

Mechanisms of Noise-induced Oscillation in Models of Biological Clock

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Yen Ting Lin

Theoretical Division and Center for Nonlinear Studies
Los Alamos National Laboratory, Los Alamos, USA

Collaborative work with Nicolas Buchler

Department of Physics, Center for Genomics and Computational Biology, Duke University, Durham, USA



Motivation

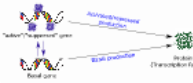
- Oscillatory dynamics are ubiquitous in biological systems: cell division/cycle, metabolism, circadian rhythms, etc.
- Many computational models were proposed for specific systems.
- Noise is an important factor in the process
 - Intrinsic noise: Temperature, light/dark cycle, heterogeneity of copy numbers of the molecules, etc.
 - Extrinsic noise: discreteness of population, volume exclusion, localization in space, etc.
- There is an ongoing debate about if noise is "beneficial" for oscillations
- Pres. noise induce oscillation. This parameter range can be used in some models.
- Gene randomly complicate the substance
- However, the arguments are often performed on model-specific manner.

Questions we ask...

- Can we propose a "simple-harmonic-oscillator"-like model to investigate these computational models, in a most simplified way to deliver generic conclusions?
- What is precisely the "noise-induced oscillation"? Are people comparing the same "animal"? We will show that there are multiple mechanisms to achieve this.
- How to analyze these mechanisms (what is the proper mathematical tool)? Can we generate hypothesis about regulating these stochastic oscillations with different mechanisms?

Prerequisite and the structure of the talk

Gene expression for each gene/TF pair: we adopted the minimal single-stage process



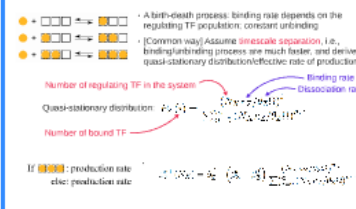
Scope of the models: titration-based oscillators: Two genes, two transcription factors. We will explore possible feedback mechanism until deterministic oscillation is possible.



We will investigate both the deterministic and stochastic models

Deterministic model: mass action kinetics

Discrete promoter site dynamics



Deterministic model: mass action kinetics

Using the effective production rate, we arrived at the two-dimensional deterministic dynamics described by the ODE:

$$\dot{x}(t) = \beta_X^{eff}(\Omega x) - \delta_X x - \alpha x y,$$

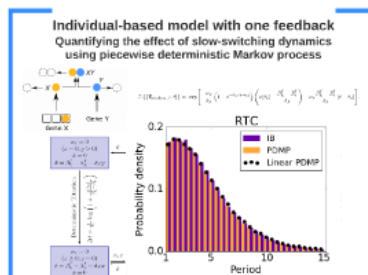
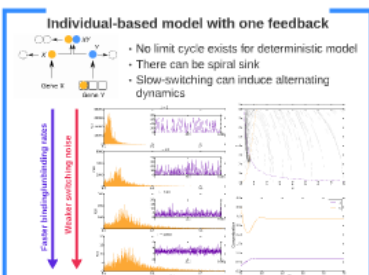
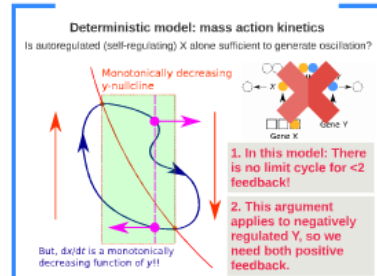
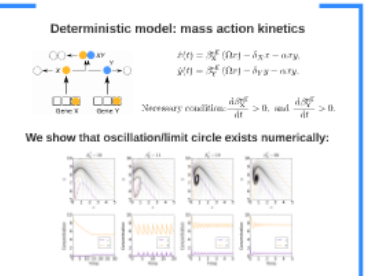
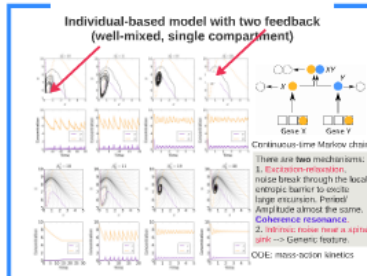
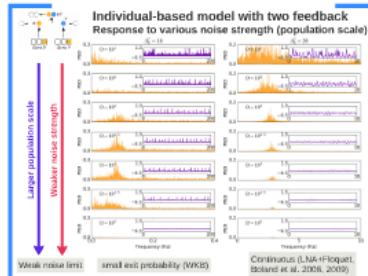
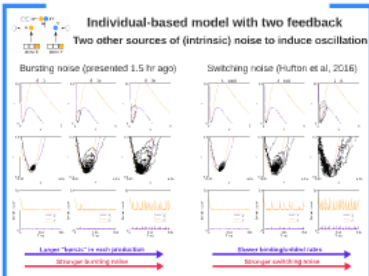
$$\dot{y}(t) = \beta_Y^{eff}(\Omega x) - \delta_Y y - \alpha x y.$$

Bifurcation criterion stated that a limit cycle does not exist when

$$\partial_x \mathcal{F} + \partial_y \mathcal{G} = -\delta_X - \delta_Y - \alpha(x+y) + \Omega \frac{d\beta_X^{eff}}{dx}$$

does not change sign on a simply connected domain (in our case, $\mathbb{R}^+ \times \mathbb{R}^+$). If a limit cycle exists, a necessary condition

$$= \frac{d\beta_X^{eff}}{dx} > 0 \Rightarrow X \text{ must be positively regulating itself.}$$



Summary

- We identified the minimal regulatory mechanisms which allows deterministic or stochastic oscillations
- We identified different mechanisms of noise-induced oscillations
- We provide mechanistic insights to explain noise-induced oscillations in the published models

Future directions

- Condense the material and write it up...
- Connections to the (abstract) stochastic phase oscillator models?
- WKB, quasi-potential, etc. for the excitable systems and comparison to generic stochastic broadening of the spiral sink?
- Noisy extrinsic signal (entrainment?)
- Some conclusion of the deterministic dynamics will not hold for systems with mRNA: is including mRNA a "correction", or we need more theory?

Very rough draft available; if you are interested: yentingl@lanl.gov

Thank you for your attention!

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 - **Intrinsic noise**: **discreteness of population**, volume exclusion, localization in space, etc
- There is an ongoing debate about if noise is "beneficial" for oscillations
 - Pros: exists noise-induced oscillation. The parameter range can be vast in some models.
 - Cons: inevitably compromise the coherence

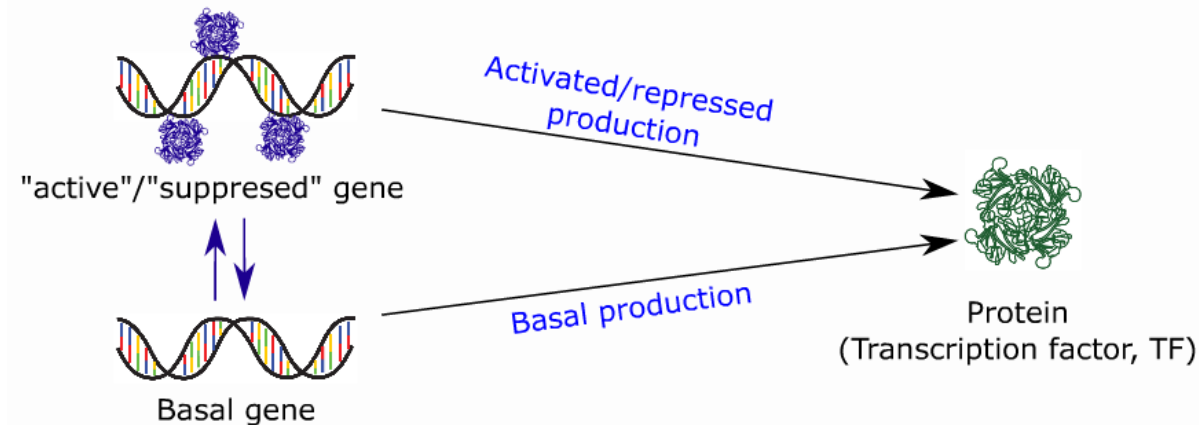
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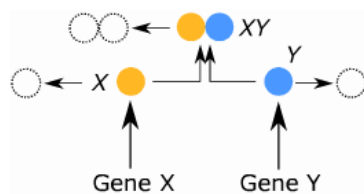
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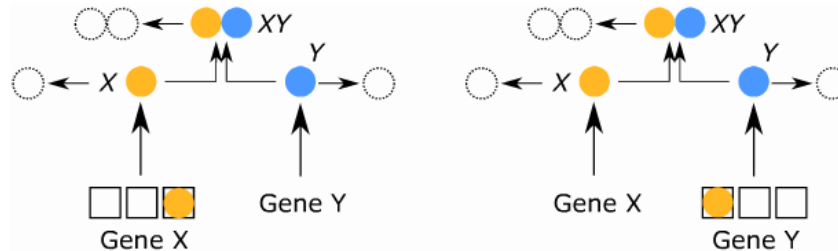


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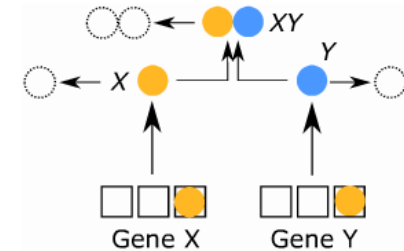
No feedback



One feedback

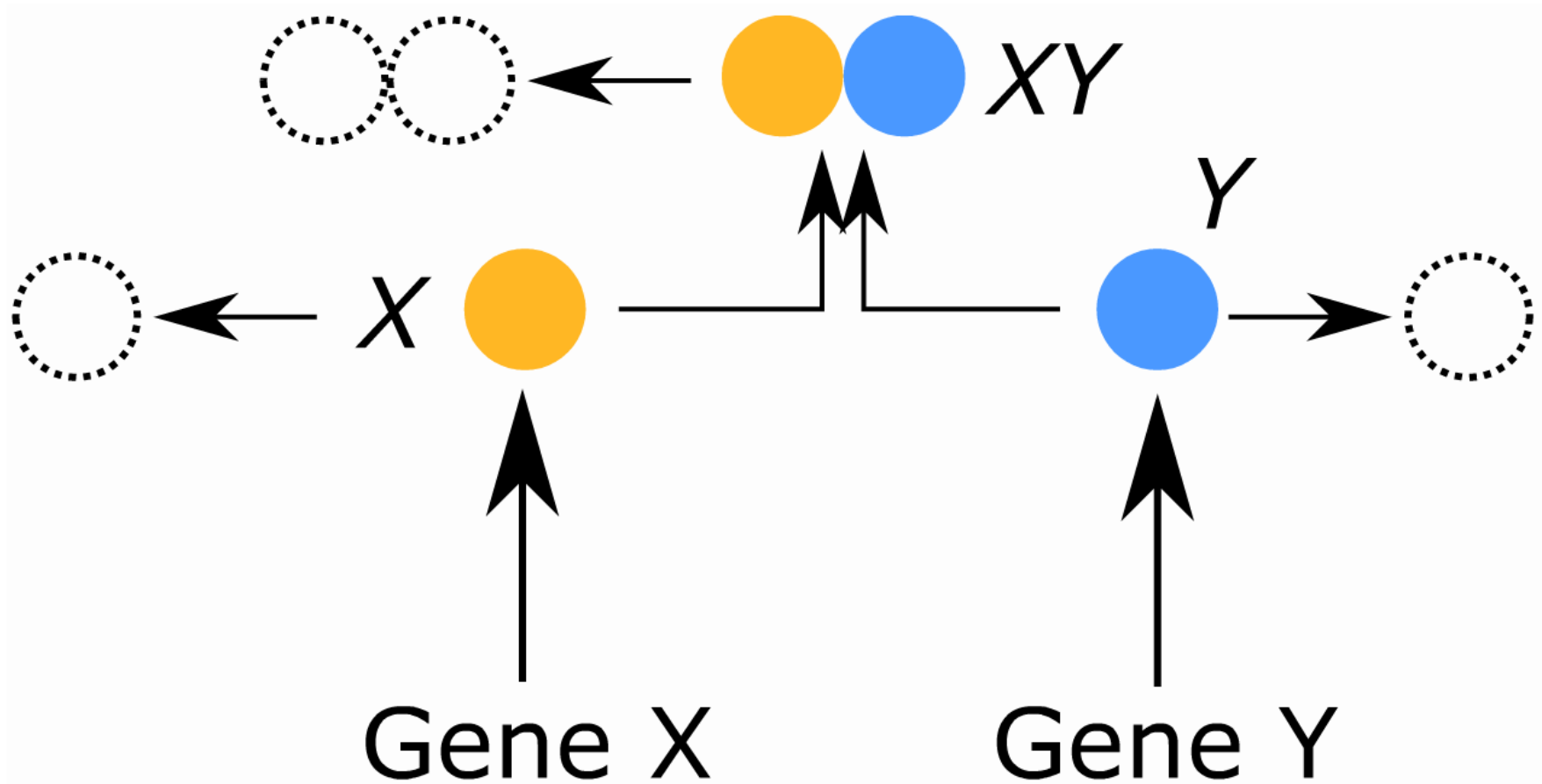


Two feedback



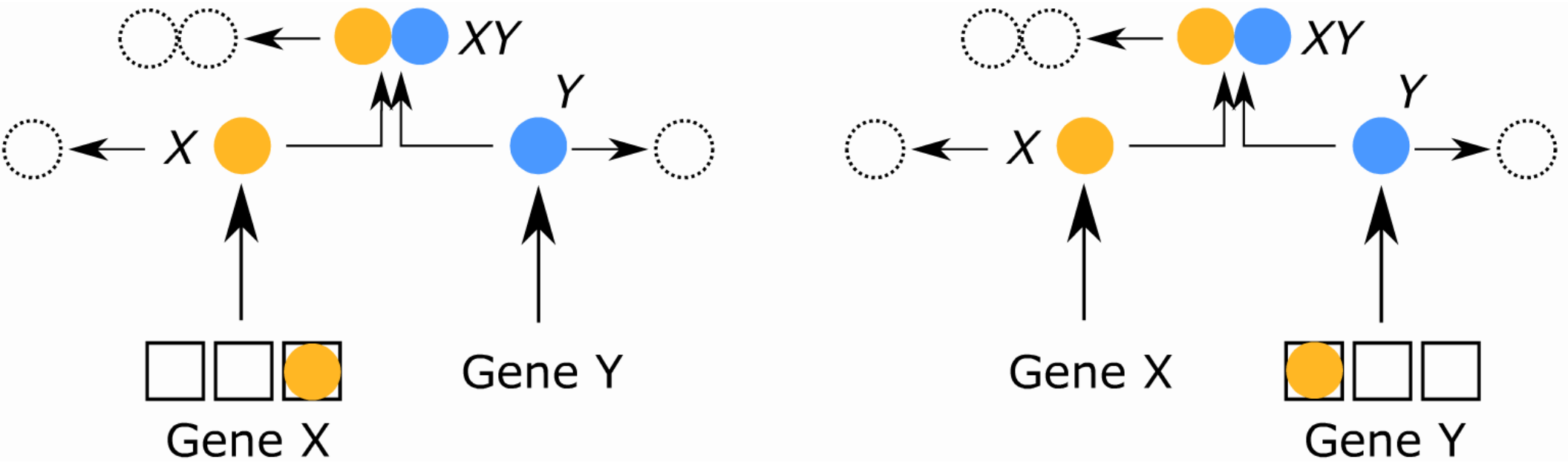
We will investigate both the **deterministic** and **stochastic** models

No feedback



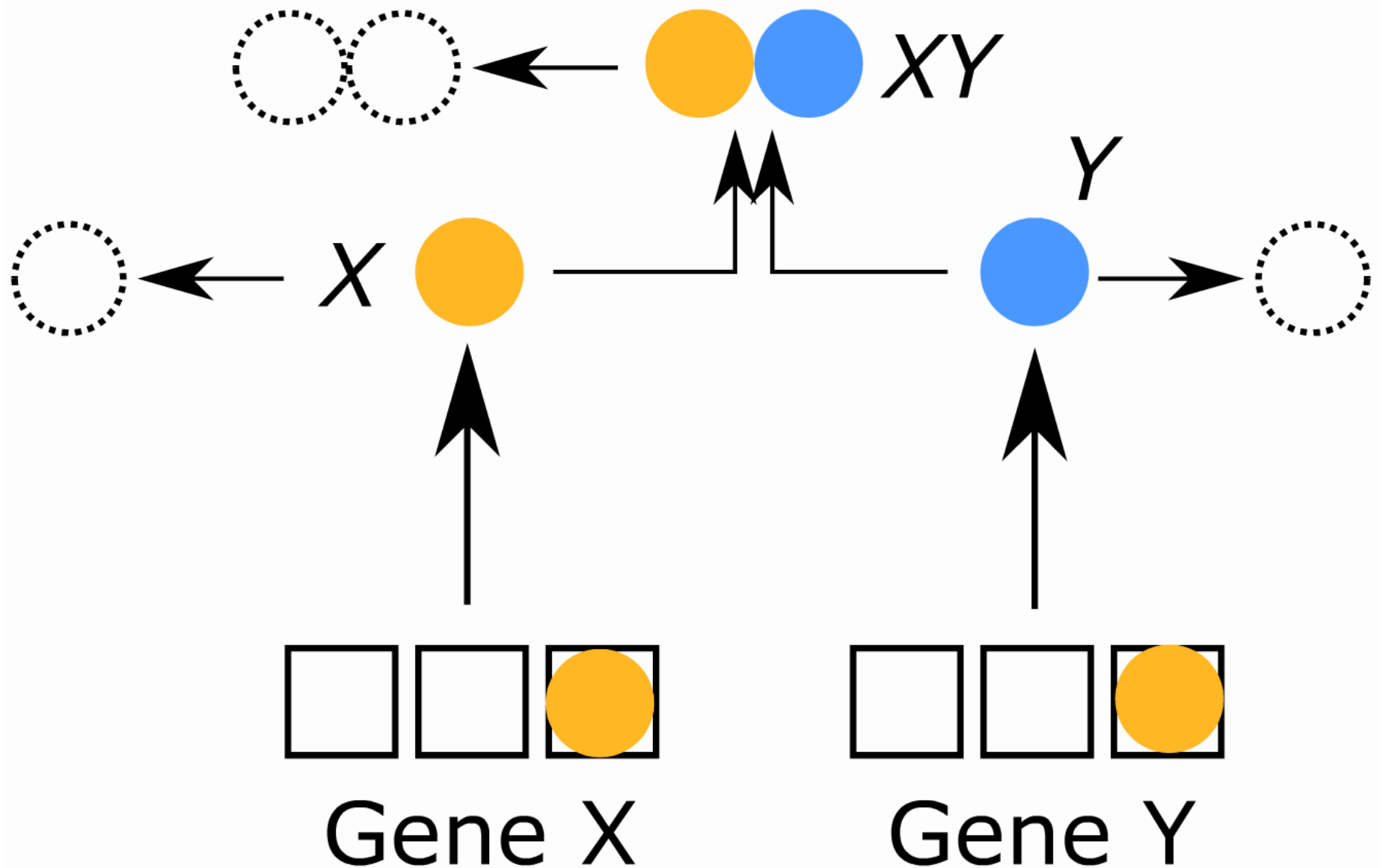
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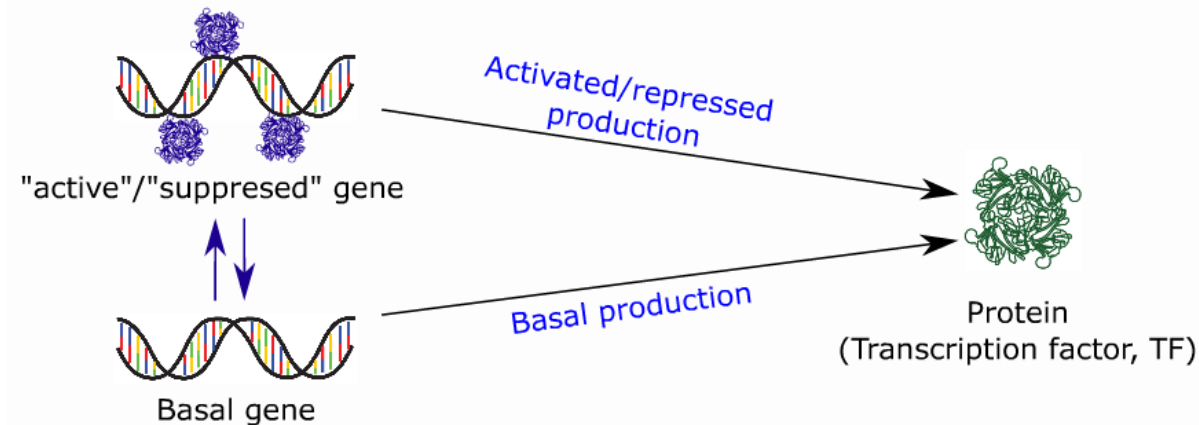
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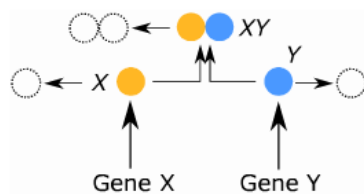
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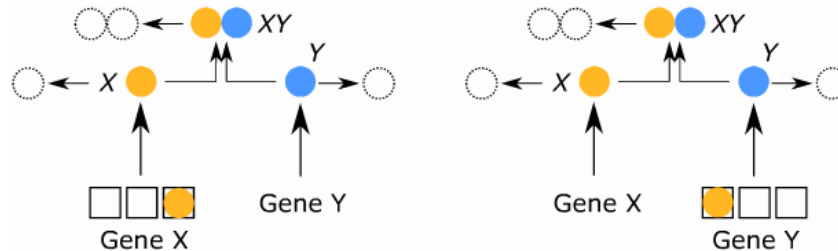


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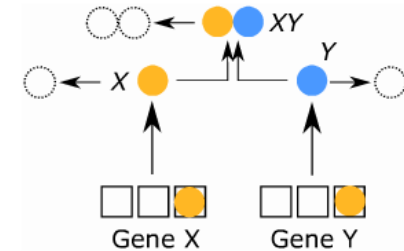
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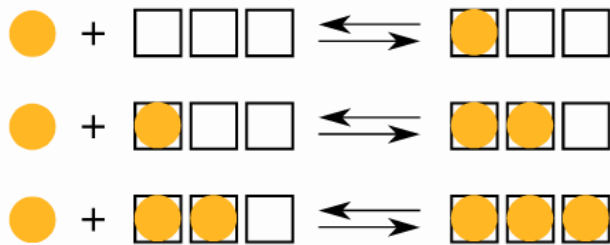
Two feedback



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Deterministic model: mass action kinetics

Discrete promoter site dynamics



- A birth-death process: binding rate depends on the regulating TF population; constant unbinding
- [Common way] Assume **timescale separation**, i.e., binding/unbinding process are much faster, and derive quasi-stationary distribution/effective rate of production

Number of regulating TF in the system \rightarrow Binding rate
 Dissociation rate \rightarrow

Quasi-stationary distribution: $P_Z(i) = \frac{(N_X \kappa_Z / \theta_Z \Omega)^i}{\sum_{m=0}^{n_Z} (N_X \kappa_Z / \theta_Z \Omega)^m}$

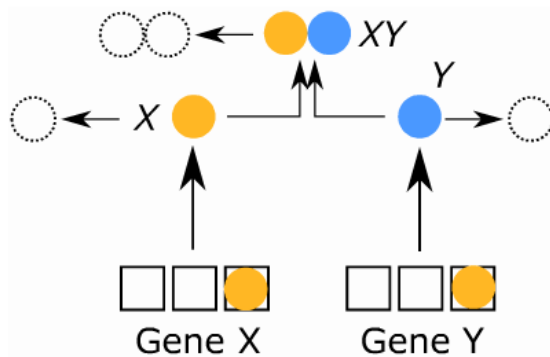
Number of bound TF \rightarrow

If : production rate β_Z^b
 else: production rate β_Z^f

$$\beta_Z^{\text{eff}}(N_X) = \beta_Z^f + \left(\beta_Z^b - \beta_Z^f \right) \frac{(N_X \kappa_Z / \theta_Z \Omega)^{n_Z}}{\sum_{m=0}^{n_Z} (N_X \kappa_Z / \theta_Z \Omega)^m}$$

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$$\partial_x \mathcal{F} + \partial_y \mathcal{G} = -\delta_X - \delta_Y - \alpha(x + y) + \Omega \frac{d\beta_X^{\text{eff}}}{dx}.$$

does not change sign on a simply connected domain (in our case, $\mathbb{R}^+ \times \mathbb{R}^+$).
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$$\Rightarrow \frac{d\beta_X^{\text{eff}}}{dx} > 0 \Rightarrow X \text{ must be positively regulating itself.}$$



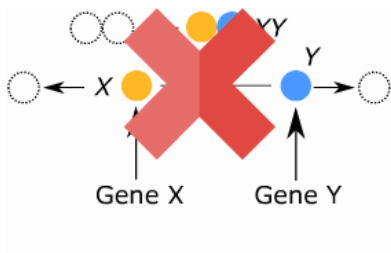
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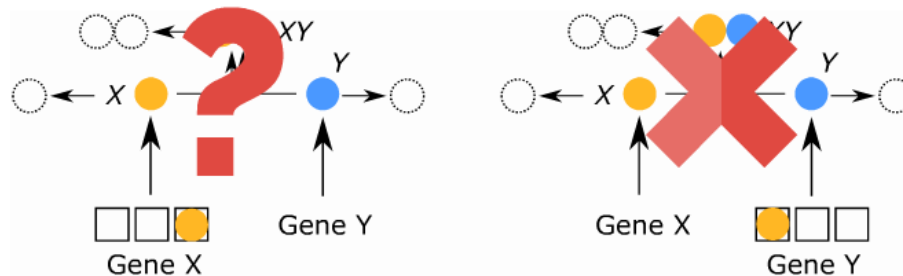
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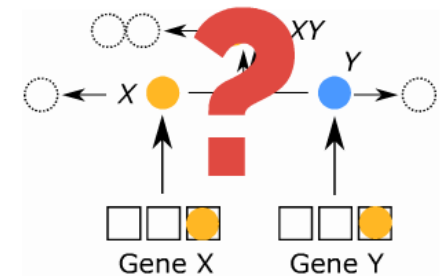
No feedback



One feedback

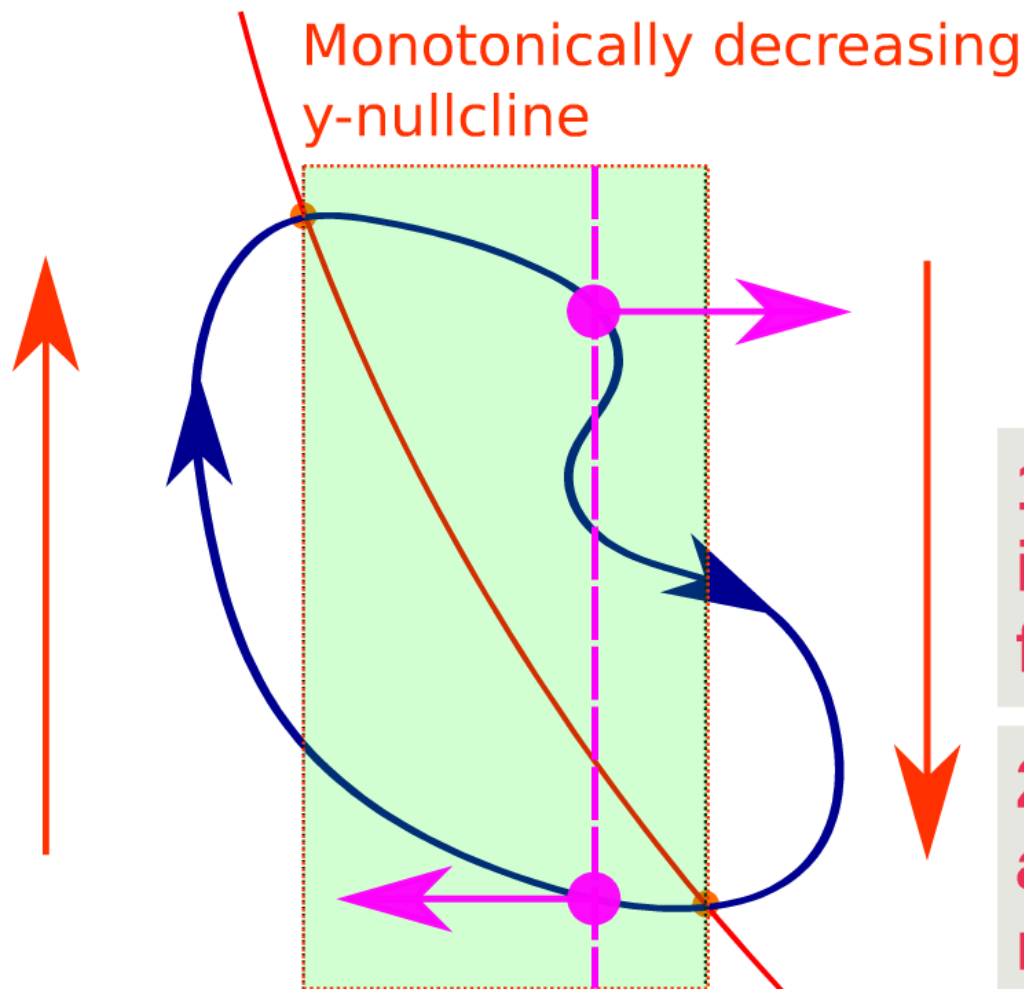


Two feedback

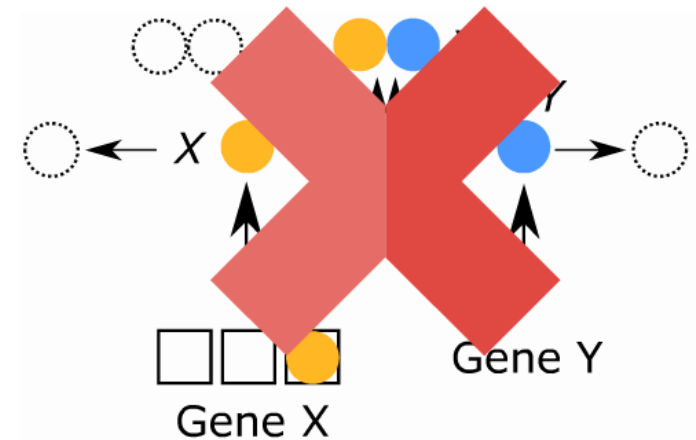


Deterministic model: mass action kinetics

Is autoregulated (self-regulating) X alone sufficient to generate oscillation?



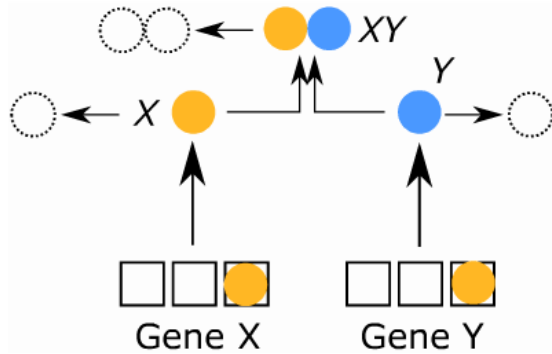
But, dx/dt is a monotonically decreasing function of y !!



1. In this model: There is no limit cycle for <2 feedback!

2. This argument applies to negatively regulated Y, so we need both positive feedback.

Deterministic model: mass action kinetics

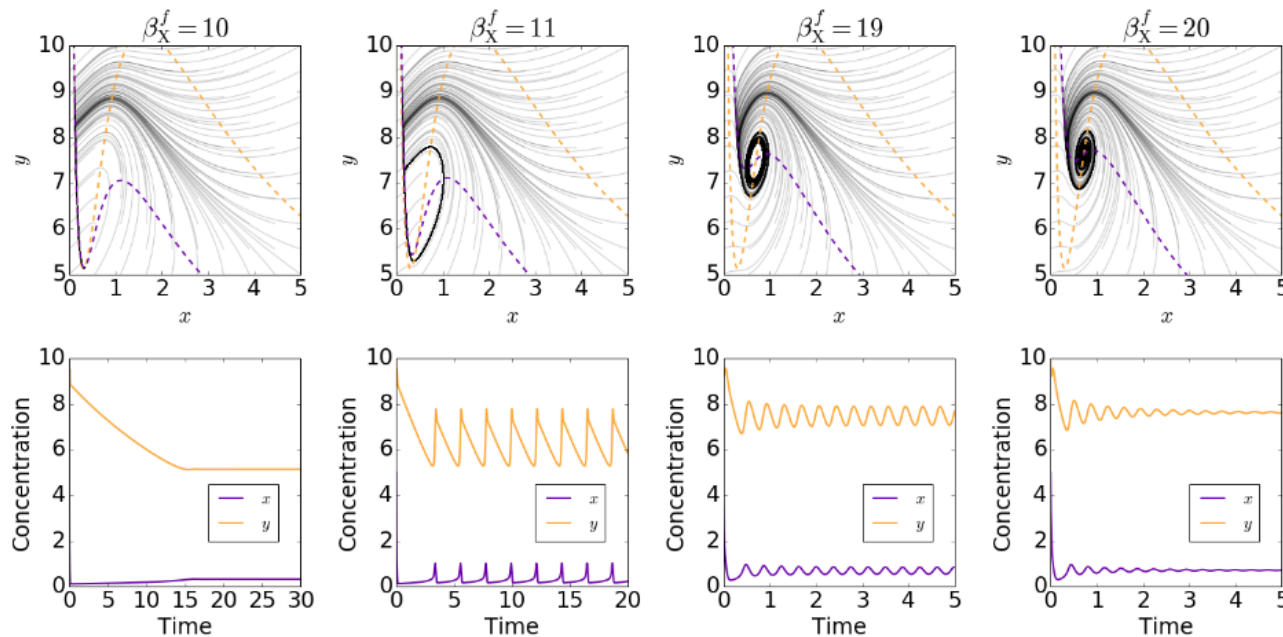


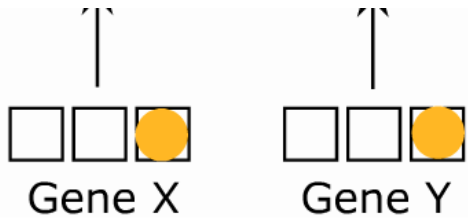
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Necessary condition: $\frac{d\beta_X^{\text{eff}}}{dt} > 0$, and $\frac{d\beta_Y^{\text{eff}}}{dt} > 0$.

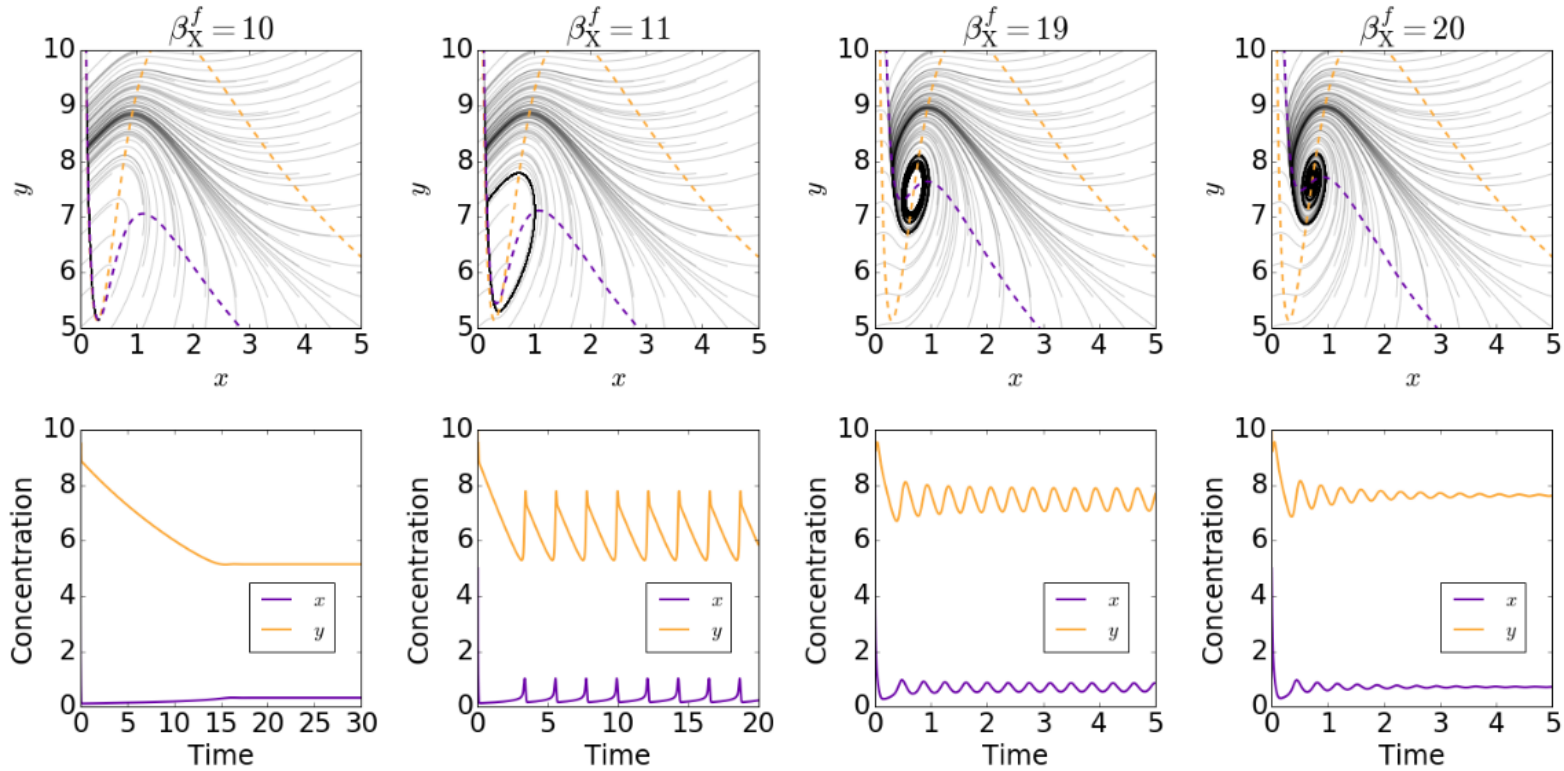
We show that oscillation/limit circle exists numerically:



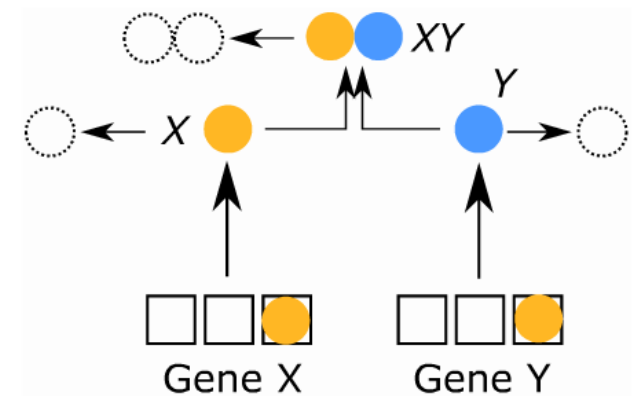
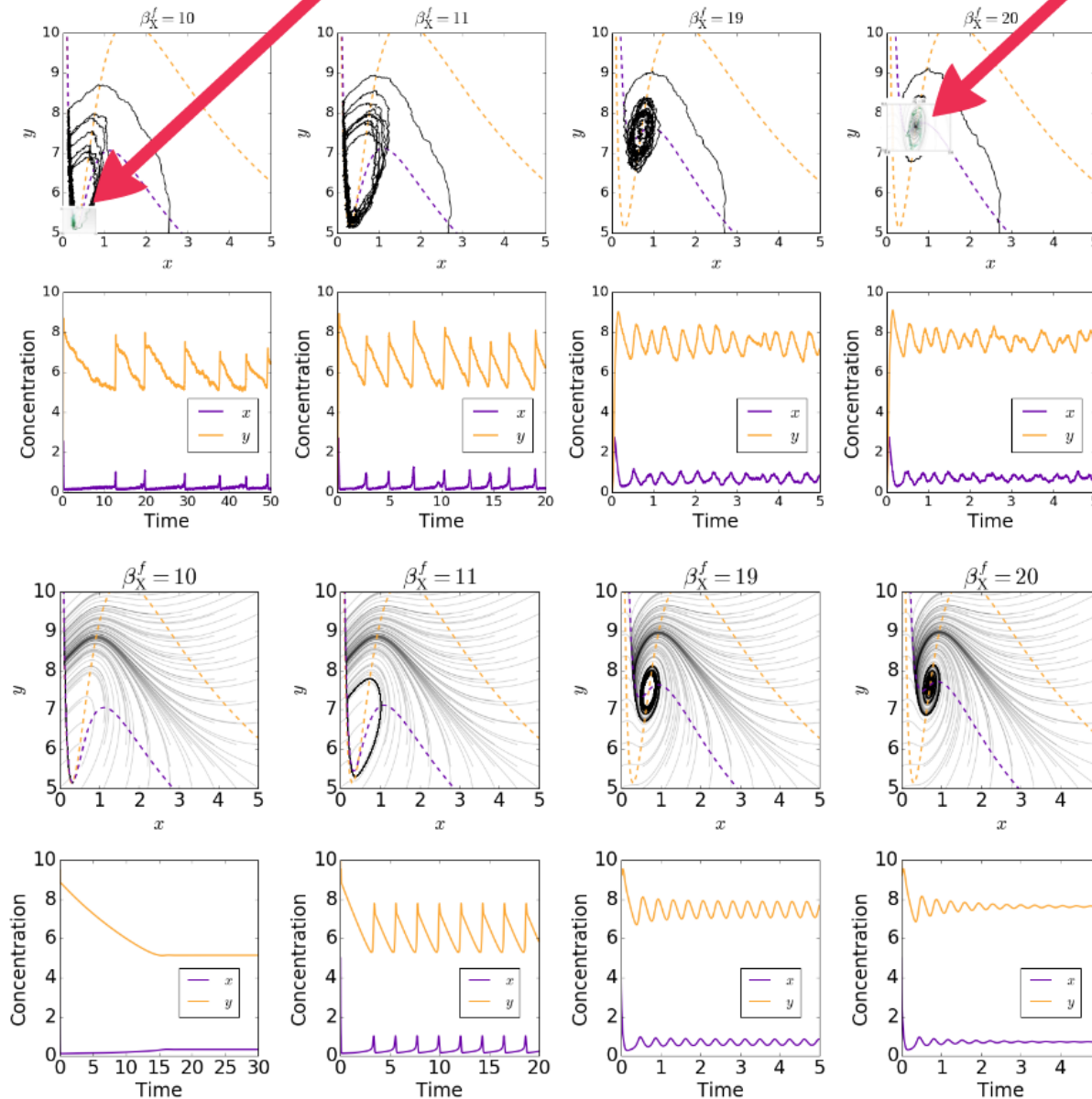


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Individual-based model with two feedback (well-mixed, single compartment)

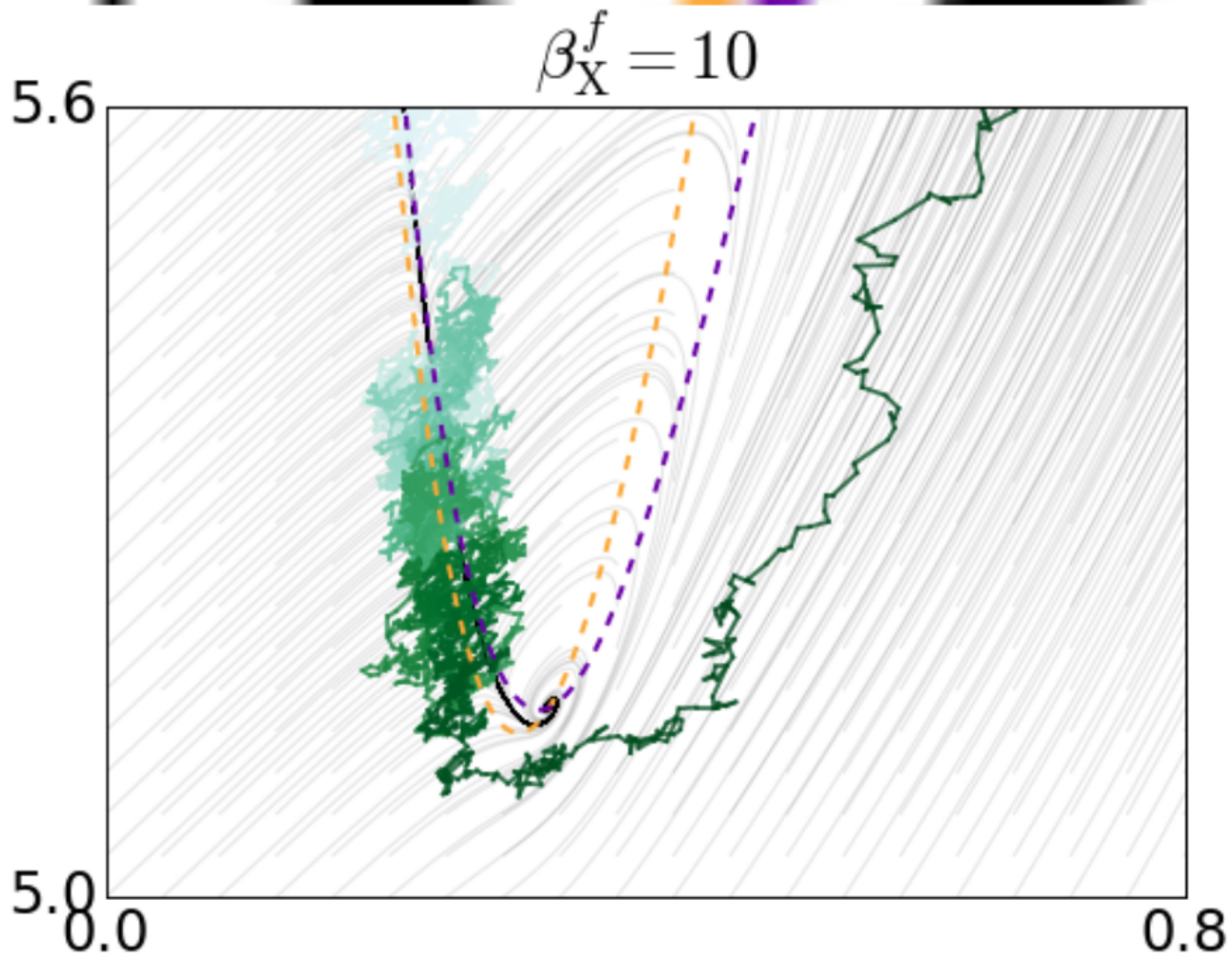


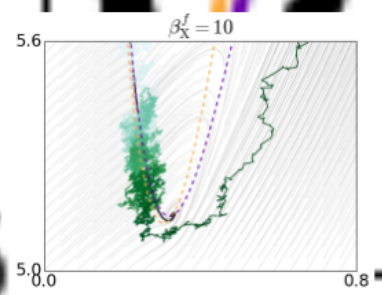
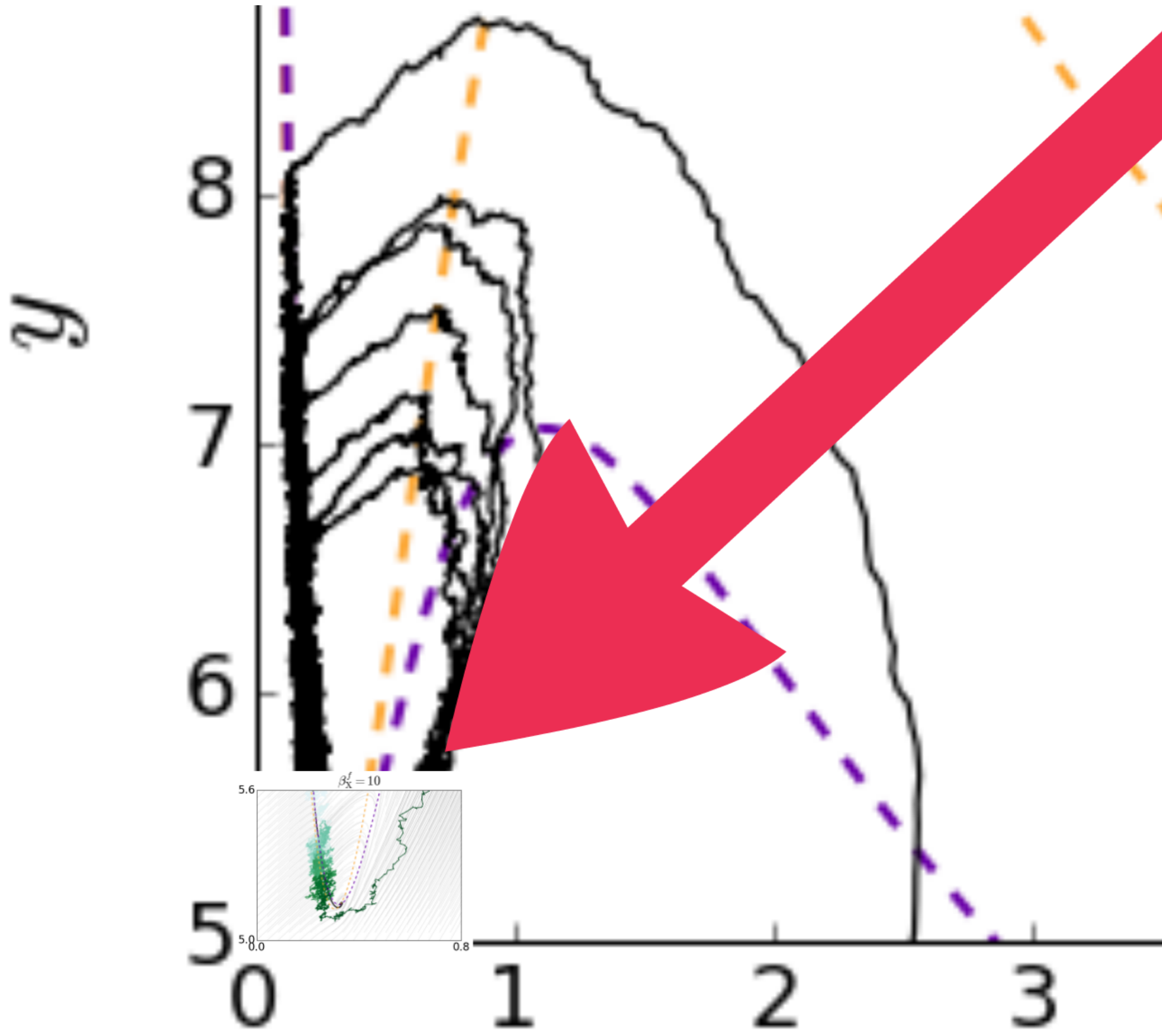
Continuous-time Markov chain

There are **two** mechanisms:

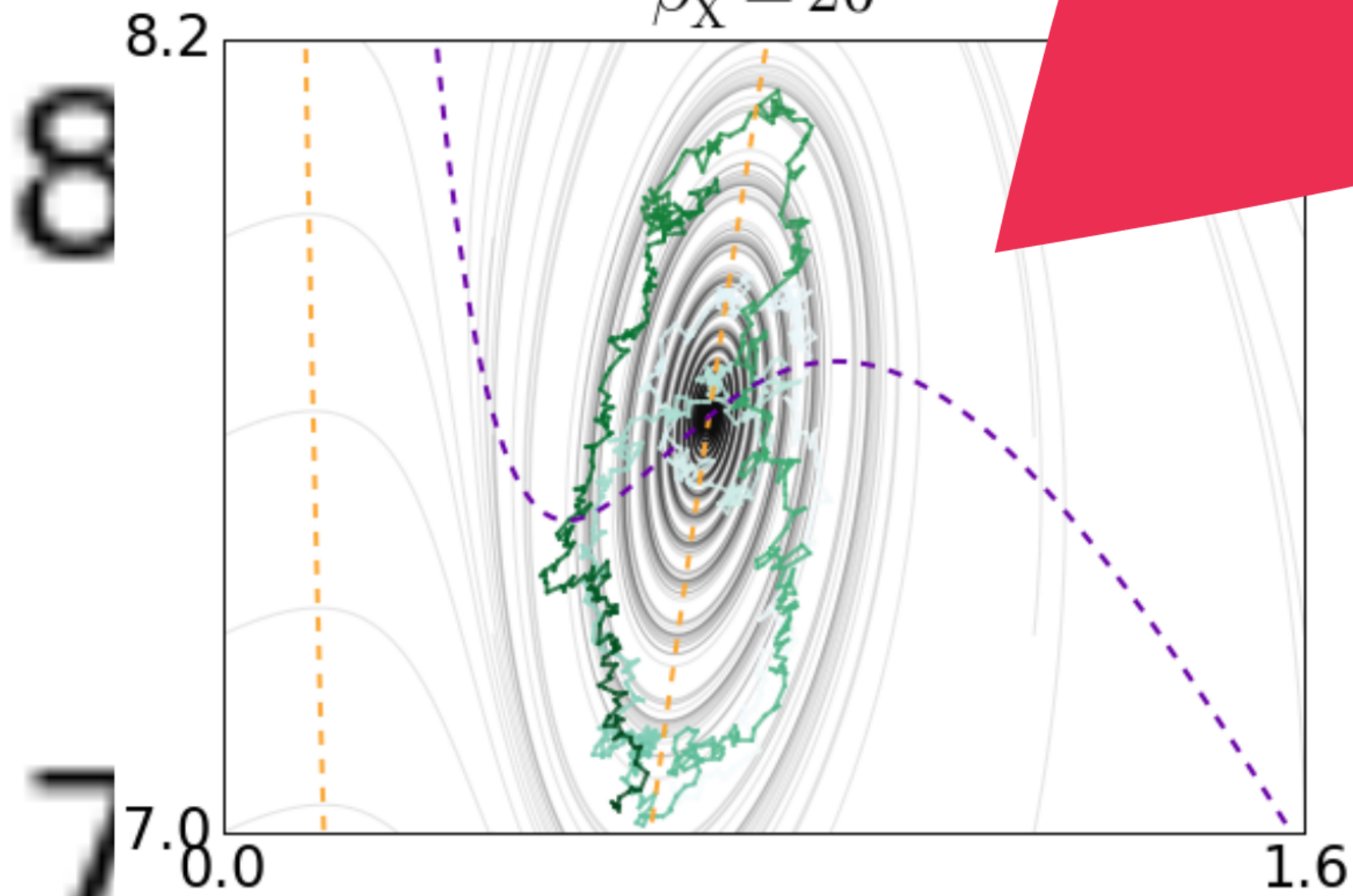
- Excitation-relaxation**, noise break through the local entropic barrier to excite large excursion. Period/Amplitude almost the same.
- Coherence resonance**.
- Intrinsic noise near a spiral sink** --> Generic feature.

ODE: mass-action kinetics

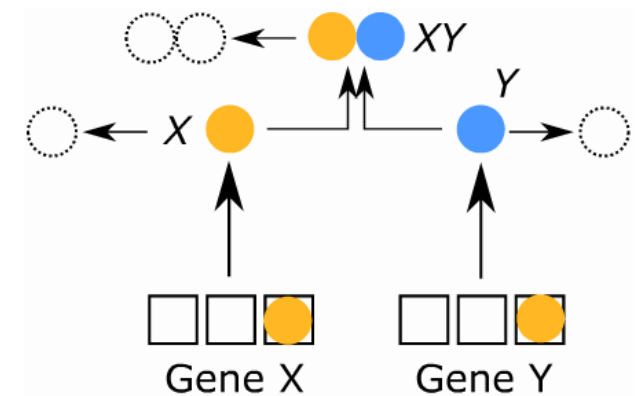
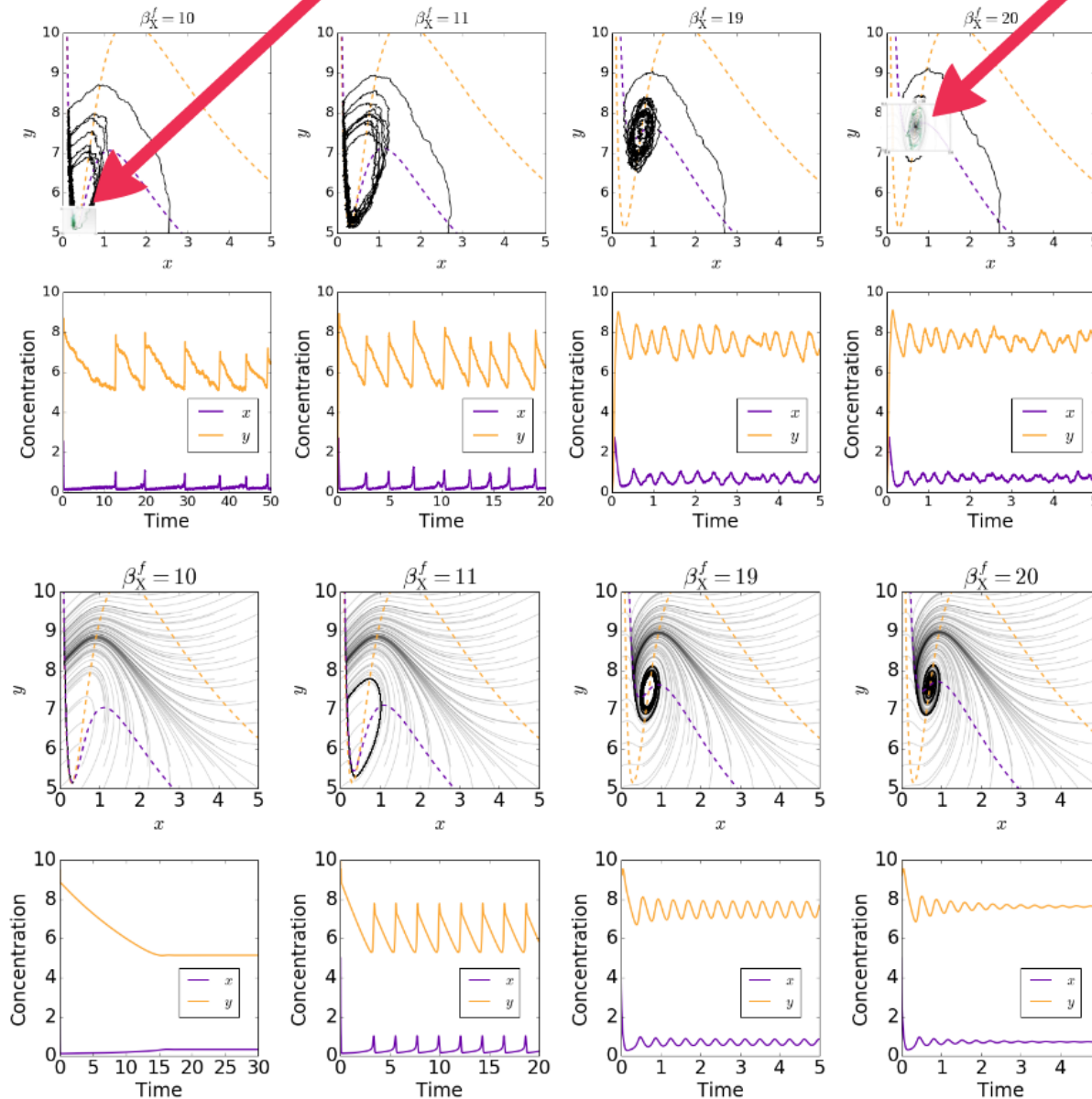




$$\beta_X^f = 20$$



Individual-based model with two feedback (well-mixed, single compartment)

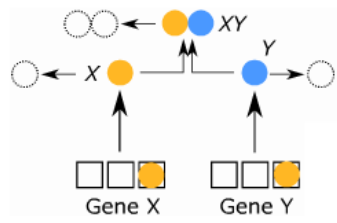


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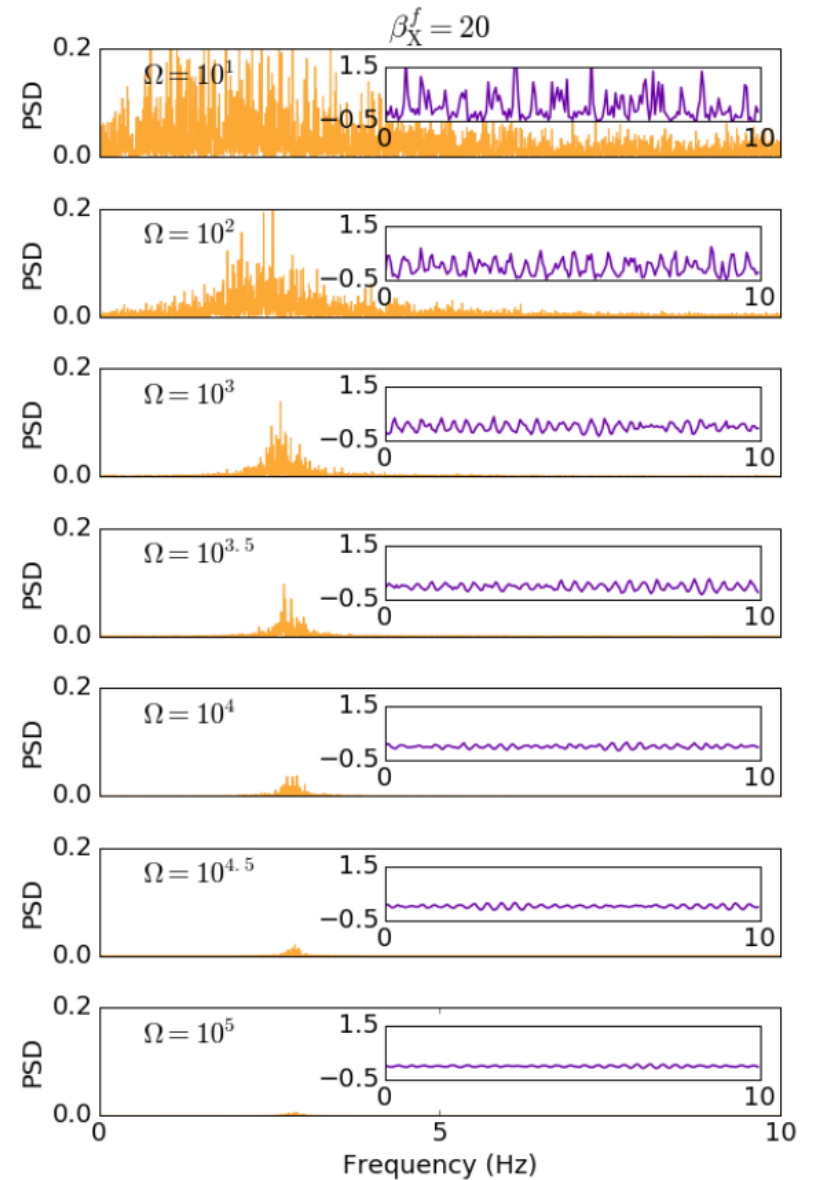
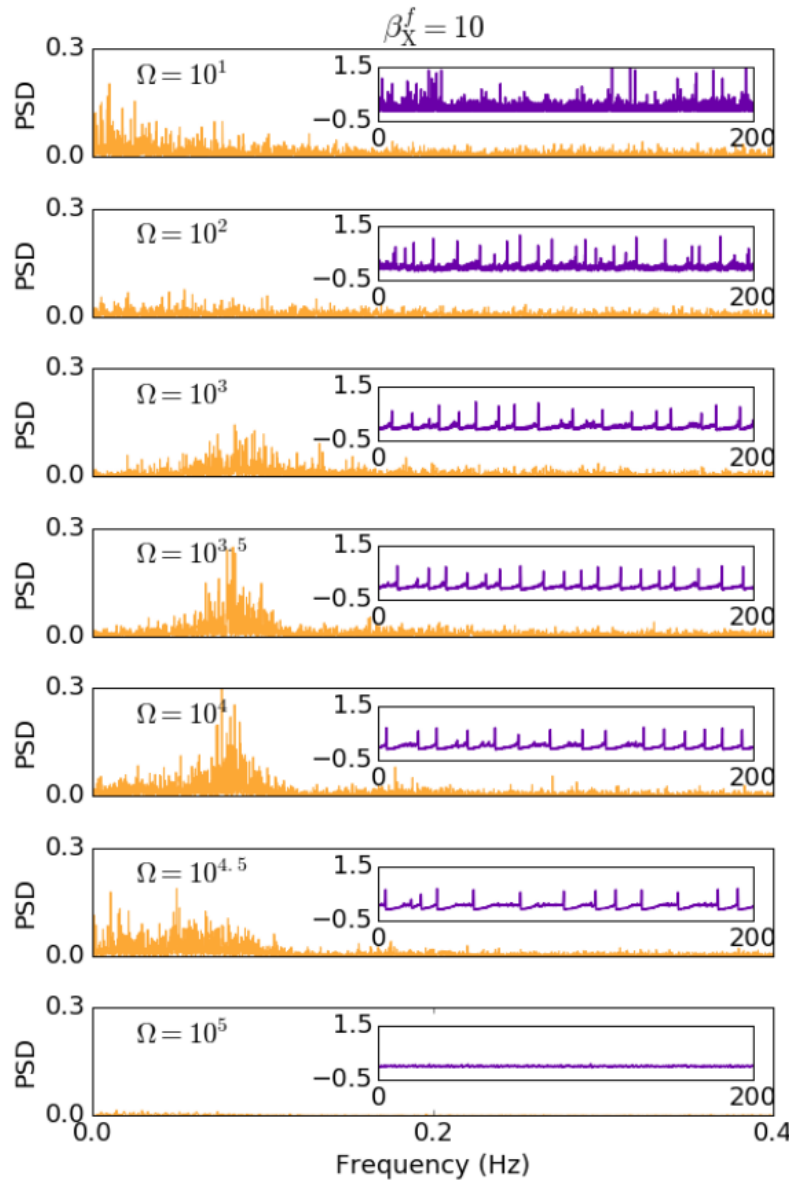


Individual-based model with two feedback

Response to various noise strength (population scale)

Larger population scale

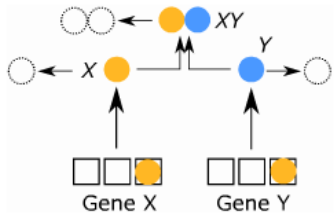
Weaker noise strength



Weak noise limit

small exit probability (WKB)

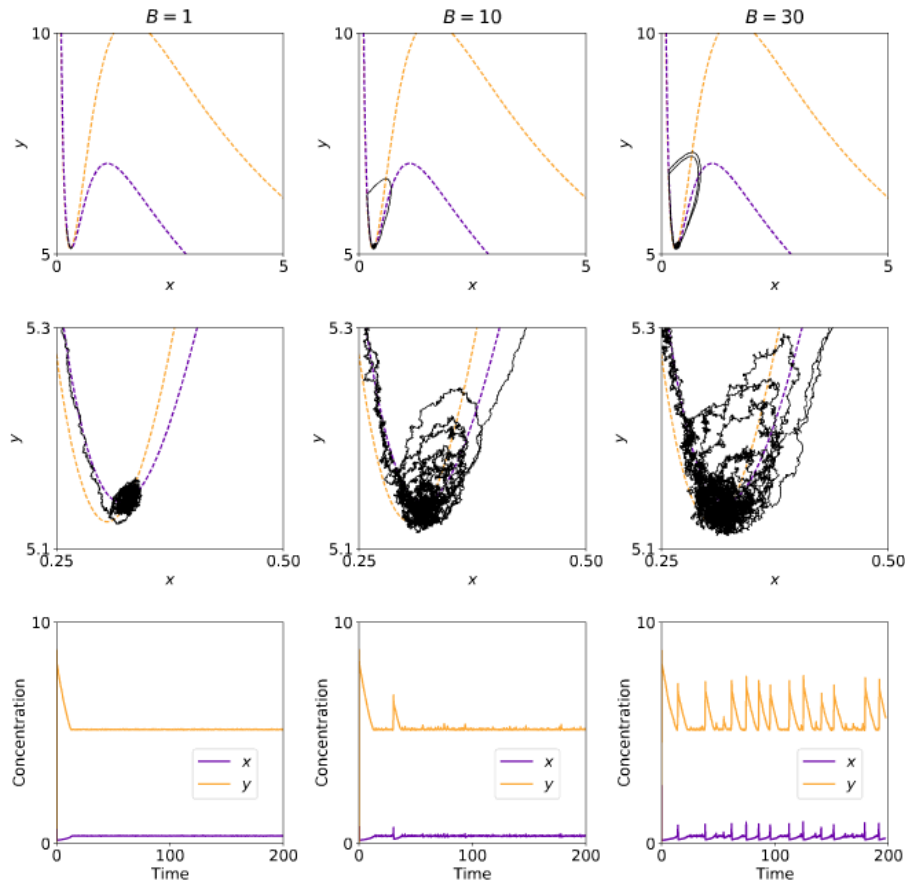
Continuous (LNA+Floquet, Boland et al. 2008, 2009)



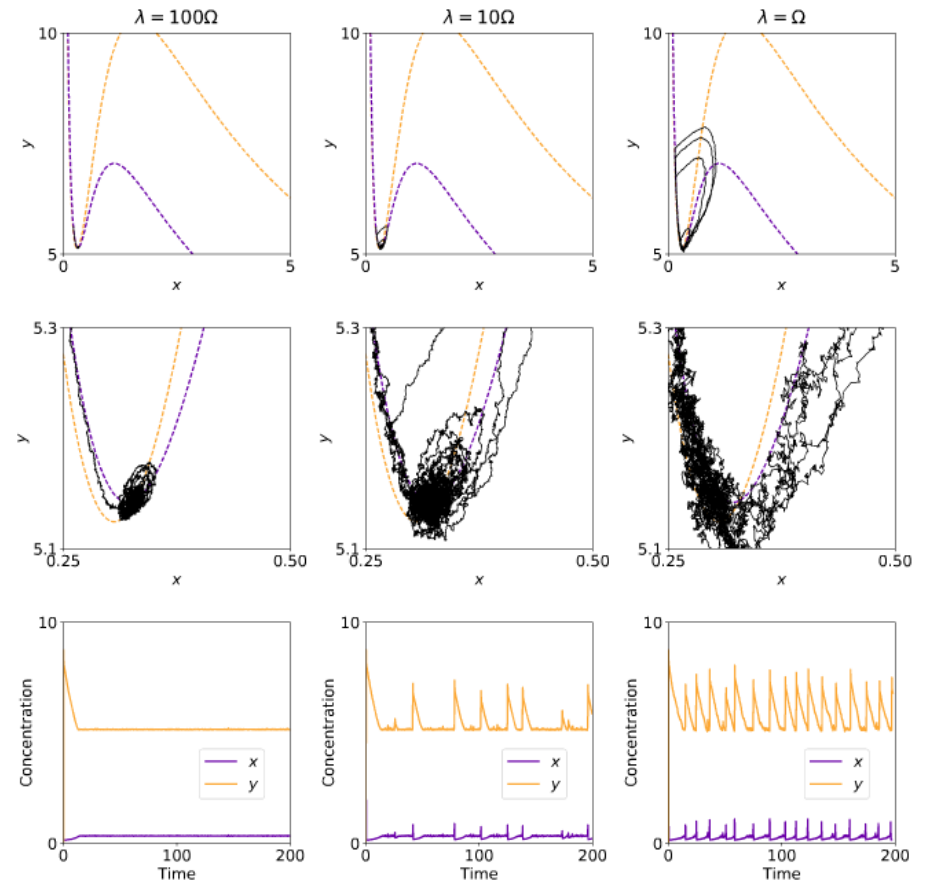
Individual-based model with two feedback

Two other sources of (intrinsic) noise to induce oscillation

Bursting noise (presented 1.5 hr ago)



Switching noise (Hufton et al, 2016)



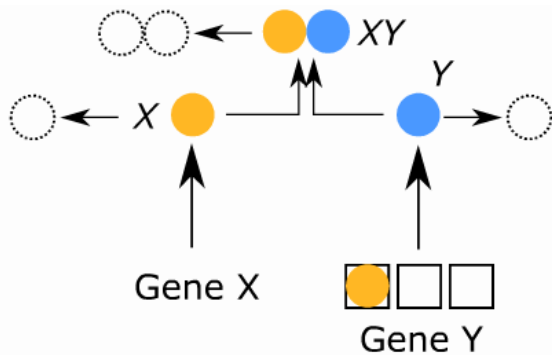
Larger "bursts" in each production

Stronger bursting noise

Slower binding/unbind rates

Stronger switching noise

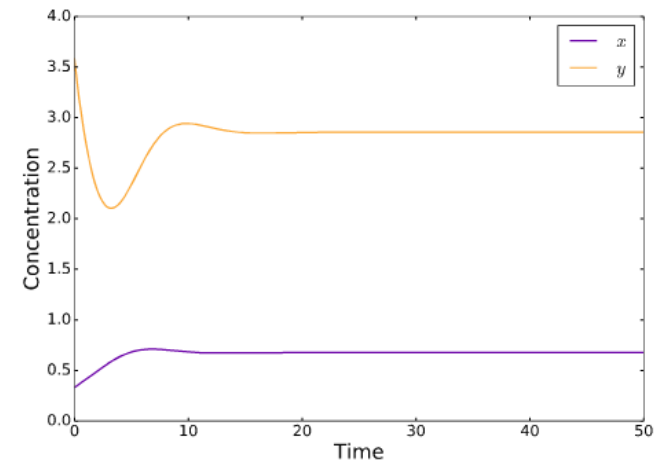
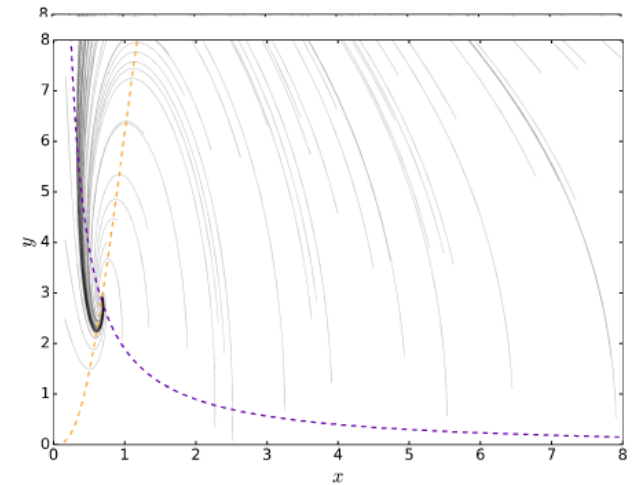
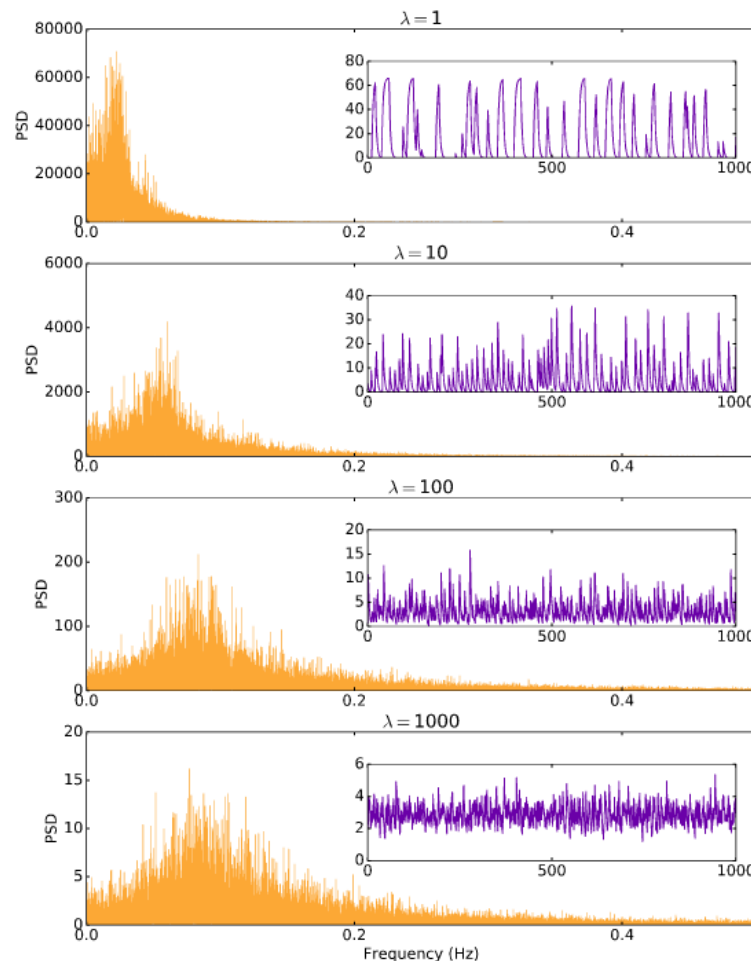
Individual-based model with one feedback



- No limit cycle exists for deterministic model
- There can be spiral sink
- Slow-switching can induce alternating dynamics

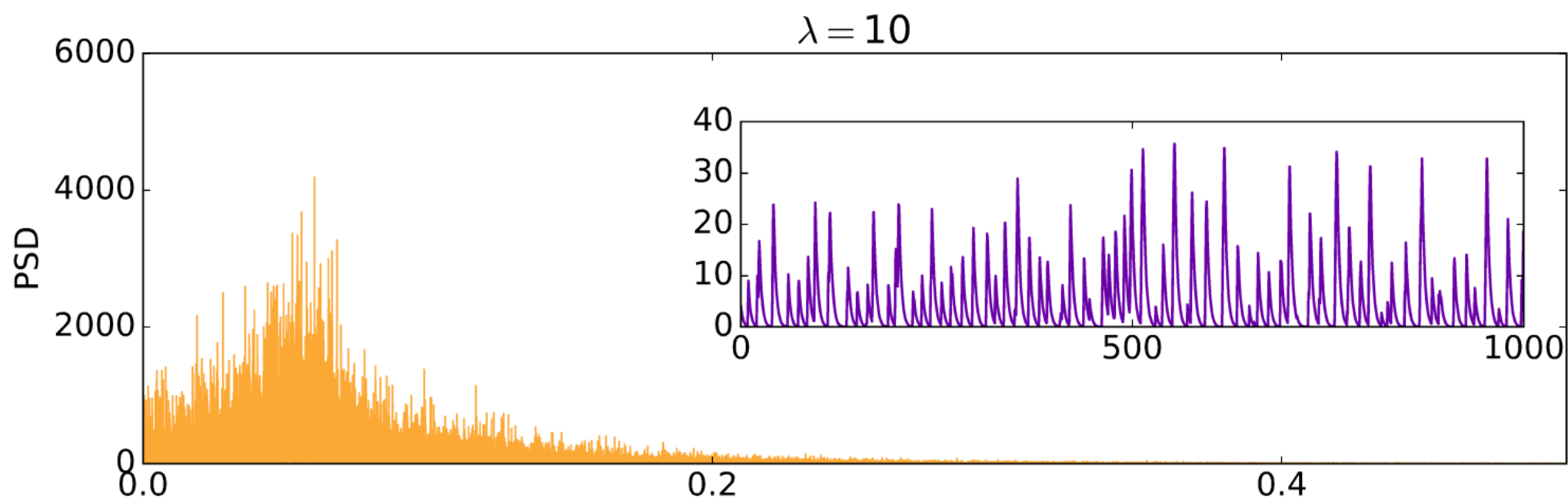
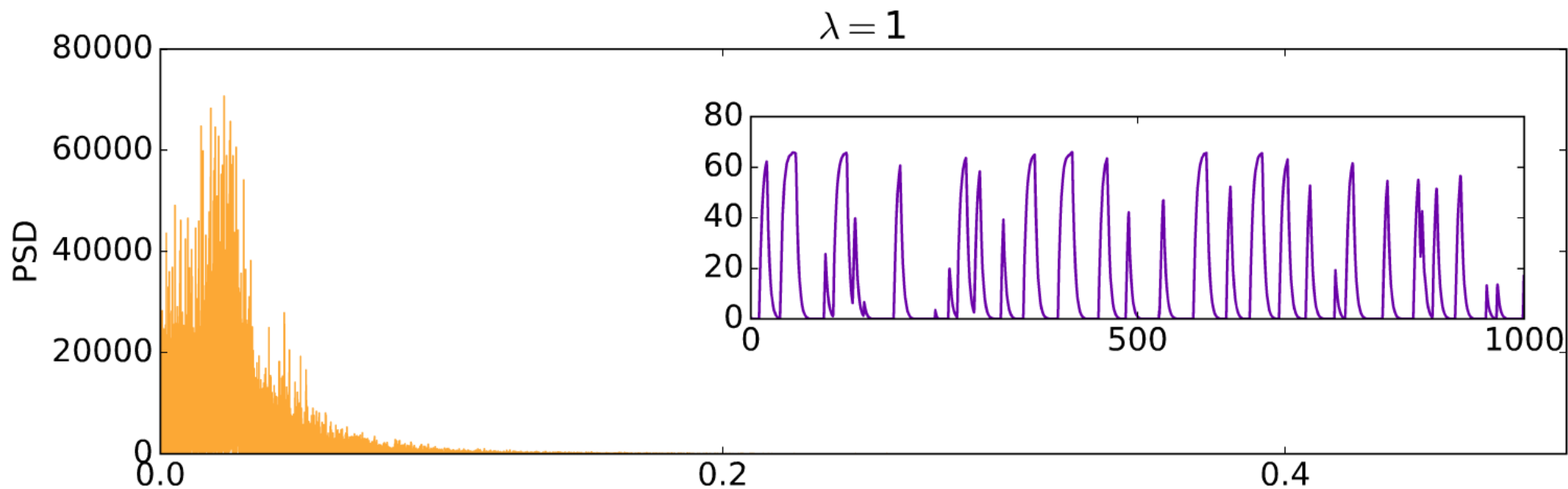
Faster binding/unbinding rates

Weaker switching noise



ene Y

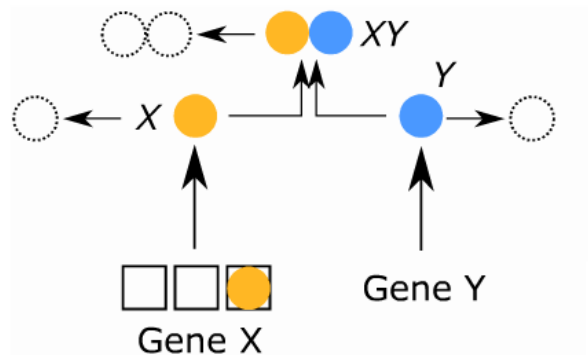
dynamics



$\lambda = 100$

Individual-based model with one feedback

Quantifying the effect of slow-switching dynamics using piecewise deterministic Markov process

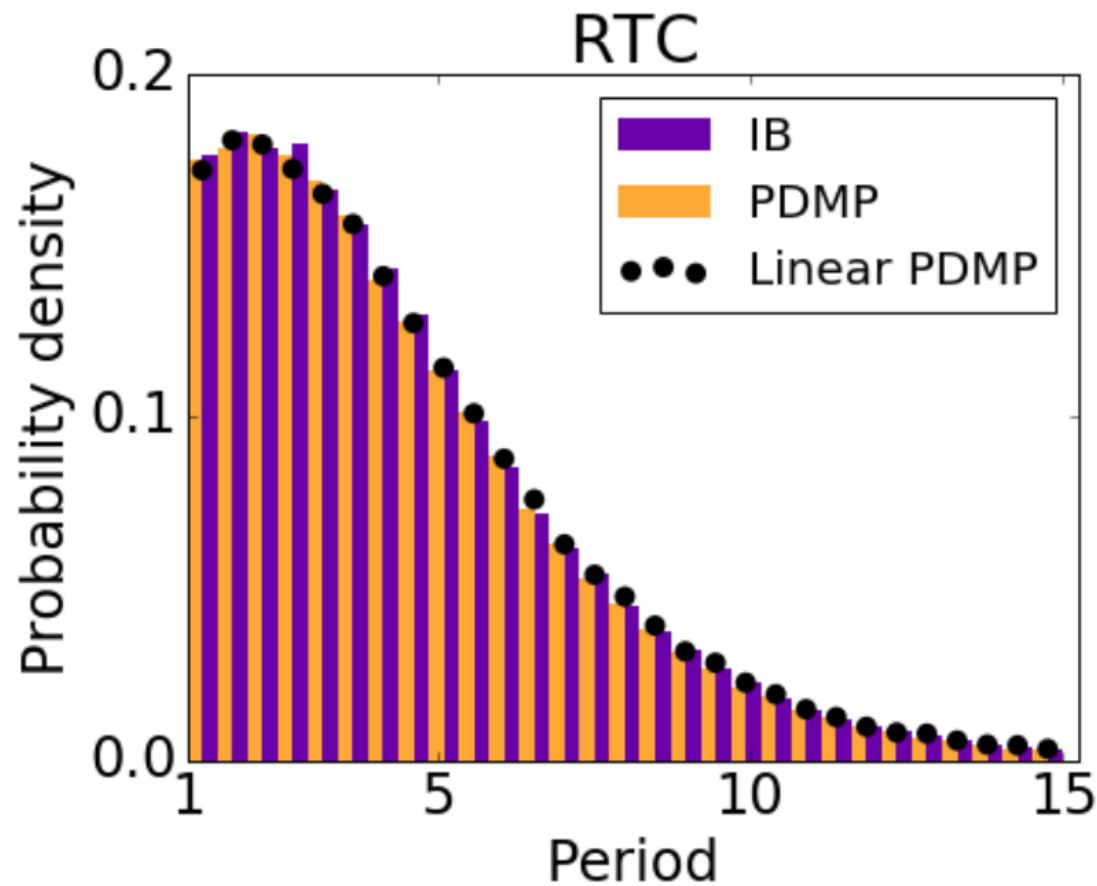


$$\mathbb{P}\{\{T_{\text{binding}} > t\}\} = \exp \left[-\frac{\kappa_Y}{\delta_X} \left(1 - e^{-\delta_X(t-t_0)}\right) \left(x(t_0) - \frac{\beta_X^f - \beta_Y^f}{\delta_X}\right) - \kappa_Y \frac{\beta_X^f - \beta_Y^f}{\delta_X} (t - t_0) \right].$$

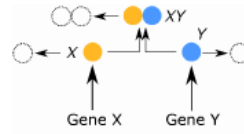
$$\begin{aligned} s_Y &= 0 \\ (x = 0, y > 0) \\ \dot{x} &= 0 \\ \dot{x} &= \beta_Y^f - \beta_X^f - \delta_Y y \end{aligned}$$

Deterministic Titration
 $\Delta t = \frac{1}{\delta_Y} \log \left(1 + \frac{y(t_0)}{\beta_X^f - \beta_Y^f} \right)$

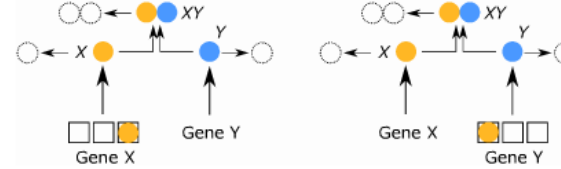
$$\begin{aligned} s_Y &= 0 \\ (x \geq 0, y = 0) \\ \dot{x} &= \beta_X^f - \beta_Y^f - \delta_X x \\ \dot{y} &= 0 \end{aligned}$$



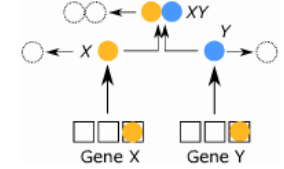
No feedback



One feedback



Two feedback



Deterministic LC
Excitation-relaxation
Spiral sink

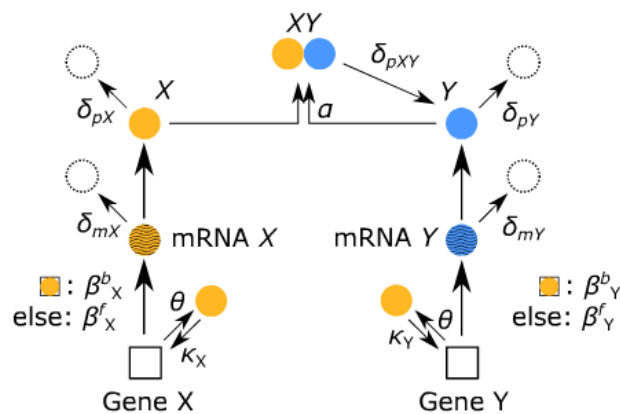


Demographic noise-induced
Burstiness induced
Slow-switching induced

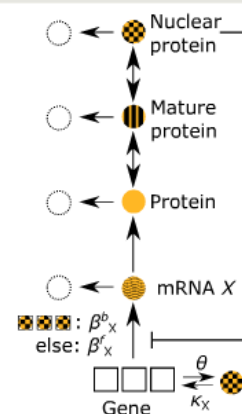


Classification of noise-induced oscillations in published models

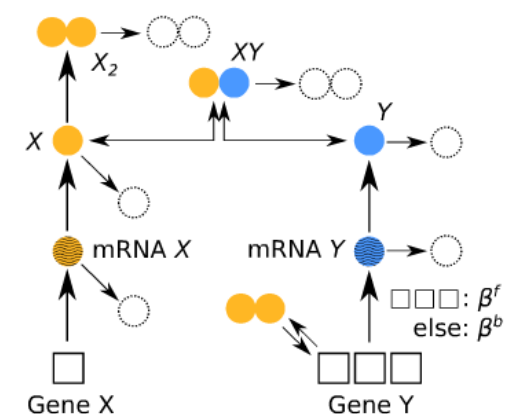
Vilar et al. PNAS 2002



Gonze et al. PNAS 2002
(+N>4 similar...)

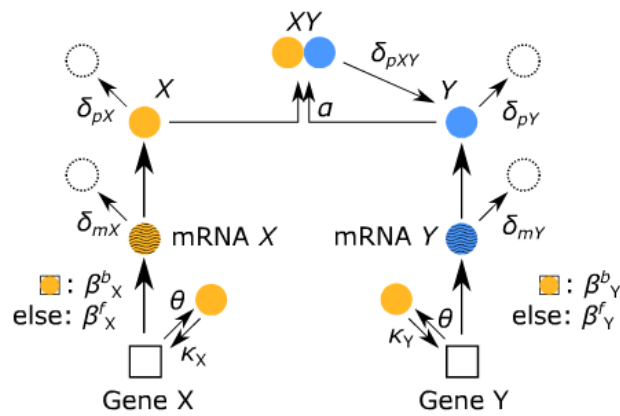


Karapetyan & Buchler, 2015

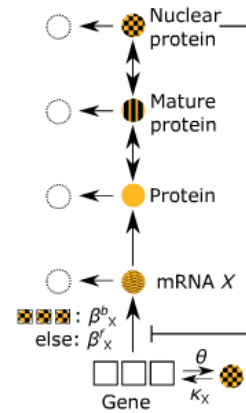


Classification of noise-induced oscillations in published models

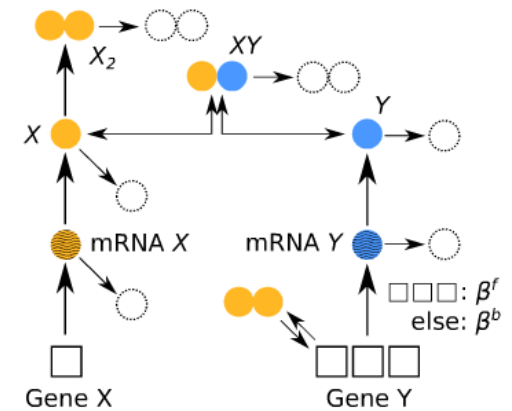
Vilar et al. PNAS 2002



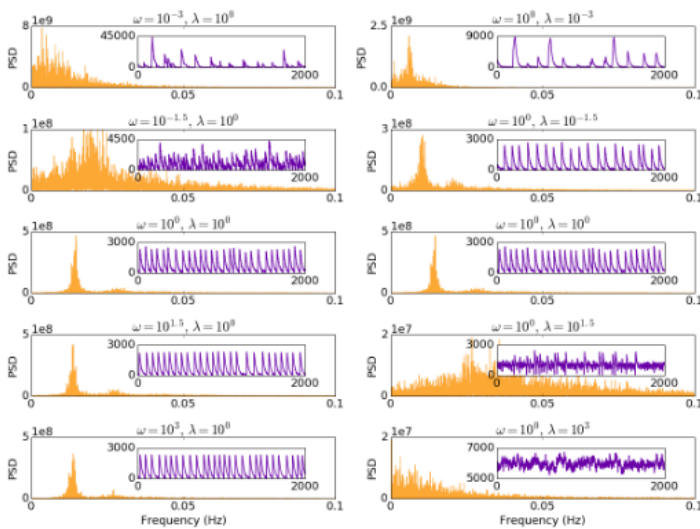
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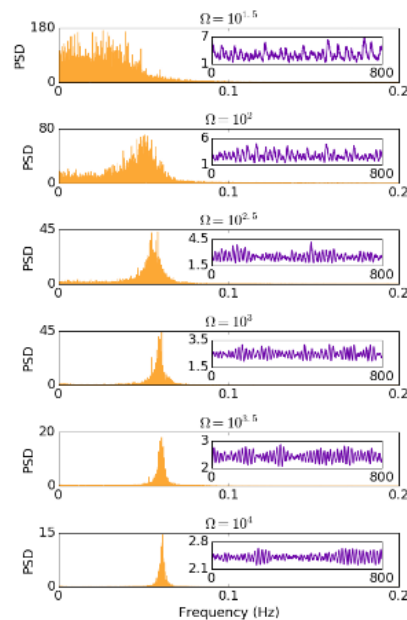
Karapetyan & Buchler, 2015



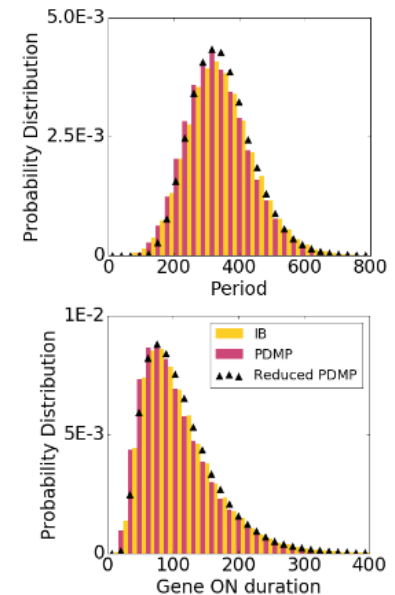
Excitation-relaxation and slow switching



Generic spiral sink



Slow switching



Summary

- We identified the minimal regulatory mechanisms which allows **deterministic** or **stochastic** oscillations
- We identified **different mechanisms** of noise-induced oscillations
- We provide mechanistic insights to explain noise-induced oscillations in the published models

Future directions

- Condense the material and write it up...
- Connections to the (abstract) stochastic phase oscillator models?
- WKB, quasi-potential, etc. for the excitable systems and comparison to generic stochastic broadening of the spiral sink?
- Noisy extrinsic signal (entrainment?)
- Some conclusion of the deterministic dynamics will not hold for systems with mRNA: is including mRNA a "correction", or we need more theory?

Mechanisms of Noise-induced Oscillation in Models of Biological Clock

SIAM Conference on Applications of Dynamical Systems
5/22/2017

Yen Ting Lin

Theoretical Division and Center for Nonlinear Studies
Los Alamos National Laboratory, New Mexico, USA

Collaborative work with Nicolas Buchler

Department of Physics, Dept. of Biology, Center for Genome & Computational Biology, Duke University, North Carolina, USA



Motivation

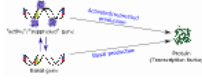
- Oscillatory dynamics are ubiquitous in biological systems: cell division cycle, metabolism, circadian rhythm, etc.
- Many computational models were proposed for specific systems.
- Noise is an important factor in the process.
- Dynamic noise: Temperature, light/dark cycle, heterogeneity of copy numbers of the molecules, etc.
- Intrinsic noise: discreteness of population, volume reduction, localization in space, etc.**
- There is an ongoing debate about if noise is "beneficial" for oscillations
- Prok. relab. noise sink and oscillator: The summer range can be vast in some models
- Copy number can compromise the conversion

Questions we ask...

- Can we propose a "simple-harmonic-oscillator"-like model to investigate these computational models, in a most simplified way to deliver generic conclusions?
- What is precisely the "noise-induced oscillator"? Are people comparing the same "animal"? We will show that there are multiple mechanisms to achieve this.
- How to analyze these mechanisms (what is the proper mathematical tool)? Can we generate hypothesis about regulating these stochastic oscillators with different mechanisms?

Prerequisite and the structure of the talk

Gene expression for each gene/TF pair: we adopted the minimal single-stage process



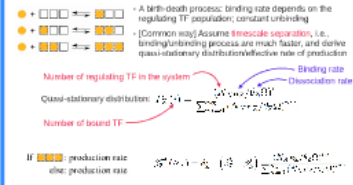
Scope of the models: Intrinsic-based oscillators: Two genes, two transcription factors. We will explore possible feedback mechanism until deterministic oscillation is possible.



We will investigate both the **deterministic** and **stochastic** models

Deterministic model: mass action kinetics

Discrete promoter site dynamics



A birth-death process: binding rate depends on the regulating TF population; constant unbinding
 (Common way) Assume timescale separation, i.e., binding/unbinding process are much faster, and derive quasi-stationary distribution/effective rate of production
 Number of regulating TF in the system
 Quasi-stationary distribution: $P(x) \propto \frac{1}{x!} \left(\frac{\lambda}{\mu}\right)^x$
 Number of bound TF
 If $\frac{d\langle x \rangle}{dt} > 0$ production rate else production rate

Deterministic model: mass action kinetics

Using the effective production rate, we arrived at the two-dimensional deterministic dynamics described by the ODE:

$$\dot{x}(t) = \beta_X^{eff}(\Omega x) - \delta_X x - \alpha_X y$$

$$\dot{y}(t) = \beta_Y^{eff}(\Omega x) - \delta_Y y - \alpha_Y$$

Brouwer's criterion stated that a limit cycle does not exist when

$$\partial_x \beta + \partial_y \alpha = -\delta_X - \delta_Y - \alpha \left(x + y + \Omega \frac{d\langle x \rangle}{dt} \right)$$

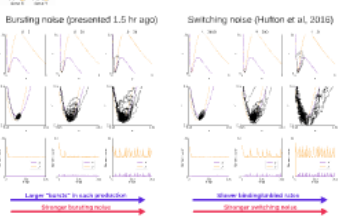
Does not change sign on a single connected domain (in our case, $\mathbb{R}^2 \times \mathbb{R}^+$). If a limit cycle exists, a necessary condition

$$\Rightarrow \frac{d\langle x \rangle}{dt} > 0 \Rightarrow X \text{ must be positively regulating itself.}$$



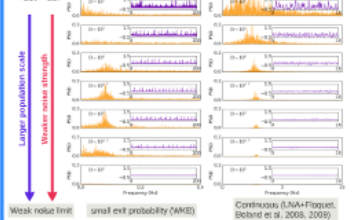
Individual-based model with two feedback

Two other sources of (intrinsic) noise to induce oscillation



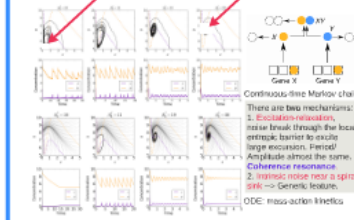
Individual-based model with two feedback

Response to various noise strength (population scale)



Individual-based model with two feedback

(well-mixed, single compartment)



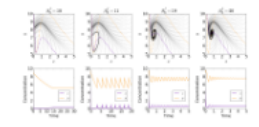
Deterministic model: mass action kinetics

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$$\dot{y}(t) = \beta_Y^{eff}(\Omega x) - \delta_Y y - \alpha_Y$$

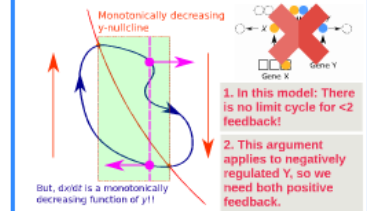
Necessary condition: $\frac{d\langle x \rangle}{dt} > 0$, and $\frac{d\langle y \rangle}{dt} > 0$.

We show that oscillation/limit cycle exists numerically:



Deterministic model: mass action kinetics

is autoregulated (self-regulating) X alone sufficient to generate oscillation?

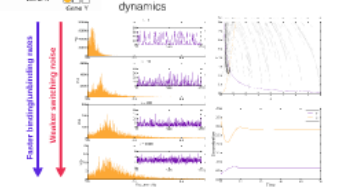


Individual-based model with one feedback

No limit cycle exists for deterministic model

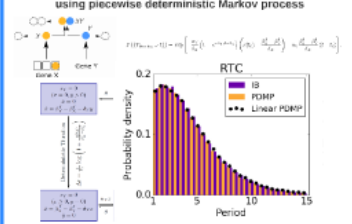
There can be spiral sink

Slow-switching can induce alternating dynamics

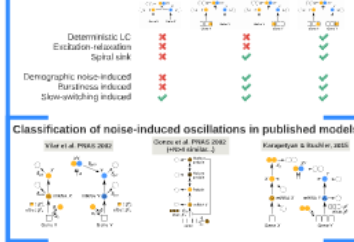


Individual-based model with one feedback

Quantifying the effect of slow-switching dynamics using piecewise deterministic Markov process



Classification of noise-induced oscillations in published models



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Very rough draft available;
if you are interested:
yentingl@lanl.gov

Thank you for your attention!