

## Averaging fast subsystems in chemical network models

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- Enzyme reaction
- Michaelis-Menten equation
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See [References](#), in particular [Anderson and Kurtz \(2015\)](#).



### Markov chains

Infinitesimal specification of a Markov chain:

$$P\{X(t + \Delta t) - X(t) = \zeta_k | \mathcal{F}_t^X\} \approx \beta_k(X(t))\Delta t$$

Generator:

$$Af(x) = \sum_k \beta_k(x)(f(x + \zeta_k) - f(x))$$

(For simplicity assume only finitely many possible jumps  $\zeta_k$ .)

$$X(t) = X(0) + \sum_k R_k(t)\zeta_k,$$

$R_k$  a counting process counting the times  $X$  takes jump  $\zeta_k$ .

$$P\{R_k(t + \Delta t) - R_k(t) = 1 | \mathcal{F}_t^X\} \approx \beta_k(X(t))\Delta t.$$

For independent unit Poisson processes  $Y_k$ , we should have

$$X(t) = X(0) + \sum_k Y_k \left( \int_0^t \beta_k(X(s)) ds \right) \zeta_k$$



## Chemical networks

Modeling counts of individual chemical species:  $X(t)$  will be the vector whose components give the numbers of molecules of each chemical species.

For a binary reaction  $S_1 + S_2 \rightarrow S_3$  or  $S_1 + S_2 \rightarrow S_3 + S_4$

$$\lambda_k(x) = \kappa'_k x_1 x_2 \quad \zeta_k = \nu'_k - \nu_k = \begin{pmatrix} 0 \\ 0 \\ 1 \\ 0 \end{pmatrix} - \begin{pmatrix} 1 \\ 1 \\ 0 \\ 0 \end{pmatrix} = \begin{pmatrix} -1 \\ -1 \\ 1 \\ 0 \end{pmatrix} \text{ or } \begin{pmatrix} -1 \\ -1 \\ 1 \\ 1 \end{pmatrix}$$

where  $\nu_k$  is the vector giving the numbers of molecules consumed and  $\nu'_k$  gives the numbers produced. For  $S_1 \rightarrow S_2$  or  $S_1 \rightarrow S_2 + S_3$ ,

$$\lambda_k(x) = \kappa'_k x_1.$$

For  $2S_1 \rightarrow S_2$ ,

$$\lambda_k(x) = \kappa'_k x_1(x_1 - 1) \quad \zeta_k = \begin{pmatrix} -2 \\ 1 \end{pmatrix}$$



## Enzyme reaction



$$X_E(t) = X_E(0) - Y_1 \left( \int_0^t \kappa_1 X_A(s) X_E(s) ds \right) + Y_2 \left( \kappa_2 \int_0^t X_{AE}(s) ds \right) \\ + Y_3 \left( \kappa_3 \int_0^t X_{AE}(s) ds \right)$$

$$X_A(t) = X_A(0) - Y_1 \left( \int_0^t \kappa_1 X_A(s) X_E(s) ds \right) + Y_2 \left( \kappa_2 \int_0^t X_{AE}(s) ds \right)$$

The remainder of the system is determined by

$$X_{AE}(t) = M - X_E(t) \quad X_B(t) = X_B(0) + X_A(0) + X_{AE}(0) - X_A(t) - X_{AE}(t)$$

$$\text{Generator: } \zeta_1 = \begin{pmatrix} -1 \\ -1 \end{pmatrix}, \zeta_2 = \begin{pmatrix} 1 \\ 1 \end{pmatrix}, \zeta_3 = \begin{pmatrix} 1 \\ 0 \end{pmatrix}$$

$$Af(x) = \kappa_1 x_A x_E (f(x + \zeta_1) - f(x)) + \kappa_2 (M - x_E) (f(x + \zeta_2) - f(x)) \\ + \kappa_3 (M - x_E) (f(x + \zeta_3) - f(x))$$



Fix  $N_0 \gg 1$ . For each species  $i$ , define the *normalized abundances* (or simply, the abundances) by

$$Z_i(t) = N_0^{-\alpha_i} X_i(t),$$

where  $\alpha_i \geq 0$  should be selected so that  $Z_i = O(1)$ . Note that the abundance may be the species number ( $\alpha_i = 0$ ) or the species concentration or something else.

The rate constants may also vary over several orders of magnitude so scale the rate constants  $\kappa'_k = \kappa_k N_0^{\beta_k}$  so that  $\kappa_k = O(1)$ .

Then

$$\begin{aligned} \kappa'_k x_i x_j &= N_0^{\beta_k + \alpha_i + \alpha_j} \kappa_k z_i z_j & \kappa'_k x_i (x_i - 1) &= N_0^{\beta_k + 2\alpha_i} \kappa_k z_i (z_i - N_0^{-\alpha_i}) \\ \kappa'_k x_i &= \kappa_k N_0^{\beta_k + \alpha_i} z_i \end{aligned}$$

Note that the exponent on  $N_0$  is  $\rho_k = \beta_k + \alpha \cdot \nu_k$ .



## A parameterized family of models

Then, noting that  $\nu_k \cdot \alpha = \sum_i \nu_{ik} \alpha_i$ ,

$$Z_i(t) = Z_i(0) + \sum_k N_0^{-\alpha_i} Y_k \left( \int_0^t N_0^{\beta_k + \nu_k \cdot \alpha} \lambda_k(Z(s)) ds \right) (\nu'_{ik} - \nu_{ik}).$$

Let

$$Z_i^N(t) = Z_i(0) + \sum_k N^{-\alpha_i} Y_k \left( \int_0^t N^{\beta_k + \nu_k \cdot \alpha} \lambda_k(Z^N(s)) ds \right) (\nu'_{ik} - \nu_{ik}).$$

Then the “true” model is  $Z = Z^{N_0}$ .



Let

$$\begin{aligned} Z_i^{N,\gamma}(t) &\equiv Z_i^N(tN^\gamma) \\ &= Z_i(0) + \sum_k N^{-\alpha_i} Y_k \left( \int_0^t N^{\gamma+\beta_k+\nu_k \cdot \alpha} \lambda_k(Z^{N,\gamma}(s)) ds \right) \zeta_{ik}. \end{aligned}$$

Equation is “balanced” if

$$\max\{\beta_k + \nu_k \cdot \alpha : \zeta_{ik} > 0\} = \max\{\beta_k + \nu_k \cdot \alpha : \zeta_{ik} < 0\}$$

If the equation is not balanced then we need

$$\gamma + \beta_k + \nu_k \cdot \alpha \leq \alpha_i \tag{1}$$

for all  $i$  and  $k$  such that  $\zeta_{ik} \neq 0$ .

The time-scale of species  $i$ :  $\gamma_i = \alpha_i - \max\{\beta_k + \nu_k \cdot \alpha : \zeta_{ik} \neq 0\}$

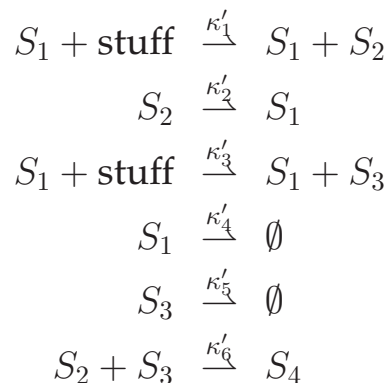


## Example: Model of a viral infection

Srivastava, You, Summers, and Yin (2002), Haseltine and Rawlings (2002), Ball, Kurtz, Popovic, and Rempala (2006)

Three time-varying species, the viral template, the viral genome, and the viral structural protein (indexed, 1, 2, 3 respectively).

The model involves six reactions,



## Stochastic system

$$X_1(t) = X_1(0) + Y_2\left(\int_0^t \kappa'_2 X_2(s) ds\right) - Y_4\left(\int_0^t \kappa'_4 X_1(s) ds\right)$$

$$X_2(t) = X_2(0) + Y_1\left(\int_0^t \kappa'_1 X_1(s) ds\right) - Y_2\left(\int_0^t \kappa'_2 X_2(s) ds\right) - Y_6\left(\int_0^t \kappa'_6 X_2(s) X_3(s) ds\right)$$

$$X_3(t) = X_3(0) + Y_3\left(\int_0^t \kappa'_3 X_1(s) ds\right) - Y_5\left(\int_0^t \kappa'_5 X_3(s) ds\right) - Y_6\left(\int_0^t \kappa'_6 X_2(s) X_3(s) ds\right)$$

$\kappa'_1$	1	$\kappa'_4$	0.25
$\kappa'_2$	0.025	$\kappa'_5$	2
$\kappa'_3$	1000	$\kappa'_6$	$7.5 \times 10^{-6}$

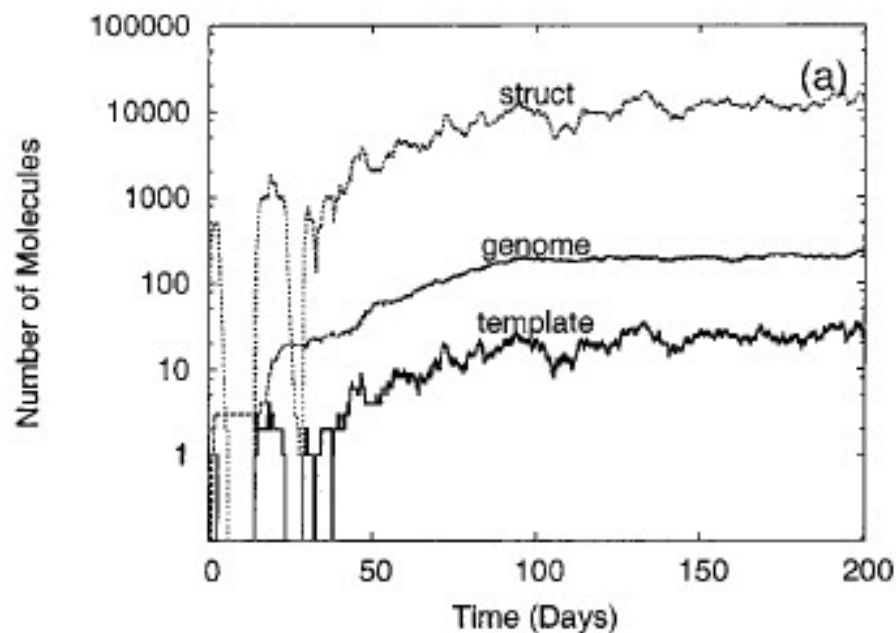


Figure 1: Simulation (Haseltine and Rawlings 2002)



## Balance equations for the viral model

$$Z_1^N(t) = Z_1(0) + N^{-\alpha_1} Y_2 \left( \int_0^t \kappa_2 N^{\beta_2 + \alpha_2} Z_2^N(s) ds \right) - N^{-\alpha_1} Y_4 \left( \int_0^t \kappa_4 N^{\beta_4 + \alpha_1} Z_1^N(s) ds \right)$$

$$Z_2^N(t) = Z_2(0) + N^{-\alpha_2} Y_1 \left( \int_0^t \kappa_1 N^{\beta_1 + \alpha_1} Z_1^N(s) ds \right) - N^{-\alpha_2} Y_2 \left( \int_0^t \kappa_2 N^{\beta_2 + \alpha_2} Z_2^N(s) ds \right) \\ - N^{-\alpha_2} Y_6 \left( \int_0^t \kappa_6 N^{\beta_6 + \alpha_2 + \alpha_3} Z_2^N(s) Z_3^N(s) ds \right)$$

$$Z_3^N(t) = Z_3(0) + N^{-\alpha_3} Y_3 \left( \int_0^t \kappa_3 N^{\beta_3 + \alpha_1} Z_1^N(s) ds \right) - N^{-\alpha_3} Y_5 \left( \int_0^t \kappa_5 N^{\beta_5 + \alpha_3} Z_3^N(s) ds \right) \\ - N^{-\alpha_3} Y_6 \left( \int_0^t \kappa_6 N^{\beta_6 + \alpha_2 + \alpha_3} Z_2^N(s) Z_3^N(s) ds \right)$$

$$\beta_2 + \alpha_2 = \beta_4 + \alpha_1$$

$$\beta_1 + \alpha_1 = (\beta_2 + \alpha_2) \vee (\beta_6 + \alpha_2 + \alpha_3)$$

$$\beta_3 + \alpha_1 = (\beta_5 + \alpha_3) \vee (\beta_6 + \alpha_2 + \alpha_3)$$

$$\beta_3 \geq \beta_5 \geq \beta_1 \geq \beta_4 \geq \beta_2 \geq \beta_6$$



## An example

$$\beta_2 + \alpha_2 = \beta_4 + \alpha_1$$

$$\beta_1 + \alpha_1 = (\beta_2 + \alpha_2) \vee (\beta_6 + \alpha_2 + \alpha_3)$$

$$\beta_3 + \alpha_1 = (\beta_5 + \alpha_3) \vee (\beta_6 + \alpha_2 + \alpha_3)$$

$$\beta_3 \geq \beta_5 \geq \beta_1 \geq \beta_4 \geq \beta_2 \geq \beta_6$$

$\beta_1$	0	$\alpha_1$	0
$\beta_2$	$-\frac{2}{3}$	$\alpha_2$	$\frac{2}{3}$
$\beta_3$	1	$\alpha_3$	1
$\beta_4$	0	$\gamma_1$	0
$\beta_5$	0	$\gamma_2$	$\frac{2}{3}$
$\beta_6$	$-\frac{5}{3}$	$\gamma_3$	0



## Scaling parameters Ball, Kurtz, Popovic, and Rempala (2006)

Each  $X_i$  is scaled according to its abundance in the system.

For  $N_0 = 1000$ ,  $X_1 = O(N_0^0)$ ,  $X_2 = O(N_0^{2/3})$ , and  $X_3 = O(N_0)$  and we take  $Z_1 = X_1$ ,  $Z_2 = X_2 N_0^{-2/3}$ , and  $Z_3 = X_3 N_0^{-1}$ .

Expressing the rate constants in terms of  $N_0 = 1000$

$\kappa'_1$	1	1
$\kappa'_2$	0.025	$2.5N_0^{-2/3}$
$\kappa'_3$	1000	$N_0$
$\kappa'_4$	0.25	.25
$\kappa'_5$	2	2
$\kappa'_6$	$7.5 \times 10^{-6}$	$.75N_0^{-5/3}$



## Normalized system

With the scaled rate constants and abundances, we have

$$Z_1^N(t) = Z_1^N(0) + Y_2 \left( \int_0^t 2.5 Z_2^N(s) ds \right) - Y_4 \left( \int_0^t .25 Z_1^N(s) ds \right)$$

$$\begin{aligned}
 Z_2^N(t) = Z_2^N(0) + N^{-2/3} Y_1 \left( \int_0^t Z_1^N(s) ds \right) - N^{-2/3} Y_2 \left( \int_0^t 2.5 Z_2^N(s) ds \right) \\
 - N^{-2/3} Y_6 \left( \int_0^t .75 Z_2^N(s) Z_3^N(s) ds \right)
 \end{aligned}$$

$$\begin{aligned}
 Z_3^N(t) = Z_3^N(0) + N^{-1} Y_3 \left( \int_0^t N Z_1^N(s) ds \right) - N^{-1} Y_5 \left( \int_0^t 2N Z_3^N(s) ds \right) \\
 - N^{-1} Y_6 \left( \int_0^t .75 Z_2^N(s) Z_3^N(s) ds \right),
 \end{aligned}$$



## Limiting system for $\gamma = 0$

Passing to the limit, we have

$$Z_1(t) = Z_1(0) + Y_2\left(\int_0^t 2.5Z_2(s)ds\right) - Y_4\left(\int_0^t .25Z_1(s)ds\right)$$

$$Z_2(t) = Z_2(0)$$

$$Z_3(t) = Z_3(0) + \int_0^t Z_1(s)ds - \int_0^t 2Z_3(s)ds$$



## Fast time-scale

Define  $Z_i^{N,\gamma}(t) = Z_i^N(N^\gamma t)$ . For  $\gamma = \frac{2}{3}$ ,

$$Z_1^{N,\gamma}(t) = Z_1(0) + Y_2\left(\int_0^t 2.5N^{2/3}Z_2^{N,\gamma}(s)ds\right) - Y_4\left(\int_0^t .25N^{2/3}Z_1^{N,\gamma}(s)ds\right)$$

$$Z_2^{N,\gamma}(t) = Z_2(0) + N^{-2/3}Y_1\left(\int_0^t N^{2/3}Z_1^{N,\gamma}(s)ds\right)$$

$$-N^{-2/3}Y_2\left(\int_0^t 2.5N^{2/3}Z_2^{N,\gamma}(s)ds\right)$$

$$-N^{-2/3}Y_6\left(N^{2/3}\int_0^t .75Z_2^{N,\gamma}(s)Z_3^{N,\gamma}(s)ds\right)$$

$$Z_3^{N,\gamma}(t) = Z_3(0) + N^{-1}Y_3\left(\int_0^t N^{5/3}Z_1^{N,\gamma}(s)ds\right) - N^{-1}Y_5\left(\int_0^t 2N^{5/3}Z_3^{N,\gamma}(s)ds\right)$$

$$-N^{-1}Y_6\left(\int_0^t .75N^{2/3}Z_2^{N,\gamma}(s)Z_3^{N,\gamma}(s)ds\right)$$





## Averaging

As  $N \rightarrow \infty$ , dividing the equations for  $Z_1^{N,\gamma}$  and  $Z_3^{N,\gamma}$  by  $N^{2/3}$  shows that

$$\begin{aligned}\int_0^t Z_1^{N,\gamma}(s) ds - 10 \int_0^t Z_2^{N,\gamma}(s) ds &\rightarrow 0 \\ \int_0^t Z_3^{N,\gamma}(s) ds - 5 \int_0^t Z_2^{N,\gamma}(s) ds &\rightarrow 0.\end{aligned}$$

The assertion for  $Z_3^{N,\gamma}$  and the fact that  $Z_2^{N,\gamma}$  is asymptotically regular imply

$$\int_0^t Z_2^{N,\gamma}(s) Z_3^{N,\gamma}(s) ds - 5 \int_0^t Z_2^{N,\gamma}(s)^2 ds \rightarrow 0.$$

It follows that  $Z_2^{N,\gamma}$  converges to the solution of (2).



## Law of large numbers

**Theorem 1** Let  $\gamma = \frac{2}{3}$ . For each  $\delta > 0$  and  $t > 0$ ,

$$\lim_{N \rightarrow \infty} P\left\{ \sup_{0 \leq s \leq t} |Z_2^{N,\gamma}(s) - Z_2^{\infty,\gamma}(s)| \geq \delta \right\} = 0,$$

where  $Z_2^{\infty,\gamma}$  is the solution of

$$Z_2^{\infty,\gamma}(t) = Z_2(0) + \int_0^t 7.5 Z_2^{\infty,\gamma}(s) ds - \int_0^t 3.75 Z_2^{\infty,\gamma}(s)^2 ds. \quad (2)$$



# Martingale approach to averaging

Kurtz (1992)

Kang and Kurtz (2013)

Let  $e_k$  be the vector in  $\mathbb{Z}^3$  with  $k$ th component 1 and other components zero. For  $\gamma = 2/3$ , the generator is

$$\begin{aligned} A^N f(z) = & N^{2/3} z_1 (f(z + N^{-2/3} e_2) - f(z)) \\ & + 2.5 N^{2/3} z_2 (f(z + e_1 - N^{-2/3} e_2) - f(z)) + N^{5/3} z_1 (f(z + N^{-1} e_3) - f(z)) \\ & + .25 N^{2/3} z_1 (f(z - e_1) - f(z)) + 2 N^{5/3} z_3 (f(z - N^{-1} e_3) - f(z)) \\ & + .75 N^{2/3} z_2 z_3 (f(z - N^{-1} e_3 - N^{-2/3} e_2) - f(z)) \end{aligned}$$

If  $f$  depends only on  $z_2$ , then

$$\lim_{N \rightarrow \infty} A^N f(z) = (z_1 - 2.5 z_2 - .75 z_2 z_3) f'(z_2).$$

If  $f$  depends only on  $z_1$  and  $z_3$ , then

$$\begin{aligned} \lim_{N \rightarrow \infty} N^{-2/3} A^N f(z) = & 2.5 z_2 (f(z_1 + 1, z_3) - f(z_1, z_3)) \\ & + .25 z_1 (f(z_1 - 1, z_3) - f(z_1, z_3)) \\ & + (z_1 - 2 z_3) \partial_{z_3} f(z_1, z_3) \end{aligned}$$



## Limiting martingale

$$\begin{aligned} & f(Z_2^N(t)) - \int_0^t (Z_1^N(s) - 2.5 Z_2^N(s) - .75 Z_2^N(s) Z_3^N(s)) f'(Z_2^N(s)) ds + \epsilon_N(t) \\ & = f(Z_2^N(t)) - \int_{[0,t] \times [0,\infty)^2} (z_1 - 2.5 Z_2^N(s) - .75 Z_2^N(s) z_3) f'(Z_2^N(s)) \Gamma^N(ds, dz_1, dz_3) + \epsilon_N(t) \\ & \Rightarrow f(Z_2(t)) - \int_0^t \int_{[0,\infty)^2} (z_1 - 2.5 Z_2(s) - .75 Z_2(s) z_3) f'(Z_2(s)) \Gamma_s(dz_1, dz_3) ds \end{aligned}$$

Need to check relative compactness of  $\{\Gamma^N\}$ .

**Exercise:** Bound the expectations of  $Z_1^N, Z_2^N, Z_3^N$  by taking expectations of the stochastic equations and throwing away the nonlinear terms.

Need to identify the limit  $\Gamma^N \Rightarrow \Gamma$ .



## Identification of $\Gamma$

For a function of  $z_1, z_2$

$$\begin{aligned}
 & N^{-2/3} f(Z_1^N(t), Z_3^N(t)) - \int_0^t N^{-2/3} A^N f(Z^N(s)) ds \\
 & \approx \int_{[0,t] \times [0,\infty)^2} \left[ 2.5 Z_2^N(s) (f(z_1 + 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + .25 z_1 (f(z_1 - 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + (z_1 - 2z_3) \partial_{z_3} f(z_1, z_3) \right] \Gamma^N(ds, dz_1, dz_3) \\
 & \Rightarrow \int_0^t \int_{[0,\infty)^2} \left[ 2.5 Z_2(s) (f(z_1 + 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + .25 z_1 (f(z_1 - 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + (z_1 - 2z_3) \partial_{z_3} f(z_1, z_3) \right] \Gamma_s(dz_1, dz_3) ds
 \end{aligned}$$

A martingale that is continuous and finite variation is a constant. Consequently, the last expression is zero.



## $\Gamma_s$ is a “stationary distribution”

Almost surely, for almost every  $s$

$$\begin{aligned}
 & \int_{[0,\infty)^2} \left[ 2.5 Z_2(s) (f(z_1 + 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + .25 z_1 (f(z_1 - 1, z_3) - f(z_1, z_3)) \right. \\
 & \quad \left. + (z_1 - 2z_3) \partial_{z_3} f(z_1, z_3) \right] \Gamma_s(dz_1, dz_3) = 0
 \end{aligned}$$



## Approximate models

We have a family of models indexed by  $N$  for which  $N = N_0$  gives the “correct” model.

Other values of  $N$  and any limits as  $N \rightarrow \infty$  (perhaps with a change of time-scale) give approximate models. The challenge is to select the  $\alpha_i$ , but once that is done, the initial condition for index  $N$  is given by

$$Z_i^N(0) = N_i^{-\alpha_i} X_i(0),$$

where the  $X_i(0)$  are the initial species numbers in the correct model.

If  $\lim_{N \rightarrow \infty} Z_i^N(\cdot N^\gamma) = Z_i^{\infty, \gamma}$ , then we should have

$$X_i(t) \approx N_0^{\alpha_i} Z_i^{\infty, \gamma}(t N_0^{-\gamma}).$$

For example, in the virus model

$$X_2(t) \approx (1000)^{2/3} Z_2^{\infty, \gamma}(t(1000)^{-2/3})$$



## Enzyme reaction

(cf. Darden (1982))

Basic assumption: A small number of enzymes interacts with a large number  $O(N)$  of substrate molecules.



$$\begin{aligned} X_E(t) &= X_E(0) - Y_1 \left( \int_0^t \kappa_1 X_A(s) X_E(s) ds \right) + Y_2 \left( \kappa_2 \int_0^t X_{AE}(s) ds \right) \\ &\quad + Y_3 \left( \kappa_3 \int_0^t X_{AE}(s) ds \right) \\ X_A(t) &= X_A(0) - Y_1 \left( \int_0^t \kappa_1 X_A(s) X_E(s) ds \right) \\ &\quad + Y_2 \left( \kappa_2 \int_0^t X_{AE}(s) ds \right) \end{aligned}$$



## Scaled model

Assume that production and dissociation reactions are fast, and the substrate is present in large numbers.  $Z_A^N = N^{-1}X_A^N$ .

$$X_E^N(t) = X_E(0) - Y_1(N \int_0^t \kappa_1 Z_A^N(s) X_E^N(s) ds) + Y_2(N \kappa_2 \int_0^t X_{AE}^N(s) ds) \\ + Y_3(N \kappa_3 \int_0^t X_{AE}^N(s) ds)$$

$$Z_A^N(t) = Z_A^N(0) - N^{-1}Y_1(N \int_0^t \kappa_1 Z_A^N(s) X_E^N(s) ds) \\ + N^{-1}Y_2(N \kappa_2 \int_0^t X_{AE}^N(s) ds)$$

$m \equiv X_E^N(t) + X_{AE}^N(t)$  does not depend on  $t$ .



## Averaging by the equation

Dividing the first equation by  $N$  and taking the limit

$$N^{-1}X_E^N(t) = N^{-1}X_E(0) - N^{-1}Y_1(N \int_0^t \kappa_1 Z_A^N(s) X_E^N(s) ds) \\ + N^{-1}Y_2(N \kappa_2 \int_0^t X_{AE}^N(s) ds) \\ + N^{-1}Y_3(N \kappa_3 \int_0^t X_{AE}^N(s) ds) \\ 0 = \lim_{N \rightarrow \infty} \int_0^t (-\kappa_1 Z_A^N(s) X_E^N(s) + \kappa_2 X_{AE}^N(s) + \kappa_3 X_{AE}^N(s)) ds$$

and

$$\lim_{N \rightarrow \infty} \int_0^t X_E^N(s) ds = \lim_{N \rightarrow \infty} \int_0^t (m - X_{AE}^N(s)) ds = \int_0^t \frac{m(\kappa_2 + \kappa_3)}{\kappa_1 Z_A(s) + \kappa_2 + \kappa_3} ds.$$



## Michaelis-Menten equation

$$Z_A^N(t) = Z_A^N(0) - N^{-1}Y_1(N \int_0^t \kappa_1 Z_A^N(s) X_E^N(s) ds) \\ + N^{-1}Y_2(N \kappa_2 \int_0^t X_{AE}^N(s) ds)$$

which converges to

$$Z_A(t) = Z_A(0) - \int_0^t \kappa_1 Z_A(s) \frac{m(\kappa_2 + \kappa_3)}{\kappa_1 Z_A(s) + \kappa_2 + \kappa_3} ds \\ + \kappa_2 \int_0^t \frac{m \kappa_1 Z_A(s)}{\kappa_1 Z_A(s) + \kappa_2 + \kappa_3} ds \\ = Z_A(0) - \int_0^t \frac{m \kappa_1 \kappa_3 Z_A(s)}{\kappa_1 Z_A(s) + \kappa_2 + \kappa_3} ds$$



## Gaussian limit

Kang, Kurtz, and Popovic (2014)

Call the limiting quadratic variation  $\int_0^t \sigma^2(Z_A(s)) ds$ .

$$\int_0^t \sigma^2(Z_A(s)) ds \\ = \int_0^t ((h(Z_A(s)) - 1)^2(\kappa_2 + \kappa_3) + (h(Z_A(s)) + 1)^2 \kappa_2 + h(Z_A(s))^2 \kappa_3) \\ \times \frac{m \kappa_1 Z_A(s)}{\kappa_1 Z_A(s) + \kappa_2 + \kappa_3} ds$$

Then

$$U^N(t) = \sqrt{N}(Z_A^N(t) - Z(t))$$

converges to the solution of

$$U(t) = U(0) + W(\int_0^t \sigma^2(Z_A(s)) ds) - \int_0^t \frac{m \kappa_1 \kappa_3 (\kappa_2 + \kappa_3)}{(\kappa_2 + \kappa_3 + \kappa_1 Z_A(s))^2} U(s) ds$$



## Diffusion approximation

Rationale: Find a sequence of diffusion processes  $D_A^N$  such that

$$\sqrt{N}(D_A^N - Z_A) \Rightarrow U.$$

So

$$D_A^N(t) = Z_A^N(0) + \frac{1}{\sqrt{N}} \int_0^t \sigma(D_A^N(s)) dW(s) - \int_0^t \frac{m\kappa_1\kappa_3 D_A^N(s)}{\kappa_2 + \kappa_3 + \kappa_1 D_A^N(s)} ds$$



## Heat shock model

The following reaction network is given as a model for the heat shock response in E. Coli by [Srivastava, Peterson, and Bentley \(2001\)](#). The analysis is from [Kang \(2012\)](#).

Reaction	Intensity	Reaction	Intensity
$\emptyset \rightarrow S_8$	$4.00 \times 10^0$	$S_6 + S_8 \rightarrow S_9$	$3.62 \times 10^{-4} X_{S_6} X_{S_8}$
$S_2 \rightarrow S_3$	$7.00 \times 10^{-1} X_{S_2}$	$S_8 \rightarrow \emptyset$	$9.99 \times 10^{-5} X_{S_8}$
$S_3 \rightarrow S_2$	$1.30 \times 10^{-1} X_{S_3}$	$S_9 \rightarrow S_6 + S_8$	$4.40 \times 10^{-5} X_{S_9}$
$\emptyset \rightarrow S_2$	$7.00 \times 10^{-3} X_{S_1}$	$\emptyset \rightarrow S_1$	$1.40 \times 10^{-5}$
$\text{stuff} + S_3 \rightarrow S_5 + S_2$	$6.30 \times 10^{-3} X_{S_3}$	$S_1 \rightarrow \emptyset$	$1.40 \times 10^{-6} X_{S_1}$
$\text{stuff} + S_3 \rightarrow S_4 + S_2$	$4.88 \times 10^{-3} X_{S_3}$	$S_7 \rightarrow S_6$	$1.42 \times 10^{-6} X_{S_4} X_{S_7}$
$\text{stuff} + S_3 \rightarrow S_6 + S_2$	$4.88 \times 10^{-3} X_{S_3}$	$S_5 \rightarrow \emptyset$	$1.80 \times 10^{-8} X_{S_5}$
$S_7 \rightarrow S_2 + S_6$	$4.40 \times 10^{-4} X_{S_7}$	$S_6 \rightarrow \emptyset$	$6.40 \times 10^{-10} X_{S_6}$
$S_2 + S_6 \rightarrow S_7$	$3.62 \times 10^{-4} X_{S_2} X_{S_6}$	$S_4 \rightarrow \emptyset$	$7.40 \times 10^{-11} X_{S_4}$



## References

- David F. Anderson and Thomas G. Kurtz. *Stochastic analysis of biochemical systems*, volume 1 of *Mathematical Biosciences Institute Lecture Series. Stochastics in Biological Systems*. Springer, Cham; MBI Mathematical Biosciences Institute, Ohio State University, Columbus, OH, 2015. ISBN 978-3-319-16894-4; 978-3-319-16895-1.
- Karen Ball, Thomas G. Kurtz, Lea Popovic, and Greg Rempala. Asymptotic analysis of multiscale approximations to reaction networks. *Ann. Appl. Probab.*, 16(4):1925–1961, 2006. ISSN 1050-5164.
- Thomas A. Darden. Enzyme kinetics: stochastic vs. deterministic models. In *Instabilities, bifurcations, and fluctuations in chemical systems (Austin, Tex., 1980)*, pages 248–272. Univ. Texas Press, Austin, TX, 1982.
- Eric L. Haseltine and James B. Rawlings. Approximate simulation of coupled fast and slow reactions for stochastic chemical kinetics. *J. Chem. Phys.*, 117(15):6959–6969, 2002.
- Hye-Won Kang. A multiscale approximation in a heat shock response model of e. coli. *BMC Systems Biology*, 6(1):1–22, 2012. ISSN 1752-0509. doi: 10.1186/1752-0509-6-143. URL <http://dx.doi.org/10.1186/1752-0509-6-143>.
- Hye-Won Kang and Thomas G. Kurtz. Separation of time-scales and model reduction for stochastic reaction networks. *Ann. Appl. Probab.*, 23(2):529–583, 2013. ISSN 1050-5164. doi: 10.1214/12-AAP841. URL <http://dx.doi.org/10.1214/12-AAP841>.
- Hye-Won Kang, Thomas G. Kurtz, and Lea Popovic. Central limit theorems and diffusion approximations for multiscale Markov chain models. *Ann. Appl. Probab.*, 24(2):721–759, 2014. ISSN 1050-5164. doi: 10.1214/13-AAP934. URL <http://dx.doi.org/10.1214/13-AAP934>.
- Thomas G. Kurtz. Averaging for martingale problems and stochastic approximation. In *Applied stochastic analysis (New Brunswick, NJ, 1991)*, volume 177 of *Lecture Notes in Control and Inform. Sci.*, pages 186–209. Springer, Berlin, 1992.



- R. Srivastava, M. S. Peterson, and W. E. Bentley. Stochastic kinetic analysis of escherichia coli stress circuit using sigma(32)-targeted antisense. *Biotechnol. Bioeng.*, 75:120–129, 2001.
- R. Srivastava, L. You, J. Summers, and J. Yin. Stochastic vs. deterministic modeling of intracellular viral kinetics. *J. Theoret. Biol.*, 218(3):309–321, 2002. ISSN 0022-5193.

