



Biochemical systems with stochastic parameter variation

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Causes of uncertainty

- Measurement uncertainty

- Biological variation

External conditions, genotype, differential expression ...

- Unknown parameters

Rough guess:

K_M values between 10^{-4} and 100 mM (Brenda database)

Modelling objectives

- Prediction with missing/uncertain knowledge:
Probabilistic statements
- Mean values in cell populations
- Bayesian parameter estimation
- Robustness studies
Which properties are (almost) determined by the model structure?

Outline

- **Choice of parameter distributions**
- **Direct computation of variable distribution**
- **Stochastic parameter fluctuations**

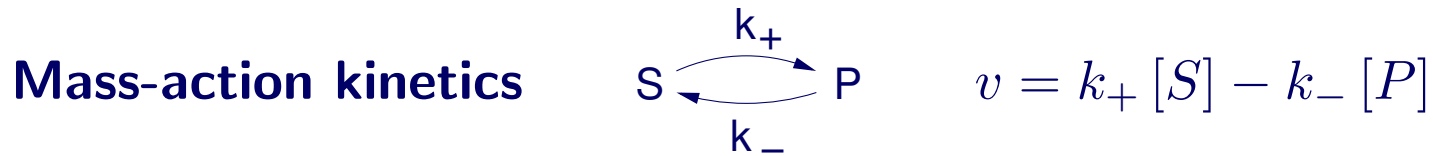
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The log-normal distribution

- Definition:
 x is log-normally distributed $\Leftrightarrow \ln x$ is normally distributed
- x, y log-normal $\rightarrow z = x^a \cdot y^b$ also log-normal
- Joint distributions of several variables:
 $\ln \mathbf{x}$ is multivariate normal with $\langle \ln \mathbf{x} \rangle, \text{cov}(\ln \mathbf{x})$
- Parameter distribution defined by
standard parameters \mathbf{p}^0 and coefficients of variation σ_{p_i}/p_i^0
 \rightarrow mean values $\langle \ln \mathbf{p} \rangle$ and variances $\text{var}(\ln \mathbf{p})$

Consistent joint distributions. Example 1



$$\begin{aligned} k_+/k_- &= q = e^{-\beta\Delta g} && (= \text{equilibrium constant}) \\ \sqrt{k_+ \cdot k_-} &= ru && (= \text{ratio} \cdot \text{enzyme activity}) \end{aligned}$$

This can be fulfilled by choosing g, r, u , and setting

$$\begin{aligned} \ln q &= -\beta\Delta g \\ \ln k_+ &= \ln u + 1/2 \ln q + \ln r \\ \ln k_- &= \ln u - 1/2 \ln q + \ln r \end{aligned}$$

Linear combination of logarithms

$$\begin{aligned} (\ln k_+, \ln k_-)^T &= \mathbf{a} \ln u + \mathbf{b} \ln q + \mathbf{c} \ln r \\ \text{with } \mathbf{a} &= \begin{pmatrix} 1 \\ 1 \end{pmatrix}, \quad \mathbf{b} = \frac{1}{2} \begin{pmatrix} 1 \\ -1 \end{pmatrix}, \quad \mathbf{c} = \begin{pmatrix} 1 \\ 1 \end{pmatrix} \end{aligned}$$

Compute mean and cov for $(\ln k_+, \ln k_-)^T$ from mean and variance for $g, \ln r, \ln u$

Consistent joint distributions. Example 2

Reversible Michaelis-Menten kinetics $v = \frac{V_{\max}^{\rightarrow}/K_M^{\rightarrow}[S] - V_{\max}^{\leftarrow}/K_M^{\leftarrow}[P]}{1 + [S]/K_M^{\rightarrow} + [P]/K_M^{\leftarrow}}$

Assumptions about parameters (with \mathbf{z} multivariate standard normal):

$$\begin{aligned} \frac{V_{\max}^{\rightarrow} K_M^{\leftarrow}}{K_M^{\rightarrow} V_{\max}^{\leftarrow}} &= q & \text{with} & \ln K_M^{\rightarrow} &= \mu_K^S + \sigma_K z_1 \\ \sqrt{V_{\max}^{\rightarrow} \cdot V_{\max}^{\leftarrow}} &= u V_{\max}^* & & \ln K_M^{\leftarrow} &= \mu_K^P + \sigma_K z_2 \\ & & & \ln V_{\max}^* &= \mu_V + \sigma_V z_3, \end{aligned}$$

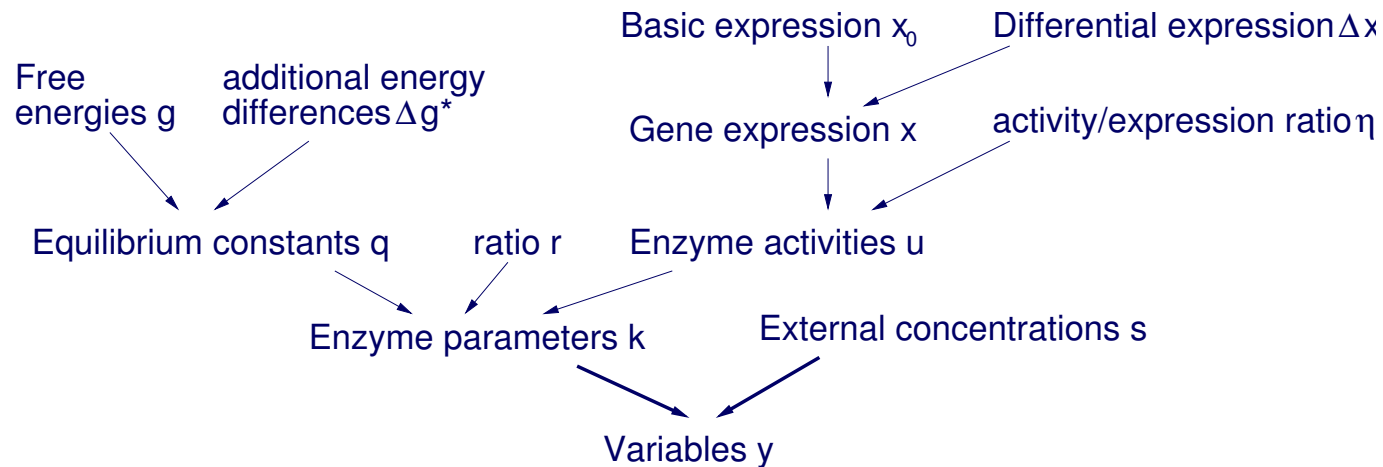
Linear combination of logarithms

$$(\ln K_M^{\rightarrow}, \ln K_M^{\leftarrow}, \ln V_{\max}^{\rightarrow}, \ln V_{\max}^{\leftarrow})^T = \mathbf{a} \ln u + \mathbf{b} \ln q + \mathbf{C} \mathbf{z} + \mathbf{d}$$

$$\text{with } \mathbf{a} = (0, 0, 1, 1)^T, \quad \mathbf{b} = (0, 0, 1/2, -1/2)^T, \quad \mathbf{d} = (\mu_K^S, \mu_K^P, \mu_V, \mu_V)^T$$

$$\mathbf{C} = \begin{pmatrix} \sigma_K & 0 & 0 \\ 0 & \sigma_K & 0 \\ 1/2\sigma_K & -1/2\sigma_K & \sigma_V \\ -1/2\sigma_K & 1/2\sigma_K & \sigma_V \end{pmatrix}$$

Scheme for dependent parameters



Distribution of all parameters: multiplicative relations yield

$$\ln \mathbf{k} = A \ln \mathbf{u} + B \ln \mathbf{q} + C\mathbf{z} + \mathbf{d}, \quad \ln \mathbf{q} = -\beta N^T \mathbf{g}$$

Compute means and covariances

$$\begin{aligned} \langle \ln \mathbf{k} \rangle &= A \langle \ln \mathbf{u} \rangle + B \langle \ln \mathbf{q} \rangle + \mathbf{d} \\ \text{cov}(\ln \mathbf{k}) &= A \text{cov}(\ln \mathbf{u}) A^T + B \text{cov}(\ln \mathbf{q}) B^T + CC^T \end{aligned}$$

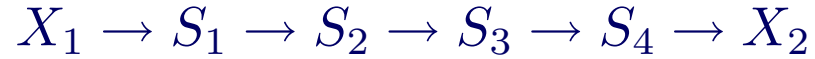
Consistent description by log-normal distributions!

Outline

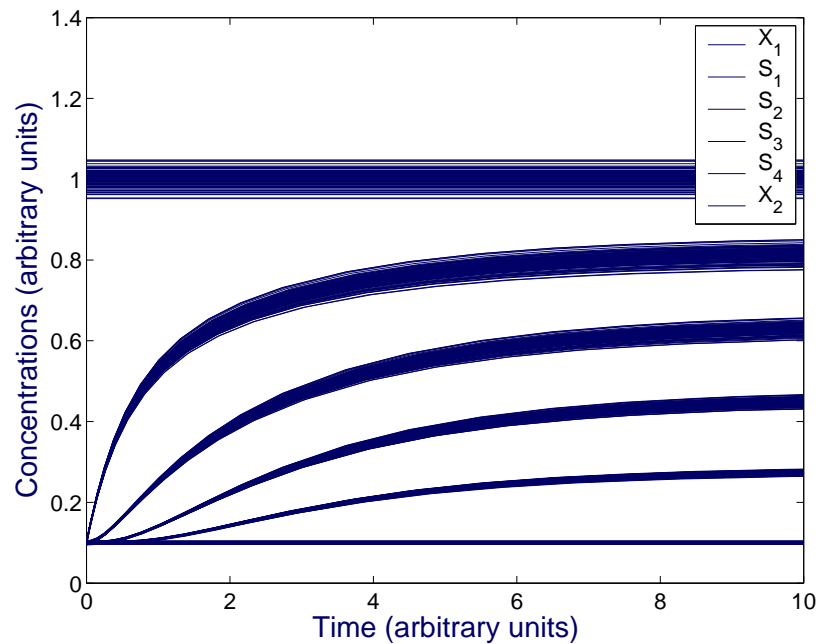
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Distribution of variables

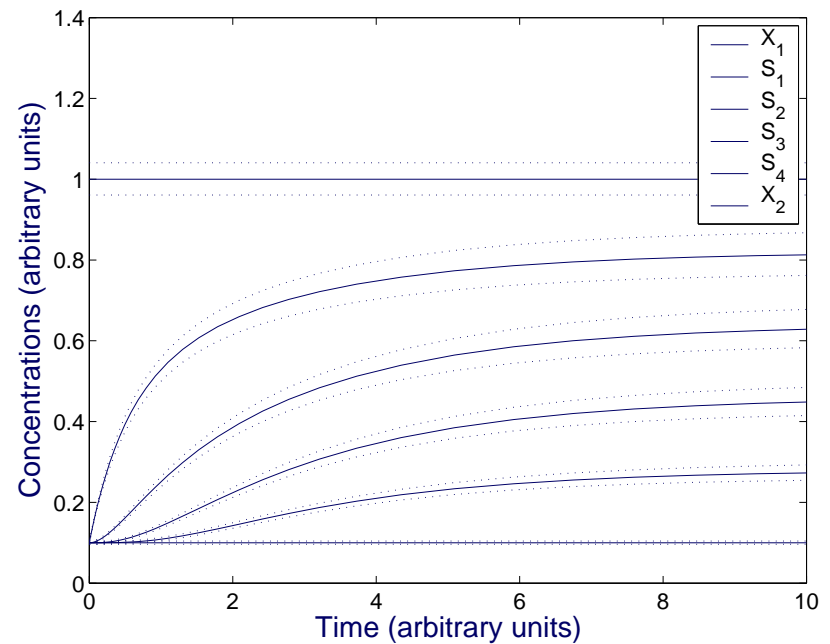
Example: Linear chain



Parameters independent log-normal



Monte Carlo simulations

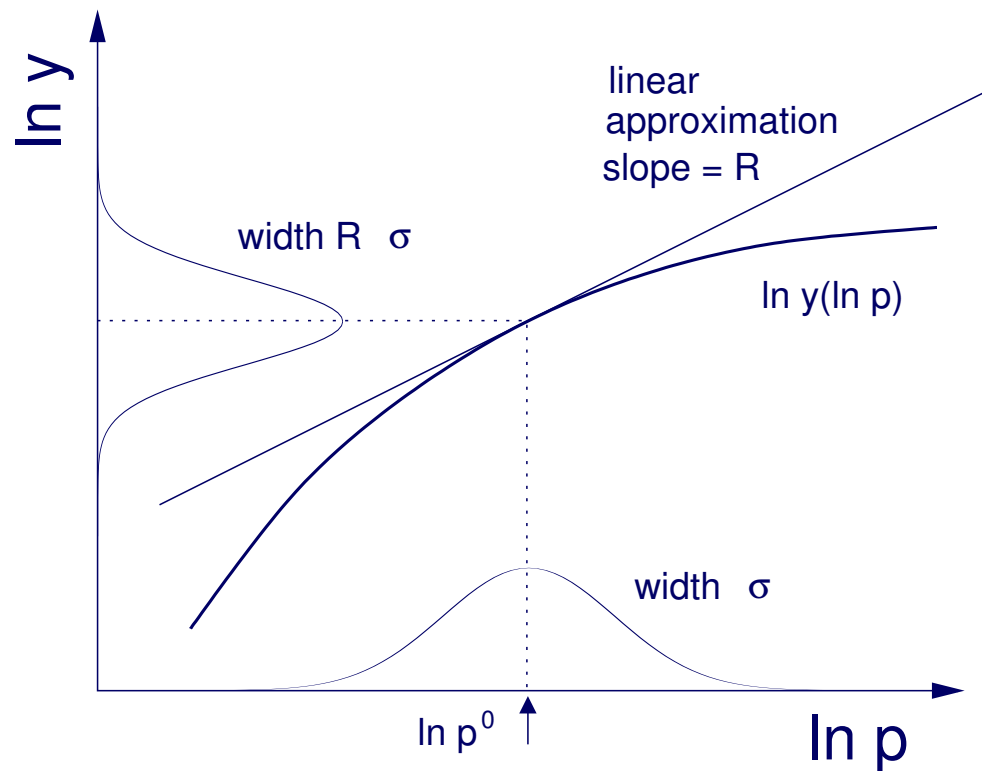


Quantiles

Expansion method: Idea

Distribution of system variable y

(steady state flux, concentration, control coefficient, time courses, oscillation characteristics, signal characteristics, ...)



Expansion method: Results

Mean and covariance of $\ln \mathbf{y}$ can be computed by

First order expansion

$$\begin{aligned}\langle \ln y_i \rangle &= \ln(y_i(\mathbf{p}^0)) \\ \text{cov}(\ln \mathbf{y}) &= \hat{R}^Y \text{cov}(\ln \mathbf{p})(\hat{R}^Y)^T\end{aligned}$$

\hat{R}^Y : (normalised) response coefficients

Log-normal parameters yield log-normal variables.

Expansion method: Results

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Second order expansion (with the shortcut $Q := \text{cov}(\ln \mathbf{p})$)

$$\begin{aligned}\langle \ln y_i \rangle &= \ln(y_i(\mathbf{p}^0)) + \frac{1}{2} \sum_{mn} \hat{R}_{lmn}^{(2)Y} Q_{jk} \\ \text{cov}(\Delta \ln y_l, \Delta \ln y_k) &= \sum_{mn} \hat{R}_{lm}^Y Q_{mn} \hat{R}_{kn}^Y \\ &\quad + \frac{1}{4} \sum_{mnr s} \hat{R}_{lmn}^{(2)Y} \hat{R}_{krs}^{(2)Y} (Q_{ms} Q_{nr} + Q_{mr} Q_{ns})\end{aligned}$$

Ratios and order relations among variables

Probabilities of **binary variables** derived from log-normal system variables y, y_1, y_2 :

$$\text{Prob}(y > a) = \Phi \left(\frac{\langle \ln y \rangle - \log a}{\text{var}(\ln y)^{1/2}} \right)$$

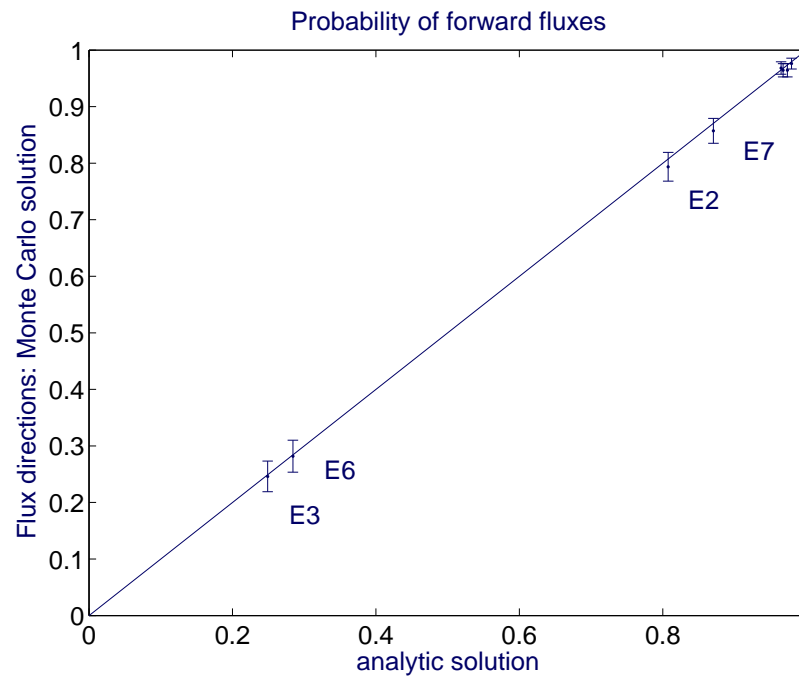
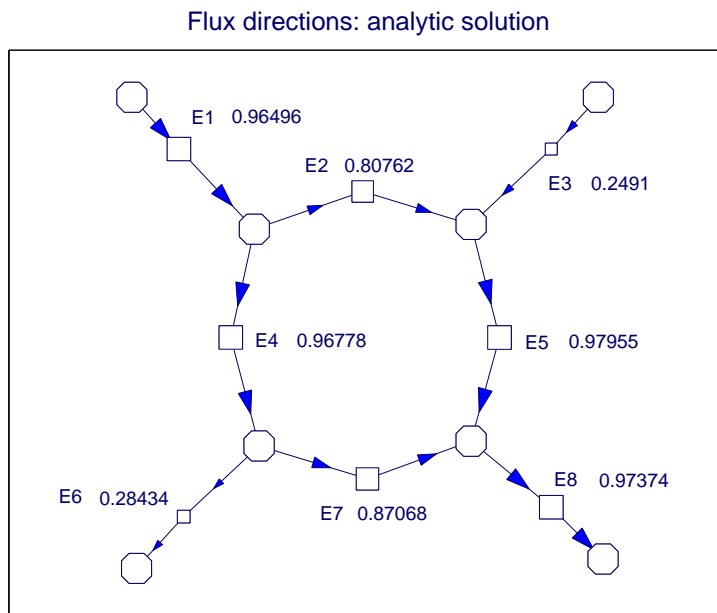
$$\text{Prob}(y_1 > y_2) = \Phi \left(\frac{\langle \ln y_1 \rangle - \langle \ln y_2 \rangle}{(\text{var}(\ln y_1) + \text{var}(\ln y_2) - 2 \text{cov}(\ln y_1, \ln y_2))^{1/2}} \right)$$

$$\text{Prob}(y_1/y_2 > a) = \Phi \left(\frac{\langle \ln y_1 \rangle - \langle \ln y_2 \rangle - \ln a}{(\text{var}(\ln y_1) + \text{var}(\ln y_2) - 2 \text{cov}(\ln y_1, \ln y_2))^{1/2}} \right)$$

where $\Phi(\cdot)$ is the cumulative density of the standard normal distribution

Example: Probability of flux directions

1. determine joint distribution of forward and backward fluxes
2. compute $\text{Prob}(\text{forward flux} > \text{backward flux})$



Other example:

Fold changes of metabolite concentrations by differential expression

Variability and sensitivity

The linear expansion yields measures of ...

- **Variability**

$$\text{cov}(\ln \mathbf{y}) \approx \hat{R}^Y \text{cov}(\ln \mathbf{p})(\hat{R}^Y)^T$$

Log-normal distributions: coefficient of variation reads $\frac{\sigma_y}{\langle y \rangle} = \sqrt{e^{\text{var}(\ln y)} - 1}$.

- **Stochastic sensitivity** (Krewski et al., 1995)

$$\begin{aligned} U(\ln y_l | \ln p_i) &:= \text{var}_a(\mathcal{E}(\ln y_l | \ln p_i = a)) \\ &\approx \left(\sum_j \hat{R}_{lj}^Y \text{cov}(\ln p_j, \ln p_i) \right)^2 / \text{var}(\ln p_i) \end{aligned}$$

- **Covariance between parameter and variable**

$$\text{cov}(\ln y_l, \ln p_i) \approx \sum_j \hat{R}_{lj}^Y \text{cov}(\ln p_j, \ln p_i)$$

Expansion method: Comments

- Many independent parameters:

First-order expansion:

For any parameter distribution, variable distributions tend to become log-normal (central limit theorem).

- Second-order expansion

... to check the quality of first-order expansion

... shows that population mean \neq results from mean parameters!

- Bayesian parameter estimation

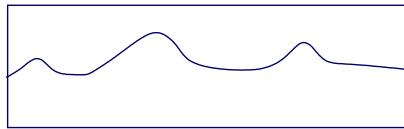
Improve parameter distributions by using experimental data

Explicit formulae for the posterior follows from the first-order expansion

Outline

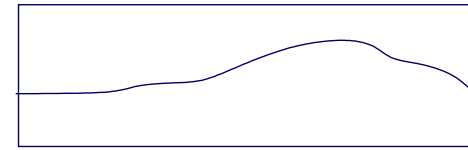
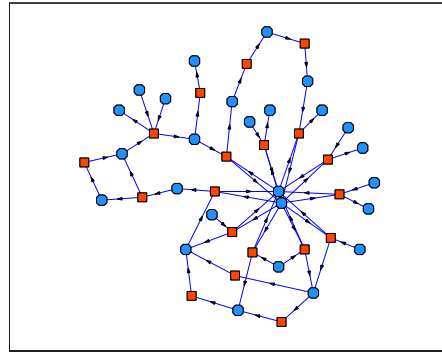
- Choice of parameter distributions
- Direct computation of variable distribution
- **Stochastic parameter fluctuations**

Time-dependent parameter perturbations



$p(t')$ →

parameter
time courses

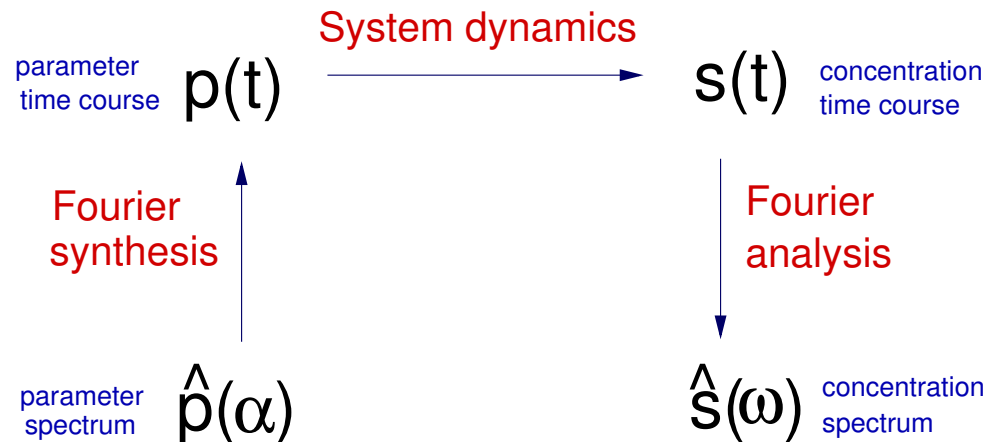


→ $s(t), j(t)$

variable
time courses

Parameter perturbation of frequency $\omega \rightarrow$ Forced oscillations of $s(t), \mathbf{j}(t)$

Spectral response coefficients



Spectral response coefficients

Definition

$$R_{lm}^S(\omega) := \frac{\delta \hat{s}_l(\hat{p}(\cdot), \omega)}{\delta \hat{p}_m(\omega)}$$

Formula

$$R^S(\omega) = -(N\epsilon^S - i\omega\mathbf{I})^{-1}N\epsilon^P$$

Spectral density of concentration fluctuations

$$\mathcal{S}^S(\omega) = R^S(\omega) \mathcal{S}^P(\omega) (R^S(\omega))^\dagger$$

Stochastic parameter fluctuations

Stochastic fluctuations due to small particle numbers:

Chemical Langevin equation (see Gillespie, 2000)

$$\frac{d}{dt} \bar{x}_i(t) = N_{ik} a_k(\bar{\mathbf{x}}(t)) + N_{ik} \sqrt{a_k(\bar{\mathbf{x}}(t))} \eta_k(t)$$

\bar{x}_i : molecule numbers, a_k : propensity functions, η_k : white standard noise

Fluctuations around steady state:

1. Linearise system at mean concentrations \mathbf{x}^0
2. Represent the fluctuations by virtual noise parameters with elasticities

$$\epsilon^P := (n \text{ liters/mol})^{-1/2} \text{diag}(\mathbf{v}(\mathbf{x}^0))^{1/2}$$

$n \text{ liters/mol} = N_A \Omega$ (n : average molecule number at 1 mol/liter)

Ω : system volume, N_A : Avogadro's constant

White Gaussian noise \rightarrow spectral density reads $\mathcal{S}^S(\omega) = R^S(\omega) (R^S(\omega))^\dagger$

Resonance

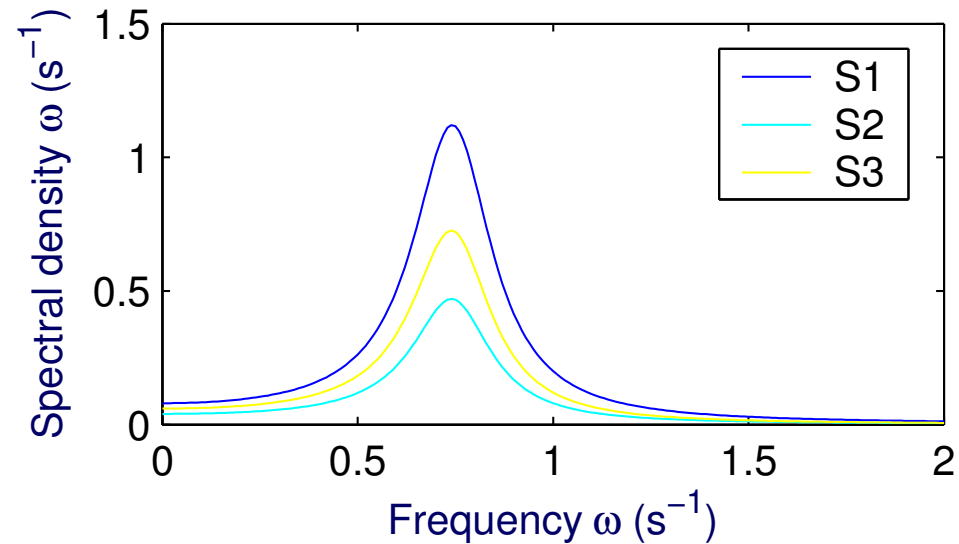
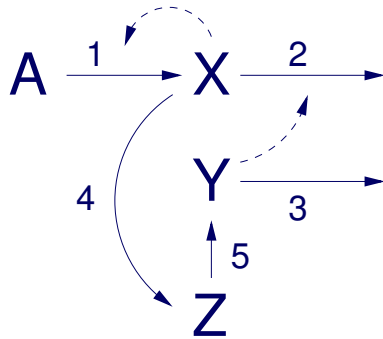
Example:

Minimal reaction system with Hopf bifurcation (Wilhelm + Heinrich, 1995)

Stable steady state:

Small volume $(3.9 \text{ nm})^3$ (≈ 100 molecules of each substance)

\Rightarrow spontaneous fluctuations of concentrations



Resonance near system's own oscillation frequency $\omega_0 \approx 0.75 \text{ s}^{-1}$

Conclusions

- **Choice of parameter distributions**

- Thermodynamic constraints → correlated parameter distributions
- Graphical scheme → easy computation of distributions
- Log-normal distribution is convenient, but not necessary

- **Monte Carlo or expansion method?**

- Monte Carlo is simple and general, but error is $\mathcal{O}(\sqrt{1/n})$
- Expansion relies on small parameter variance

But: Analytical formulae, showing relations to model structure

- **Thanks to ...**

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