## Metabolic and regulatory networks

## **Computer practical course**

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Day 1

## **Exercice 1**

$$v = \frac{v_{\text{max}} \cdot S}{(K_m + S)(1 + \frac{S}{K_i})}$$

```
% Defining symbols
syms v(S) vmax S Km Ki

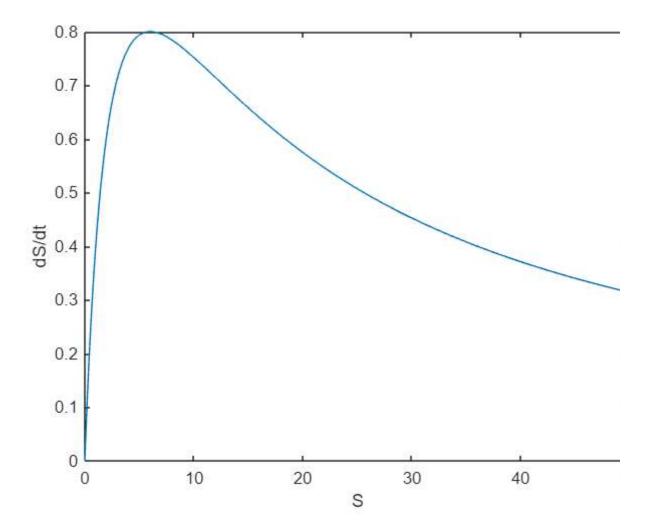
% Excluding negative values for S since negative concentrations are
% biologically irrelevant
assume(S >= 0)

% Defining the differential equation
v(S) = (vmax*S)/((Km+S)*(1+S/Ki))
```

$$v(S) = \frac{S \text{ vmax}}{\left(\frac{S}{\text{Ki}} + 1\right) (\text{Km} + S)}$$

a) Plotting for  $K_m = 9$ ,  $v_{\text{max}} = 5$ ,  $K_i = 4$ 

```
fplot(subs(v, {Km,vmax,Ki},{9,5,4}), [0,50])
xlabel('S');
ylabel('dS/dt')
```



b) Find S such that  $\frac{dS}{dt}$  is maximal for the given parameters

→from the plot it is apparent that there is only one extreme which is the maximum

c) Find S such that  $\frac{dS}{dt}$  is maximal for any parameters

```
solve(diff(v) == 0, S)
```

Warning: Solutions are only valid under certain conditions. To include parameters and condition ans =  $\sqrt{Ki\,Km}$ 

## **Exercice 2**

a) Perform a regression analysis for hypothtical experimental data by fitting the Michaelis-Mentenequation

v\_values = 1×8 0.0550 0.1210 0.2400 0.4250 0.4560 0.5220 0.5840 0.8050

Km 2.9063

vmax 0.9402

Goodness of Fit:

R-square 0.9620

b) Perform a linear regression analysis in a Lineweaver-Burk-plot

rec\_S\_values = 1./S\_values

rec\_S\_values = 1×8

10.0000 3.3333 1.0000 0.5000 0.3448 0.2439 0.1429 0.1000

rec\_v\_values = 1./v\_values

rec\_v\_values = 1×8

18.1818 8.2645 4.1667 2.3529 2.1930 1.9157 1.7123 1.2422

Km 0.9871

vmax 0.5862

Goodness of Fit

R-square 0.9907

- → We observe huge differences in the resulting values
- c) Excliding an outlier

Excluding the last measuring point changes the results the most, so it is the least consistent with the rest of the data

For a):

Km 1.6853

vmax 0.7312

Goodness of Fit

R-square 0.9912

For b):

Km 1.4838

vmax 0.7051

Goodness of Fit

R-square 0.9831