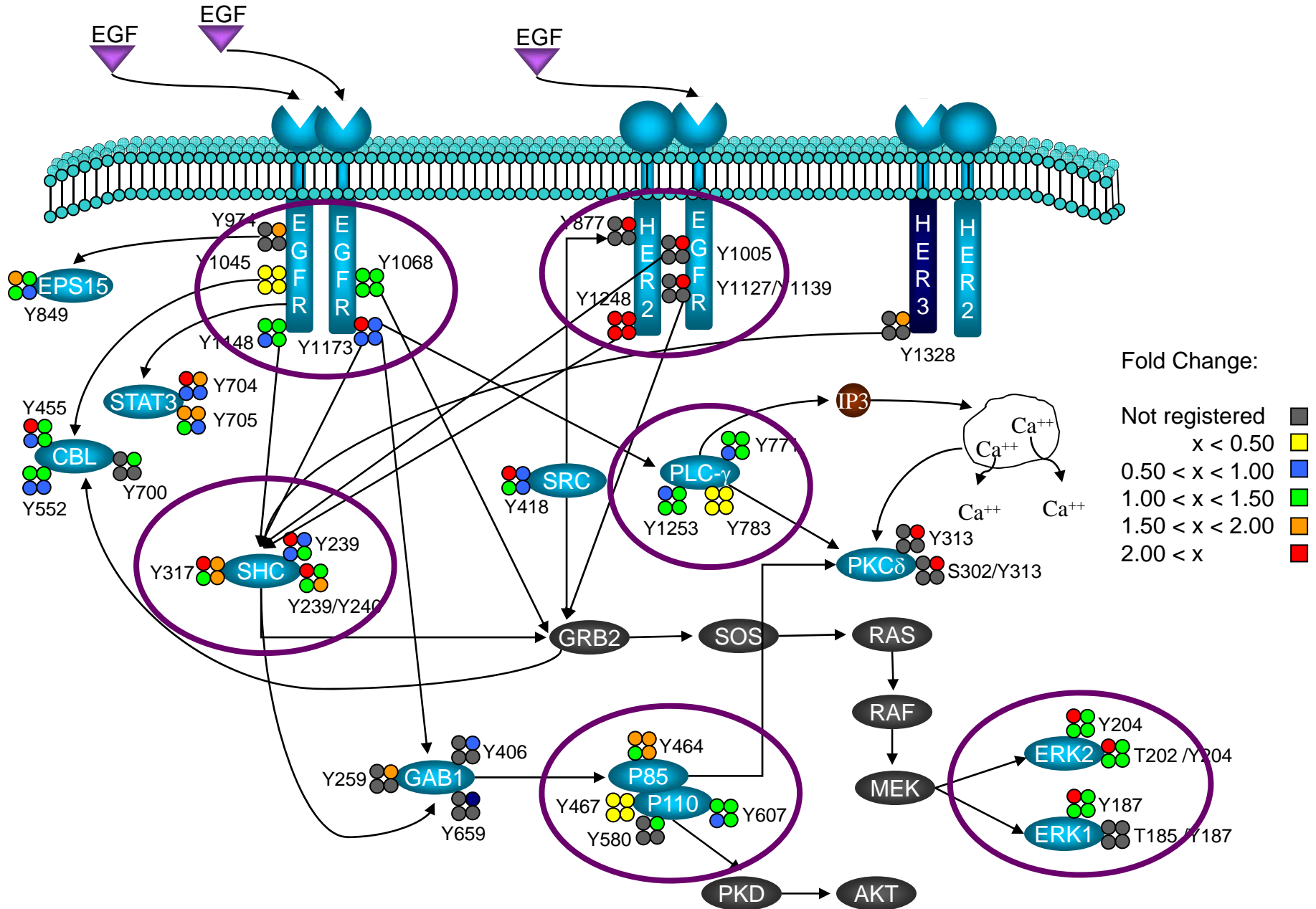


Live-cell imaging

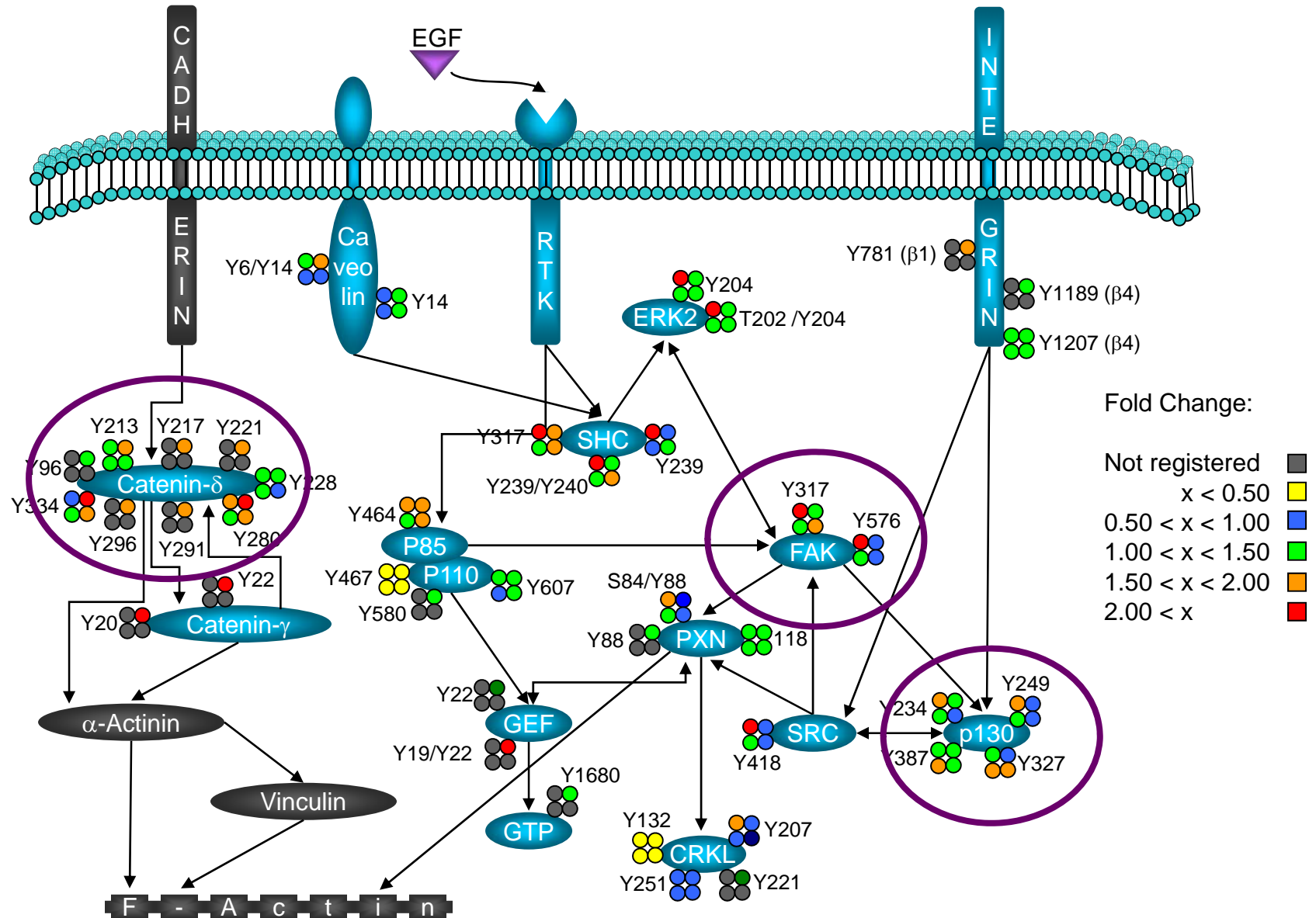
*Example Problem:
HER2 Over-Expression (e.g., breast cancer)*



HER2 Over-Expression Effect on EGF-Induced Signaling - A



HER2 Over-Expression Effect on EGF-Induced Signaling - B

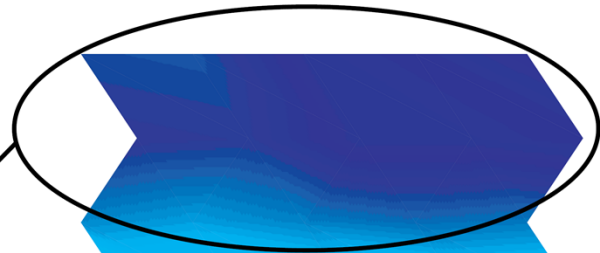


Neuronal Self-Organizing Maps (4 cell states, 4 time-points)

-- elucidates consistent dynamic modules

ERK2|T|Y|185/187
 P38 A|Y|182
 paxillin|S|Y|84/88
 PTRF|Y|308
 PZR|Y|263
 EGFR|Y|1173
 SHC|Y|239
 SHC|Y|317
 C18 of 11|Y|297
 ERK1|Y|204
 ERK2|Y|187
 STAT3-1|Y|705
 STAT3-2|Y|704
 Ack|Y|857
 EGFR|Y|1068
 EGFR|Y|1148
 SHIP-2|Y|986
 SHC|Y|Y|239/240
 An A2|Y|23
 An A2|Y|29
 TfR|Y|20
 Caveolin 1|Y|14
 Dsc3a|Y|818
 SCF38 m2|Y|20
 Rin1|Y|36

c1

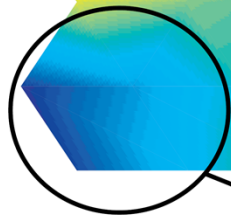


FAK|Y|397
 FAK|Y|576
 IGF1R|Y|1161
 RPTPa|Y|798
 GIT1|Y|545
 IGF1R|Y|1165
 paxillin|Y|118
 PI3KR2|Y|464
 RAI3|Y|347

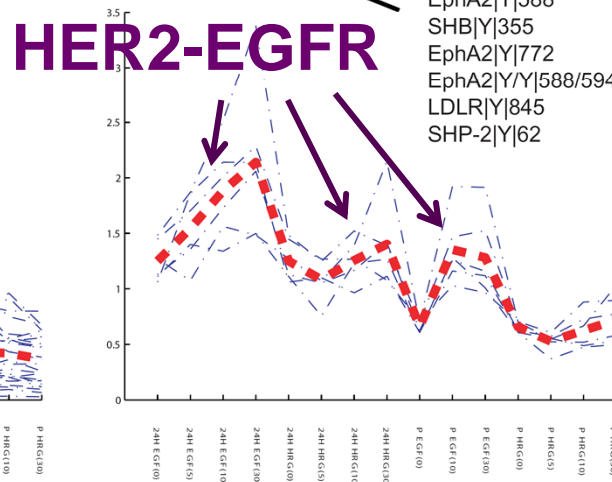
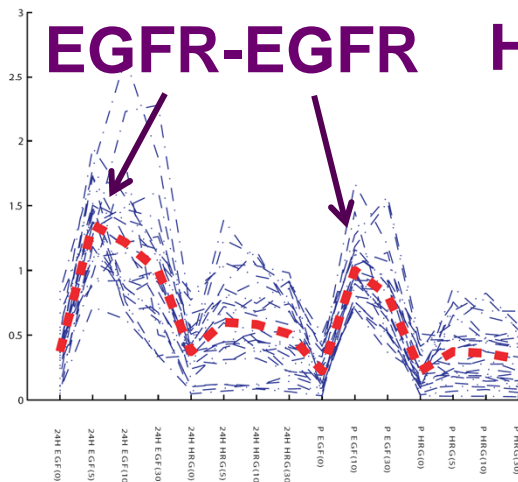
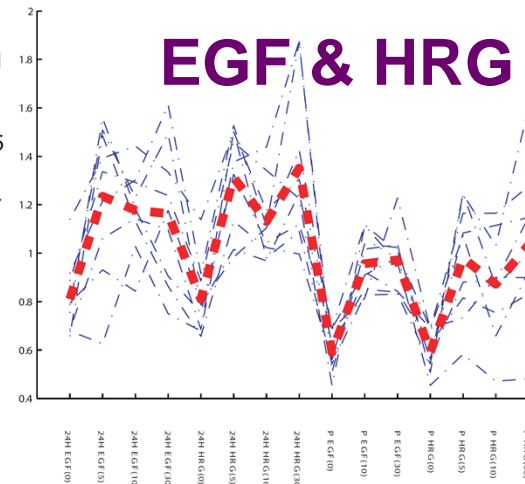
c4



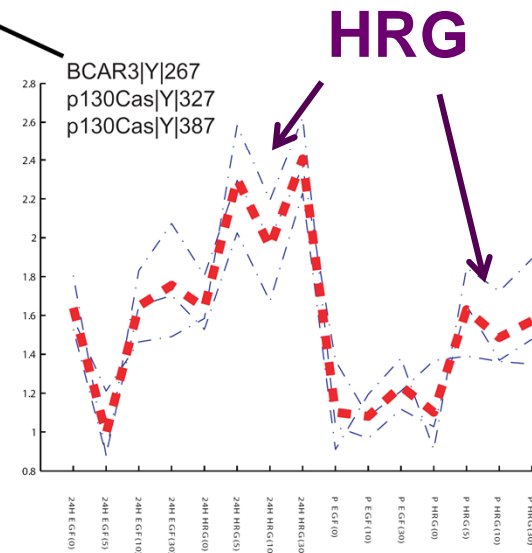
c3



c2



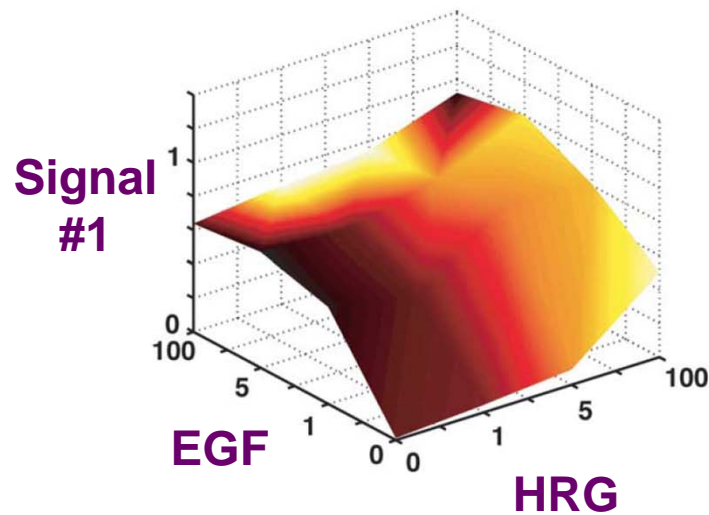
EphA2|Y|588
 SHB|Y|355
 EphA2|Y|772
 EphA2|Y|Y|588/594
 LDLR|Y|845
 SHP-2|Y|62



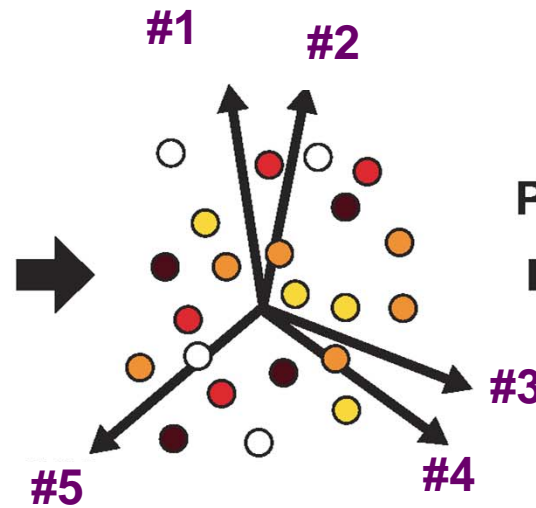
BCAR3|Y|267
 p130Cas|Y|327
 p130Cas|Y|387

Principal Component / Partial Least-Square Regression

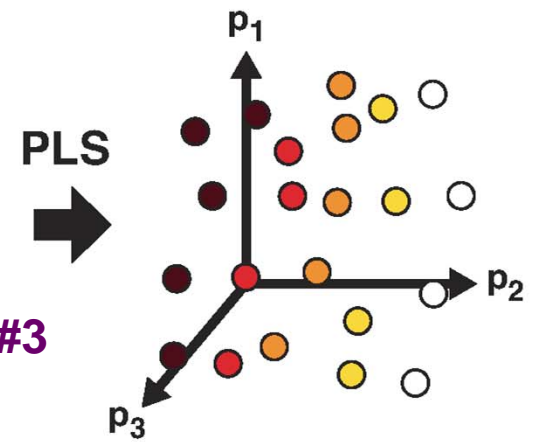
-- elucidates key signal combinations governing responses



Cue space



Signaling space

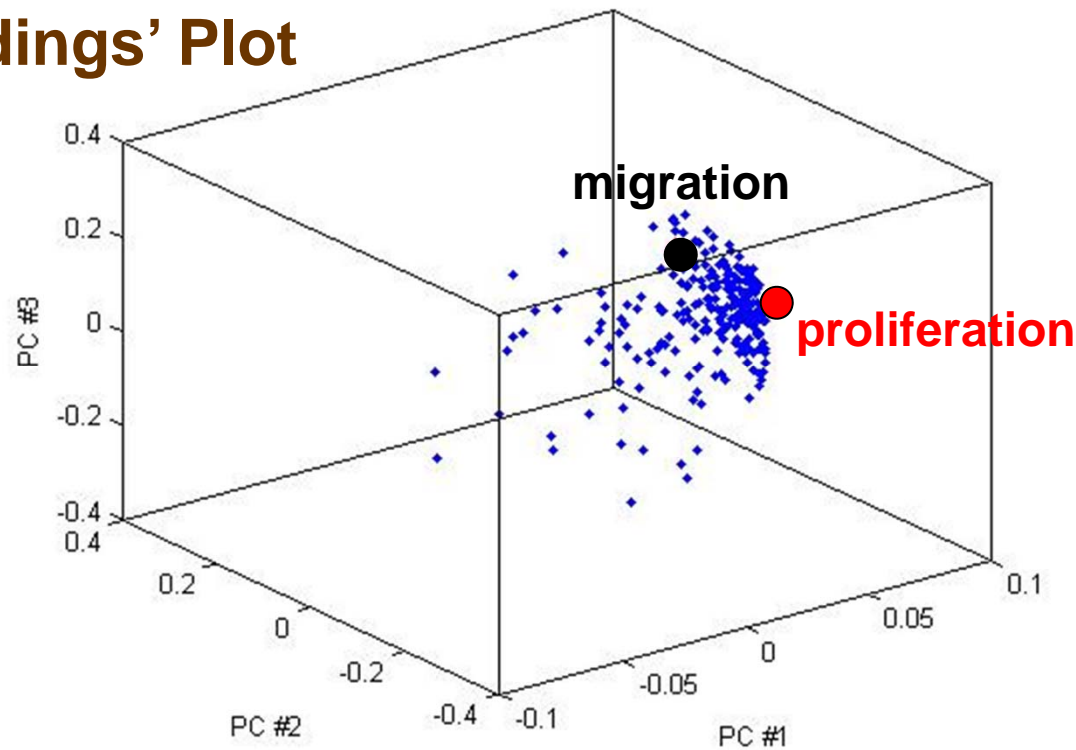


Principal component space



*Determination of Key Signals Governing
Enhanced Proliferation and Migration Arising
from HER2 Overexpression
-- Prospective Prediction of Drug Effects*

PLSR 'Loadings' Plot



Modeling Chain Underway for Prediction of Effects of Gene Mutations and Drug Effects on Tumor Cells

