

Metabolic control analysis with temporal parameter fluctuations

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Abstract

Metabolic response coefficients describe how dynamic properties of metabolic systems - like steady state concentrations - respond to small, time-independent changes of the kinetic parameters. We extend this concept to temporal parameter fluctuations and define spectral response coefficients that relate Fourier components of concentration time courses to Fourier components of the underlying parameters. The spectral response coefficients describe forced oscillations of the concentrations and fluxes, caused by small harmonic oscillations of single parameters. They depend on the driving frequency and comprise the relative phases and amplitudes. To illustrate the basic idea, only first-order response coefficients for metabolic concentrations are addressed on this poster.

Example: Glycolysis model

We start with the glycolysis model from Hynne et al. (2001) (see [3]) at a stable steady state (low external glucose concentration Glcx0=5.0).

The storage reaction (parameter k_{22}) is perturbed by a harmonic oscillation of frequency $\alpha = 2\pi/(10 \text{ min}) \Rightarrow$ Forced oscillations of all concentrations.

Spectral response coefficients of concentrations and fluxes

Temporal response to an oscillatory perturbation (top left diagram)

 $\Delta k22$

Metabolic control analysis of steady states

Stable metabolic system: Each choice of the kinetic parameters (enzyme activities, Michaelis constants, etc.) leads to certain steady state concentrations



Expand the stationary concentrations S_l after a parameter change $\Delta \mathbf{p}$:

Metabolic response coefficients (see [2]):

 $R_{lm}^{\rm S} := \frac{\partial S_l(\mathbf{p})}{\partial p_m}$

 v_k : kinetics functions

$$S_l(\mathbf{p}^0 + \Delta \mathbf{p}) \approx S_l(\mathbf{p}^0) + \sum_m R_{lm}^{\mathrm{S}} \Delta p_m$$

Computation:

 Consider the stationarity condition $0 = \dot{\mathbf{x}} = \mathbf{N}\mathbf{v}(\mathbf{x}, \mathbf{p})$ $\mathbf{R}^{\mathrm{S}} = -(\mathbf{N}\epsilon_{\mathrm{s}})^{-1}\mathbf{N}\epsilon_{\mathrm{p}}$ \circ Differentiation by ${f p}$ yields

with stoichiometric matrix N and reaction elasticities $(\epsilon_s)_{kl} := \frac{\partial v_k}{\partial x_l}, \quad (\epsilon_p)_{km} := \frac{\partial v_k}{\partial p_m}$







 1^{st} order 2^{nd} order $0^{
m th}$ order Exact solution

Time courses for general perturbations can be computed via Fourier synthesis.

Propagation of stochastic fluctuations

Stochastical fluctuations due to small particle numbers can be described by the chemical Langevin equation (see [1])

$$\frac{\mathbf{d}}{\mathbf{d}t}\bar{x}_{i}(t) = N_{ik} a_{k}(\bar{\mathbf{x}}(t)) + N_{ik} \sqrt{a_{k}(\bar{\mathbf{x}}(t))} \eta_{k}(t)$$

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

After linearisation around the mean concentrations \mathbf{x}^0 , the fluctuations

Spectral response coefficients

Parameter fluctuations:

Instead of parameter vectors, we now consider vectorial time courses



Describe the time courses by their Fourier spectrum...



Computation:

- Consider the system equation
- Fourier transformation yields

 $\dot{\mathbf{x}} = \mathbf{N}\mathbf{v}(\mathbf{x}(t), \mathbf{p}(t))$ $\mathbf{i}\omega\hat{\mathbf{s}}(\omega) = \mathbf{N}\mathbf{j}(\omega)$ **j**: fluxes $\mathbf{R}^{\mathrm{S}}(\omega,\alpha) = -(\mathbf{N}\epsilon_{\mathrm{s}} - \mathrm{i}\omega\mathbf{I})^{-1}\mathbf{N}\epsilon_{\mathrm{p}}\delta(\omega - \alpha)$

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can be described by virtual noise parameters with elasticities

$$\epsilon_{\mathrm{p}} := (n \operatorname{\mathsf{liters}/\mathsf{mol}})^{-1/2} \operatorname{\mathsf{diag}}(\mathbf{v}(\mathbf{x}^0))^{1/2}$$

n: average molecule number at 1 mol/liter, n liters/mol = $N_{
m A}\Omega$ Ω : system volume $N_{\rm A}$: Avogadro's constant

Concentration fluctuations are quantified by their spectral densities:

 $\mathcal{S}^{\mathrm{S}}(\omega) = R^{\mathrm{S}}(\omega) \ R^{\mathrm{S}^{\dagger}}(\omega) = n^{-1} \left(\mathbf{N}\epsilon_{\mathrm{s}} - \mathsf{i}\omega \right)^{-1} \mathbf{N} \operatorname{diag}(\mathbf{v}^{*}) \ \mathbf{N}^{T} (\mathbf{N}\epsilon_{\mathrm{s}} + \mathsf{i}\omega)^{-1^{T}} \cdot \operatorname{liters/mol.}$

Example: Minimal reaction system (3 substances) with Hopf bifurcation [6]

Stable steady state in a cubic volume $(3.9 \text{ nm})^3$ containing, on average, n = 100 molecules of each substance \Rightarrow Spontaneous fluctuations of concentrations





Note the resonance - below the bifurcation- near the oscillation frequency $\omega_0 \approx 0.75 \text{ s}^{-1}$ of the bifurcation point.

Not shown here...

\circ Differentiation by p yields

• The spectral response coefficients are complex and frequency-dependent. \circ They describe the response to a harmonic oscillation of a parameter.

- They may show resonance near Hopf bifurcations.
- \circ To account for conservation relations \Rightarrow Restriction to independent metabolites (see [5])

Based on the same idea, it is also straightforward to compute (see [4])

• Response coefficients for fluxes

• Second-order response coefficients

• Control coefficients (1^{st} and 2^{nd} order)

• Summation and connectivity theorems

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[1] D. T. Gillespie. The chemical Langevin equation. J. Chem. Phys., 113(1):297–306, 2000. [2] R. Heinrich and S. Schuster. *The regulation of cellular systems*. Chapman & Hall, 1996. [3] F. Hynne, S. Danø, and P.G. Sørensen. Full-scale model of glycolysis in Saccharomyces cerevisiae. *Biophys. Chem.*, 94:121–163, 2001. [4] W. Liebermeister. Metabolic response to temporal parameter fluctuations in biochemical networks. submitted to J. Theor. Biol. [5] C. Reder. Metabolic control: a structural approach. J. theor. Biol., 135:175-201, 1988. [6] T. Wilhelm and R. Heinrich. The smallest chemical reaction systems with Hopf-bifurcation. J. Math. Chem., 17:1–14, 1995.