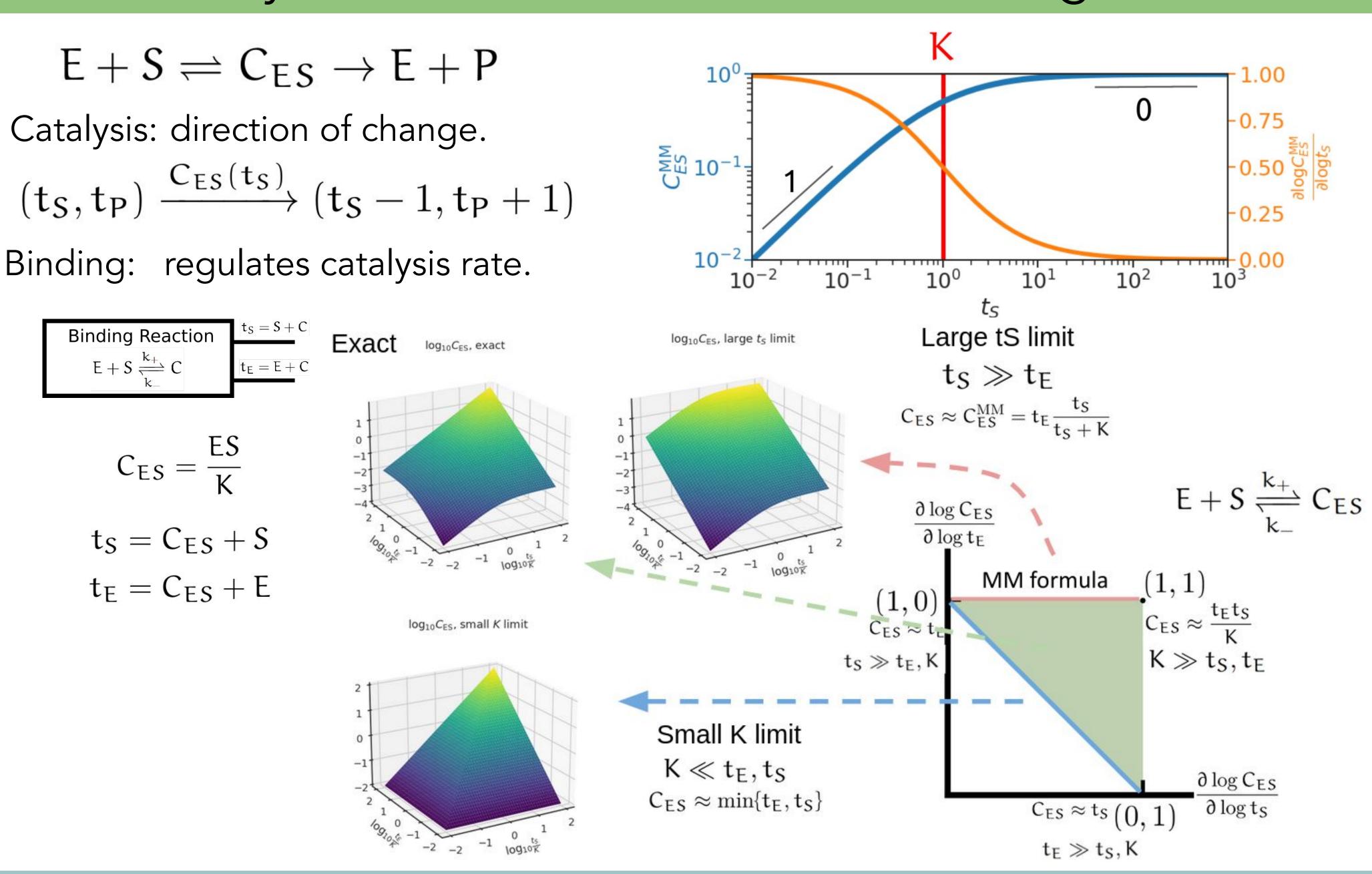
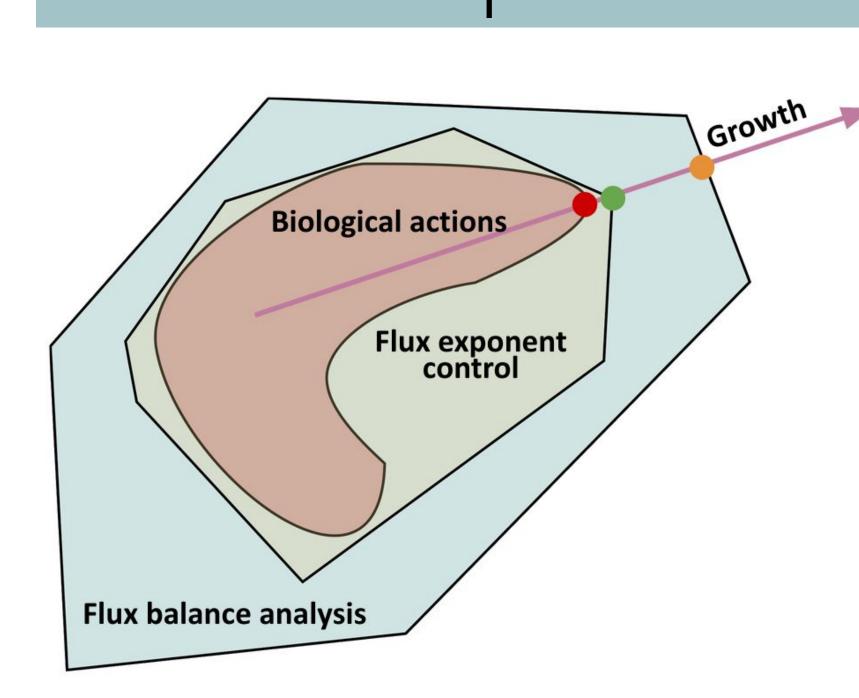
## Rule-based Systems Theory for Regulation in Networks of Biomolecules, Microbial Cells and Populations

John C. Doyle with students: Carmen Amo Alonso, Jing Shuang (Lisa) Li, Fangzhou (Fang) Xiao

## Theory Foundation: Full Profile of Bioregulation



## Flux Exponent Control for Dynamic Metabolism



Metabolism: known stoichiometry, unknown flux.

$$\dot{\mathbf{x}} = \mathbf{S}\mathbf{v}(\mathbf{x})$$

Constraint based methods: flux is optimal for some goal.

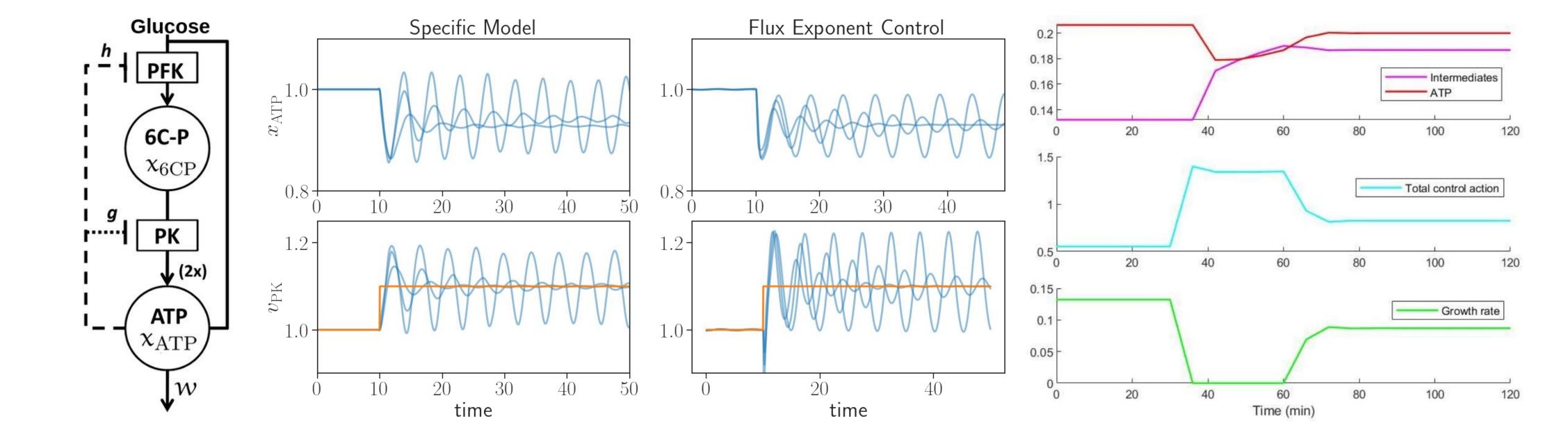
$$\dot{\mathbf{x}} = \mathbf{S}\mathbf{u}, \quad \mathbf{u}(\mathbf{t}) \in \mathcal{U}$$
 $\mathbf{v}(\mathbf{x}) = \arg \max_{\mathbf{u}} \mathbf{G}(\mathbf{u}, \mathbf{x})$ 

Flux Balance Analysis: steady state

$$\mathbf{v}(\mathbf{x}) = \arg \max_{\mathbf{u}} \mathbf{c}^{\mathsf{T}} \mathbf{u}$$
  
 $\dot{\mathbf{x}} = 0 = \mathbf{S}\mathbf{u}, \quad \mathbf{u} \in \mathcal{U}_{\text{static}}$ 

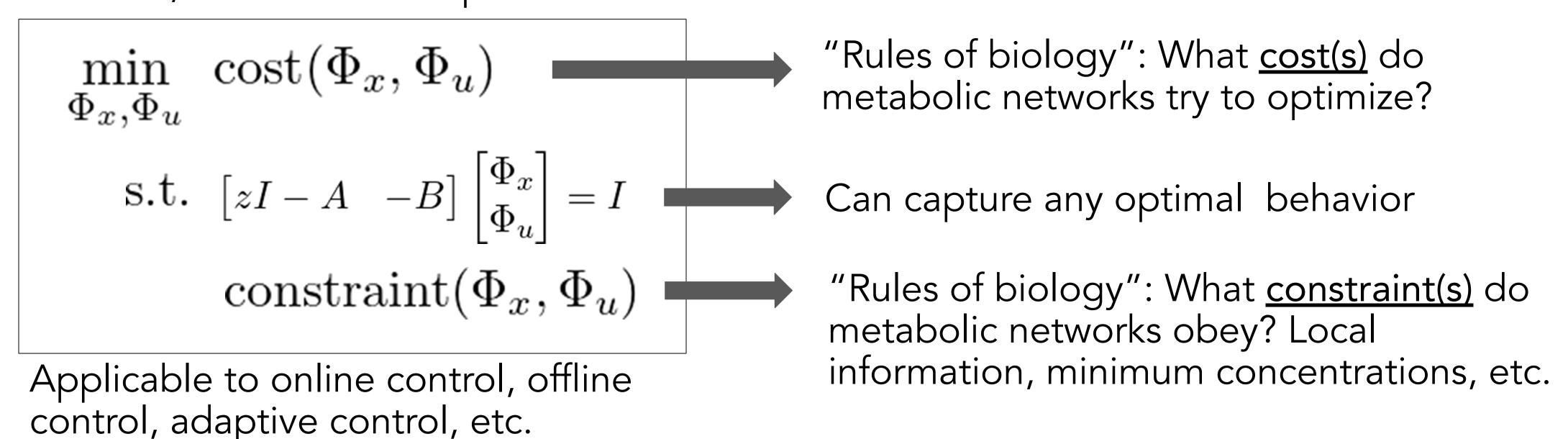
Flux Exponent Control: dynamic

$$\begin{split} \mathbf{v}(\mathbf{x}) &= e^{\mathbf{A} \log \mathbf{x} + \mathbf{B} \mathbf{u}^*}, \ \mathbf{u}^* = \arg \max_{\mathbf{u}} \mathsf{G}(\mathbf{x}, \mathbf{u}) \\ \dot{\mathbf{x}} &= \mathbf{S} e^{\mathbf{A} \log \mathbf{x} + \mathbf{B} \mathbf{u}}, \ \mathbf{u} \in \mathcal{U}_{dynamic} \end{split}$$



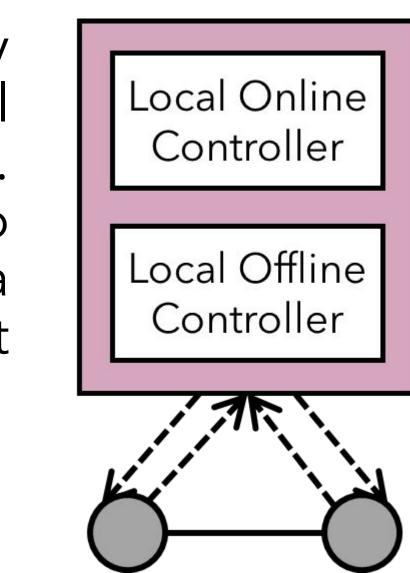
## System Level Synthesis for Dynamic Metabolic Modeling

Scalable, distributed computation



Fluxes are influenced by information on local metabolite concentrations.

Use local controllers to generate optimal fluxes for a given cost and constraint



Online control: Model predictive control. Plan optimal time-trajectories of fluxes and metabolites in response to large changes

Offline control: react to small perturbations

Use optimal control to determine optimal time-trajectories of fluxes and metabolites. Metabolites are in grey, controllers (which control flux) are in green. Not all controllers are shown.

