

Modelling Biochemical Reaction Networks

Introductory lecture:

What to model?

Why?

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Recommended reading

- ▶ Fall, Marland, Wagner and Tyson, chapter 1

A bit of philosophical background

Popper: **Falsification** of hypotheses drives science forward.

Modelling as machinery for the falsification of mechanistic hypotheses

- ▶ We start with some observations we are trying to explain.
- ▶ Someone generates a hypothesis for a **mechanism** for the phenomenon.
- ▶ Mechanistic hypotheses can be converted to mathematical models.
- ▶ Does the model replicate the observations that the hypothesis was meant to explain?
- ▶ Does the model make any new predictions that could be tested experimentally?

Other reasons to make mathematical models

Discrimination between rival models

Exploration of phenomena not readily studied experimentally

▶ Exploration of **parameter space**

Reduction of a phenomenon to its essentials for further study

(Re)engineering of a process

Exploration of possible interventions

Biochemistry as a multiscale discipline

- ▶ Biochemical processes depend on and affect phenomena over a wide range of spatial and temporal scales
- ▶ Some relevant length scales:
 - Chemical bonds: 10^{-10} m
 - Macromolecular dimensions: 10^{-9} – 10^{-8} m
 - Length of a bacterium or of a mitochondrion: 10^{-6} m
 - Red blood cell diameter: 10^{-5} m
 - Neuron length: 10^{-3} – 1 m
- ▶ Some relevant time scales:
 - Time for Na^+ to transit through a channel: 10^{-8} s
 - Macromolecular conformational changes: 10^{-7} – 10^{-3} s
 - Transcription, translation: 10^1 – 10^4 s
 - Circadian rhythm: 10^5 s

Number of molecules

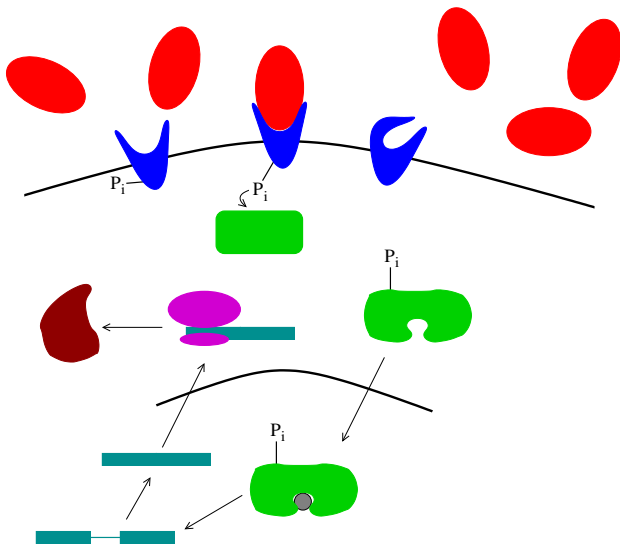
- ▶ Suppose that $[X] = 10 \mu\text{mol/L}$.
- ▶ How many molecules of X do we have?

V/L	Example	N_X
10^{-16}	Axon terminal	600
10^{-15}	Bacterium	6000
10^{-14}	Yeast cell	60 000
10^{-12}	Mammalian cell	6×10^6

Modelling biochemical systems

- ▶ You can't model everything completely.
- ▶ Many choices to make:
 - ▶ Is a qualitative model OK or do you want quantitative agreement?
 - ▶ Which physical part of the system (subcellular compartment, cell, group of cells, etc.) do you want to model?
 - ▶ Do you need to take the spatial dimension into account explicitly?
 - ▶ Do you need to explicitly model diffusive transport?
 - ▶ Is it OK to just treat the system as a set of coupled compartments?
 - ▶ What range of time scales do you need to cover?
 - ▶ What biochemical processes do you need to include?
At what level of detail?
 - ▶ Number of molecules:
continuous description (many molecules)
or stochastic (statistical; few molecules)?

Level of biochemical detail



This course

- ▶ Focus on **kinetics**
- ▶ Both differential equation (continuous) and stochastic models covered
- ▶ Compartmental descriptions of spatial effects only
- ▶ Emphasis on selecting the particular interactions to model, and the level of description required

Some central questions (some of which may not be resolved in this course)

- ▶ How do you decide if you have a “good” model?
- ▶ Past a certain level of complexity, we tend to rely heavily on computation.
How do we know if the results of a computation are correct?
- ▶ Since kinetic parameters are often difficult to get, is it OK just to get the right structure for a model?