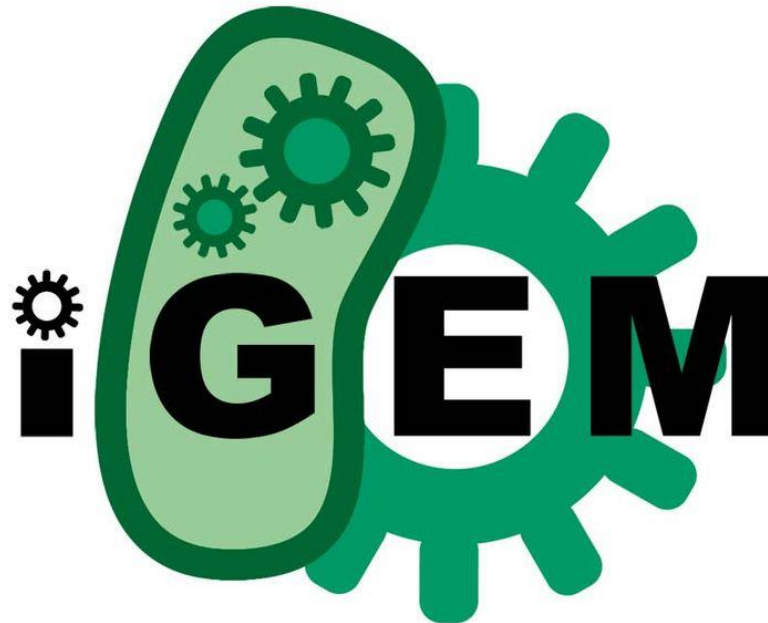


Deterministic model

iGEM2013



Why do we need it?



- To see how the mean of the population will act.
- To understand how the system behaves.
- To find out what need to be changed.
- To improve the circuit.
- To find the parameters.
-
-

How do we do it?

- ➔ Before learning how a deterministic model is done is important to understand a few concepts...

Optimization

Screening

Assumptions

Michaelis Menten

Reaction rate

Hill's equation

Equilibrium

Cooperation

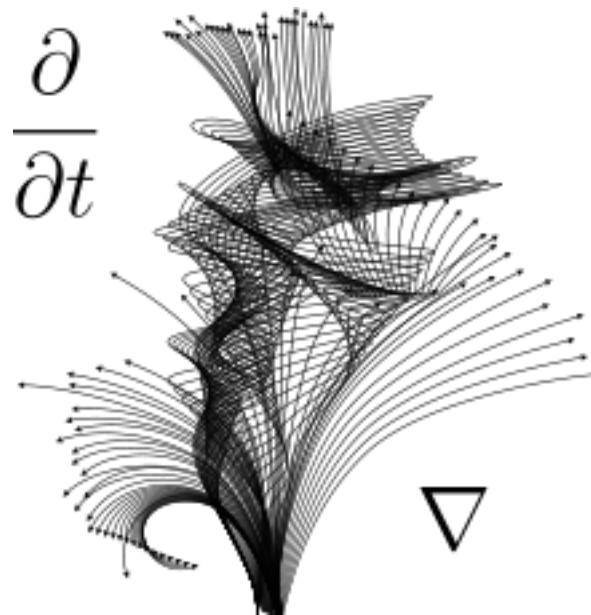
Hysteresis

Sensitivity analysis

Mass action law

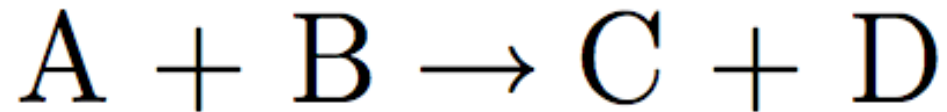
How do we do it?

- Finding an expression in terms of time and concentration usually is not easy.
- Expressions that define the change in time of the substances are easier to find.



How do we do it?

Change in time of the concentration of the molecules:

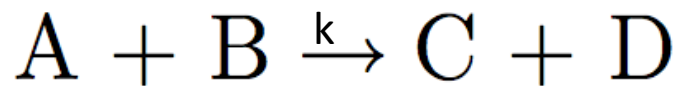


Reaction rate

$$\frac{dA}{dt}, \frac{dB}{dt}, \frac{dC}{dt}$$

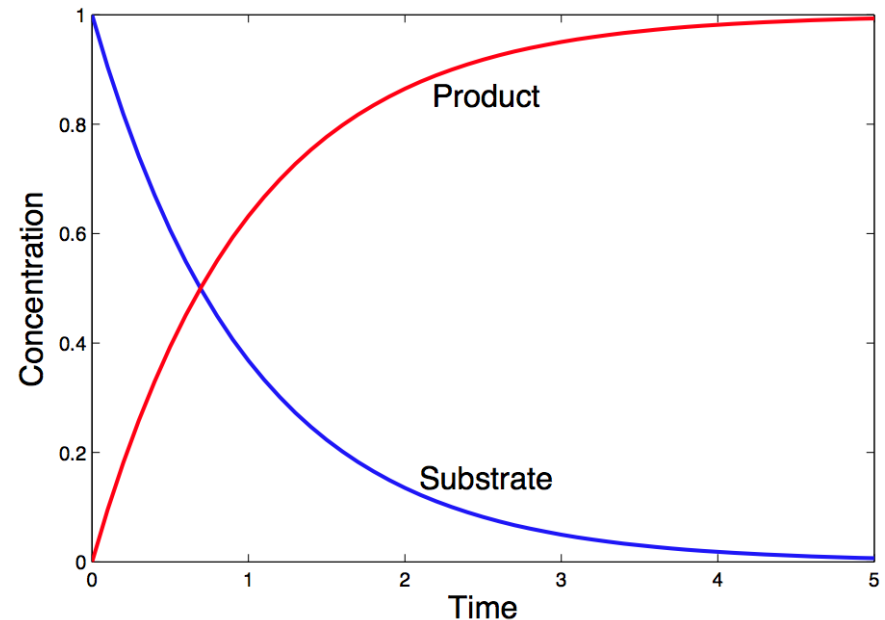
How do we do it?

Mass action law



$$\frac{dA}{dt} = \frac{dB}{dt} = -kAB$$

$$\frac{dC}{dt} = \frac{dD}{dt} = kAB$$



How do we do it?

Mass action law

More than one
reaction...

EXAMPLES!

How do we do it?

Michaelis Menten

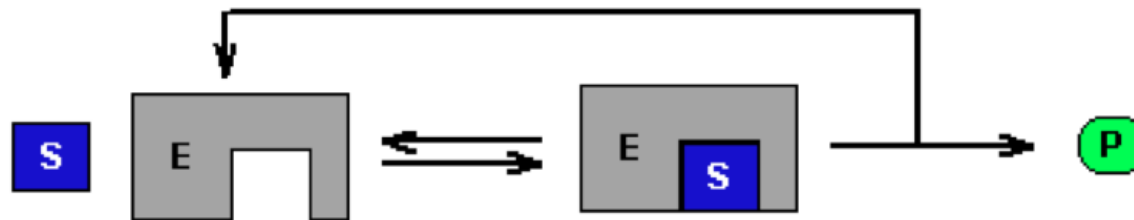
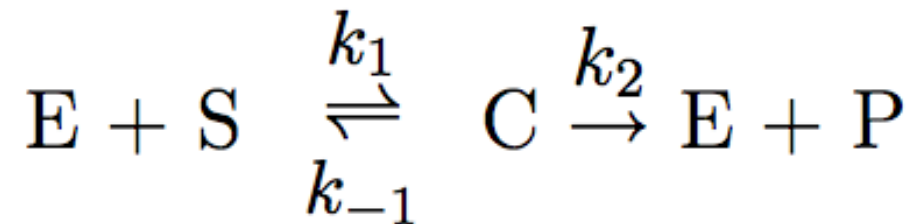


Figure 8: Michaelis-Menten mechanism.



How do we do it?

Michaelis Menten

$$\left\{ \begin{array}{l} \frac{dS}{dt} = -k_1ES + k_{-1}C \\ \frac{dE}{dt} = -k_1ES + k_{-1}C + k_2C \\ \frac{dC}{dt} = k_1ES - k_{-1}C - k_2C \\ \frac{dP}{dt} = k_2C \end{array} \right.$$

How do we do it?

Assumptions...

$$C = \frac{E_T S}{\frac{k_{-1}}{k_1} + S}$$

$$\left\{ \begin{array}{l} \frac{dS}{dt} = -k_1 ES + k_{-1} C \\ \frac{dE}{dt} = -k_1 ES + k_{-1} C + k_2 C \\ \frac{dC}{dt} = k_1 ES - k_{-1} C - k_2 C \\ \frac{dP}{dt} = k_2 C \end{array} \right.$$

How do we do it?

Michaelis Menten

$$\frac{dP}{dt} = k_2C = V_{max} \frac{S}{K_S + S}$$

Hill's equation

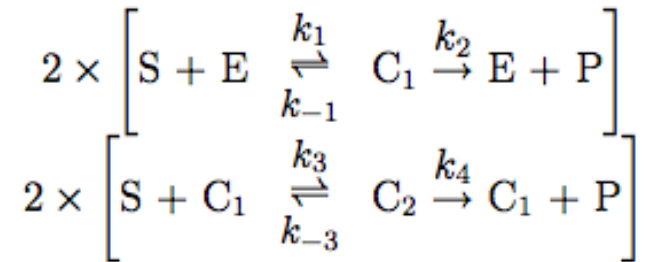
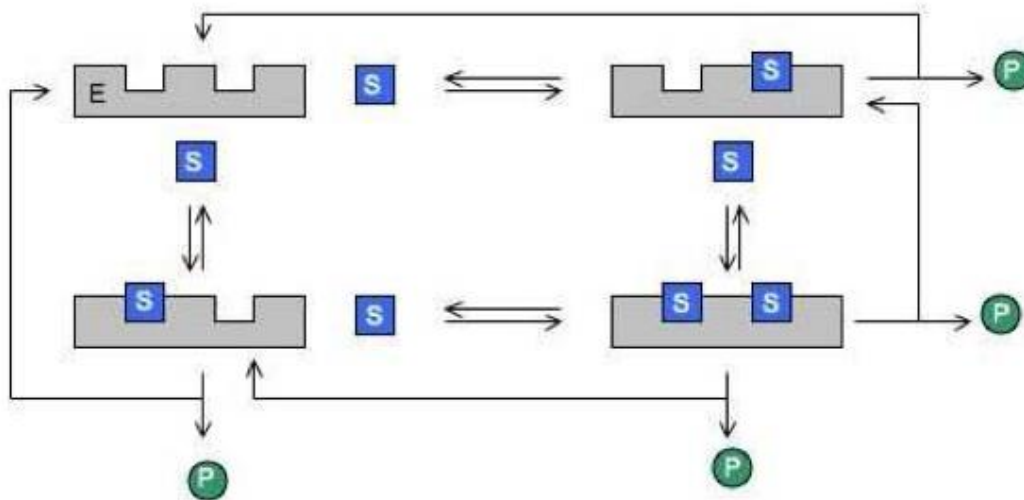
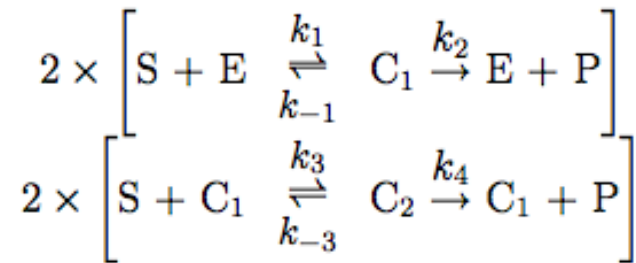


Figure 20: Enzyme with two binding sites: mechanism.

Cooperation

Hill's equation



$$\left\{ \begin{array}{l} \frac{dS}{dt} = 2(-k_1SE + k_{-1}C_1 - k_3SC_1 + k_{-3}C_2) \\ \frac{dC_1}{dt} = 2(k_1SE - (k_{-1} + k_2)C_1 - k_3SC_1 + (k_{-3} + k_4)C_2) \\ \frac{dC_2}{dt} = 2(k_3SC_1 - (k_{-3} + k_4)C_2) \end{array} \right.$$

Cooperation

How do we do it?

Assumptions...

1. $k_1, k_{-1} \gg k_2$
2. Steady state $\frac{dC_1}{dt} = \frac{dC_2}{dt} = 0$
3. $E_t = E_{\text{free}} + E_{c1} + E_{c2}$

How do we do it?

Assumptions...

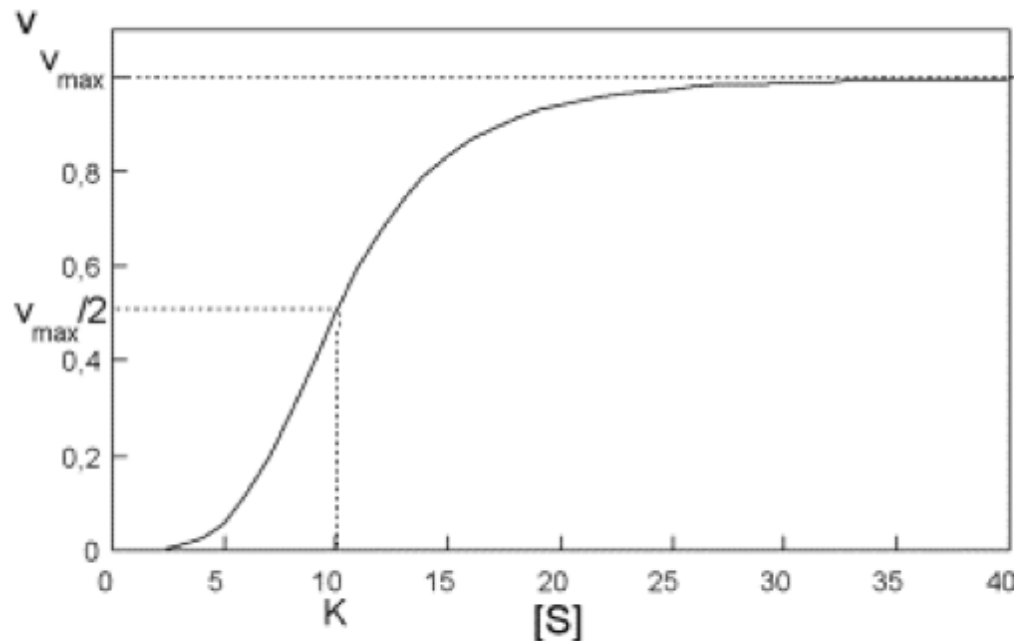
4. $k_1 = k_3 = k_+$
 $k_{-1} = k_{-3} = k_-$
 $k_2 = k_4 = k_p$

Binding sites independent.



How do we do it?

Hill's equation



$$v = V_{max} \frac{S^2}{K + S^2}$$

Figure 22: Hill kinetics.

How do we do it?

Hill's equation

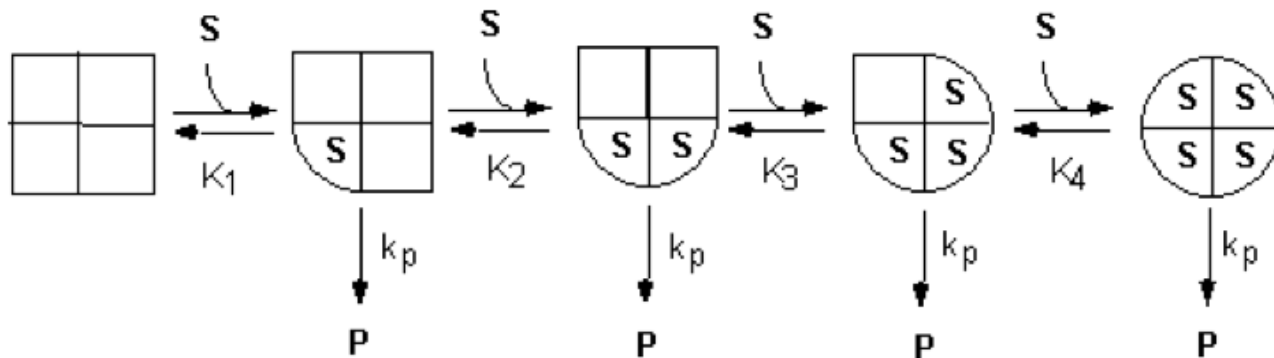
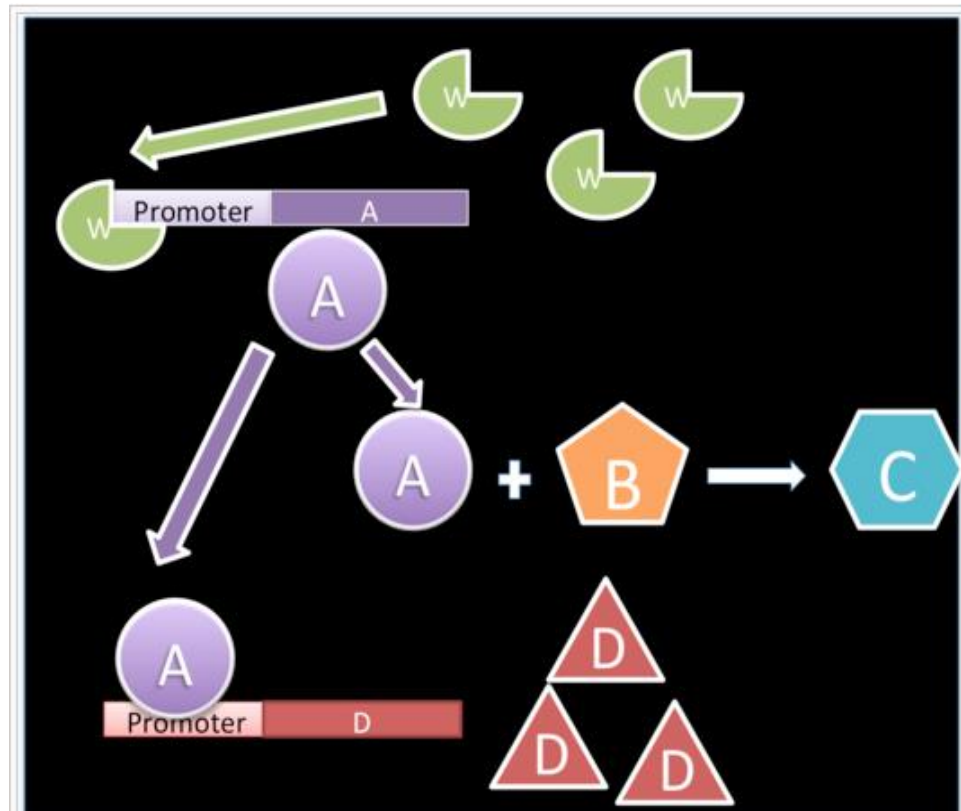


Figure 21: Cooperativity: mechanism.

$$v = V_{max} \frac{S^n}{K^n + S^n}$$

How do we do it?

Genetic Circuits



Accumulation = Input - Output + Production - Consumption

How do we do it?

THE PROBLEM:
THE PARAMETERS!!!!



Our strategy for choosing parameters

There are four main steps:

**Objective
function**

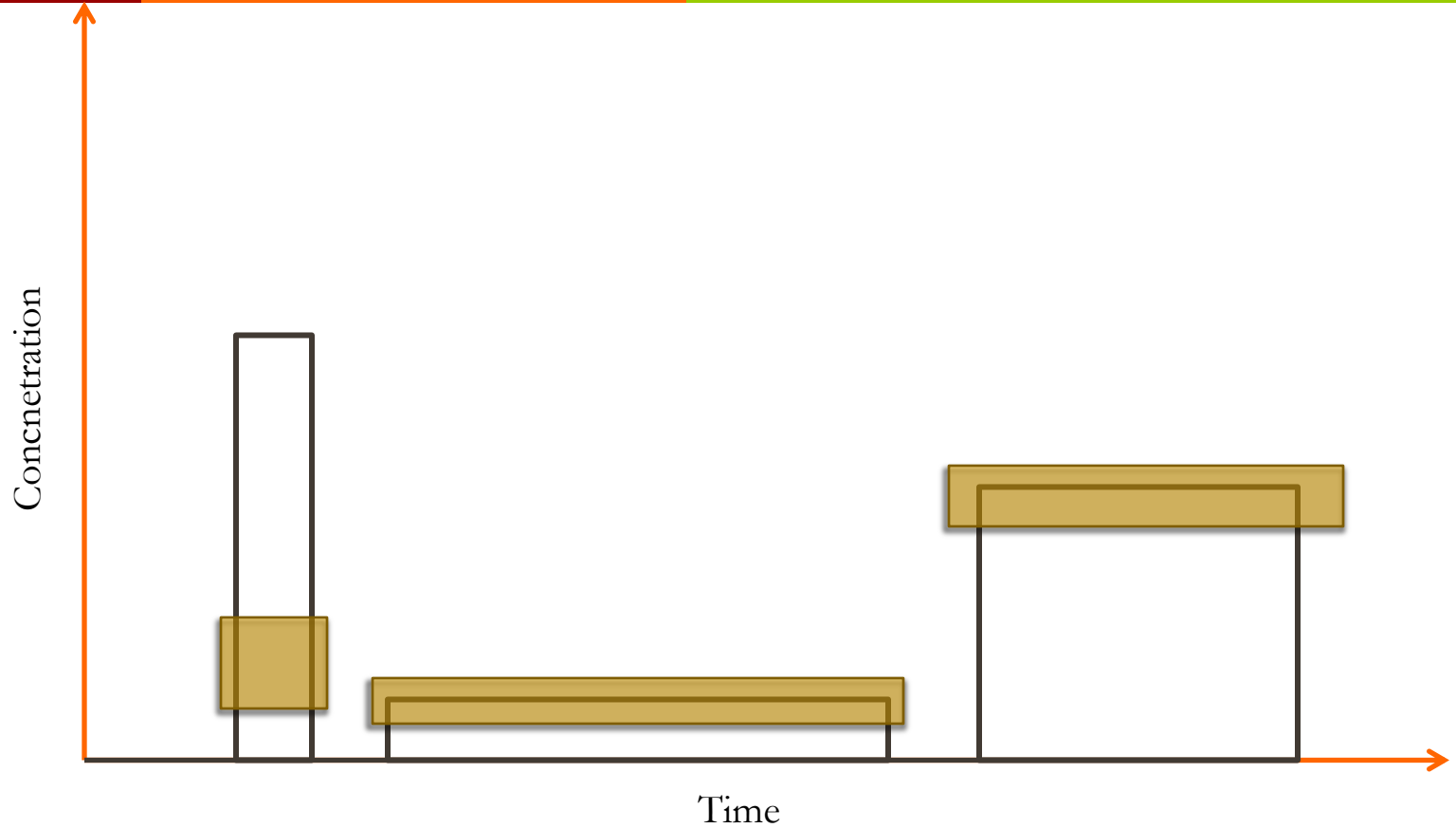
Optimization

**Sensitivity
Analysis**

Screening



Which set of parameters do we choose?



Introduction

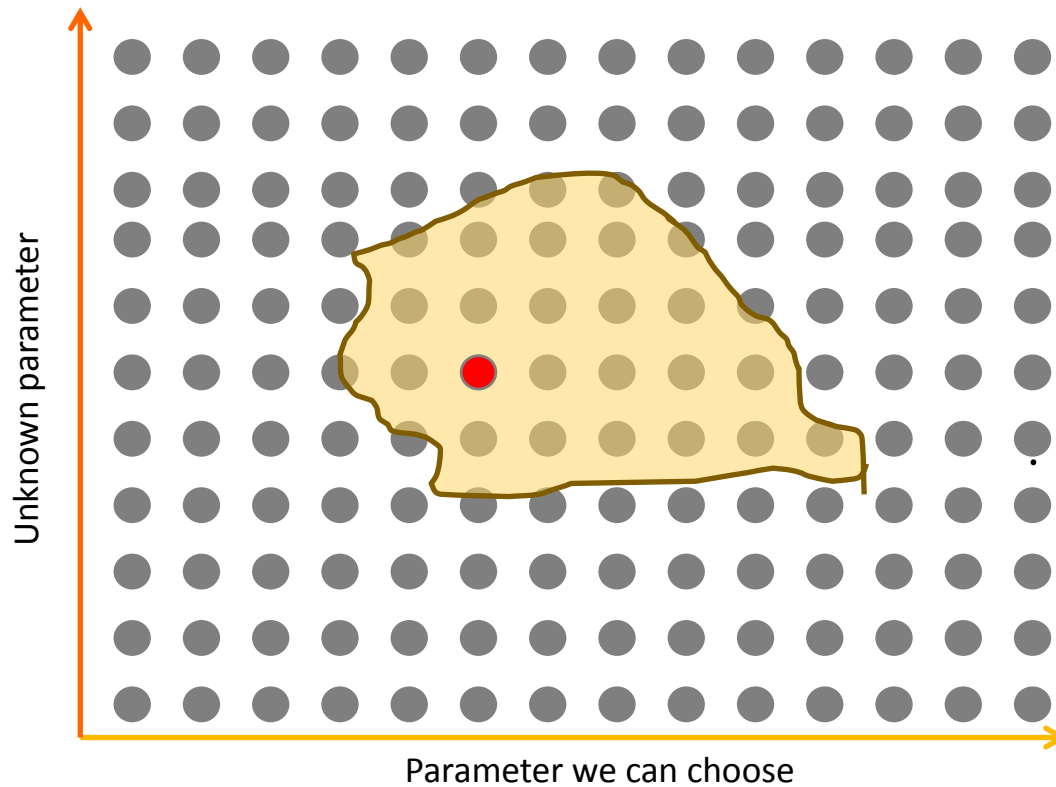
Models

Laboratory

Human Practices



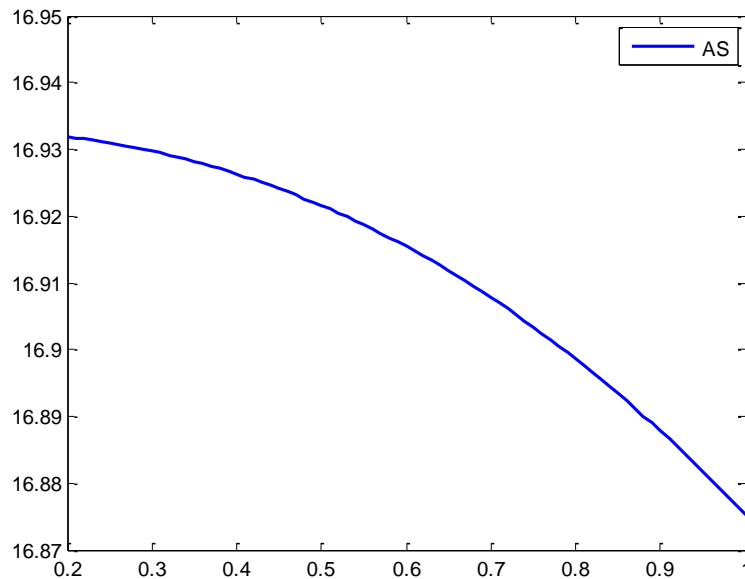
A point within the area



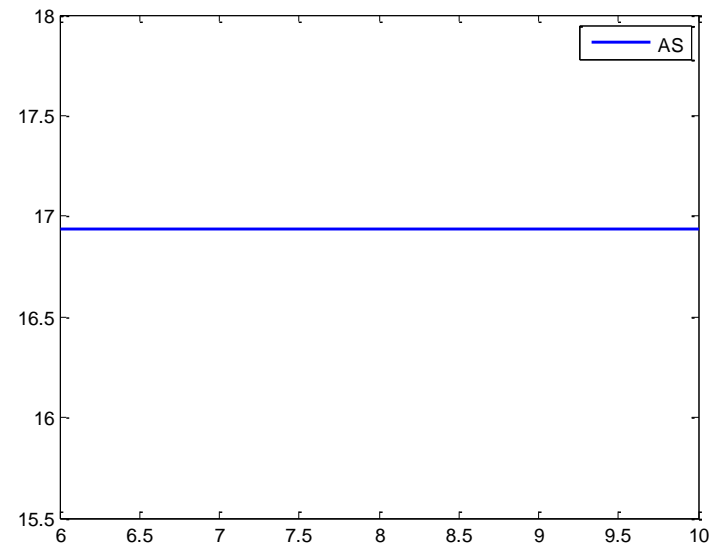


Sensitivity analysis

How does each parameter affect the model?



A sensitive parameter:
CI Hill constant

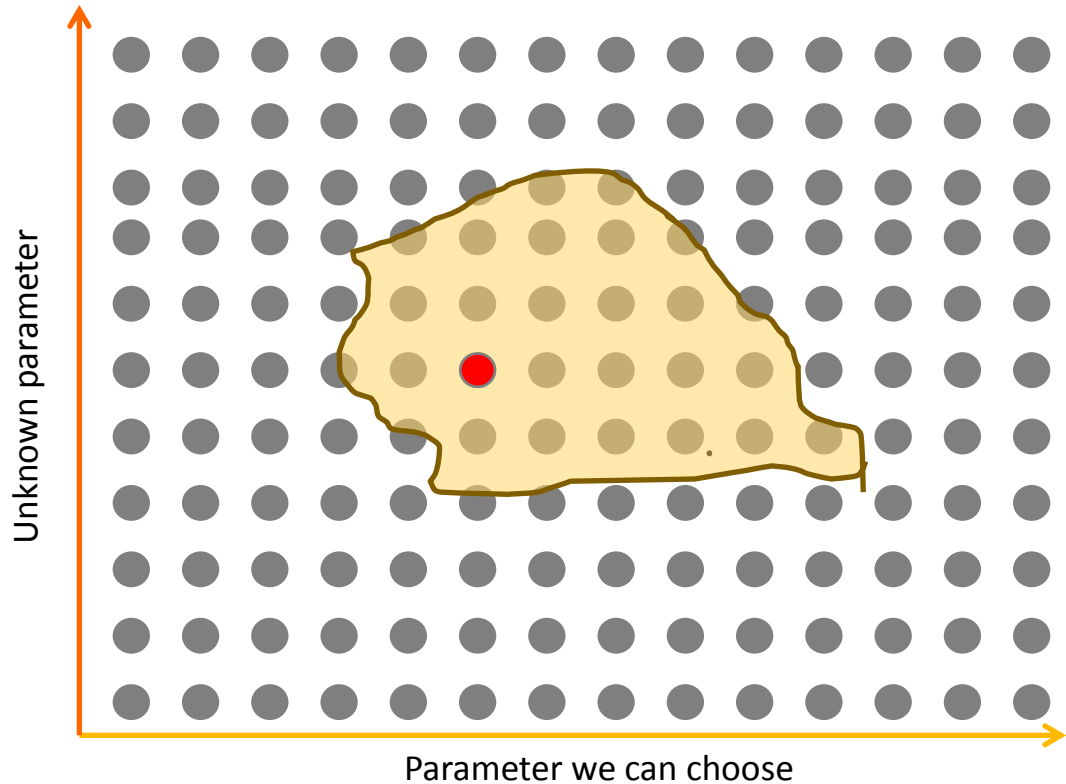


A non sensitive parameter:
Phosphorylation kinetic constant



The screening

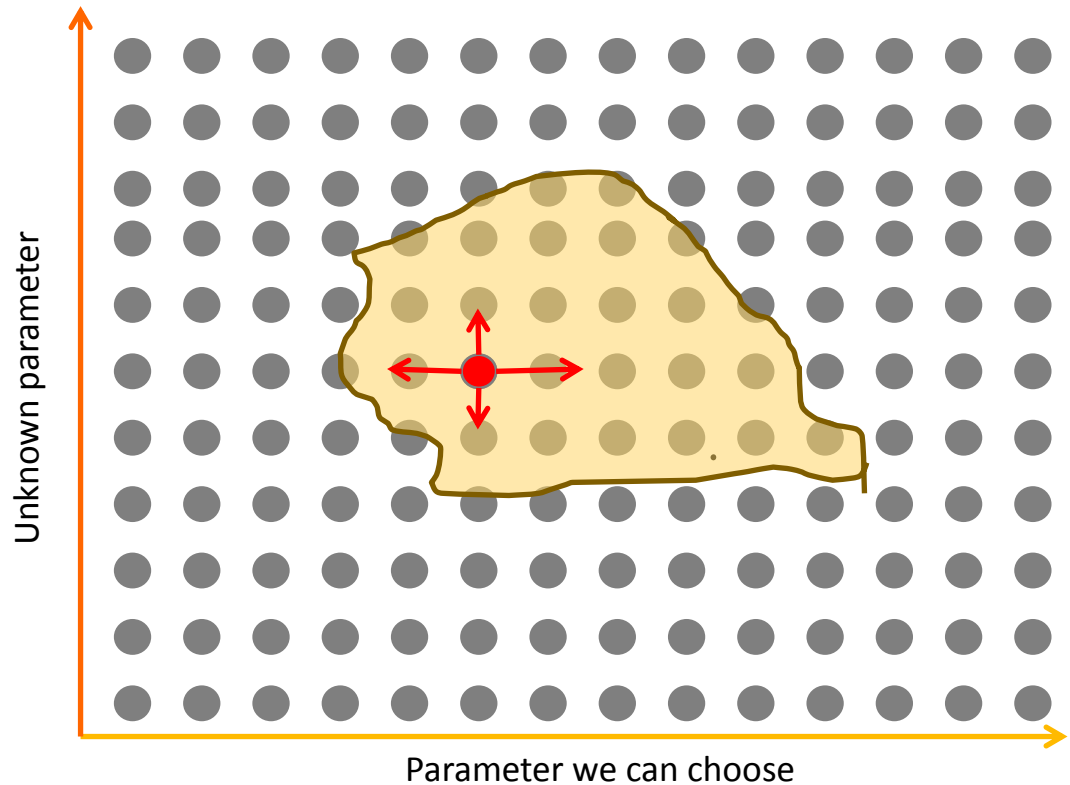
Look for the area that represents the sets of parameters that have the expected response.





The screening

Look for the area that represents the sets of parameters that have the expected response.





The screening

