

Dynamics of gene regulatory circuits

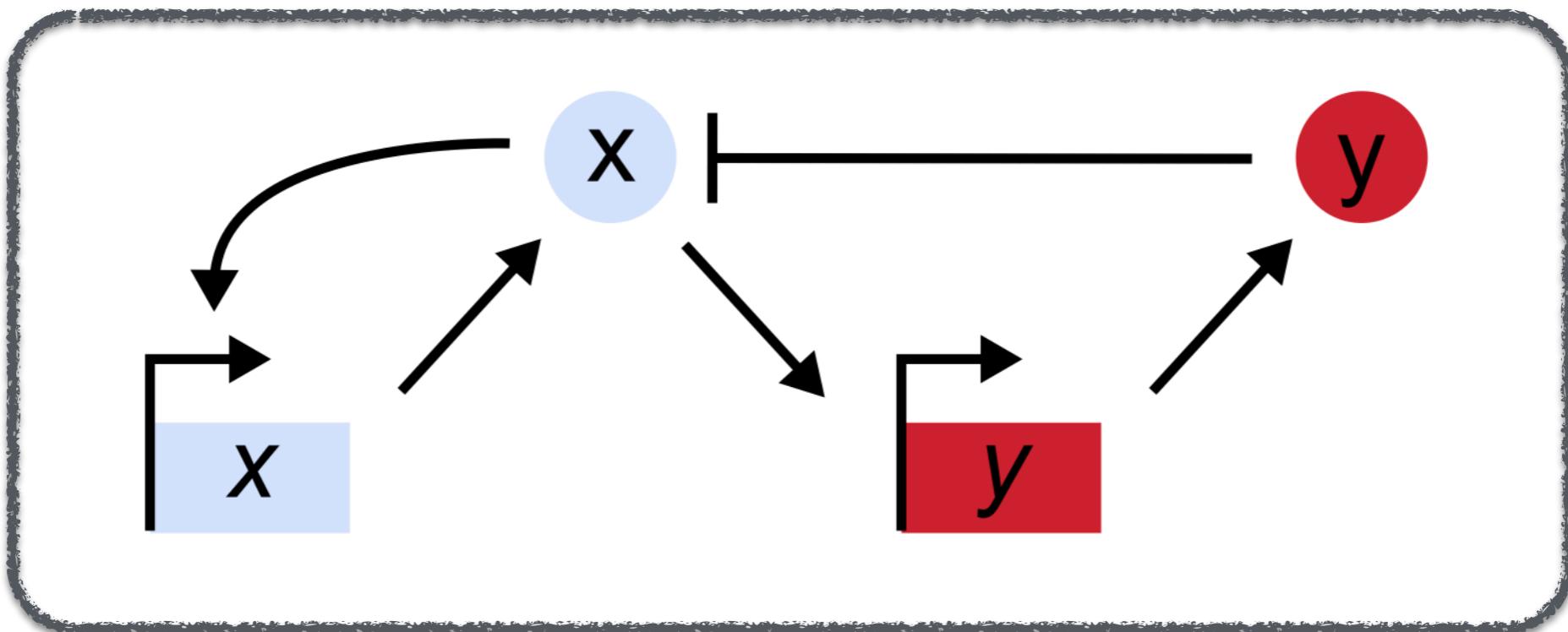
Jordi Garcia Ojalvo

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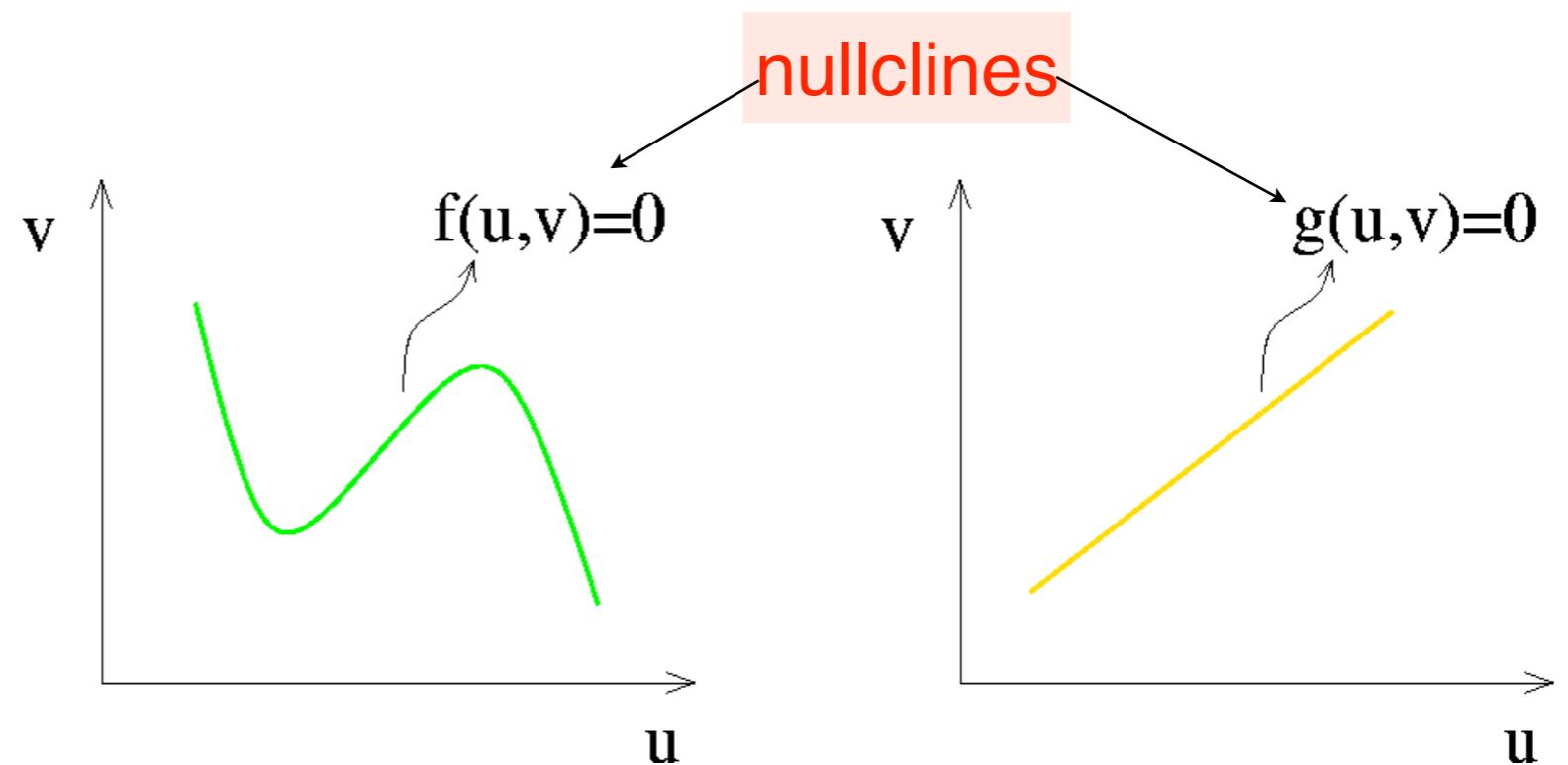
Combining a positive and a negative feedback



$$\frac{dx}{dt} = a_1 + \frac{b_1 x^n}{K_1^n + x^n} - gxy - d_1x$$
$$\frac{dy}{dt} = a_2 + \frac{b_2 x^m}{K_2^m + x^m} - d_2y$$

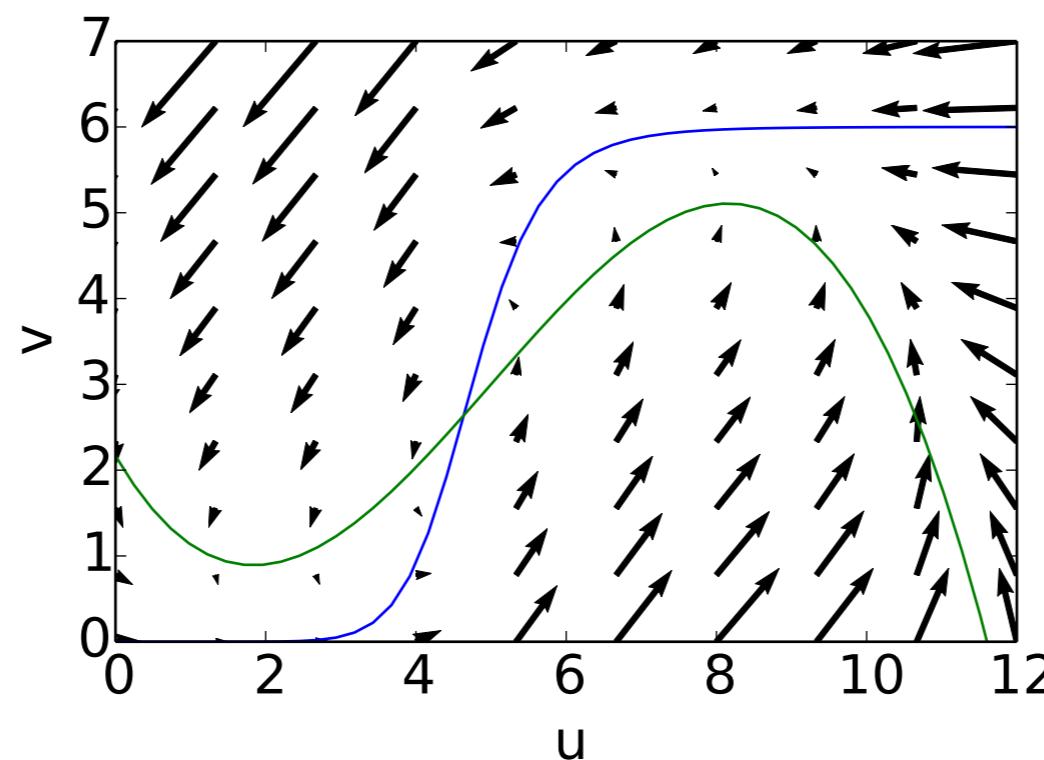
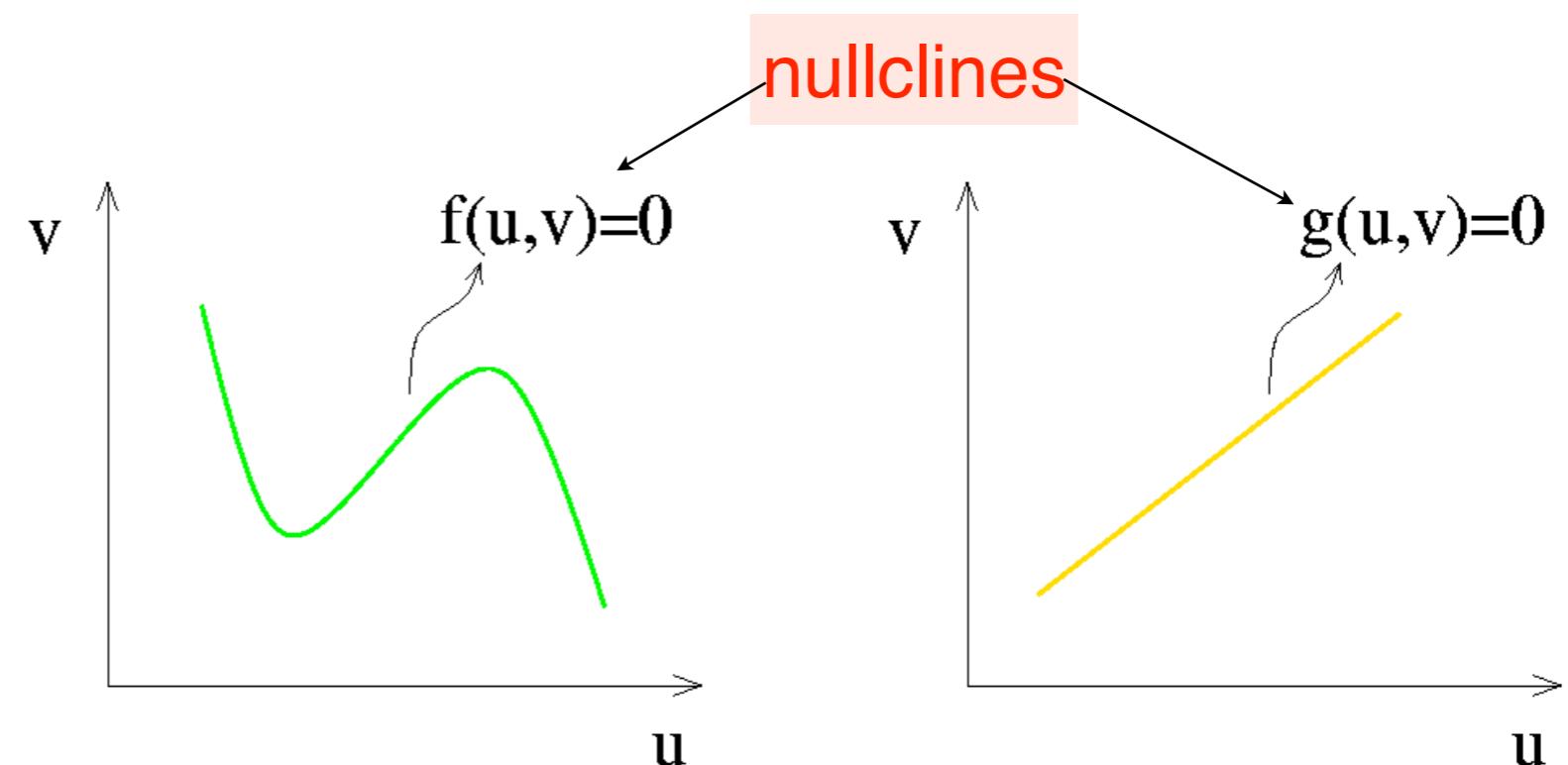
Two degrees of freedom: phase plane

$$\begin{aligned}\tau_u \dot{u} &= f(u, v) \\ \tau_v \dot{v} &= g(u, v)\end{aligned}$$



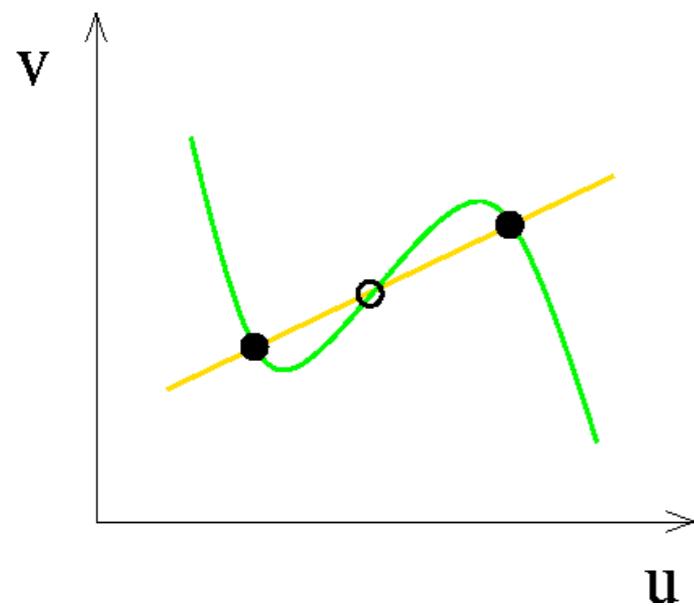
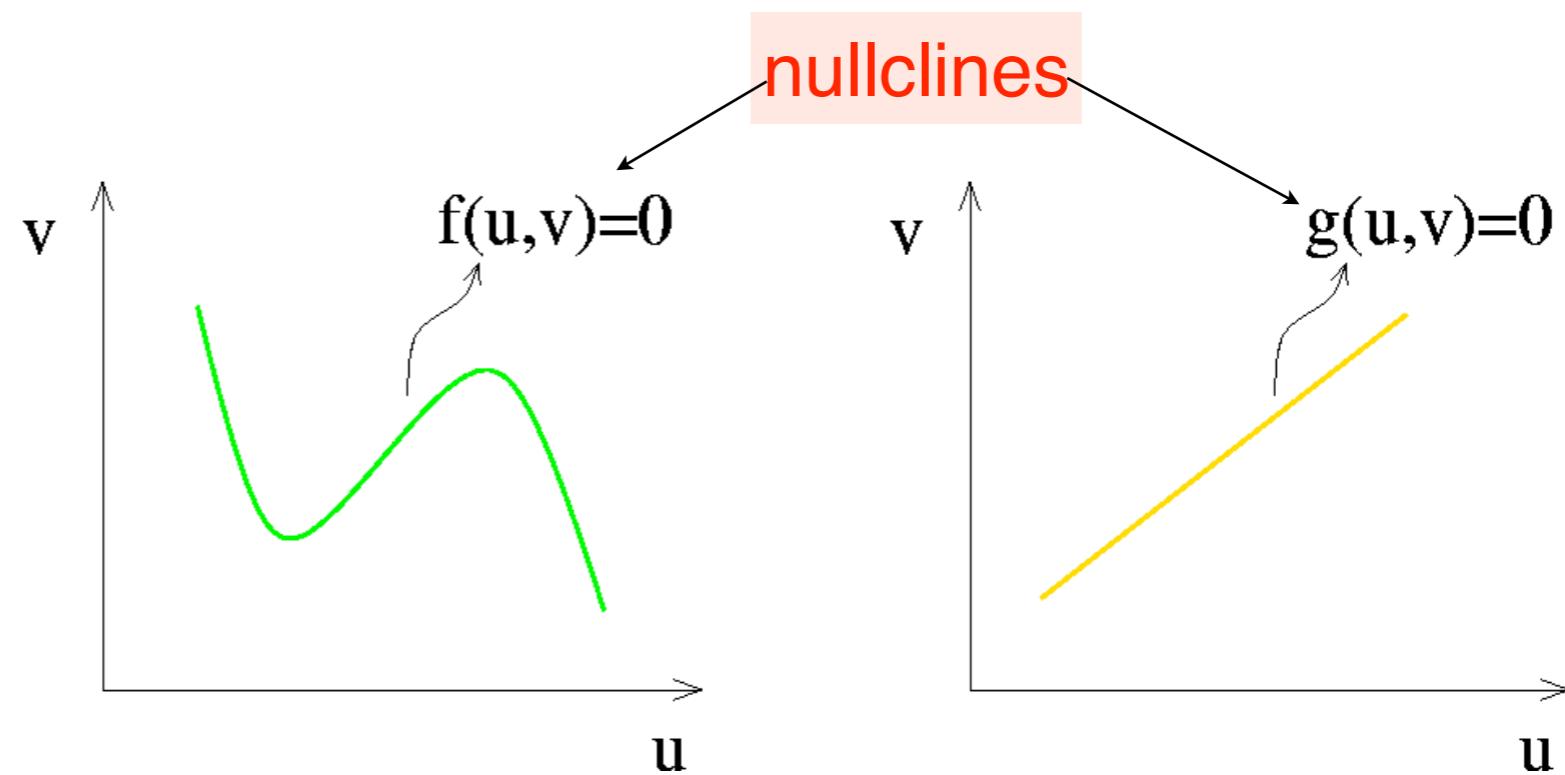
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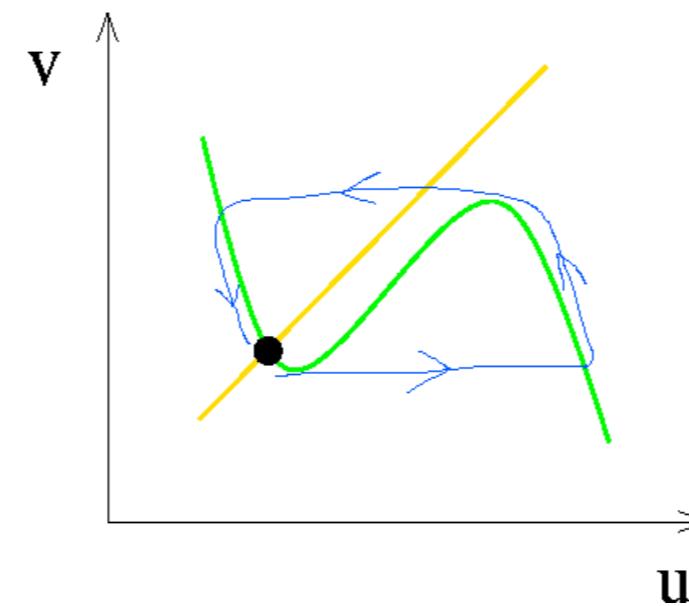


Two degrees of freedom: phase plane

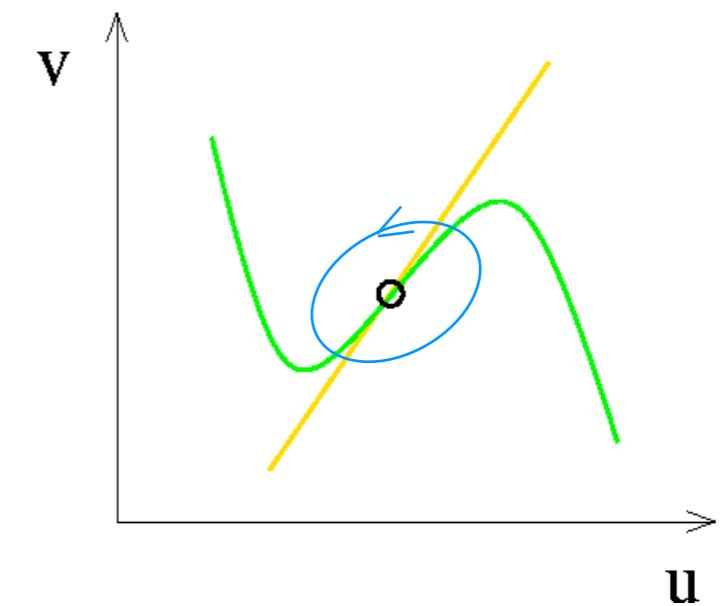
$$\begin{aligned}\tau_u \dot{u} &= f(u, v) \\ \tau_v \dot{v} &= g(u, v)\end{aligned}$$



BISTABLE

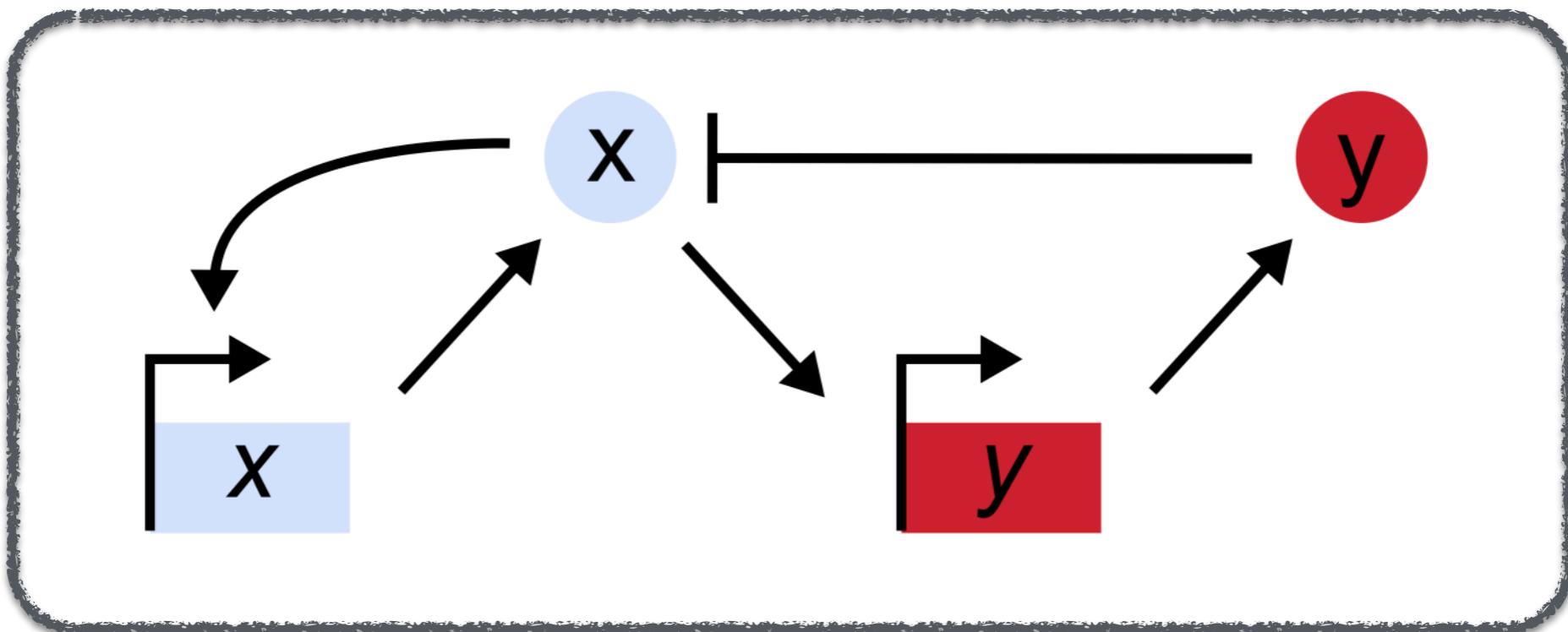


EXCITABLE



OSCILLATORY

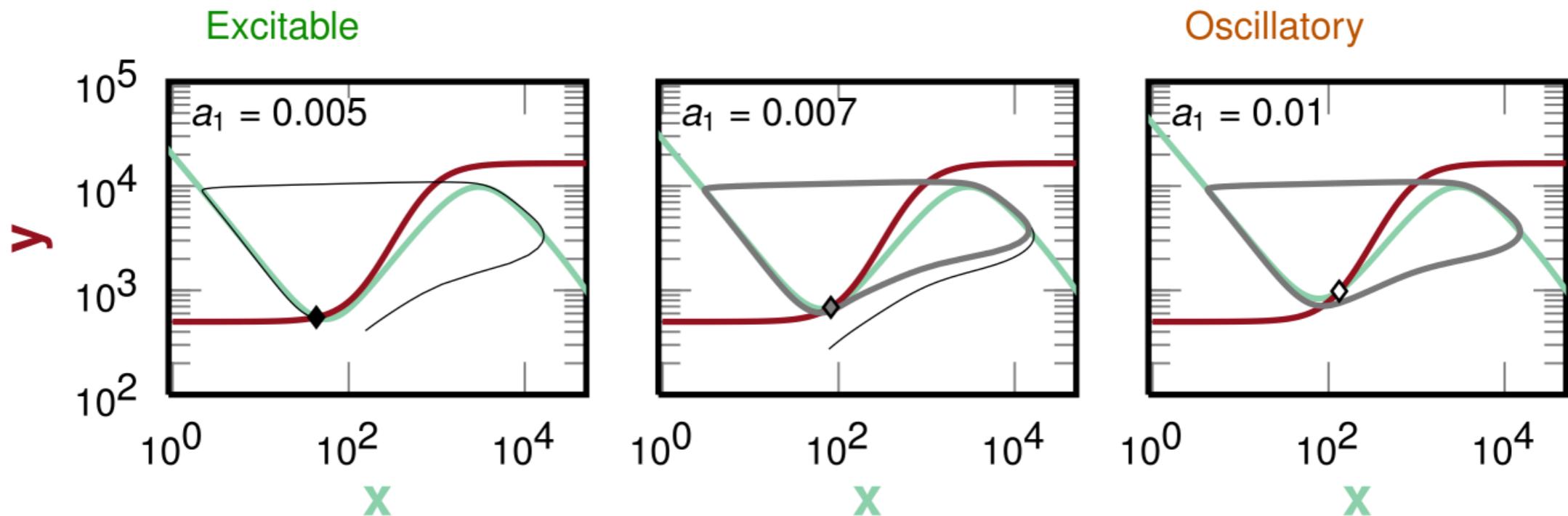
Combining a positive and a negative feedback



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$$\frac{dy}{dt} = a_2 + \frac{b_2 x^m}{K_2^m + x^m} - d_2y$$

From oscillations to pulses

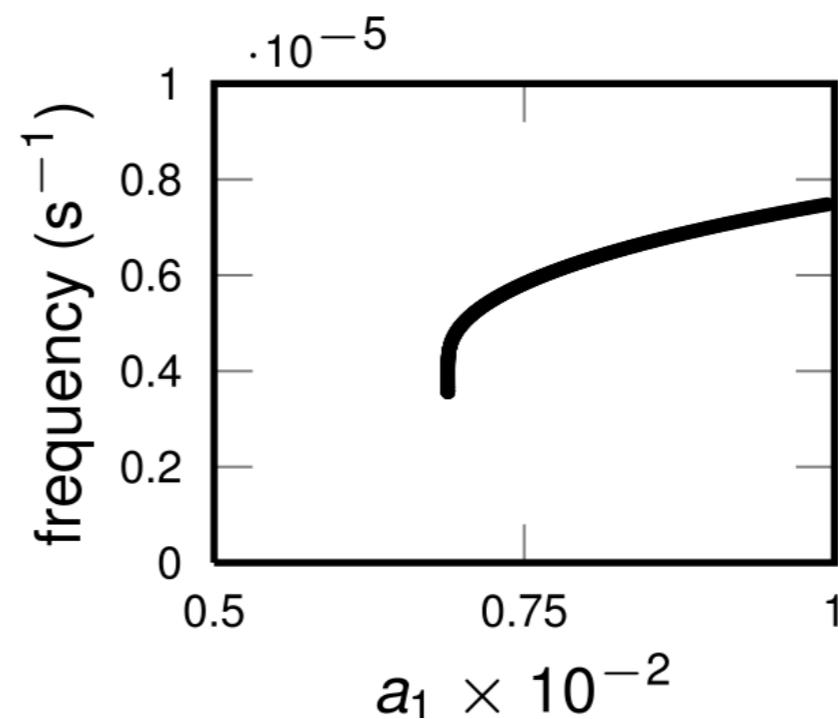
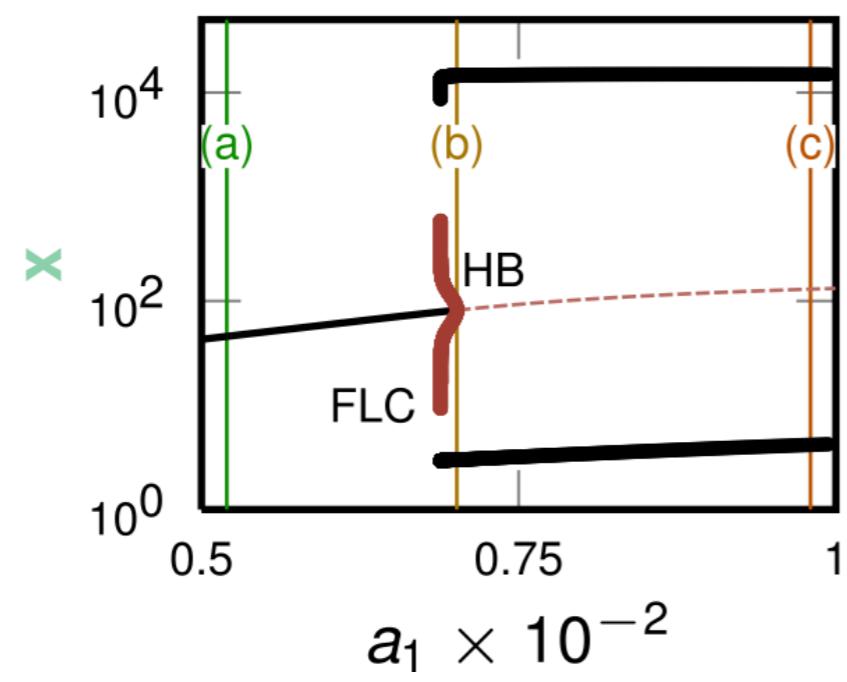
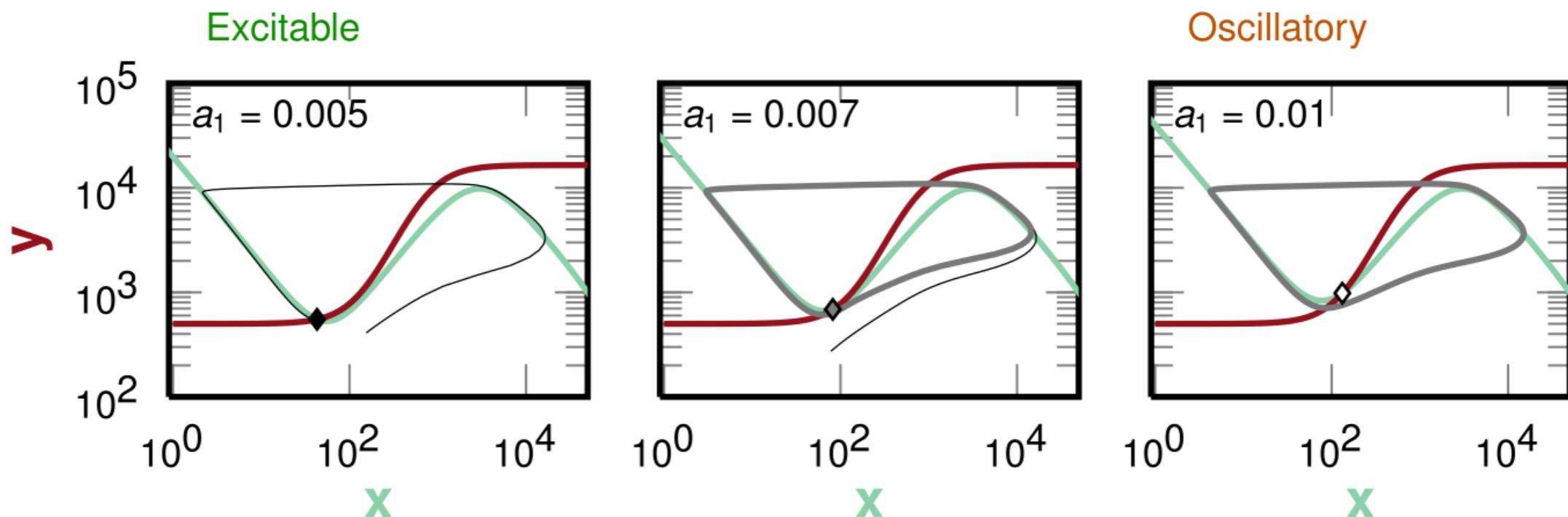
— $dx/dt = 0$ — $dy/dt = 0$ — trajectory
— limit cycle



$$\frac{dx}{dt} = \cancel{a_1} + \frac{b_1 x^n}{K_1^n + x^n} - gxy - d_1 x$$
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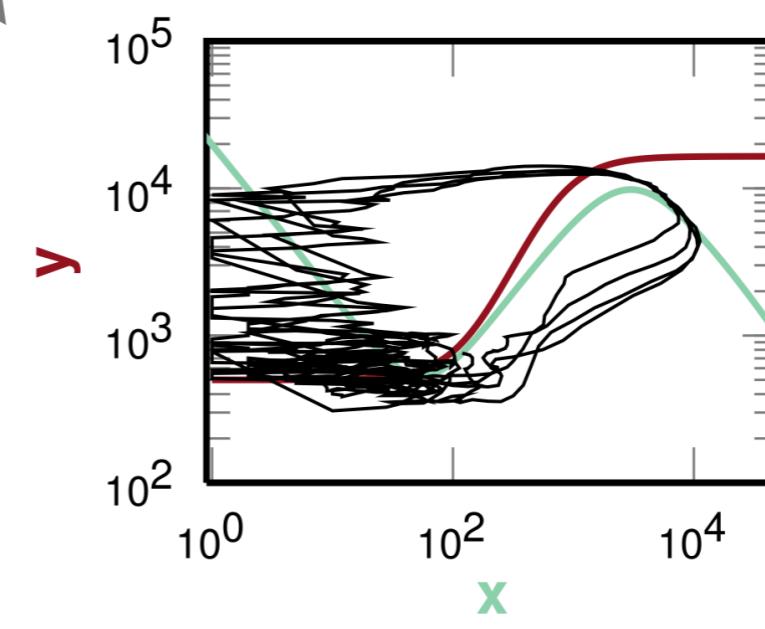
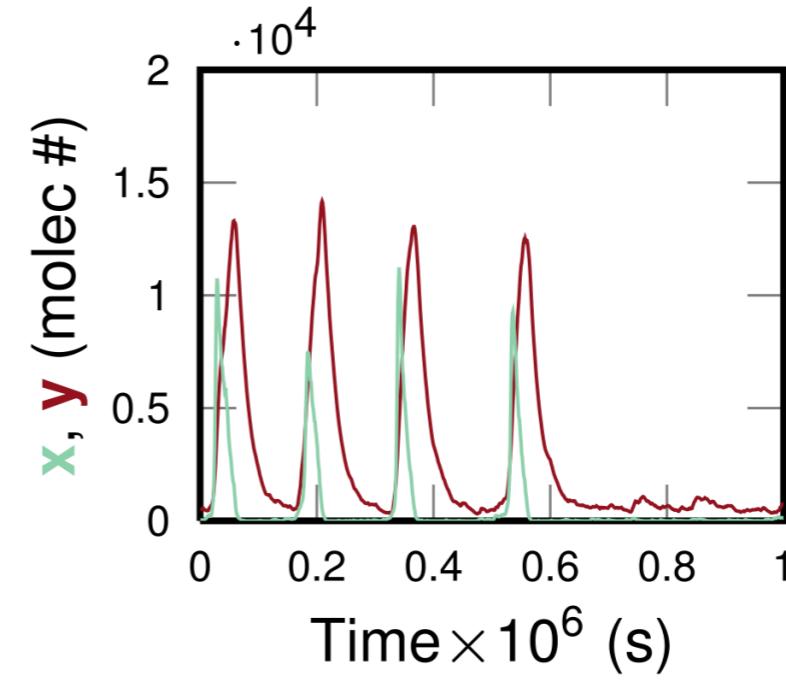
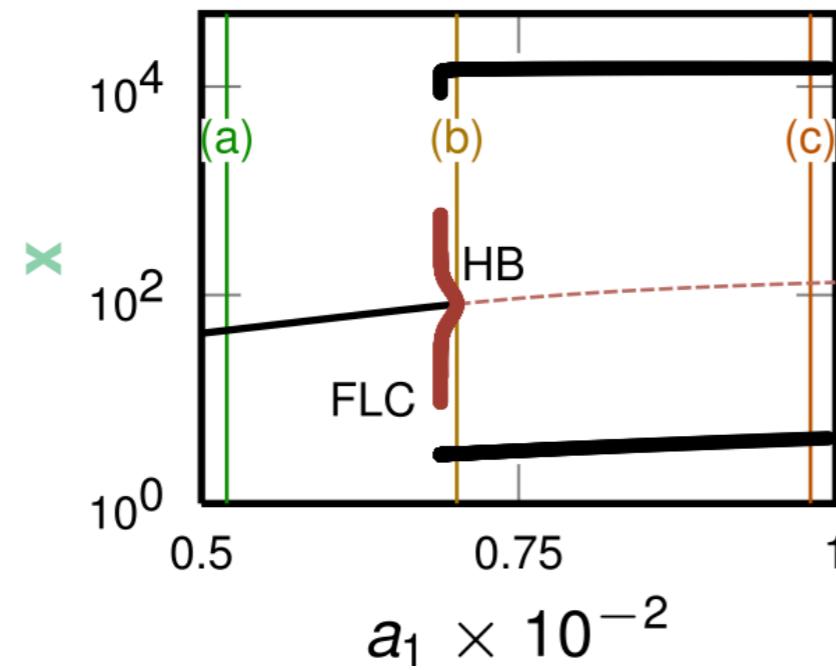
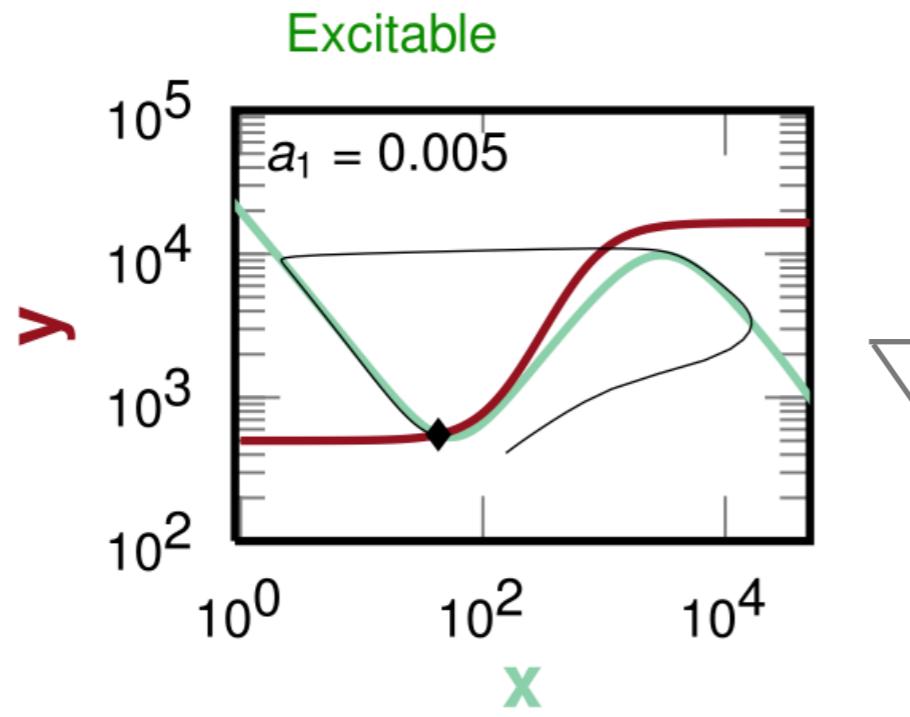
Bifurcation analysis

— $dx/dt = 0$ — $dy/dt = 0$ — trajectory
— limit cycle



Excitability

— $dx/dt = 0$ — $dy/dt = 0$ — trajectory
— limit cycle



ADVANCED SUMMER SCHOOL ON APPLIED DYNAMICS IN SYSTEMS AND SYNTHETIC BIOLOGY

EXERCISES - GENE REGULATORY CIRCUITS

subsequent decrease of α back down to 0 does not cause expression to turn off: the cell responds *irreversibly* to a pulse stimulus in α .

3. Consider the following activator-repressor model discussed in class:

$$\begin{aligned}\frac{dx}{dt} &= a_1 + \frac{b_1 x^n}{K_1^n + x^n} - gxy - d_1 x \\ \frac{dy}{dt} &= a_2 + \frac{b_2 x^m}{K_2^m + x^m} - d_2 y\end{aligned}$$

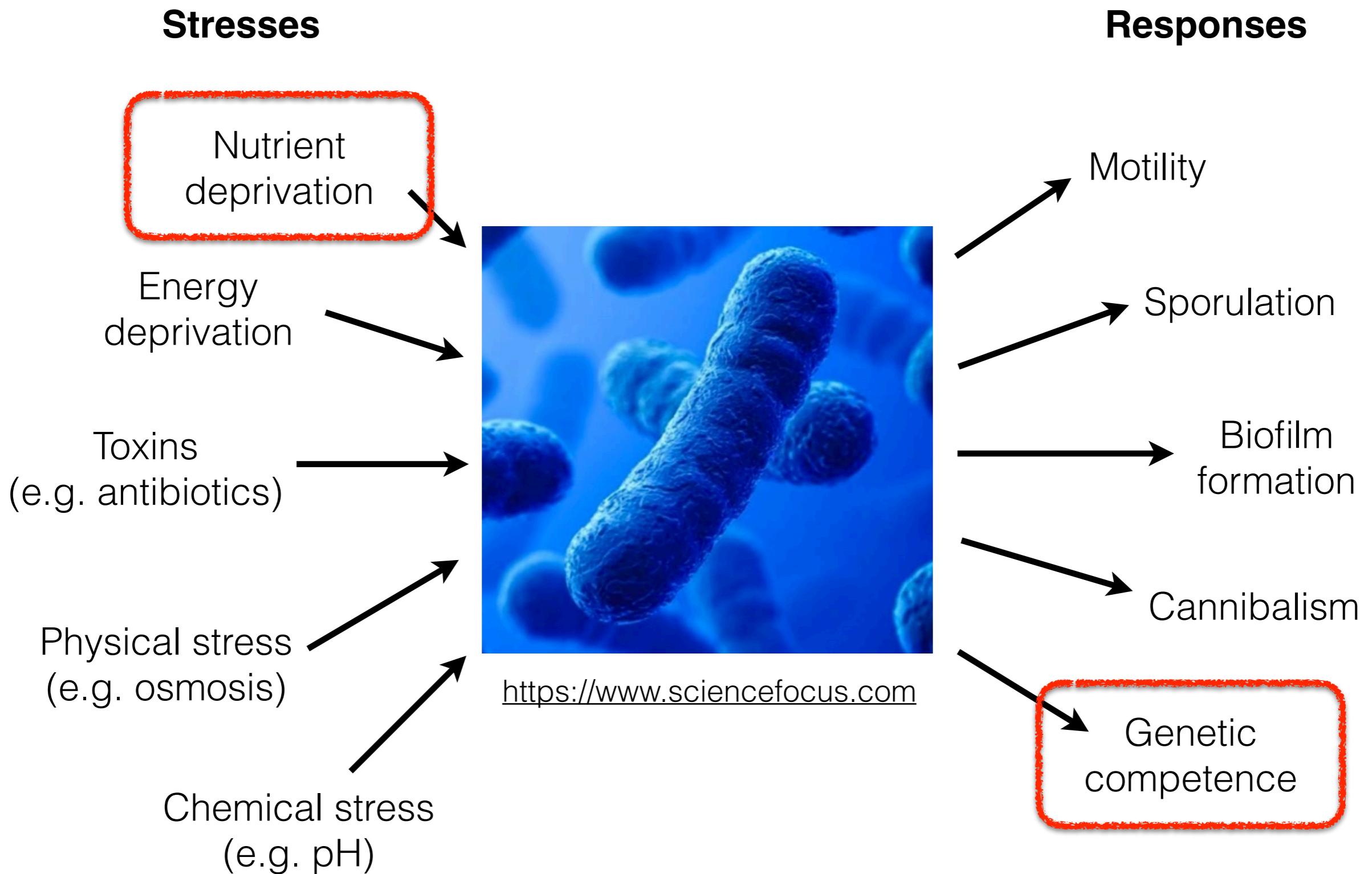
Use the tools that you have learned in the different courses of this school to analyze this dynamical system in the plane, as the basal expression rate a_1 of x varies, as shown in the slides. Use these values for the other parameters to perform simulations if necessary: $a_2=0.025$ nM/s, $b_1=15$ nM/s, $b_2=0.8$ nM/s, $d_1=d_2=5 \cdot 10^{-5}$ s $^{-1}$, $g=2.5 \cdot 10^{-7}$ nM $^{-1}$ s $^{-1}$, $K_1=3000$ nM, $K_2=750$ nM, and $n=m=2$.

4. The following model describes genetic competence in *B. subtilis*:

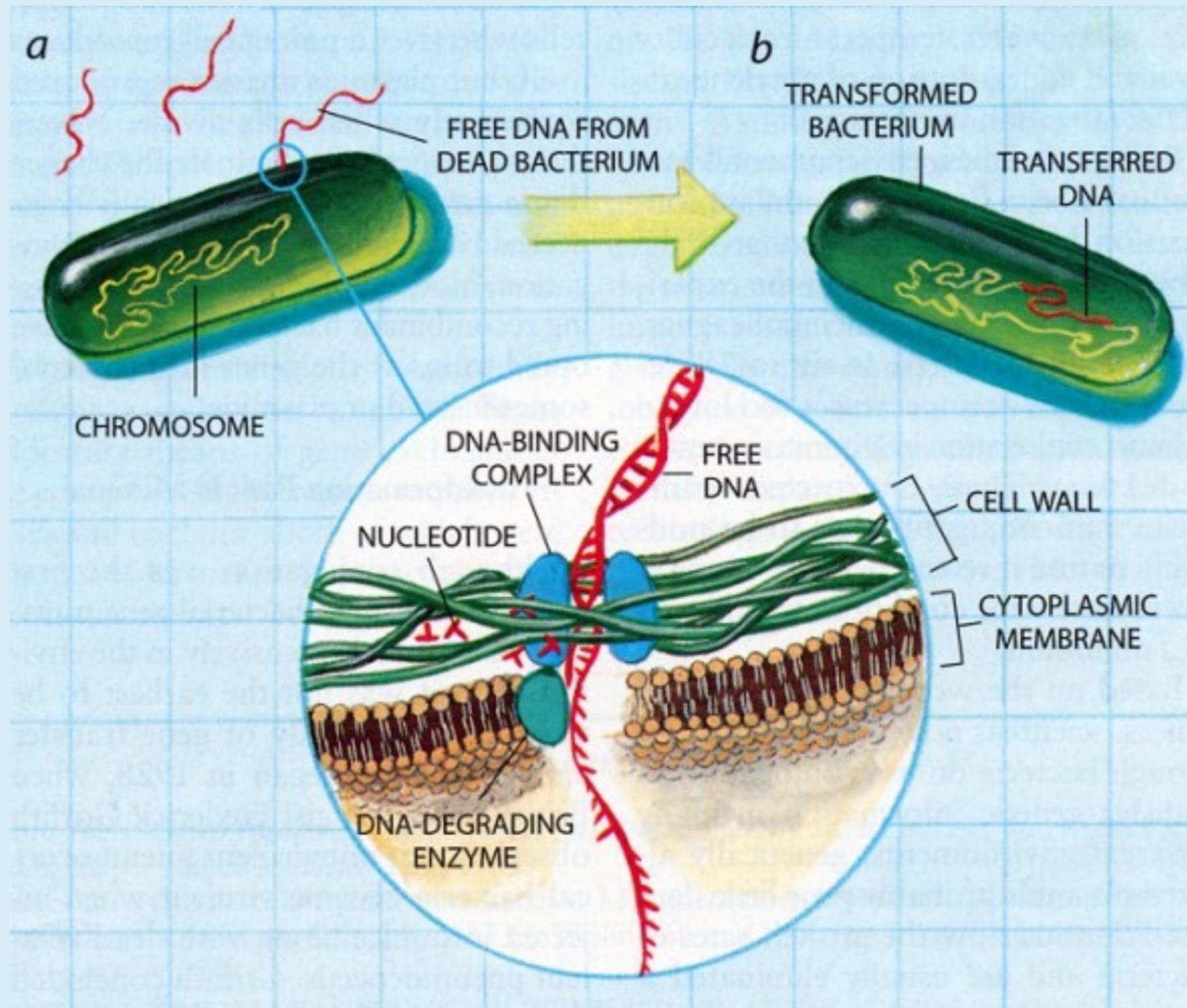
Dynamics of gene regulatory circuits

1. Gene circuit dynamics
2. Dissecting a genetic circuit
3. Noise in genetic circuits

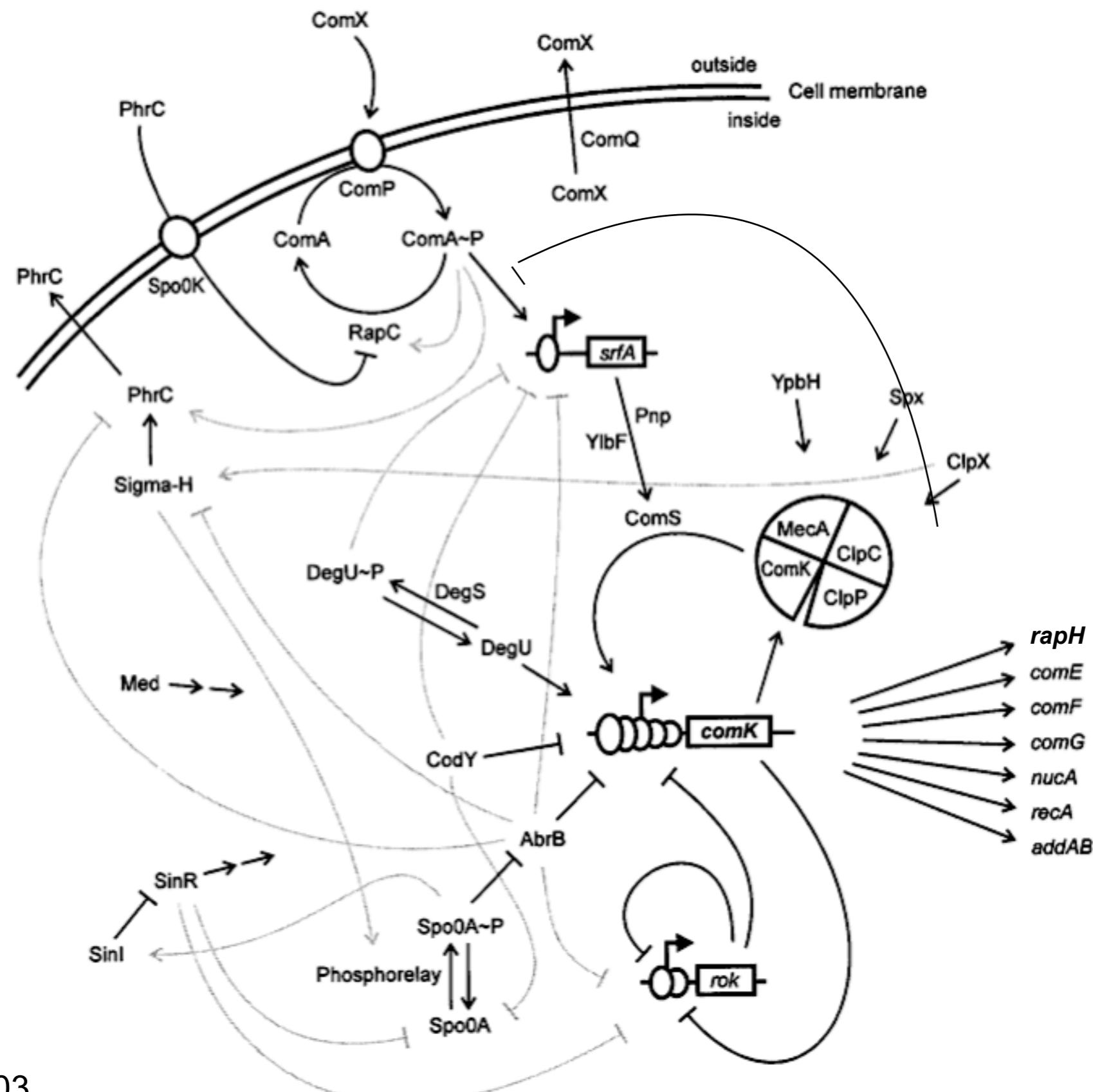
Stress responses of bacteria



Genetic competence



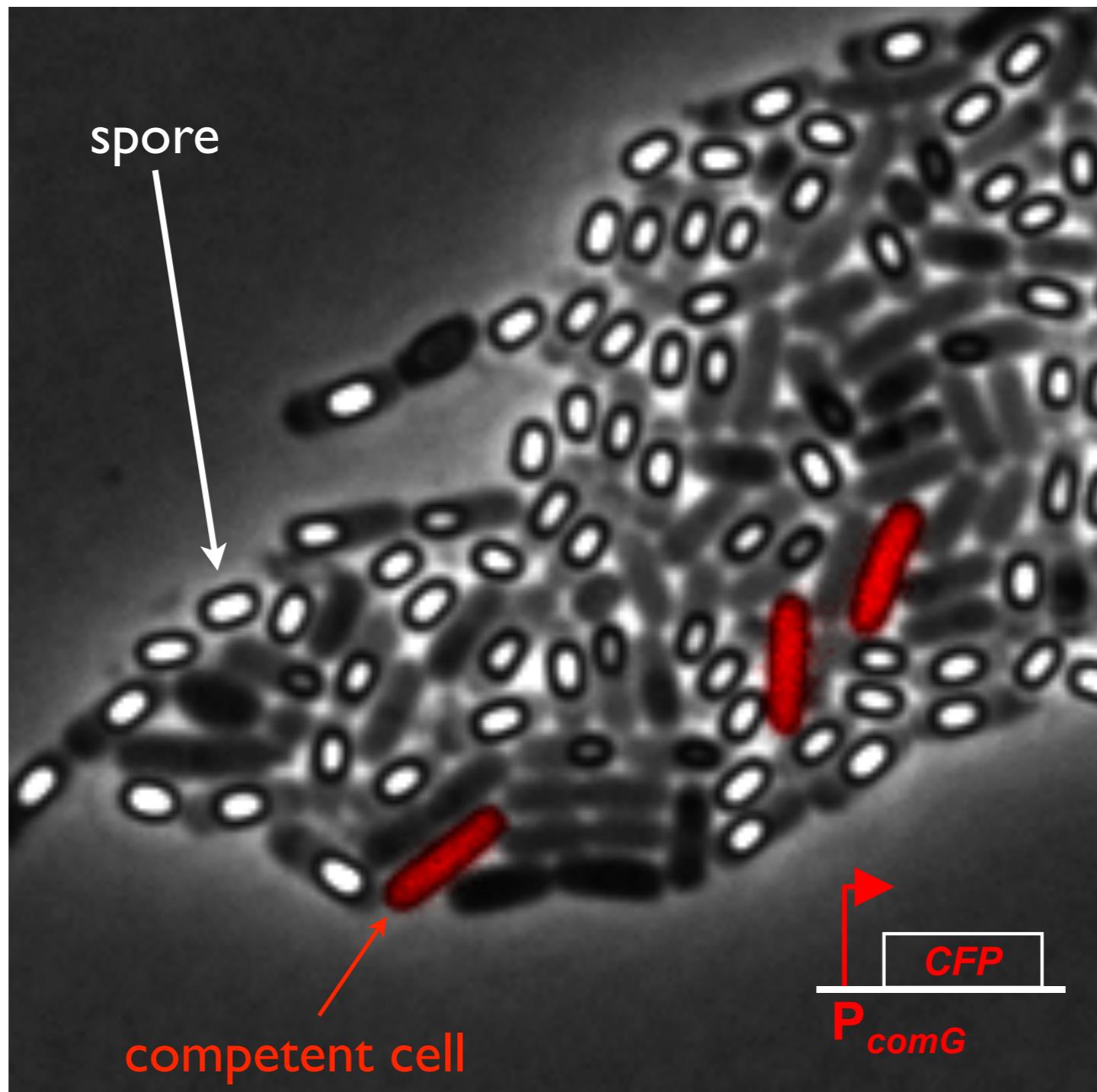
Gene network underlying genetic competence



adapted from
Hamoen et al,
Microbiology 2003

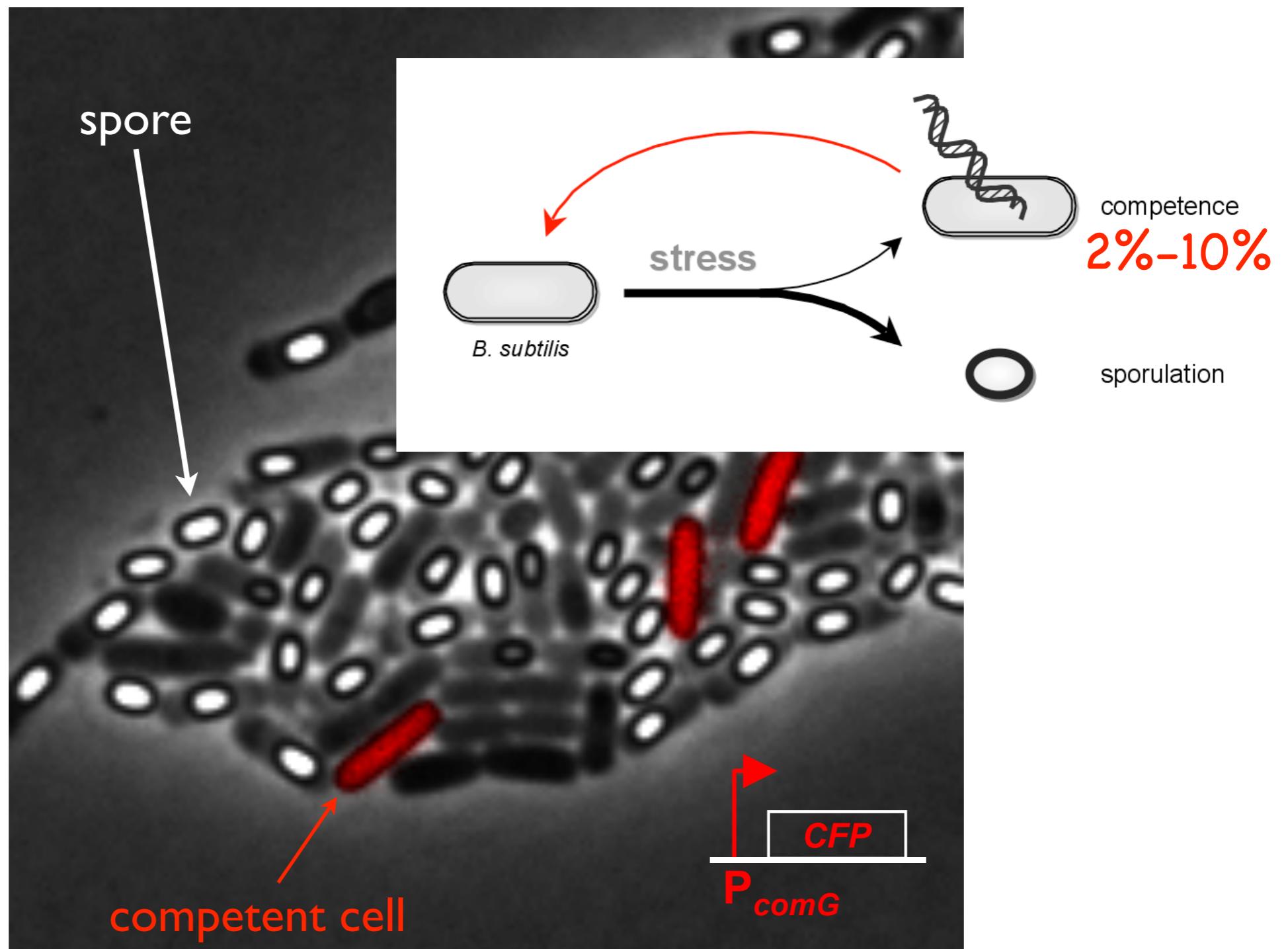
Stress response of bacteria

B. subtilis under low nutrient conditions



Stress response of bacteria

B. subtilis under low nutrient conditions

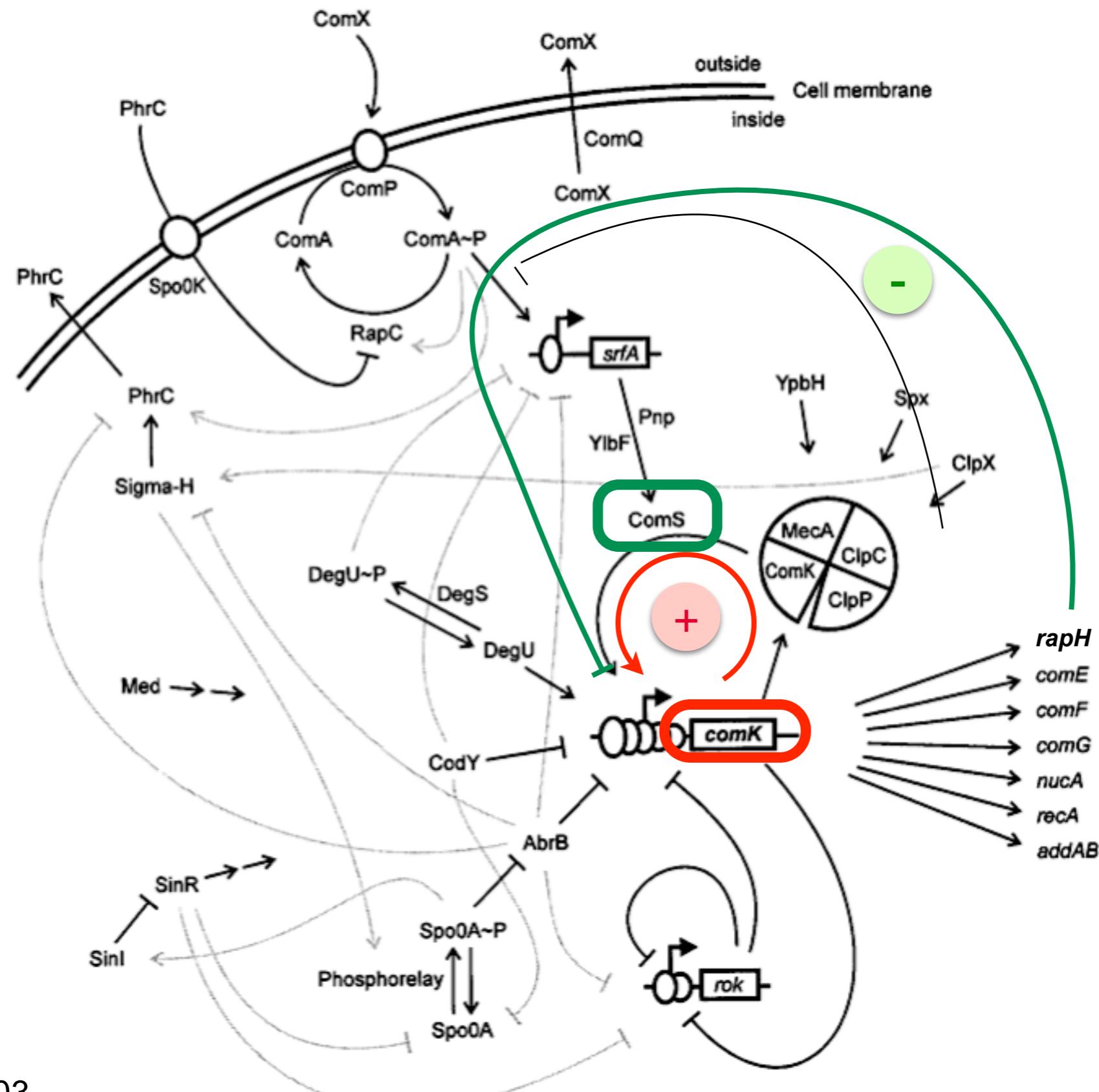




◀ cfp
 P_{comG}

◀ yfp
 P_{comS}

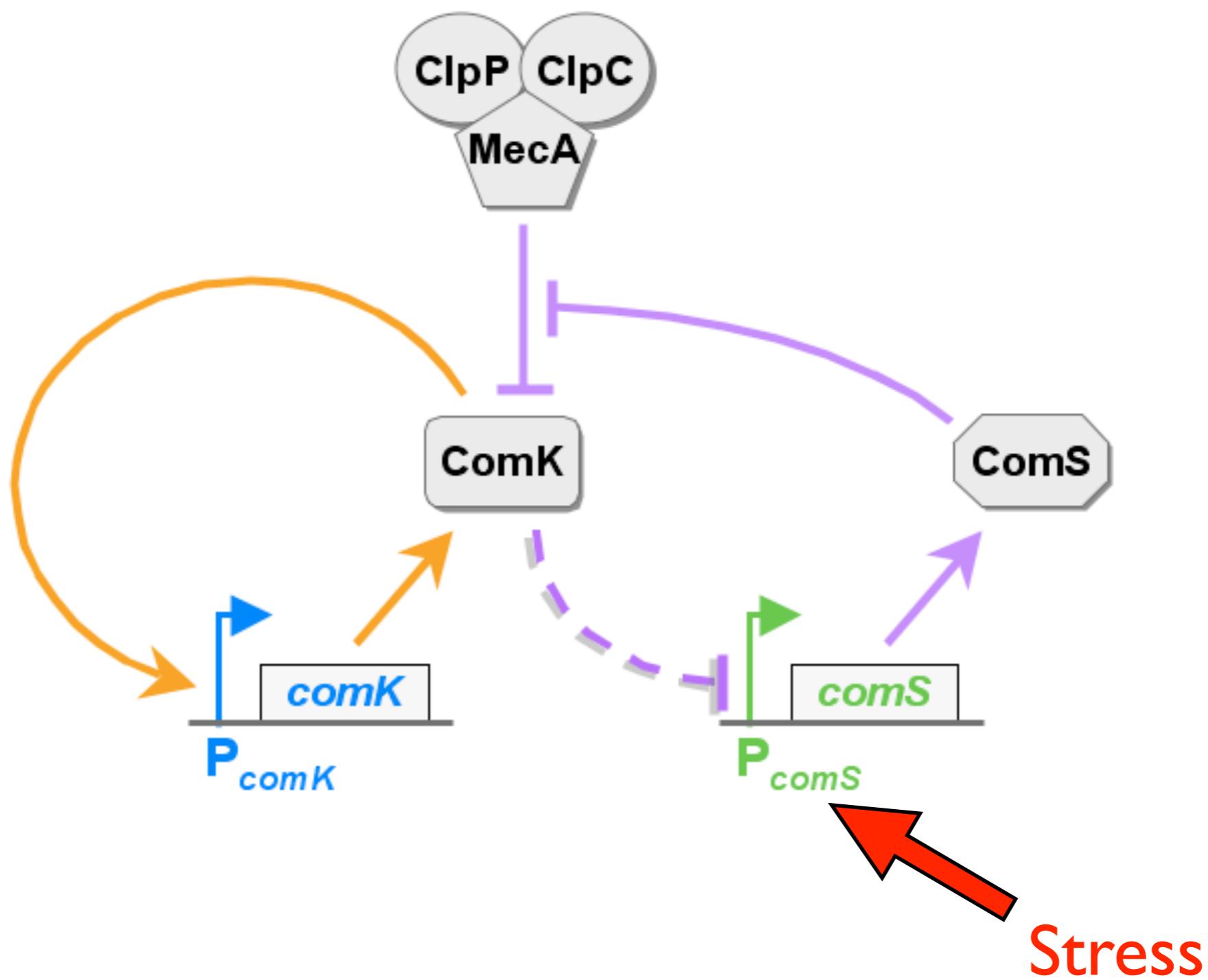
Molecular network underlying genetic competence

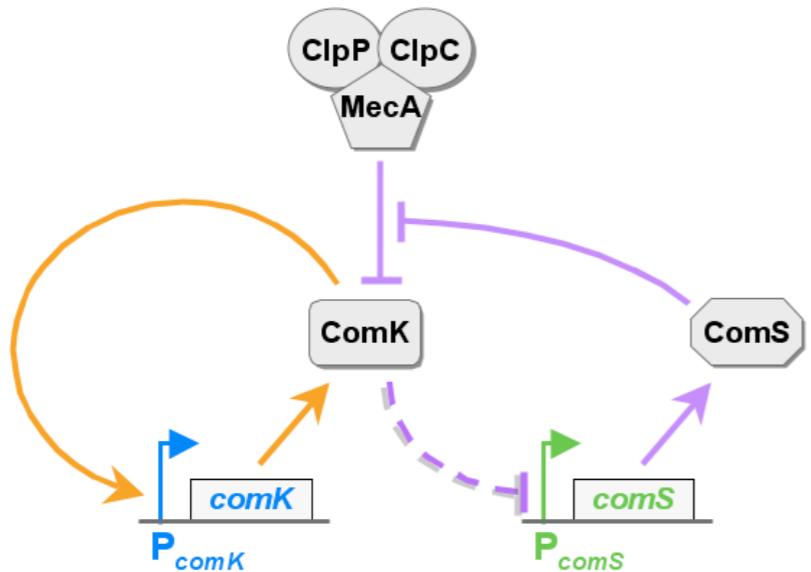


adapted from
Hamoen et al,
Microbiology 2003

stress
sensor
master
regulator

A genetic circuit for genetic competence





Competitive degradation:



$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \gamma_a M_f K + \gamma_{-a} M_K - \delta_k K$$

$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \gamma_b M_f S + \gamma_{-b} M_S - \delta_s S$$

$$\frac{dM_K}{dt} = -(\gamma_{-a} + \gamma_1) M_K + \gamma_a M_f K$$

$$\frac{dM_S}{dt} = -(\gamma_{-b} + \gamma_2) M_S + \gamma_b M_f S$$

$$M_f + M_K + M_S = M_{\text{total}} = \text{constant}$$

- Adiabatic elimination of M_k and M_s :

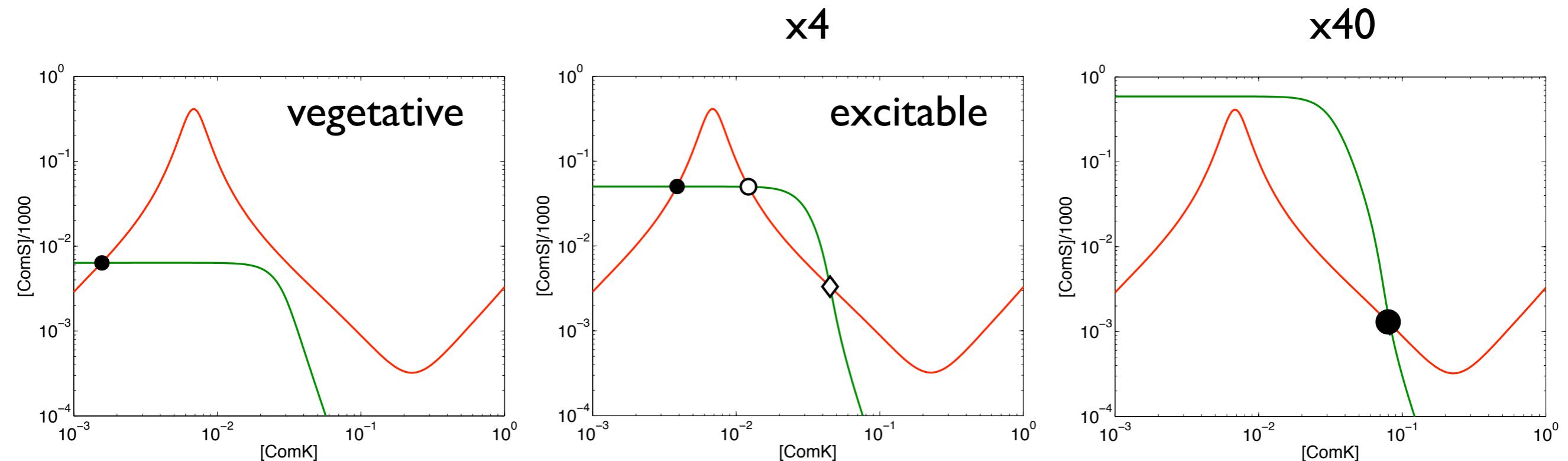
$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K$$

$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

stress

- Nulcline plots for increasing stress (β_s):

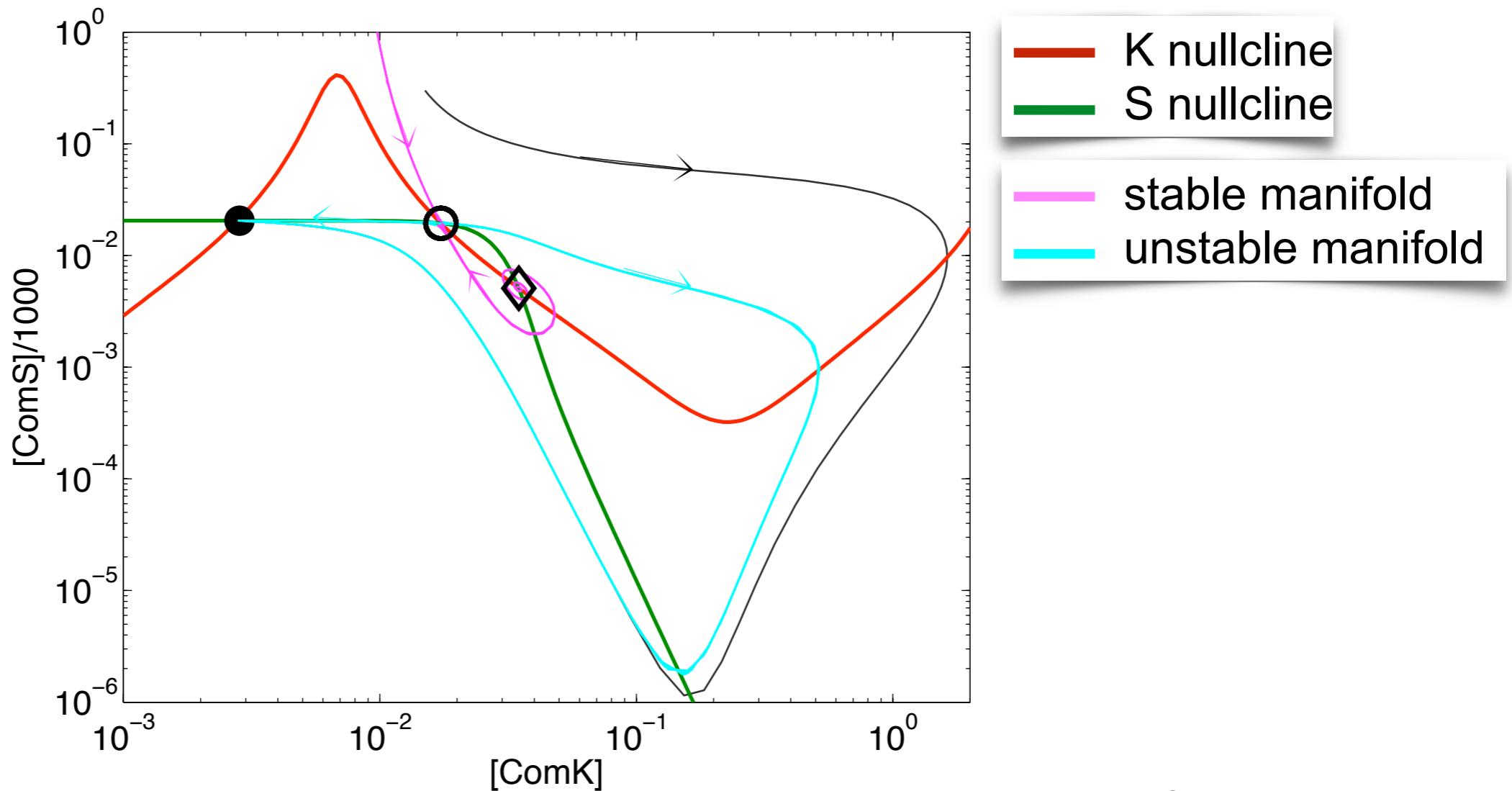
β_s



$$\begin{aligned} \frac{dK}{dt} &= \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K \\ \frac{dS}{dt} &= \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S \end{aligned}$$

- stable fixed point
- saddle
- ◇ unstable node/focus

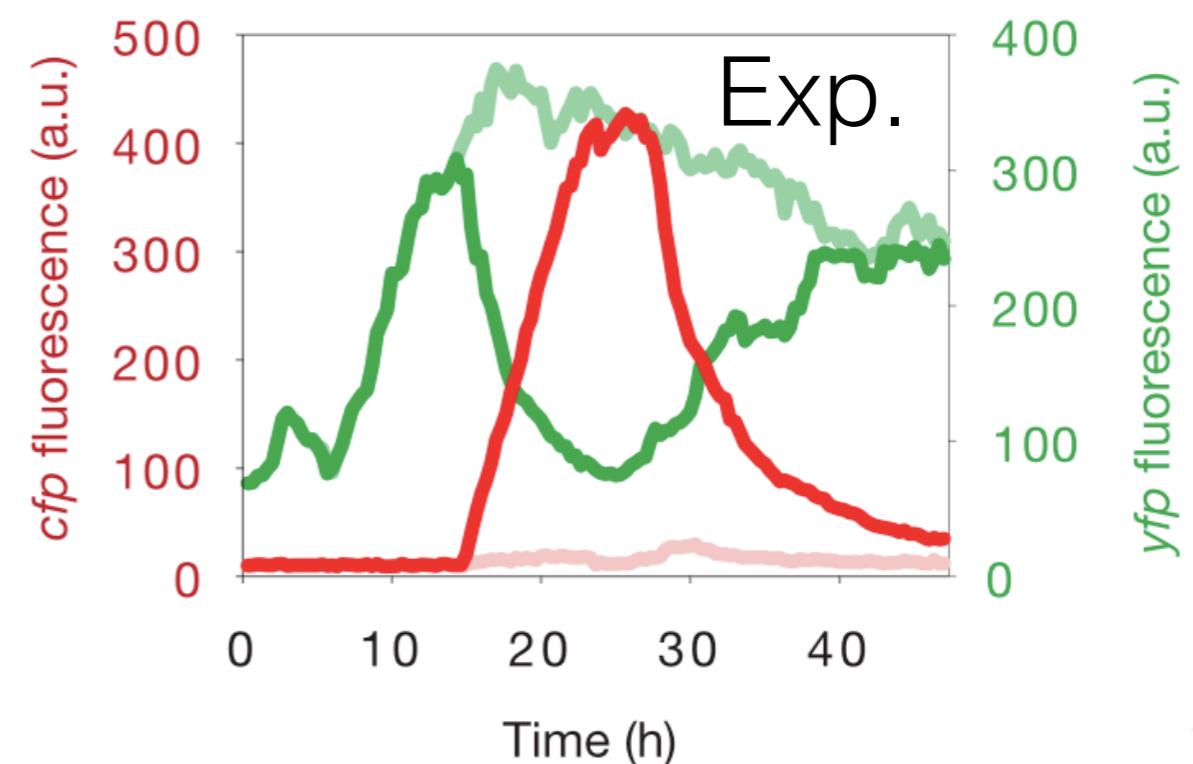
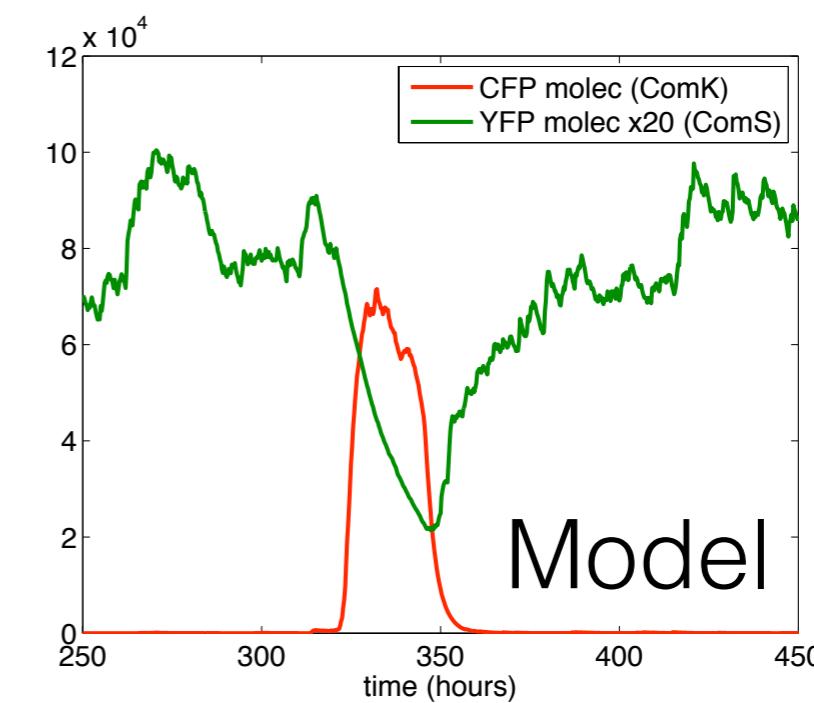
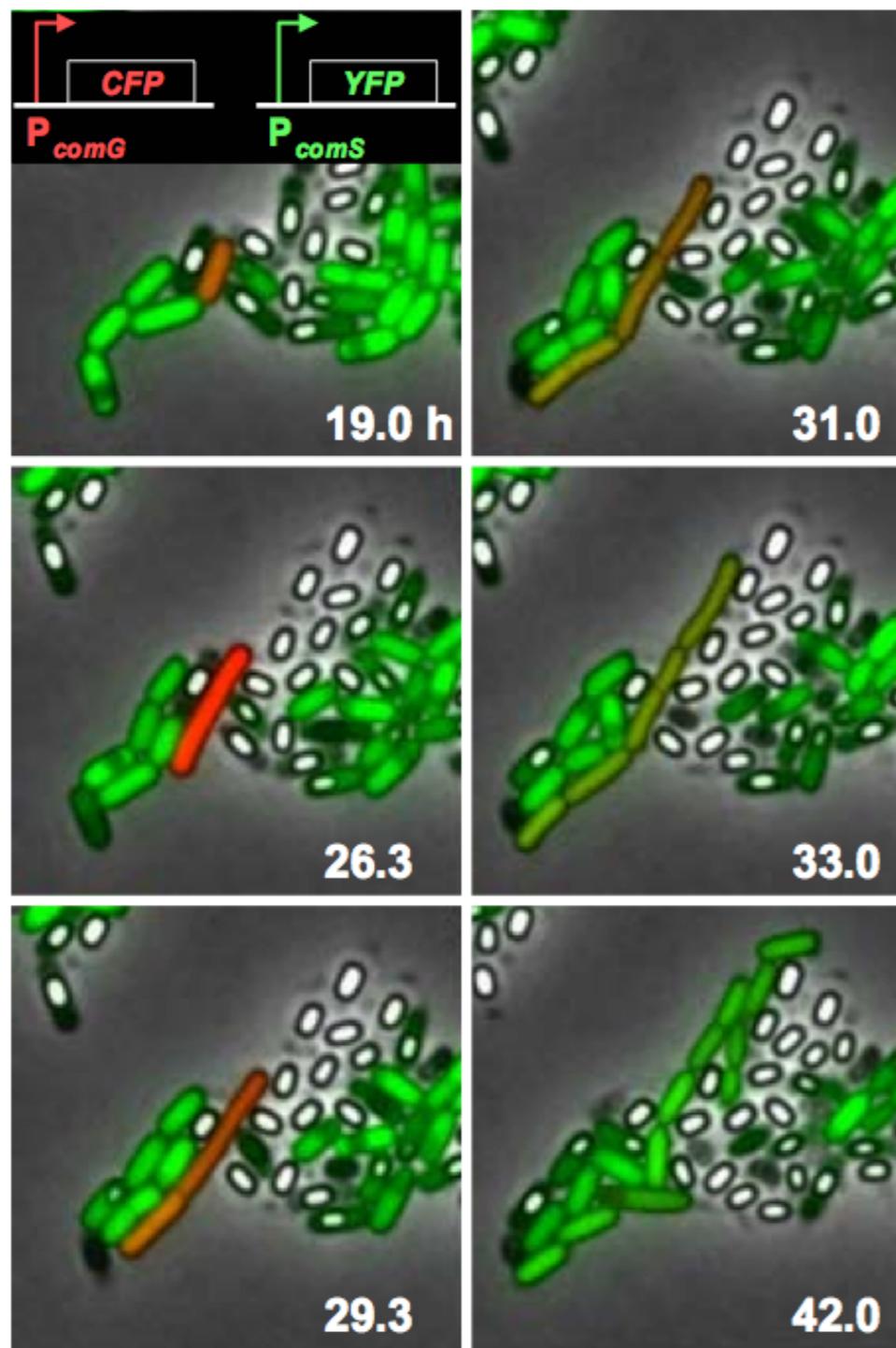
Excitable dynamics of competence



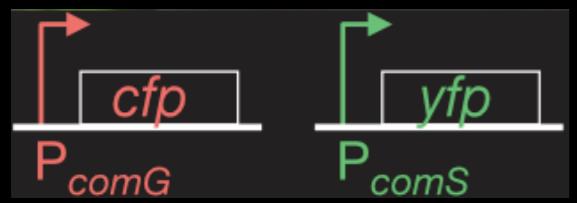
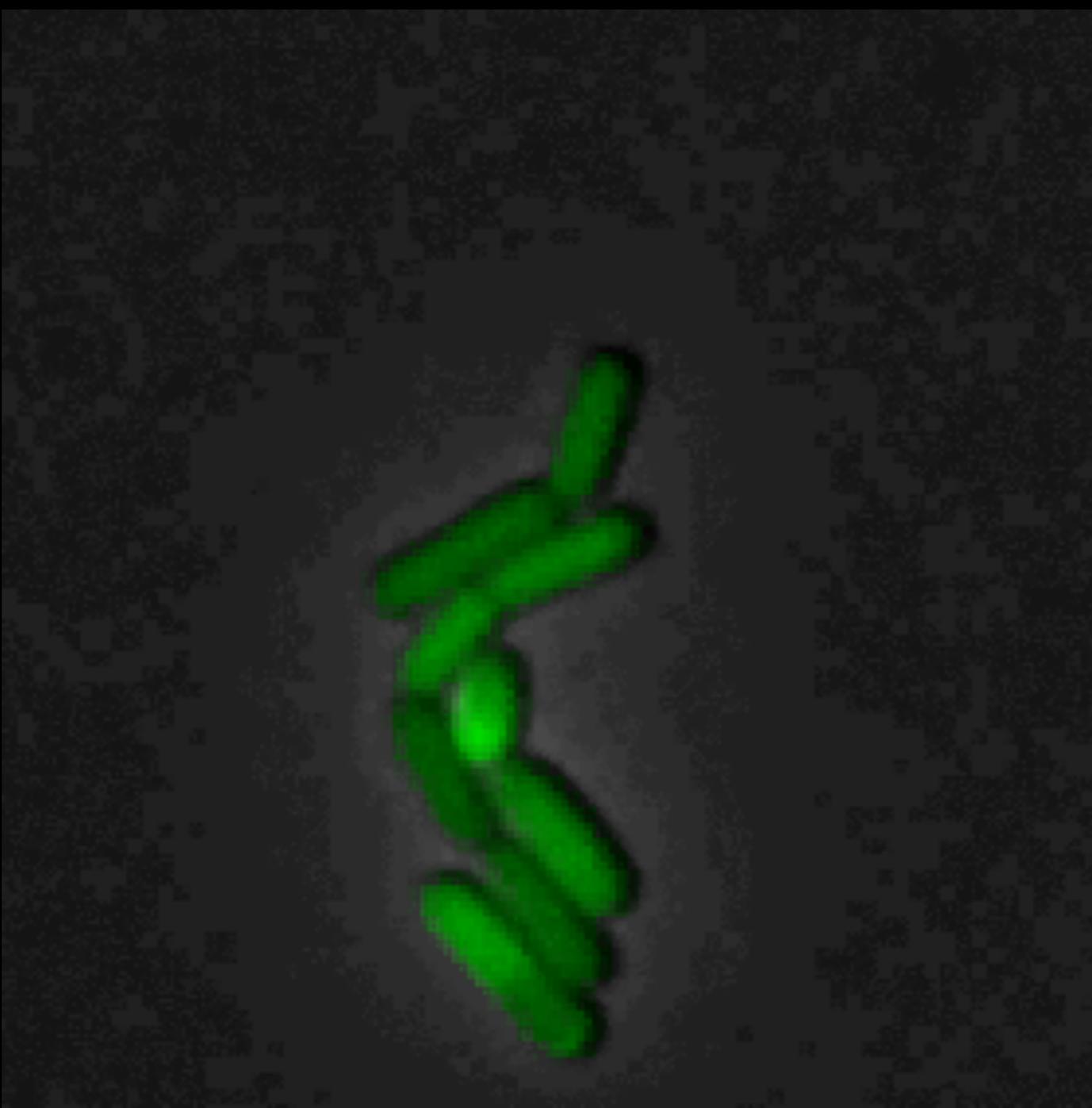
$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K$$

$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

Testing the model: ComK-ComS anticorrelation



Süel, JGO, Lieberman & Elowitz, Nature (2006)



Bifurcation analysis of the competence system

$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K$$
$$\frac{dS}{dt} = \alpha_s - \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

Two experimentally controllable parameters (by means of an **IPTG-inducible promoter**):

α_k : basal production of **ComK** protein

α_s : basal production of **ComS** protein

ADVANCED SUMMER SCHOOL ON APPLIED DYNAMICS IN SYSTEMS AND SYNTHETIC BIOLOGY

EXERCISES - GENE REGULATORY CIRCUITS

4. The following model describes genetic competence in *B. subtilis*:

$$\begin{aligned}\frac{dK}{dt} &= \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K \\ \frac{dS}{dt} &= \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S\end{aligned}$$

Again, use the tools that you have learned in the different courses of this school to analyze this dynamical system in the plane, as the parameters α_k , α_s and β_s vary (one at a time or in pairs). Reproduce and study the response of the system in the different dynamical regimes studied in class. Consider the following baseline parameter values:

Parameter	Value	Parameter	Value	Parameter	Value
α_k	0.00875 molec/s	γ_k , γ_s	0.001 s^{-1}	k_k	5000 molec
α_s	0.0004 molec/s	δ_k , δ_s	10^{-4} s^{-1}	k_s	833 molec
β_k	7.5 molec/s	Γ_k	25000 molec	n	2
β_s	0.06 molec/s	Γ_s	20 molec	p	5

Bifurcation analysis of the competence system

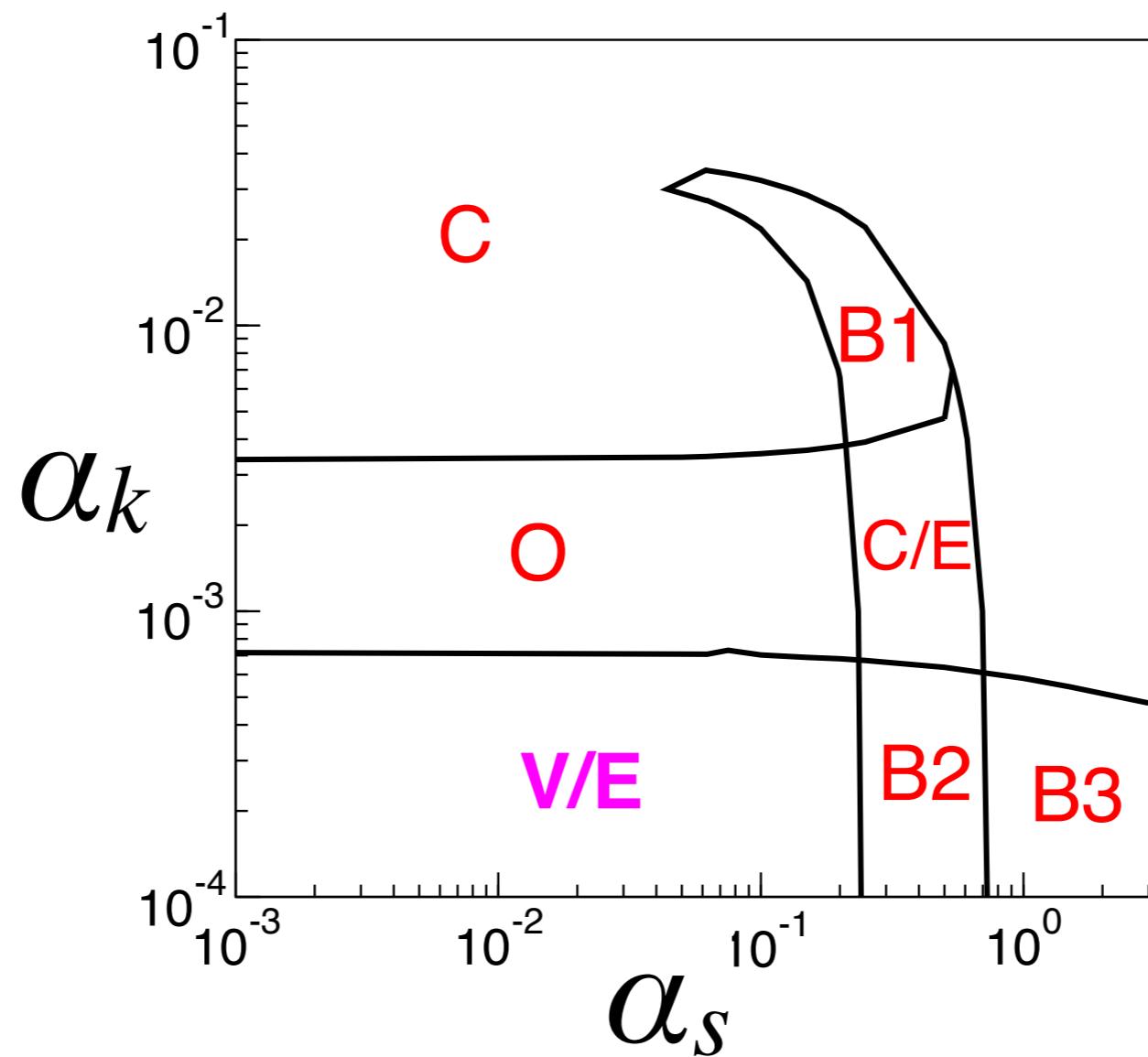
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$$\frac{dS}{dt} = \alpha_s - \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

Two experimentally controllable parameters (by means of an **IPTG-inducible promoter**):

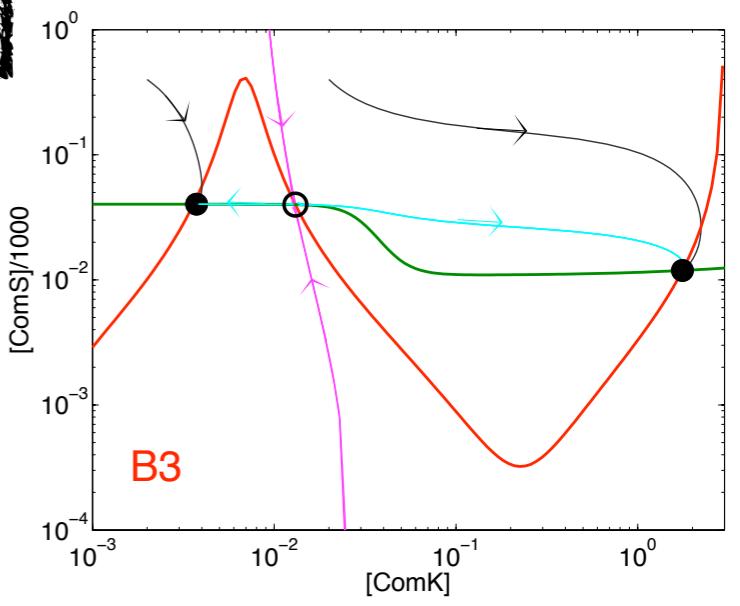
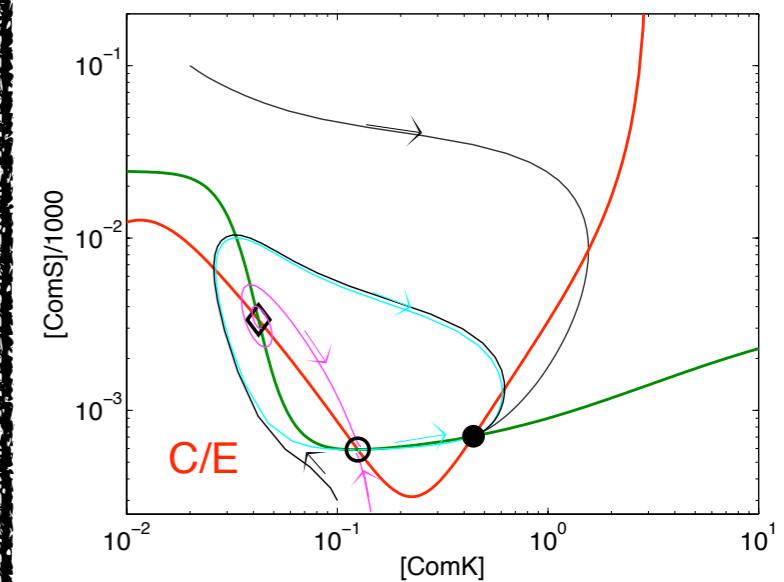
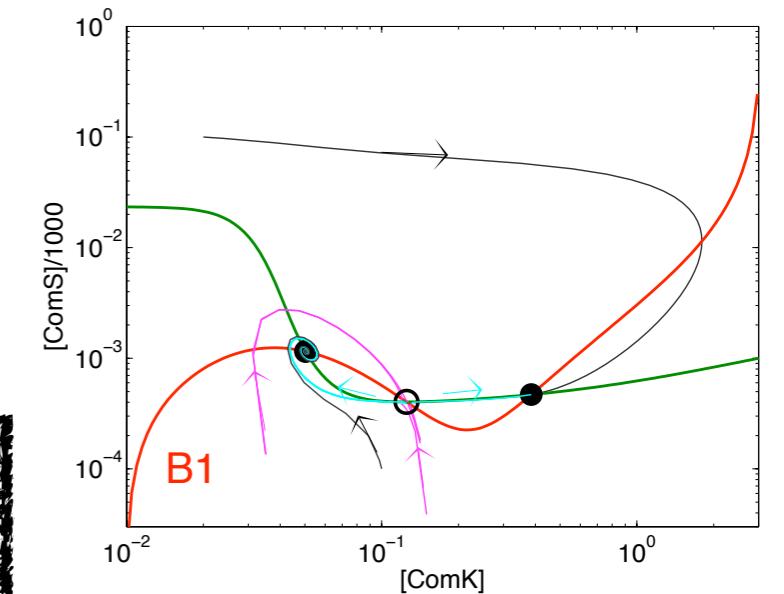
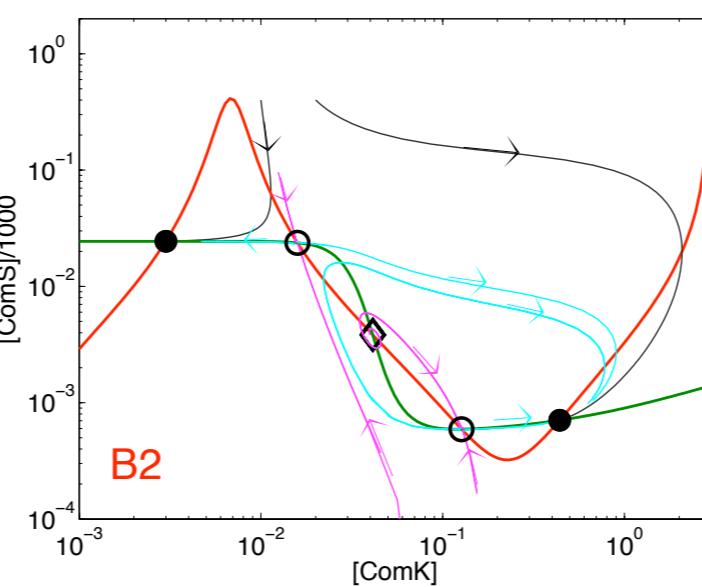
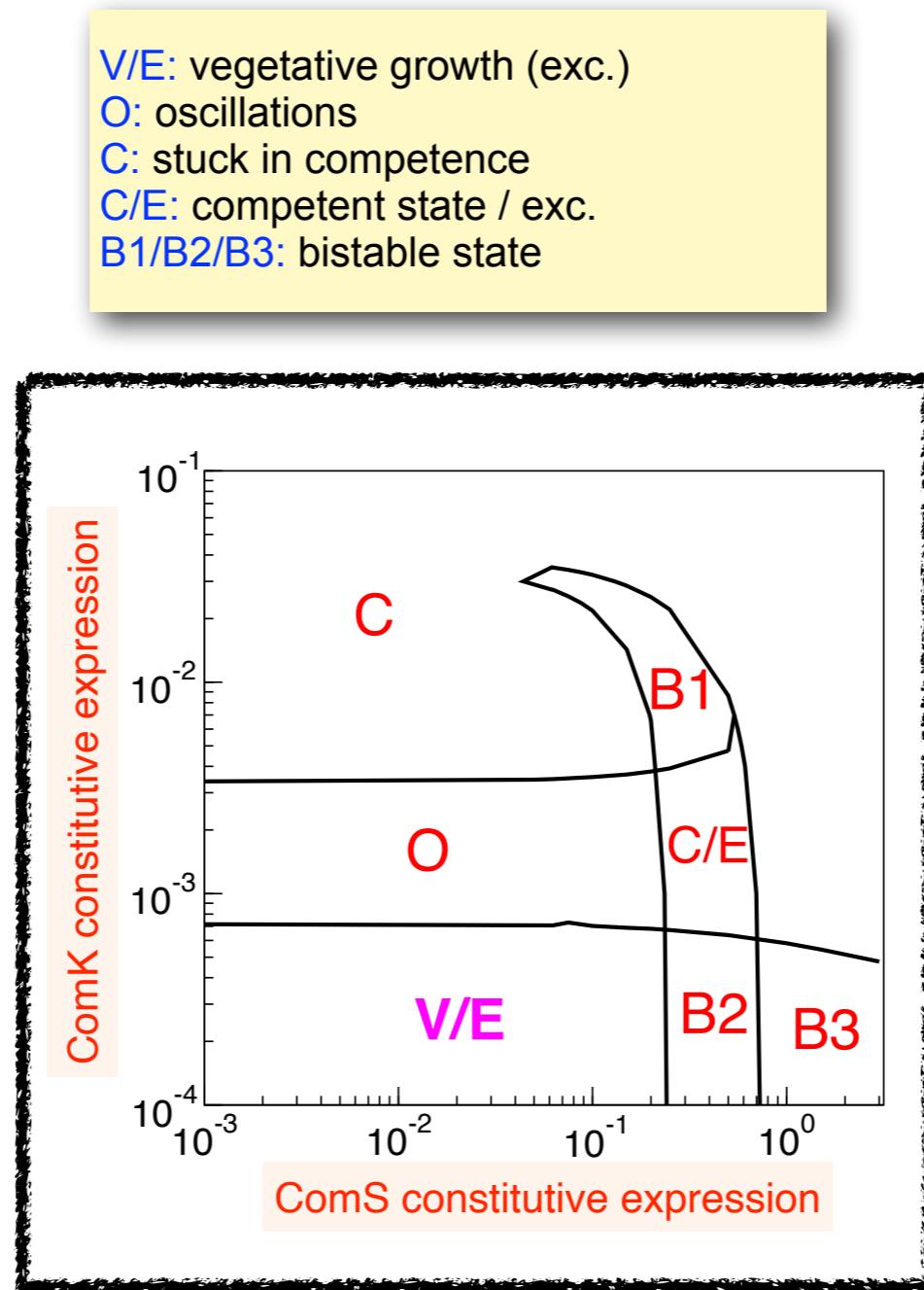
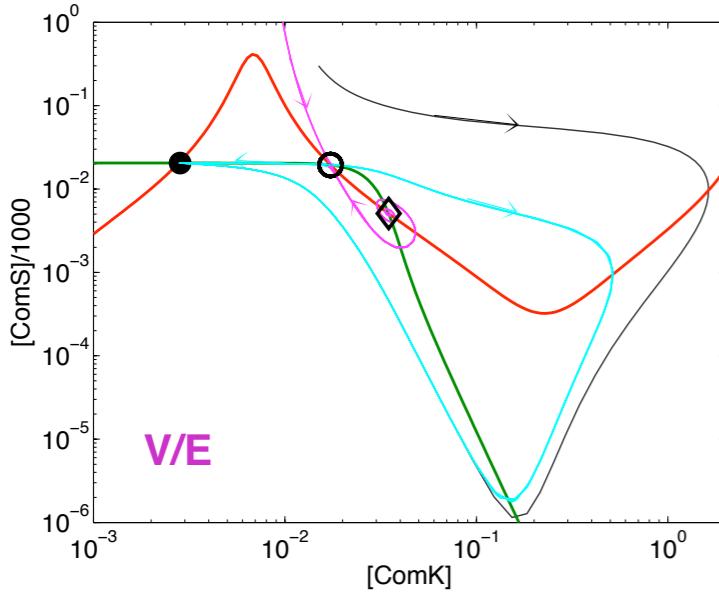
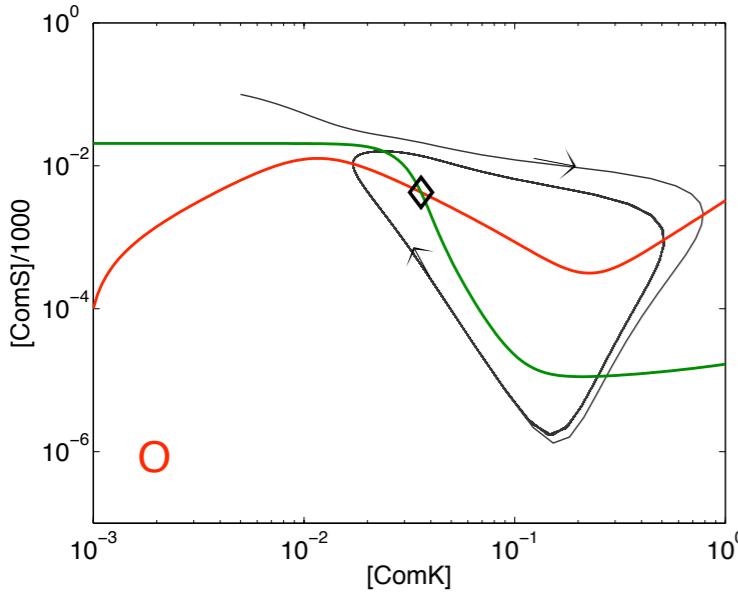
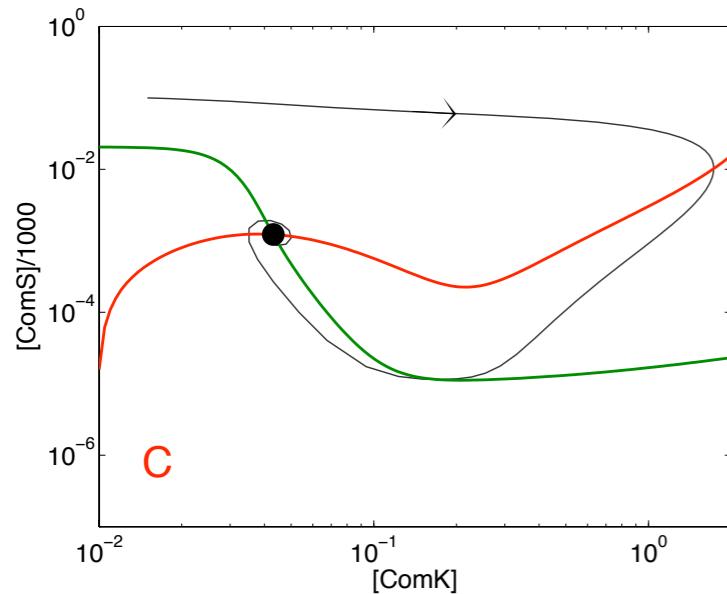
α_k : basal production of **ComK** protein

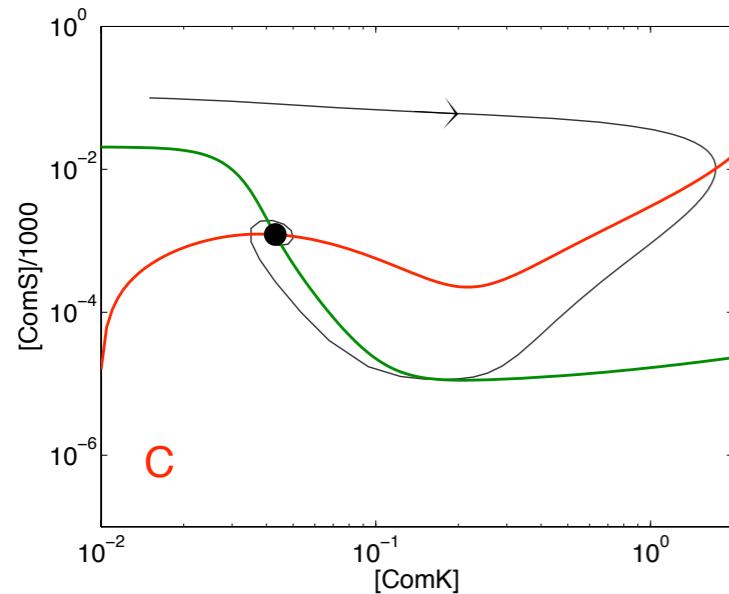
α_s : basal production of **ComS** protein

Bifurcation analysis of the competence system

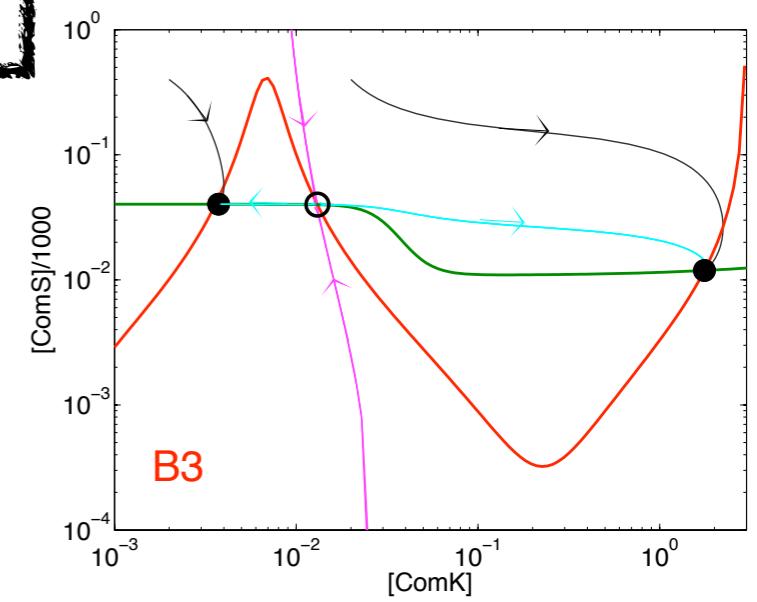
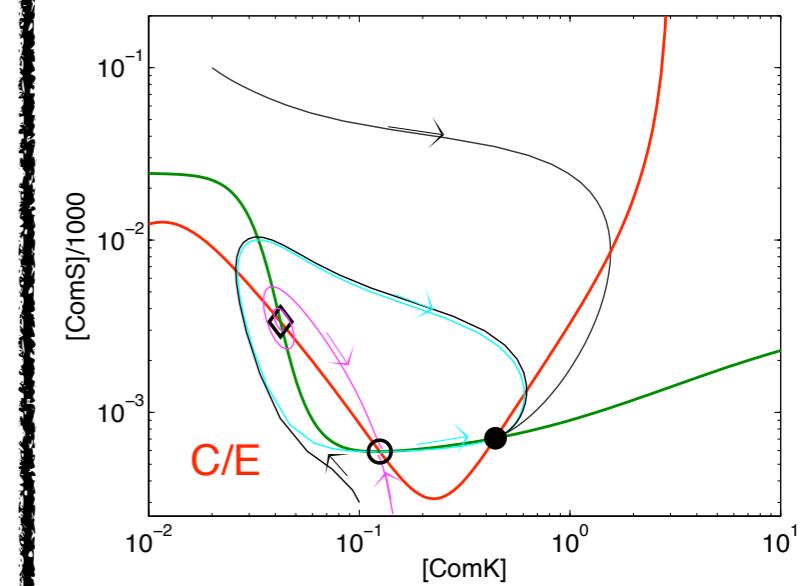
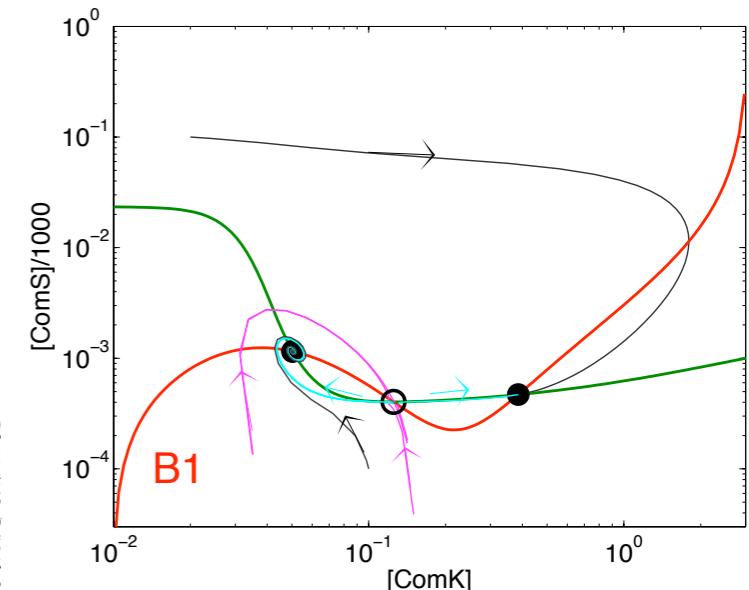
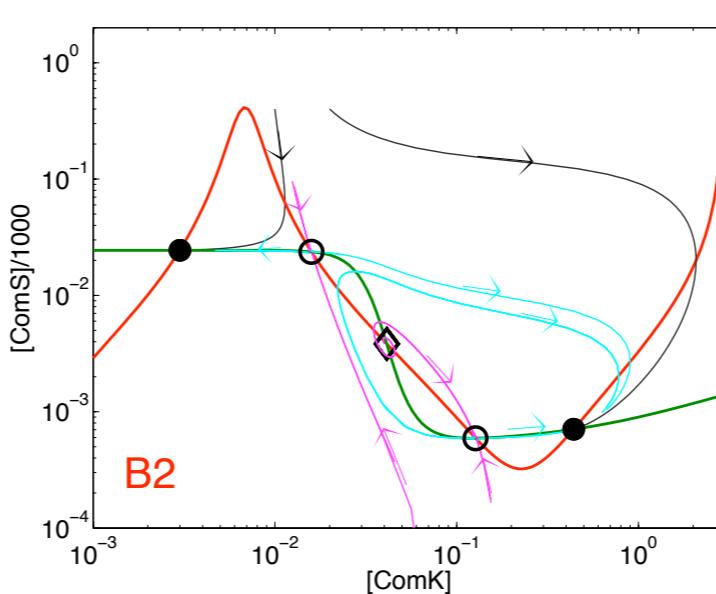
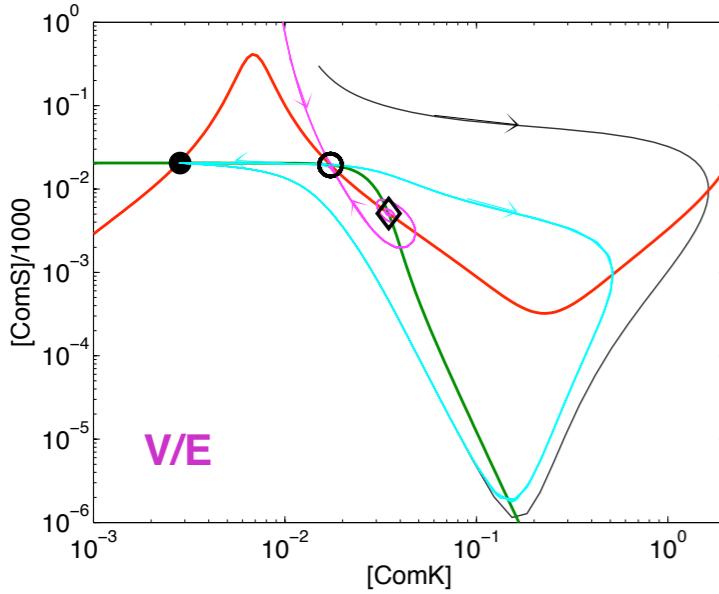
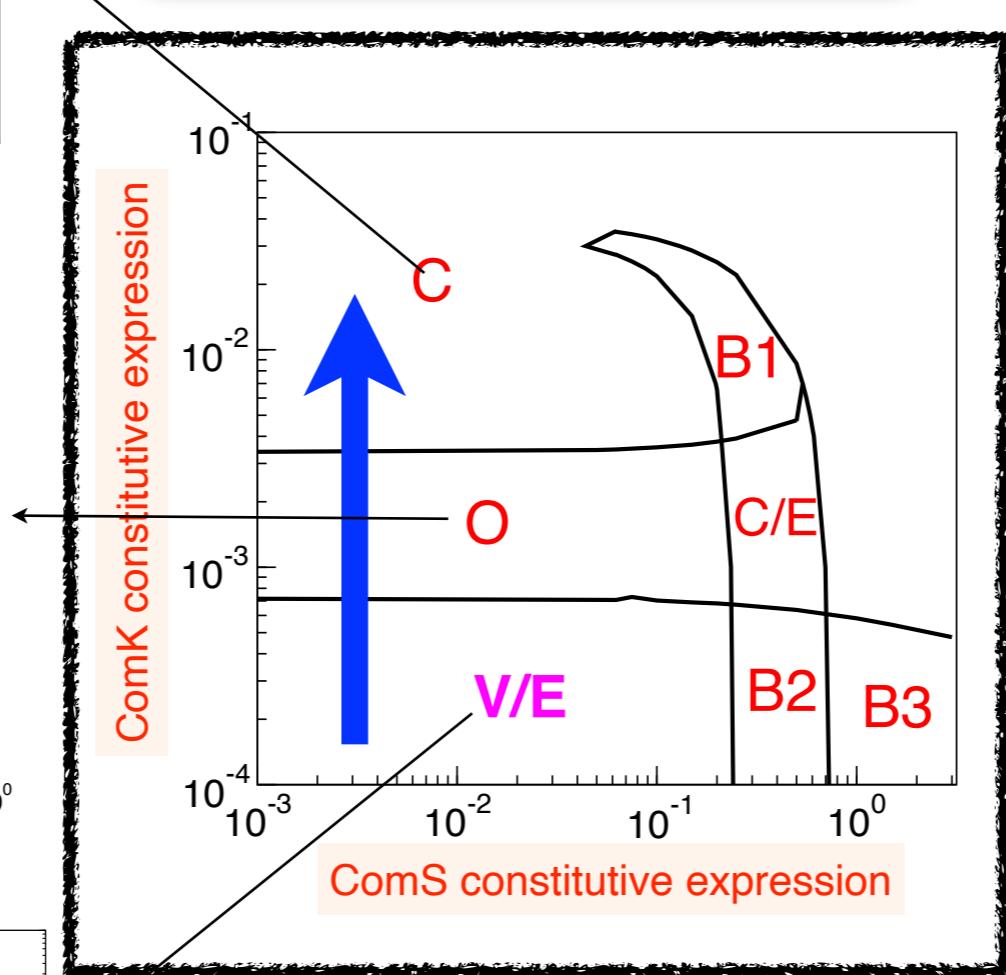
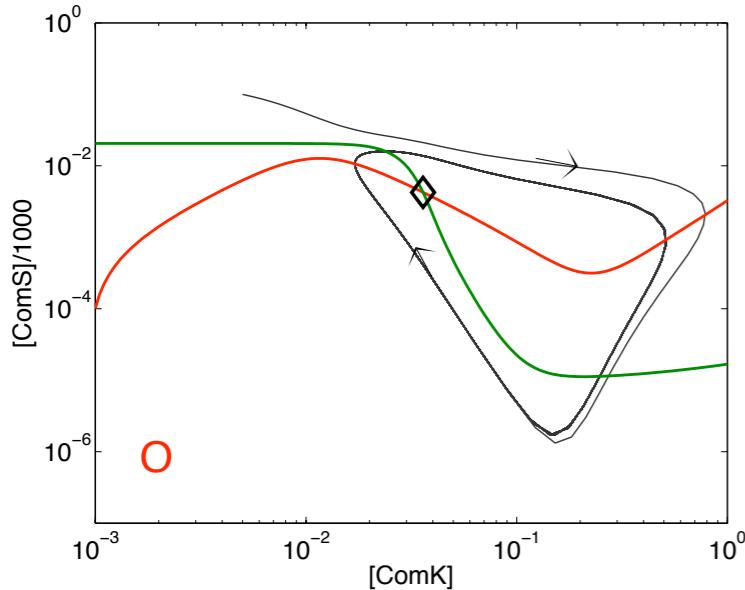


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$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

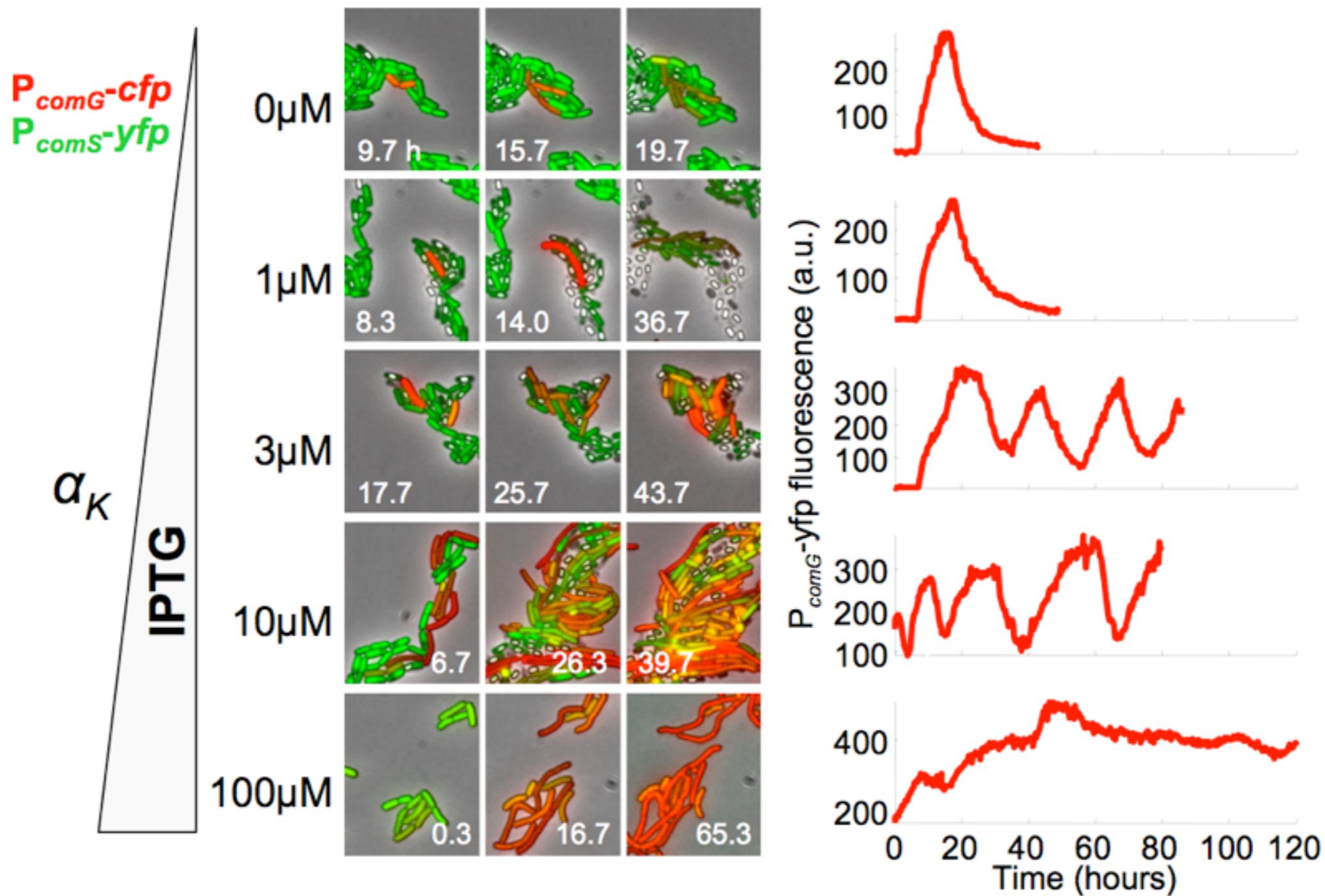


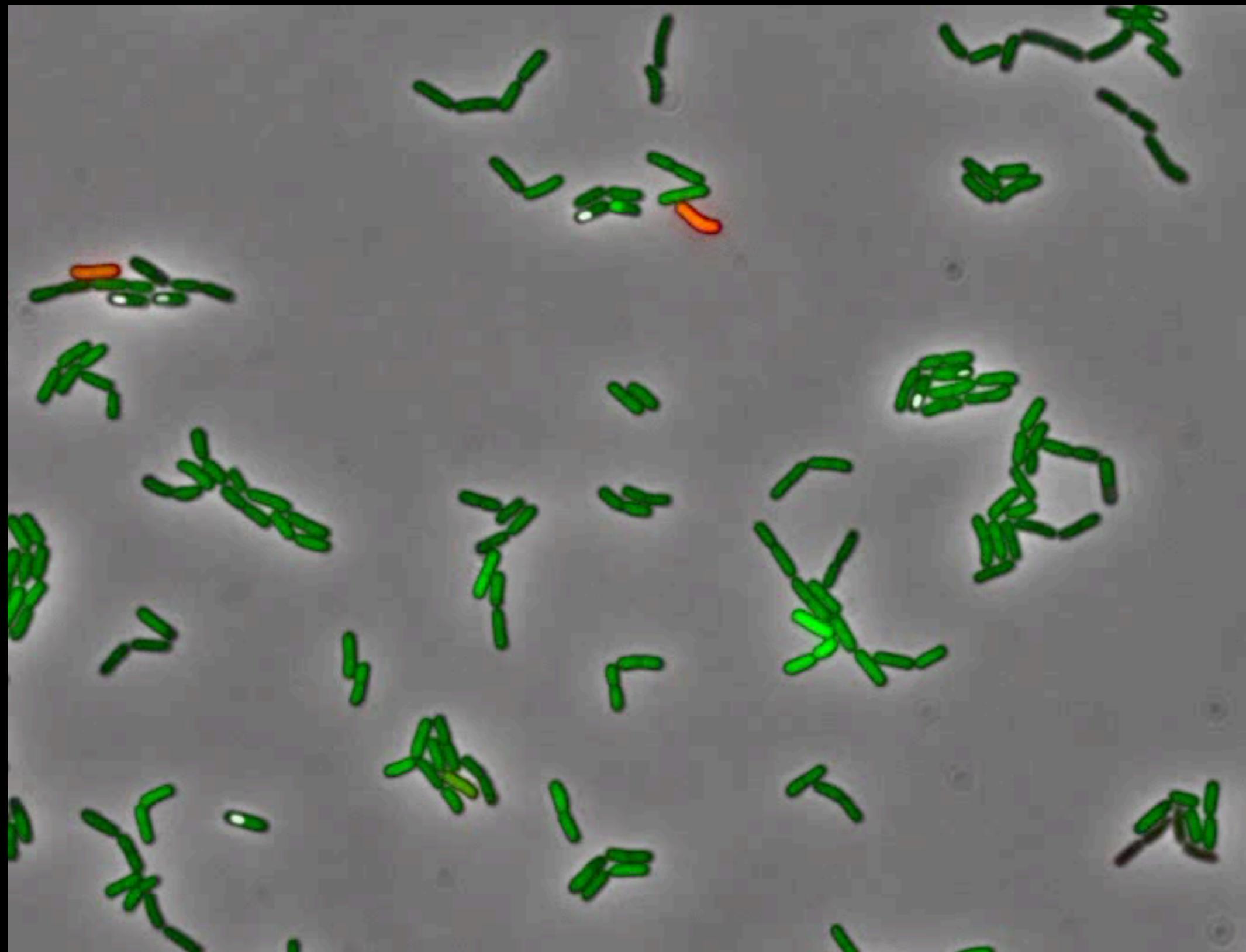


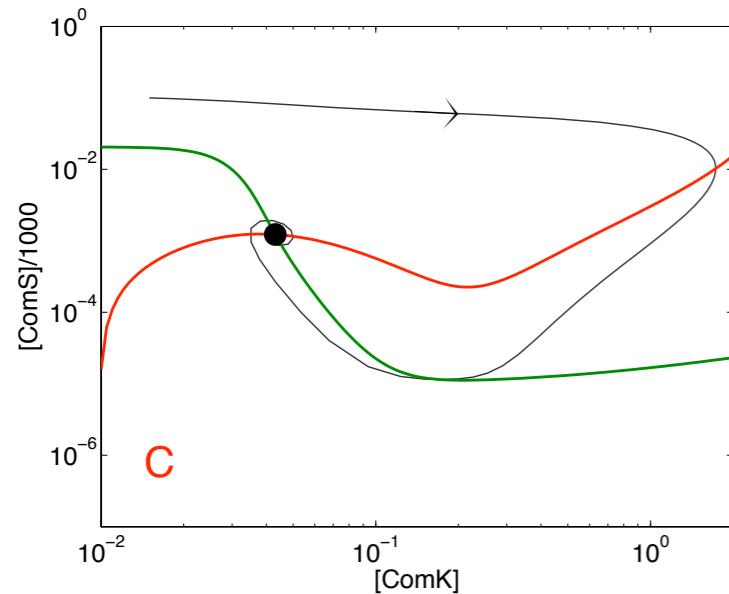
V/E: vegetative growth (exc.)
 O: oscillations
 C: stuck in competence
 C/E: competent state / exc.
 B1/B2/B3: bistable state



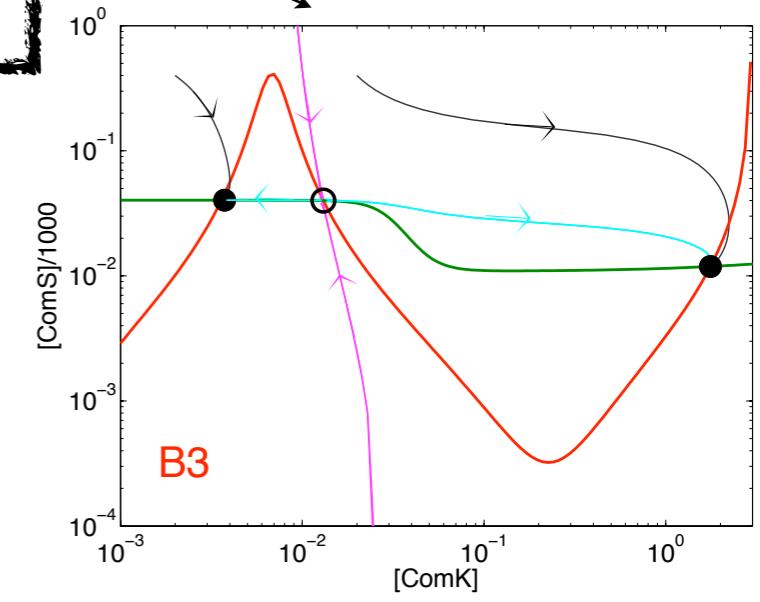
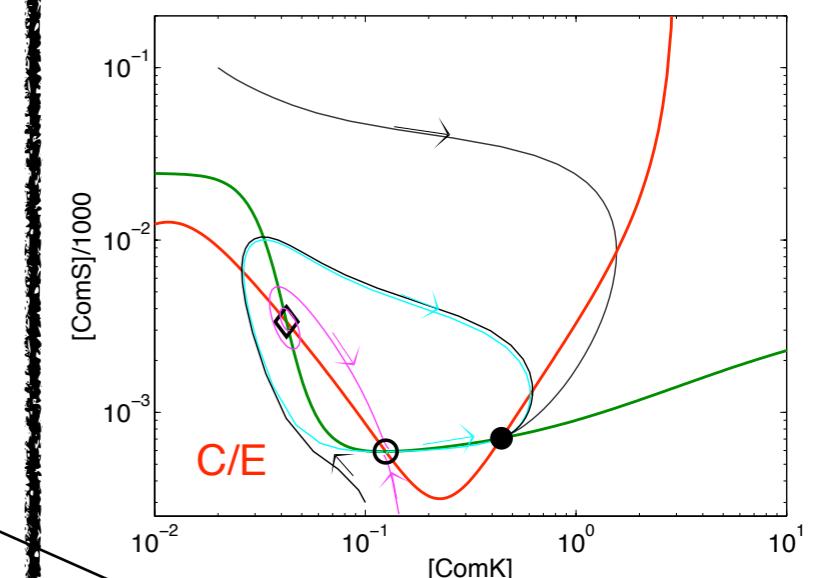
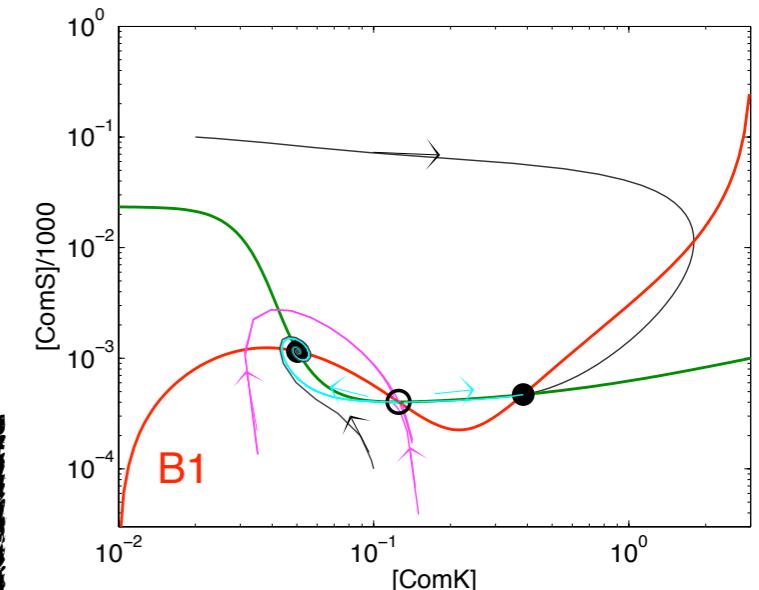
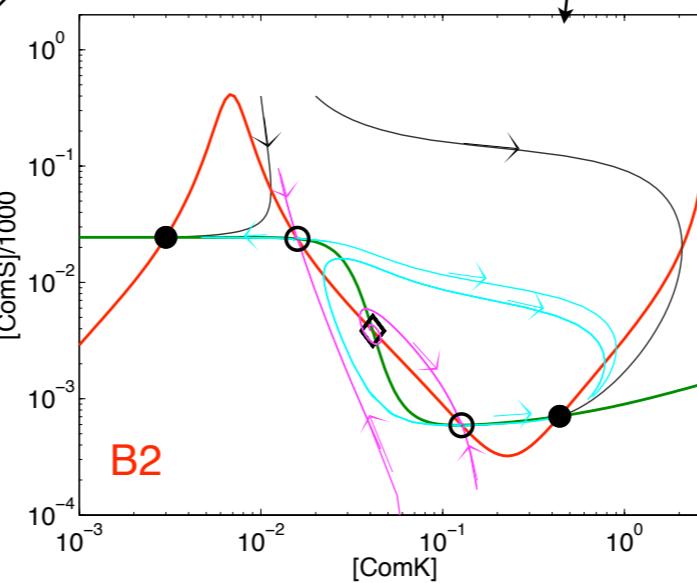
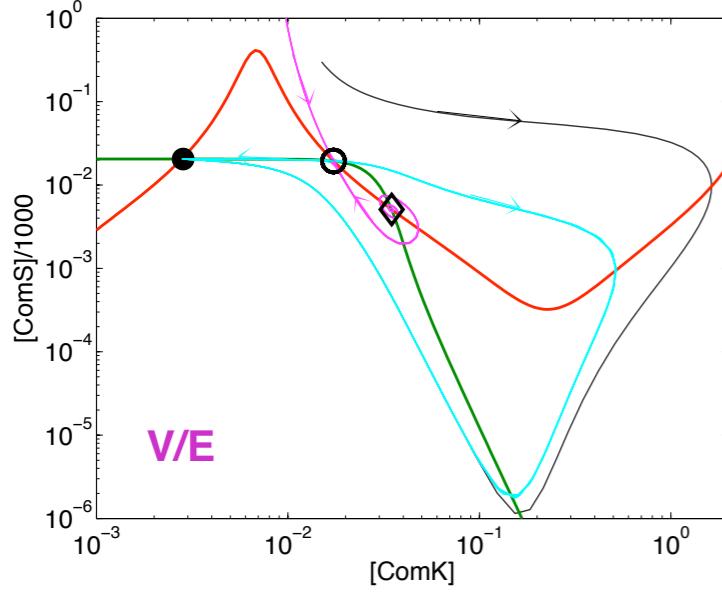
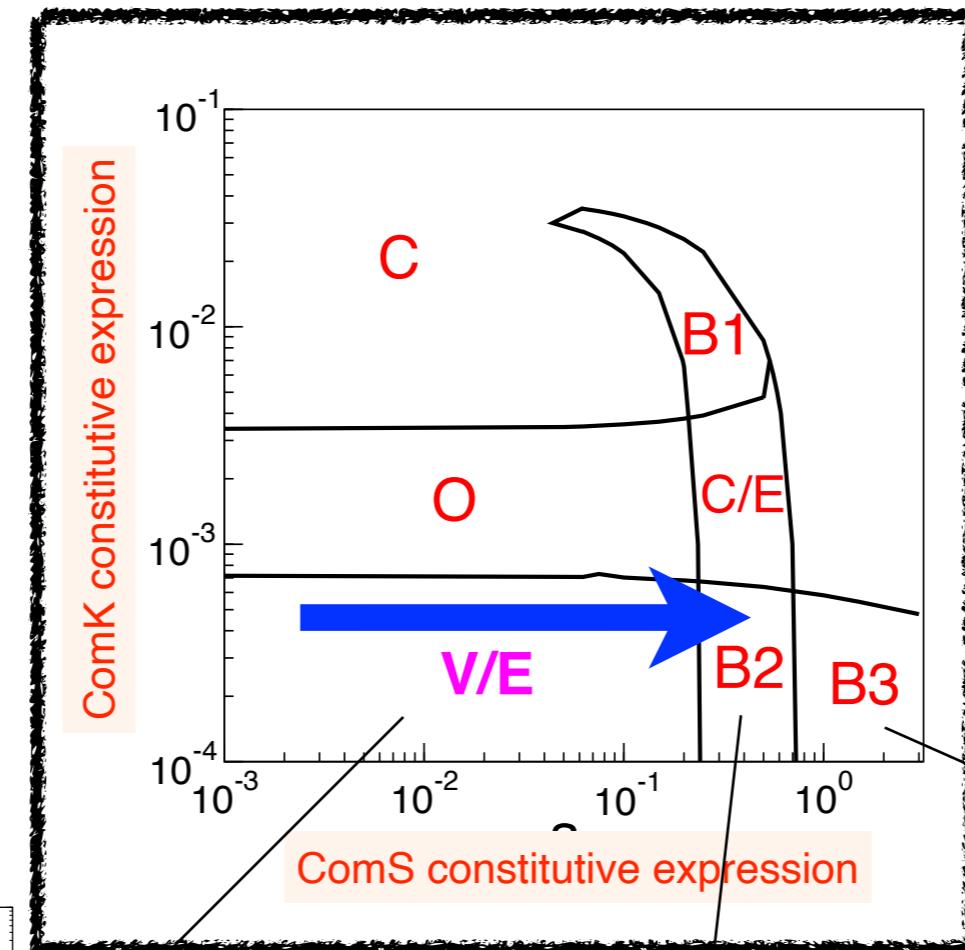
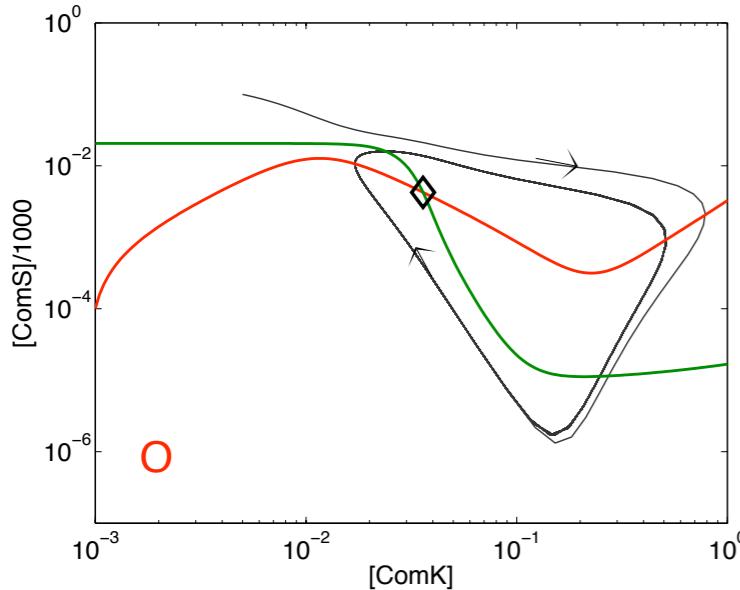
- Adding an additional copy of the *comK* gene under the control of an IPTG-inducible promoter



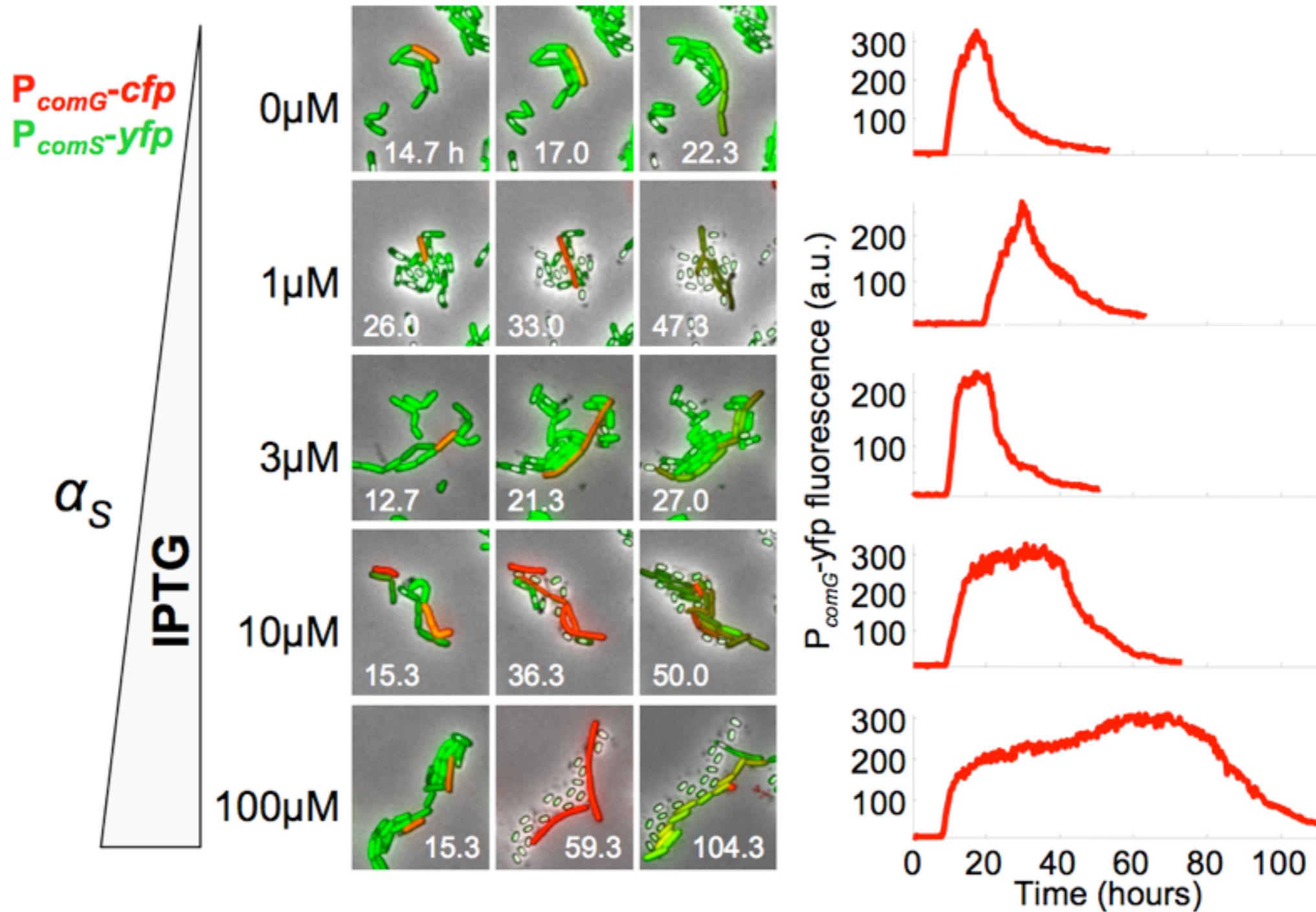




V/E: vegetative growth (exc.)
O: oscillations
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C/E: competent state / exc.
B1/B2/B3: bistable state



- Adding an additional copy of the *comS* gene under the control of an IPTG-inducible promoter





Dynamics of gene regulatory circuits

1. Gene circuit dynamics
2. Dissecting a genetic circuit
3. Noise in genetic circuits

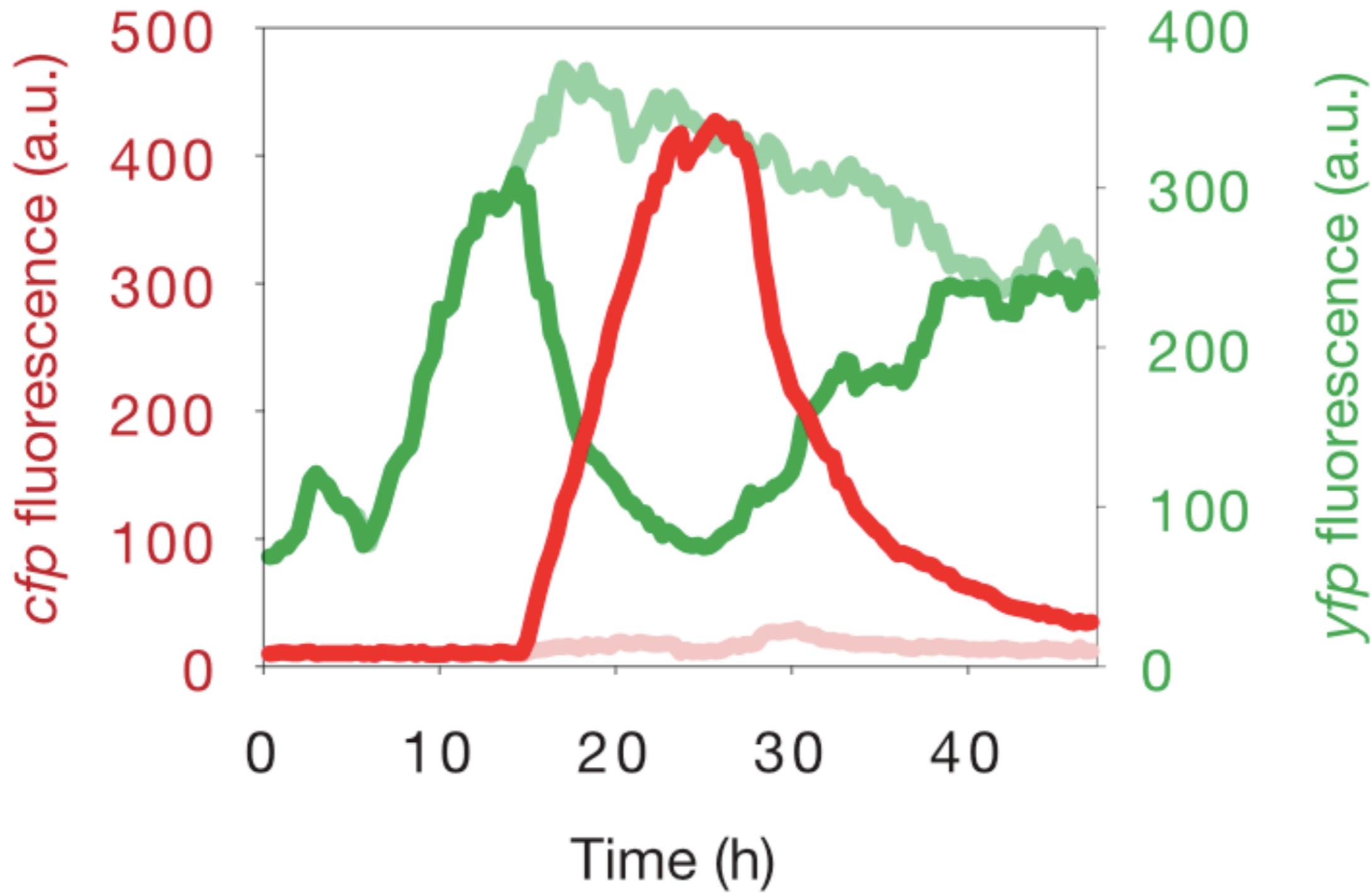


◀ cfp
 P_{comG}

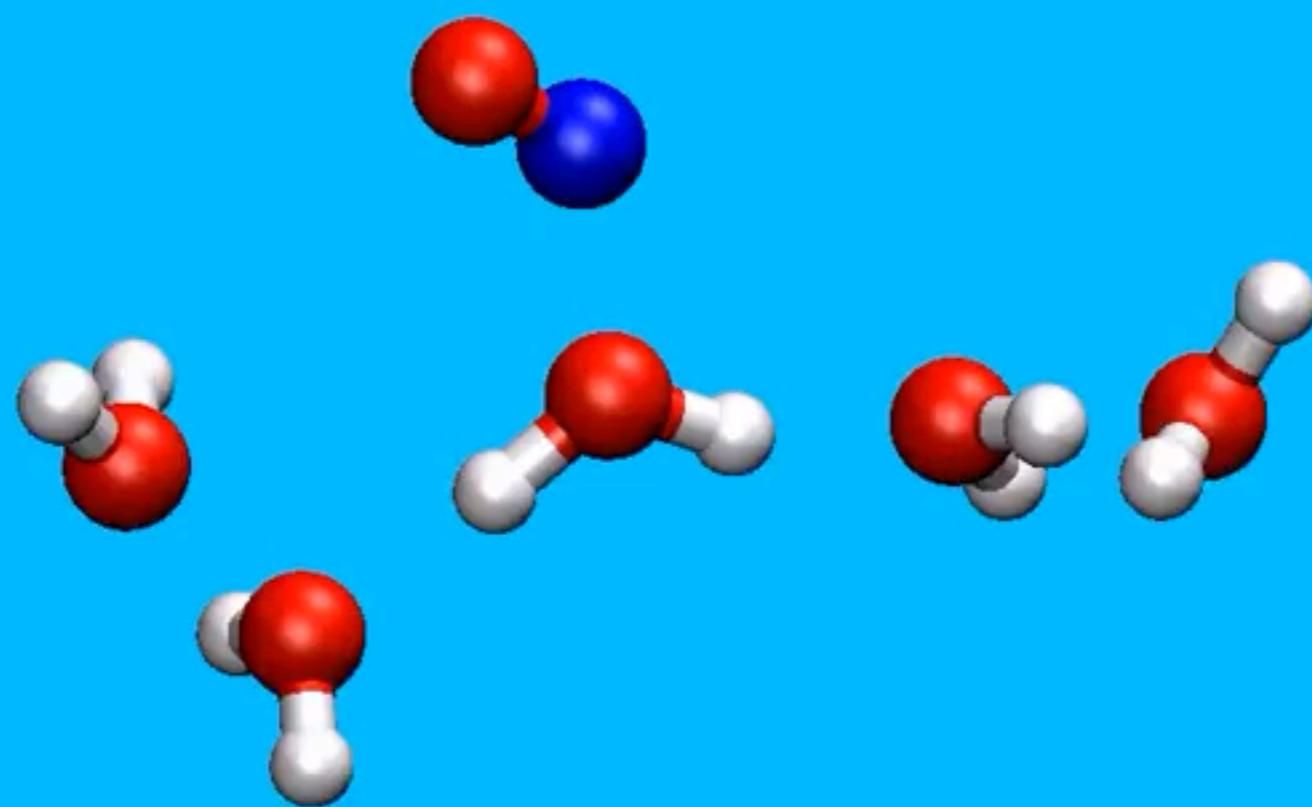
◀ yfp
 P_{comS}

An example of stochastic decision making

- Dynamics of two sister cells:



198.00 ps



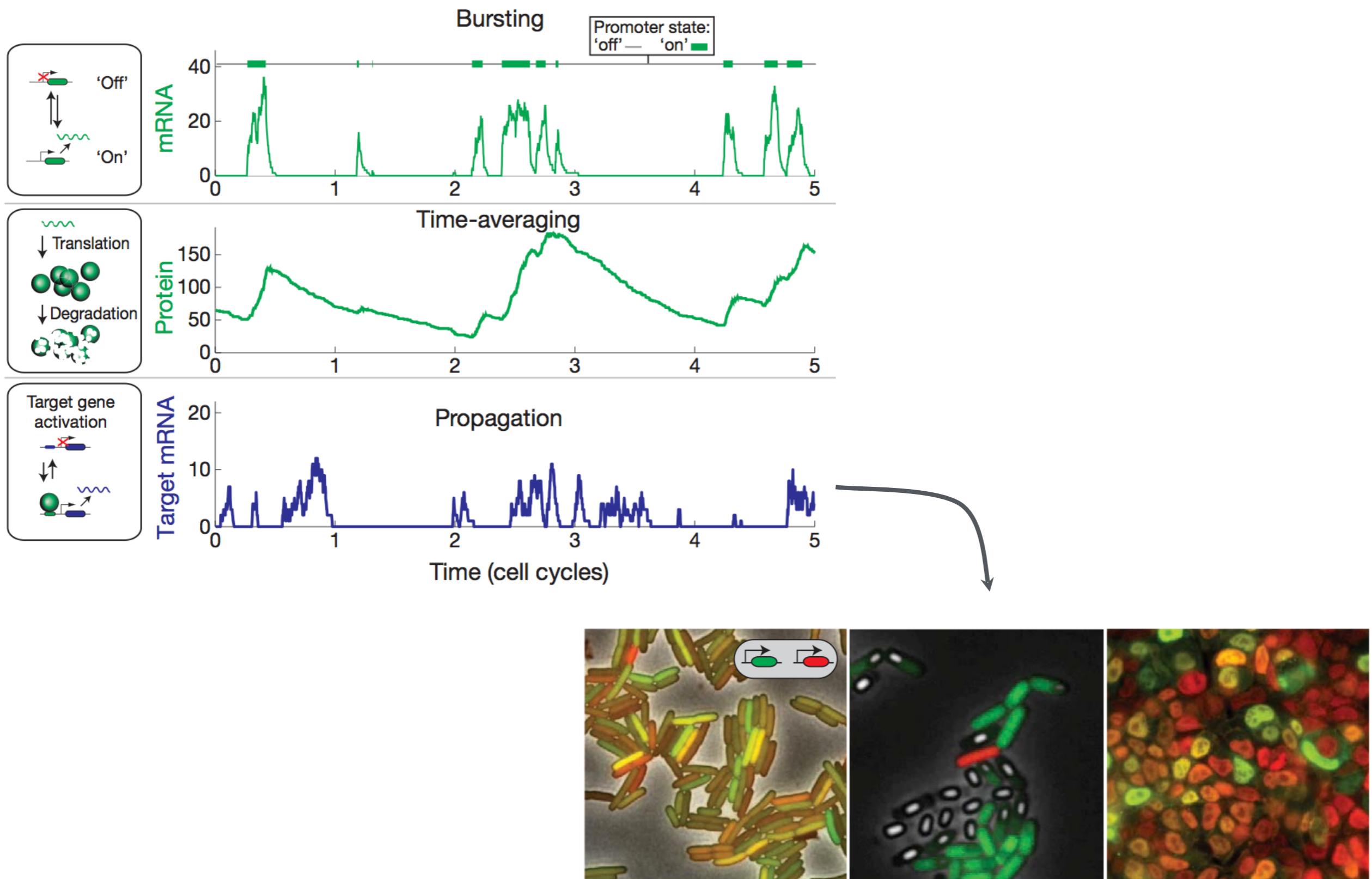
Courtesy: Proceedings of the
National Academy of Sciences

<https://www.youtube.com/watch?v=FhJ-1AOHRy0>

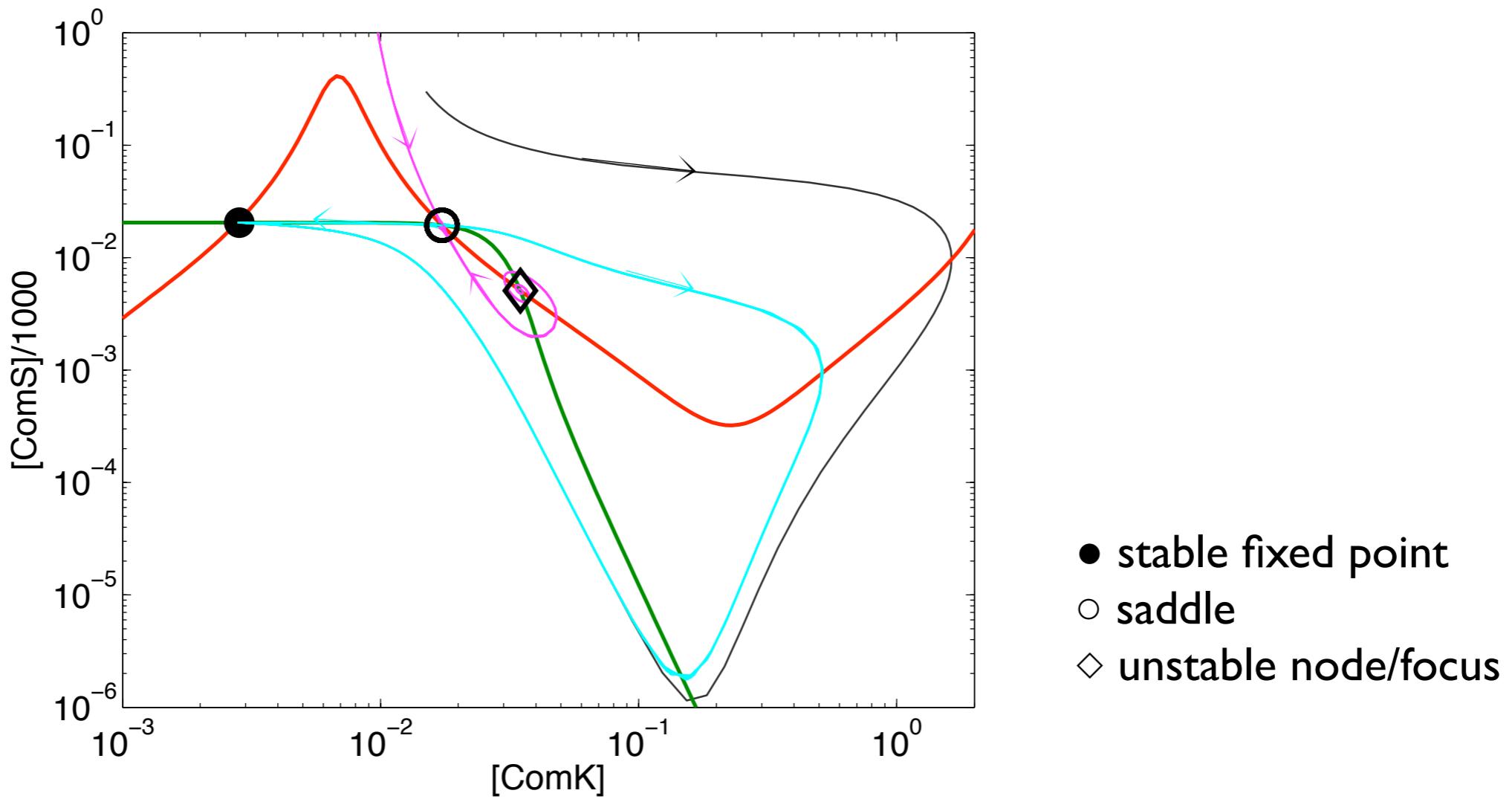
The crowded interior of cells



Noise propagates down gene circuit cascades

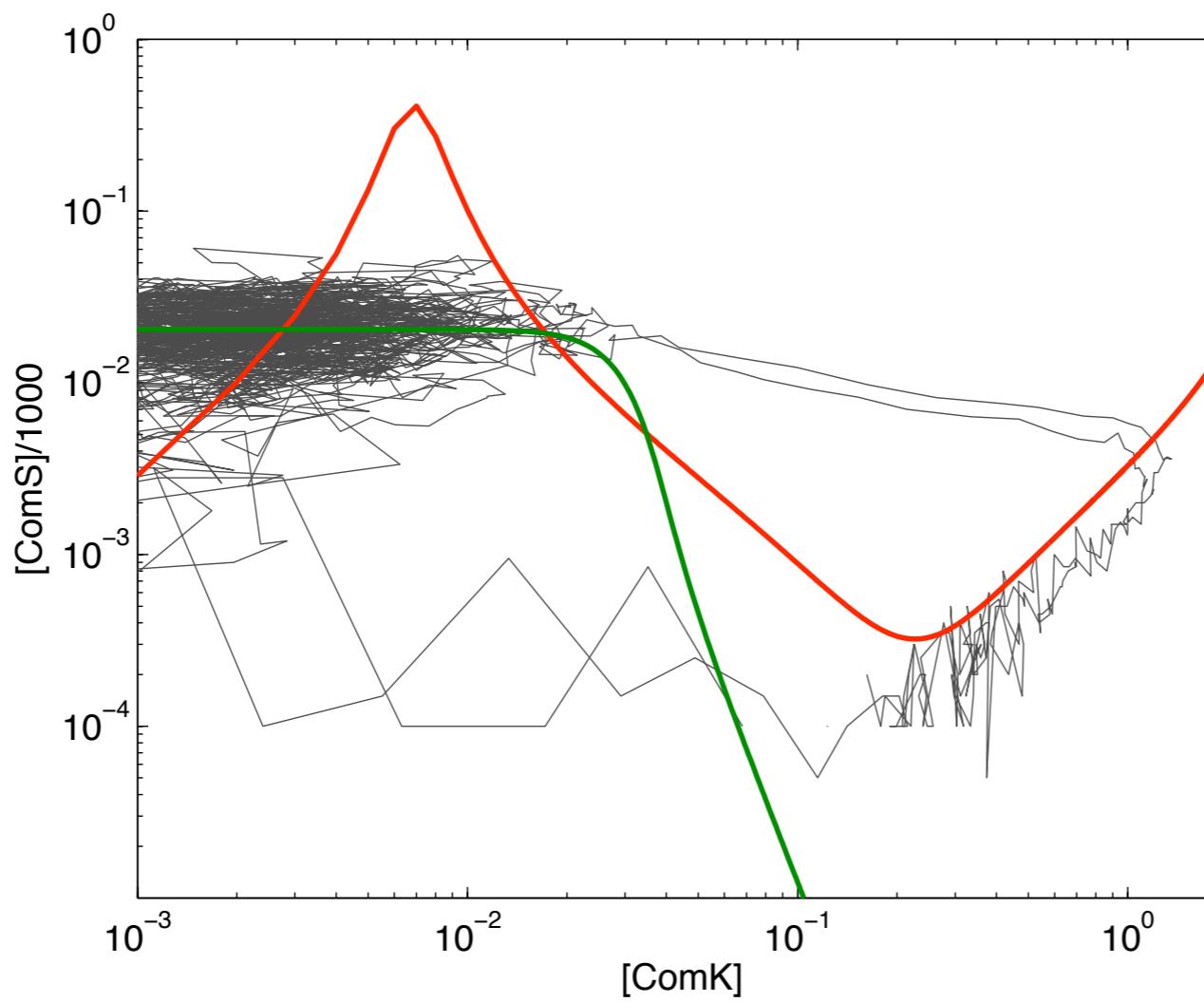


The competence circuit is an excitable system



$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K$$
$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

The competence circuit is an excitable system



- stable fixed point
- saddle
- ◇ unstable node/focus

$$\frac{dK}{dt} = \alpha_k + \frac{\beta_k K^n}{k_k^n + K^n} - \frac{\gamma_k K}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_k K$$
$$\frac{dS}{dt} = \alpha_s + \frac{\beta_s}{1 + (K/k_s)^p} - \frac{\gamma_s S}{1 + \frac{K}{\Gamma_k} + \frac{S}{\Gamma_s}} - \delta_s S$$

Dynamics of gene regulatory circuits

- ✓ 1. Gene circuit dynamics
- ✓ 2. Dissecting a genetic circuit
- ✓ 3. Noise in genetic circuits

Thank you!

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