Vorlesung "Modellierung von Zellprozessen" Aufgabenblatt 1: Modellierung

WS 2009/2010, Dozent: Wolfram Liebermeister

1. Small network structure

Draw the the reaction system

 $\begin{array}{c} P_1+S_2 \stackrel{v_1}{\leftrightarrow} S_1+S_3\\ S_1+S_3 \stackrel{v_2}{\leftrightarrow} P_2+S_2 \end{array}$

as a network, write down its stoichiometric matrix, and determine potential fluxes in stationary state. Are there linear conservation relations? (v_i : reaction rate, S_j : internal (variable) compounds, P_k : external (fixed) compounds)

2. From the ODEs back to the reaction

Consider the following system of differential equations. Try to find out a reaction scheme that corresponds to these reactions. Are there cases of activation or inhibition?

$$\frac{dM_1^-}{dt} = -k_1 \cdot M_1^- \cdot \frac{1}{(1+M_3^+)} + k_2 \cdot M_1^+ \\
\frac{dM_1^+}{dt} = +k_1 \cdot M_1^- \cdot \frac{1}{(1+M_3^+)} - k_2 \cdot M_1^+ \\
\frac{dM_2^-}{dt} = -k_3 \cdot M_2^- \cdot M_1^+ + k_4 \cdot M_2^+ \\
\frac{dM_2^+}{dt} = +k_3 \cdot M_2^- \cdot M_1^+ - k_4 \cdot M_2^+ \\
\frac{dM_3^-}{dt} = -k_5 \cdot M_3^- \cdot M_2^+ + k_6 \cdot M_3^+ \\
\frac{dM_3^+}{dt} = +k_5 \cdot M_3^- \cdot M_2^+ - k_6 \cdot M_3^+$$

3. The repressilator The repressilator (M. Elowitz and S. Leibler, Nature 2000) is a genetic circuit consisting of three proteins, each inhibiting the production of the following protein in a circle. Consider the following kinetic equations for synthesis and degradation of the proteins:

with i = 1, 2, 3 and l(1) = 3, l(2) = 1, l(3) = 2. The reaction rate vector reads $v = (v_1^{\text{syn}}, v_2^{\text{syn}}, v_3^{\text{syn}}, v_1^{\text{degr}}, v_2^{\text{degr}}, v_3^{\text{degr}})^{\text{T}}$.

- (a) Write down the stoichiometric matrix and the differential equations for the protein levels x_i .
- (b) Consider a stationary state with $x_1 = x_2 = x_3 = \bar{x}$. Calculate the elasticity matrix ε^{S} . (Hint: it is not necessary to compute the value of \bar{x} .)
- (c) Compute the Jacobian matrix by the formula $M = N\varepsilon^{S}$.
- (d) How could you decide based on M whether the stationary state is stable?