

# Large deviations for models in Systems Biology

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## Stochastic processes in systems biology

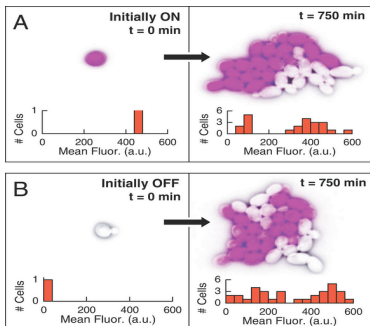
- chemical reaction networks describing basic cellular processes
- stochasticity arises from **molecular interactions and environmental noise**
- **different amounts of molecular abundances** and interaction rate magnitudes implies some stochastic features are essential and persistent in the long-term dynamics

## Rare events in systems biology

- ★ departure from typical behaviour ( $\pm$  st.dev. fluctuations)
- ★ in **bistable systems** leads to transitions to new stable state
- ★ deviations can arise from **intrinsic stochasticity** of the system (rather than from external perturbations)

## Bistable examples in systems biology

- ★ competing positive-negative feedback, enzymatic futile cycle, de/phosphorylation cycle, bistable repressible switch



**yeast cells switch between expressing and nonexpressing states**  
(Kaufmann, Yang, Mettetal, and van Oudenaarden, *PLOS Bio.* 2007)

- **cell growth+division**: rare events have many trials to occur  
not rare on the level of the cell population

## Stochastic models for reaction network system

- continuous-time Markov chains and their rescaled versions
- multi-scale properties can lead to various limiting processes: jump diffusions with state-dependent coefficients

$$X_t = X_0 + \int_0^t \mu(X_s) ds + \int_0^t \sigma(X_s) dW_s + \sum_{s \leq t} \Delta X_s$$

jumps have rate  $\lambda(X)$  and jump measure  $\int |y| \nu(X, dy) < \infty$

- reflection of  $X_s \in [0, b]$  at the boundaries  $0, b \leq \infty$

$$V_t = V_0 + X_t + L_t - U_t$$

$L_t$  and  $U_t$  are local times at the lower and upper boundary

- intrinsic constraints in reaction systems give processes with reflection  
(Leite and Williams, [www.math.ucsd.edu](http://www.math.ucsd.edu) 2017)

## Examples

► **self-regulated gene expression**

w/ **protein bursts** (due to protein translation+transcription):

$$X_t = X_0 + \int_0^t \left( \frac{c_0^\pm + c_{0*}^\pm X_s}{c_1^\pm + c_2^\pm X_s} - c_3 X_s \right) ds \\ + \epsilon \int_0^t \left( \frac{c_0^\pm + c_{0*}^\pm X_s}{c_1^\pm + c_2^\pm X_s} \left[ \frac{c_4^\pm + c_{4*}^\pm X_s}{(c_1^\pm + c_2^\pm X_s)^2} + c_5^\pm \right] + c_3 X_s \right) dW_s + \delta \sum_{s \leq t} \Delta X_s$$

► **enzymatic (Michaelis-Menten) kinetics**

w/ **substrate bursts** (due to cell division):

$$X_t = X_0 - \int_0^t \frac{X_s}{c_1 + c_2 X_s} ds + \epsilon \int_0^t \frac{c_3 X_s}{(c_4 + c_5 X_s)^2} dW_s + \delta \sum_{s \leq t} \Delta X_s$$

## Short-term behaviour

on a finite time interval  $t \in [0, T]$

when noise contribution  $(\epsilon, \delta)$  is small

- ★ **FLLN**:  $(X_t)_{[0, T]} \rightarrow (x_t)_{[0, T]}$  with  $dx_t = \mu(x_t)dt$
- ★ **FCLT**:  $\frac{1}{\epsilon}(X_t - x_t)_{[0, T]} \Rightarrow (U_t)_{[0, T]}$  with  $U$  Gaussian
- ★ **PLDP**:  $\mathbf{P}\left(\sup_{t \in [0, T]} |X_t - \tilde{x}_t| \leq \epsilon\right) \approx e^{-\epsilon I((\tilde{x}_t)_{[0, T]})}$

(Freidlin and Wentzell; Anderson and Orey - for reflected processes)

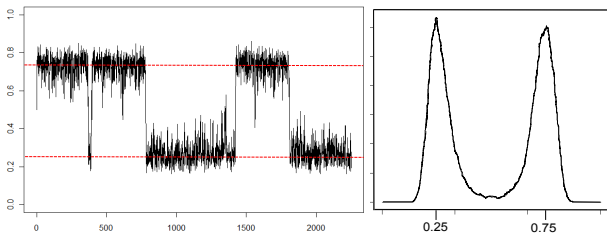
calculating the rate function  $I(\tilde{x})$

- from limiting non-linear exponential operator:

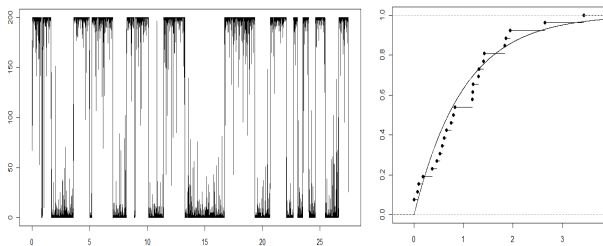
$$Hf = \lim_{\epsilon \rightarrow 0} \log \mathbf{E}\left[e^{\frac{1}{\epsilon} f((X_t)_{[0, T]})}\right], \quad I(\tilde{x}) = \sup_{f \in C_b([0, T])} [f(\tilde{x}) - Hf]$$

- operator  $H$  has additional term on the boundary for reflection

(Feng and Kurtz 'Large Deviations for Stochastic Processes' 2006)



**sample path and occupation measure for  $\delta \ll \epsilon^2$**



**sample path and switching times for  $\delta \gg \epsilon^2$  (same system)**

*(McSweeney and Popovic, Ann.Appl.Prob. 2014)*

## Long-term behaviour

on long time intervals  $t \rightarrow \infty$

with **sizeable contribution** from diffusion and jumps

- ★ assuming  $V_t$  is **ergodic**, stationary measure  $\pi$ , gives average:

$$\frac{1}{t} \int_0^t f(V_s) ds \xrightarrow{t \rightarrow \infty} \mathbf{E}_\pi[f(x)]$$

- ★ fluctuations for  $f$  with  $\mathbf{E}_\pi[f(x)] = 0$  can be calculated using solution  $g$  of the **Poisson equation**:

$$f(x) = \mu(x)g'(x) + \frac{1}{2}\sigma^2(x)g''(x) + \lambda(x) \int (g(\delta(x, y)) - g(x))\nu(x, dy), \quad g'(0) = g'(b) = 0$$

where  $V_s = \delta(V_{s-}, \Delta X_s)$  if  $\Delta X \neq 0$

for  $\delta(x, y) = 0 \mathbf{1}_{x+y \leq 0} + (x+y) \mathbf{1}_{x+y \in (0, b)} + b \mathbf{1}_{x+y \geq b}$



## Generalized occupation time

- consider the **Additive process** - generalization of local time

$$\Lambda_t = \int_0^t f(V_s) ds + \sum_{s \leq t} \tilde{f}(V_{s-}, \Delta X_s) + f_0 L_t^c + f_b U_t^c$$

$f$  bdd,  $\tilde{f}(x, 0) = 0$ ,  $\int e^{\theta \tilde{f}(x, y)} \nu(x, dy) < C \forall x$ ,  $f_0, f_b \in \mathbb{R}_+$

$L^c, U^c$  are the continuous parts of local times at 0,  $b$

- e.g.  $\Lambda_t = L_t$  with  $f = 0$ ,  $\tilde{f} = -[x + y]^-, f_0 = 1, f_b = 0$

this gives the **fraction of time** a protein is **not expressed in**, or fraction of time enzymatic substrate is at **extremely low level**

## Gaertner-Ellis theorem

for stochastic process on  $\mathbb{R}^d$

- ▶ Suppose

$$\frac{1}{t} \mathbf{E}[e^{\theta \Lambda_t}] \xrightarrow{t \rightarrow \infty} \psi(\theta)$$

exists for  $\psi$  such that

- ▶  $0 \in \text{int}(D_\psi)$ ,  $D_\psi = \{\theta : \psi(\theta) < \infty\}$
- ▶  $\psi$  is lower semi-convex, differentiable on  $\text{int}(D_\psi)$
- ▶  $D_\psi = \mathbb{R}$  or  $\lim_{\theta \rightarrow \partial D_\psi} |\psi'(\theta)| = \infty$

Then  $\frac{\Lambda_t}{t}$  satisfies the **Large Deviation Principle**  
with 'speed'  $t$  and 'good' convex **rate function**

$$\psi^*(a) = \sup_{\theta \in \mathbb{R}} [a\theta - \psi(\theta)]$$

## Integro-differential equation

each fixed  $\theta \rightarrow$  pair  $(u_\theta(x), \psi_\theta)$

- ▶ Suppose there exist  $u_\theta(x) \in \mathcal{C}_{\geq 0}^2$  and  $\psi_\theta \in \mathbb{R}$  satisfying the **integro-differential equation**

$$0 = \mu(x)u'_\theta(x) + \frac{1}{2}\sigma^2(x)u''_\theta(x) + (\theta f(x) - \psi_\theta)u_\theta(x) \\ + \lambda(x) \int (e^{\theta \tilde{f}(x,y)} u(\theta, \delta(x,y)) - u_\theta(x)) \nu(x, dy)$$

subject to **boundary conditions**

$$u_\theta(0) = 1, u'_\theta(0) = -f_0\theta, u_\theta(b) = 1, u'_\theta(b) = f_b\theta$$

- ★ the integral term makes it somewhat more difficult to **solve** the differential equation boundary problem **numerically**

## Martingale argument

- ▶ Then (under assumptions on  $\mu, \sigma, \lambda, \nu$ )

$$M_t(\theta) = e^{\theta \Lambda_t} u_\theta(V_t)$$

is a **martingale** ( $u_\theta$  is bounded) and for each  $\theta$

$$\frac{1}{t} \log \mathbf{E}[e^{\theta \Lambda_t}] \xrightarrow[t \rightarrow \infty]{} \psi_\theta$$

- ▶ Assuming  $\psi(\theta) := (\psi_\theta)_{\theta \in \mathcal{D}_\psi}$  satisfies **Gaertner-Ellis theorem**

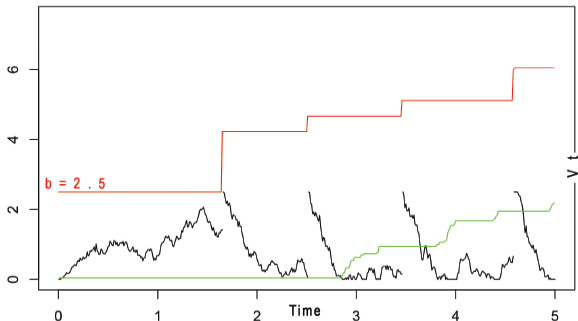
$$\frac{1}{t} \log \mathbf{P}[\Lambda_t \geq at] \xrightarrow[t \rightarrow \infty]{} \psi(\theta^*) - \theta^* a,$$

for any  $\theta^* > 0$  such that  $\psi \in C^1$  around  $\theta^*$ ,  $a = \partial_\theta \psi(\theta^*)$

- ★ **numerically** solving for  $\psi \Rightarrow$  ‘verifying’ needed assumptions only numerically  $\Rightarrow$  estimating **probabilities of rare events**

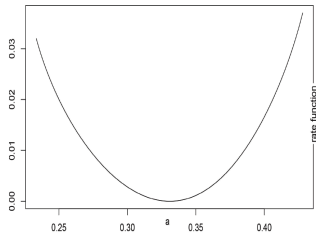
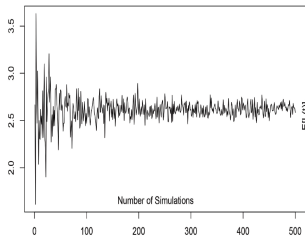
## Simulations

- ▶  $V_t$  is jump-diffusion reflected at: 0 and  $b = 2.5$  with drift and diffusion:  $\mu(x) = 2.62(1.61 - x)$ ,  $\sigma(x) = 0.62\sqrt{x}$   
jump rate and measure:  $\lambda(x) = x$ ,  $\nu(x, dy) = \frac{1}{2}\mathbf{1}_{0.72} + \frac{1}{2}\mathbf{1}_{2.87}$



$U_t$  = local time at  $b$  and  $L_t$  = local time at 0

►  $\Lambda_t = L_t$  is local time at 0



$$\frac{1}{t} \mathbf{E}[L_t] \sim \min_a \psi^*(a)$$

LDP rate function  $\psi^*$

| $t = 1000$ |                |        |             |                        |
|------------|----------------|--------|-------------|------------------------|
| $\theta$   | $\psi(\theta)$ | $a$    | $\psi^*(a)$ | $P(\Lambda_t \geq at)$ |
| 0          | 0              | 0.3401 | 0           | 1                      |
| 0.1        | 0.0346         | 0.3555 | 0.0008      | 0.4493                 |
| 0.2        | 0.0708         | 0.3653 | 0.0023      | 0.1003                 |
| 0.3        | 0.1075         | 0.3697 | 0.0034      | 0.0334                 |
| 0.4        | 0.1446         | 0.3714 | 0.0040      | 0.0183                 |
| 0.5        | 0.1818         | 0.3721 | 0.0043      | 0.0136                 |
| 0.6        | 0.2190         | 0.3724 | 0.0044      | 0.0123                 |
| 0.7        | 0.2563         | 0.3725 | 0.0045      | 0.0111                 |
| 0.8        | 0.2935         | 0.3726 | 0.0045      | 0.0111                 |
| 0.9        | 0.3308         | 0.3726 | 0.0046      | 0.0101                 |
| 1          | 0.3680         | 0.3726 | 0.0046      | 0.0101                 |