Synthetic Biology

Assignment #01

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0.1 The model

The model for a two protein network can be given by the system of equations:

$$\frac{dx}{dt} = f(y) - d_1 x \tag{0.1a}$$

$$\frac{dy}{dt} = g(x) - d_2y \tag{0.1b}$$

where x and y are the concentration of the two proteins, f and g are "repression functions" and d_1 , d_2 are positive constant describing the protein decay.

$$f(y) = \frac{k_1}{1 + [Y]^n} \tag{0.2a}$$

$$g(x) = \frac{k_2}{1 + |X|} \tag{0.2b}$$

The level of repression of [X] in a "mixed" circuit model is described by a Hill function [Cherry and Adler, 2000], which models cooperativity of binding between two different molecular species. Cooperativity magnitude is described by the Hill coefficient n. Using the conditions for a change in the number of steady states derived from nullclines analysis:

$$d_1[X] = \frac{k_1}{1 + [Y]^n} * \frac{1}{d_1} = \frac{k'_1}{1 + [Y]^n}$$
(0.3a)

$$y_1 = \sqrt[n]{\frac{k_1'}{d_1[X]} - 1} \tag{0.3b}$$

$$d_2[Y] = \frac{k_2}{1 + [X]} * \frac{1}{d_2} = \frac{k_2'}{1 + [X]^n}$$
(0.3c)

$$y_2 = \frac{k_2'}{1 + [X]} \tag{0.3d}$$

with $k'_1 = \frac{k_1}{d_1}$ and $k'_2 = \frac{k_2}{d_2}$.

The equations 0.3a and 0.3d are solved as functions of f(x) in the form y = f(x).

0.2 **Nullcline plots**

The analysis performed in graphs 0.1 shows the existence of multiple steady states depending on which parameters are chosen. Multiple steady states are the necessary condition to have a switch. The purpose of the switch is to obtain a consisting change in the concentration of a certain protein. The graph 0.1b shows transverse nullclines.

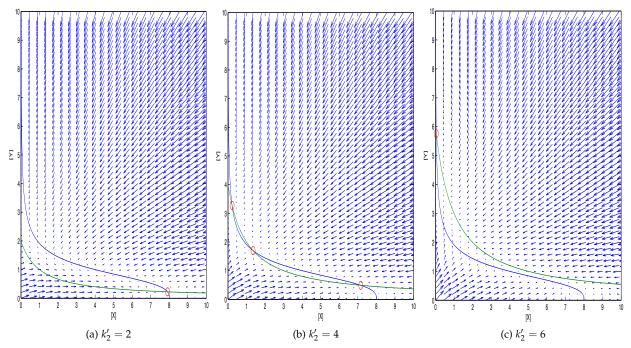


Figure 0.1: Nullclines plots for the parametres n=3, $k_1'=8$, $k_2'=\{2,4,6\}$

Intersection of functions lines are circled in red. These points represent the steady state at different values of the parameter k_2 . In the graph 0.1a we can distinguish only one point at high concentration of protein Y as well as in the graph 0.1c where the steady state is reached at high concentration of X. The major difference is visible in the graph 0.1b where we clearly confirm a possible switch, due to the presence of three steady state points.

0.3 Analytic expressions for k'_1 and k'_2

According to Matlab solve function for the equations:

$$e1: x = \frac{k_1'}{(1+y^3)} \tag{0.4a}$$

$$e2: y = \frac{k2}{(1+x)}$$

$$e3: \delta(\frac{k_1}{(1+y^3)}, y) * \delta(\frac{k_2}{(1+x)}, x) = 1$$
(0.4b)

$$e3: \delta(\frac{k_1}{(1+y^3)}, y) * \delta(\frac{k_2}{(1+x)}, x) = 1$$
 (0.4c)

we retrieve three solutions:

$$k_1' = \frac{(3*x^2)}{(2*x-1)} \tag{0.5a}$$

$$k_2' = \left(\frac{(x+1)}{(2*x-1)}\right)^{\frac{1}{3}} * (x+1)$$
 (0.5b)

$$y = \left(\frac{(x+1)}{(2*x-1)}\right)^{\frac{1}{3}} \tag{0.5c}$$

0.4 Bifurcation diagram

In bifurcation diagram the region delimited by the graph is the bistable region. Only in this region switches are possible. From this diagram one can see clearly that $k_2'=2$ is completely outside the bistable region, whereas $k_2'=4$ lies inside. This corresponds to the nullclines analyses in plot 0.1. There we could find a possible switch for $k_2'=4$ but not for $k_2'=2$. Even though $k_2'=6$ seems to be in the bistable region here, we can see from graph 0.1c that there are no possible switches.

0.5 synthbio_script_01.m

```
%## Synth Bio Project 01 ##
  %# Author: Stephanie Fingerhuth, Lorenzo Gatti
  %# Date: 09-Mar-2012 v.0.0.1
  %# Update: 10-Mar-2012 v.0.1.1
  %# Usage:
  %# Parametres:
  %# Output:
10
  %# TODO: Add comments [DONE]
12
  13
14
  15
17
  \mbox{\ensuremath{\mbox{\%}}} Solving equation with given parametres:
  % n=3, k1p = 8; k2p = {2,4,6}
20
  % Initialization of variables
  n = 3;
  k1 = 8;
k2 = [2 4 6];
23
24
  x_val = [0:0.01:10];
25
26
27
  % Defining the range of X and Y
29
  [vf_x, vf_y] = meshgrid(0:0.45:15,0:0.20:10);
30
  \mbox{\ensuremath{\mbox{\%}}} Plotting solutions on a graph
31
32
  for i=1:length(k2)
33
34
      % Plotting the vector field
35
      quiver(vf_x, vf_y,dx(k1, n, vf_x, vf_y), dy(k2(i), vf_x, vf_y), 2);
36
37
38
      \mbox{\ensuremath{\mbox{\%}}} Find intersection points for the defined functions (X,Y)
39
      [xi, yi] = polyxpoly(x_val, nx(k1, n, x_val), x_val, ny(x_val, k2(i)));
41
42
      \mbox{\ensuremath{\mbox{\%}}} Plotting the nulclines
      plot(x_val, nx(k1, n, x_val), x_val, ny(x_val, k2(i)));
43
44
      % Plot intersection of graphs
plot(xi, yi, 'or', 'MarkerSize',10);
45
46
48
      xlabel('[X]');
      ylabel('[Y]');
49
50
51
      % Setting the new axis
      axis([0 max(x_val) 0 10]);
52
53
54
55
      % Export graph in a pdf file
print('-dpdf', ['plot_B' num2str(i)]);
56
57
58
60
  % Problem [C]
  62
63
  % Defining the equations to solve
64
65
  e2 = 'y<sub>\u=\uk2</sub>/(1+x)';
e3 = 'diff(k1/(1+y^3),<sub>\uk2</sub>)*diff(k2/(1+x),x)<sub>\u=1</sub>';
68
69
70
  % Solving system of equations
  S = solve(e1, e2, e3, 'y', 'k1', 'k2');
72
73
74
75
  % Retriving first solution for each variable
```

```
76 | k1 = S.k1(1);
   k2 = S.k2(1);
   y = S.y(1);
79
   \% Defining the range of X
81
   x = [0:0.01:5];
82
83
84
   % Solve the equations in the range of x
   k1_solv = x.*subs(k1);
86
87
   k2\_solv = x.*subs(k2);
88
89
   % Problem [D]
   92
93
94
   % Plotting of the solutions from the previous step
   plot(k1_solv, k2_solv);
xlabel('[k1]');
ylabel('[k2]');
95
96
97
98
99
   \mbox{\ensuremath{\mbox{\%}}} Setting the new axis
100
101
   axis([0 max(k1_solv) 0 max(k2_solv)]);
102
   % Export graph in a pdf file
104 print('-dpdf', 'plot_D0');
```

Listing 1: ./synthbio_script_01.m

0.6 dx.m

```
function [ dx ] = dx( k1, n, x, y )

dx = (k1./(1+(y.^n))) - x;
end
```

Listing 2: ./dx.m

0.7 dy.m

```
function [ dy ] = dy( k2, x, y )

dy = (k2./(1+x))-y;
end
```

Listing 3: ./dy.m

0.8 nx.m

```
function [y] = nx(k1, n, x)

y = nthroot((k1-x)./x, n);
end
```

Listing 4: ./nx.m

0.9 ny.m

```
function [ y ] = ny( x, k2 )

y = k2./(1+x);
end
```

Listing 5: ./ny.m

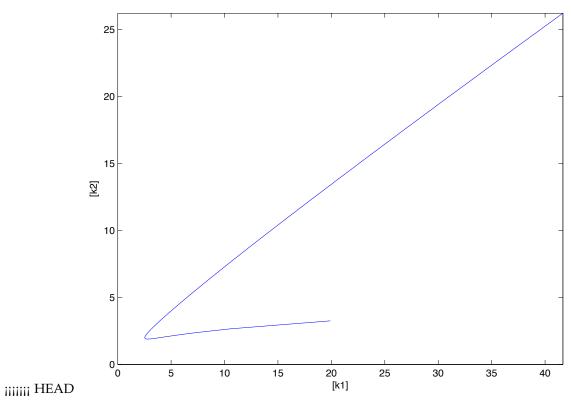


Figure 0.2: Values of the parameter k'_1 and k'_2 needed for a working switch for particular values of the exponent n in the Hill function.

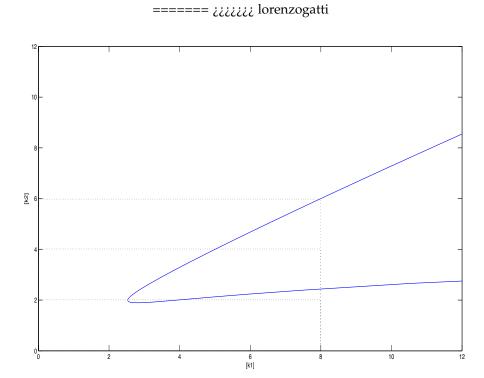


Figure 0.3: Values of the parameter k'_1 and k'_2 needed for a working switch for particular values of the exponent n in the Hill function.

Bibliography

Joshua L Cherry and Frederick R Adler. How to make a biological switch. *Journal of Theoretical Biology*, 203(2):117 – 133, 2000. ISSN 0022-5193. doi: 10.1006/jtbi.2000.1068. URL http://www.sciencedirect.com/science/article/pii/S0022519300910683.