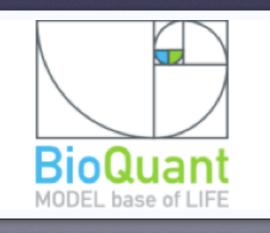
## In Silico Systems Biology

# Parameter Estimation with COPASI

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## Introduction

- When starting a modeling project usually many parameters of the model are not known
- How can I find out about the effect of parameter changes in the model?
  - Sensitivities / Metabolic control analysis
  - Parameter scans
- How can I find out about parameter values?

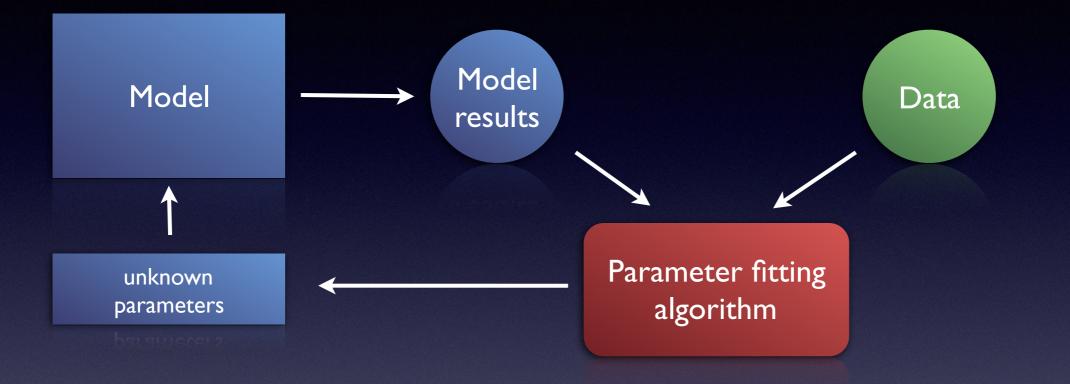
## Introduction

- simple approach: try to design an experiment for measuring the specific parameter
  - typically in vitro experiment
  - e.g. for rate constants: put different amounts of substrate in a test tube and measure how fast the reaction proceeds
- Problems:
  - often not possible, different from *in vitro* conditions

## Introduction

- Systems biology approach: adapt a complete model to experimental data
  - indirect method: use the model to find out about a parameter by measuring something else
  - can also be used to answer more complex questions, such as model identification
  - more difficult

## Basic idea of parameter fitting



• Change the parameters of a model so that it ,,best" fits the data

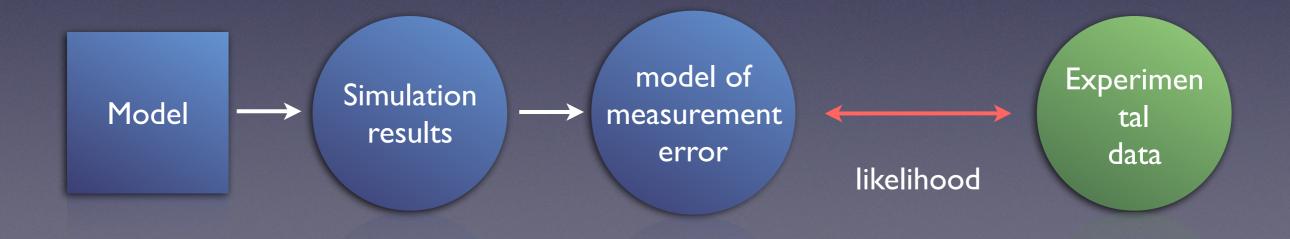
## Criteria for parameter fitting

- What is the "best" fit for a given set of data?
  - This is a mathematically difficult question
  - Fortunately, heuristically there are simple solutions.

- Required information:
  - Model
  - Knowledge about measurement process

## Maximum Likelihood

- In principle, if we know the model and the measurement process, we can calculate the probability that the the measurements would be the result of a simulation of the model.
- If this probability is high, the model is good.



## Maximum Likelihood

- To do parameter fitting, we need an algorithm that changes the unknown parameters so that the likelihood becomes maximal.
- This is mathematically very nice, but the probality is difficult to calculate in realistic cases.

## Least squares method

- If we make some assumptions about the measurement errors, we can find a rather simpler criterion:
- Assumption: Error follows a normal distribution, measurement error is uncorrelated
- Leads to Criterion: Likelihood is maximal when the difference (as defined on the next slide) between measurements and simulation results is minimal
  - This is easy to calculate but not the most general case.

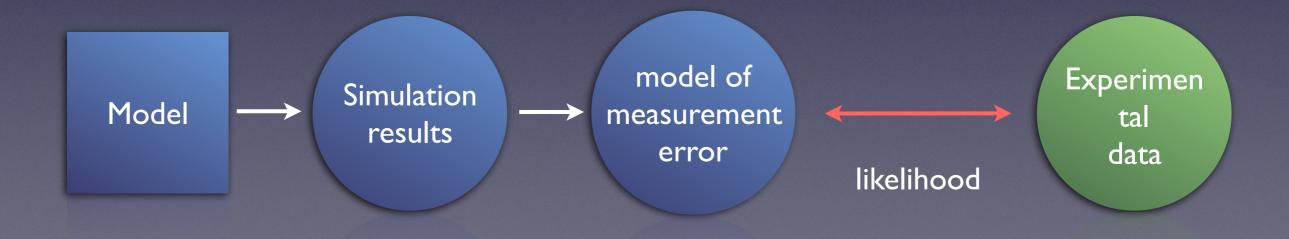
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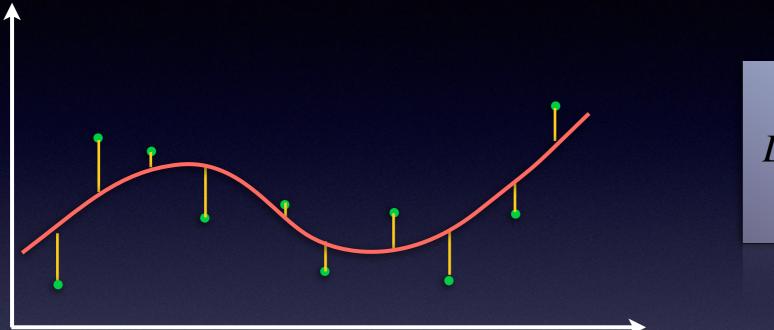
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  - This is easy to calculate but not the most general case. But it still works.

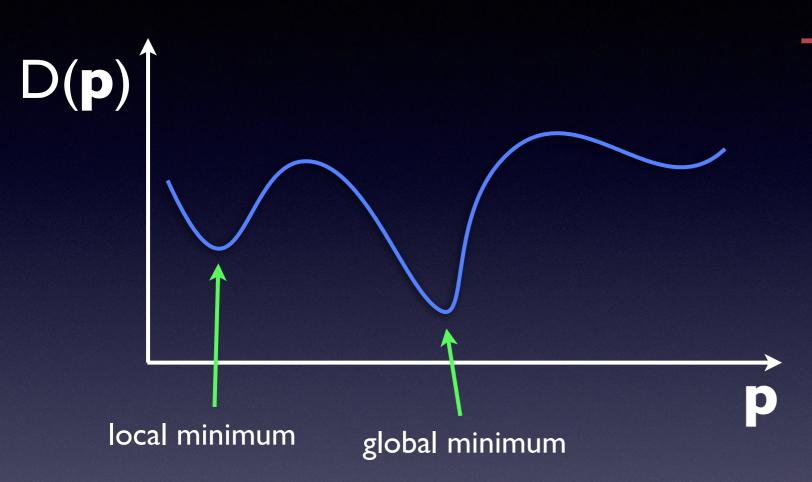
## Least squares distance measure



$$D(p) = \sum_{i=1}^{N} (x_i - y_i(p))^2$$

#### $x_i$ : measured values for time $t_i$ $y_i(p)$ : simulated values for time $t_i$ , parameter p

## The target function



## The function is usually high dimensional!

- This is the function that needs to be minimized.
- usually has many local minima

## Parameter space

- For a complete specification of the parameter fitting problem we still need to specify the unknown parameters:
- List of parameters with allowed range of values
- These parameters span an M-dimensional space, the *parameter space*.
- One specific set of parameters corresponds to a point in parameter space

## Optimization problem

- We now need a way to find the set of parameter values (a point in parameter space) for which the distance D is minimal (the best fit).
- A systematic scan of the parameter space is not possible when the dimensionality is large (many unknown parameters)
- Example: 10 parameters with 10 values each: 10<sup>10</sup> evaluations. Even if we can do 100 simulations/s, it would take 3 years.

## Optimization algorithms

- In general it is very difficult to find the parameter values for which D(p) is minimal
- It can be shown that there is no optimal optimization algorithm for all cases (this means there is no way to decide which one is best for a given problem)
- This means you should always try several algorithms for difficult parameter estimation problems.

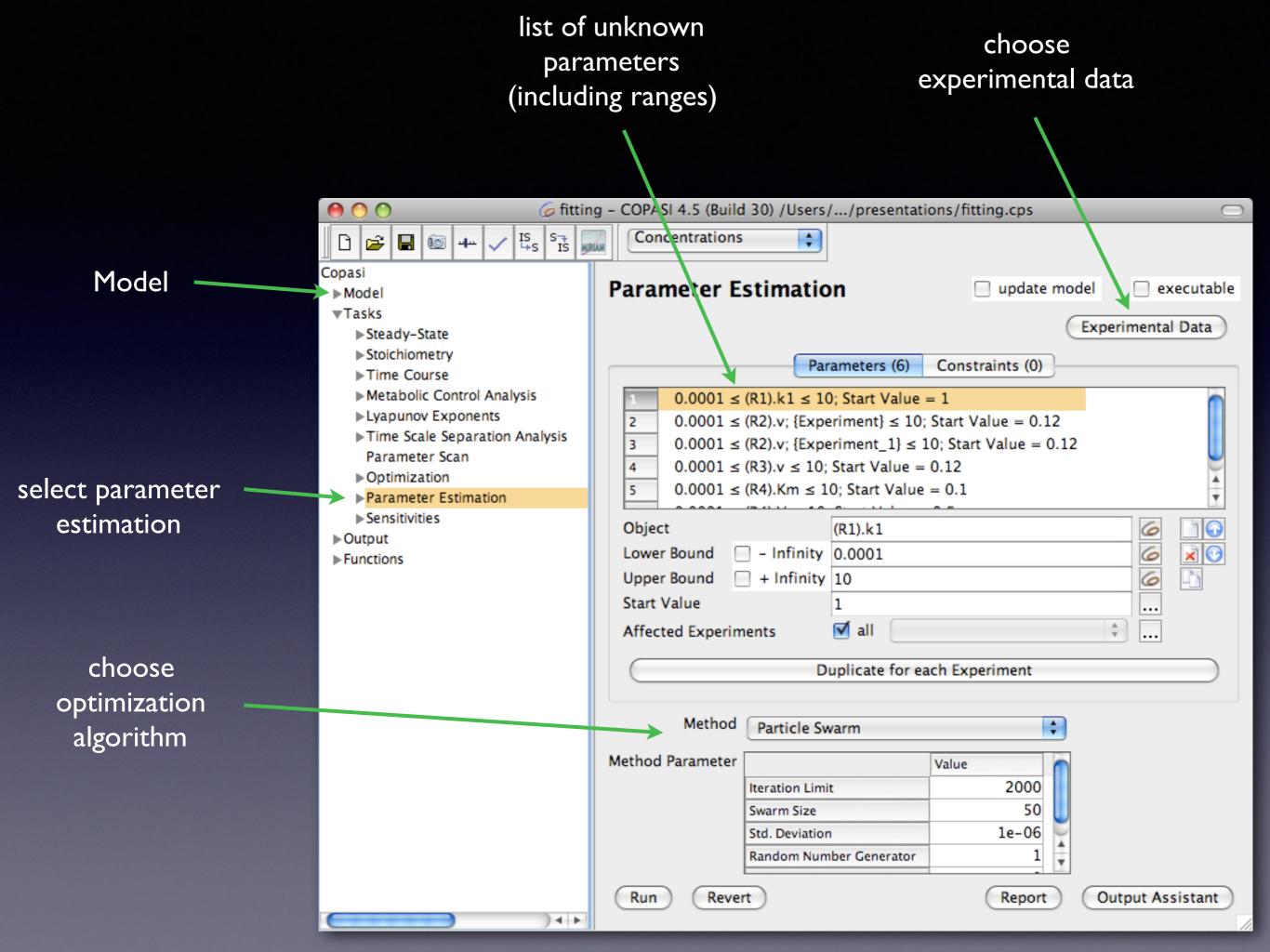
## **Optimization Algorithms**

- Based on derivatives
  - Steepest descent
  - Newton
  - Levenberg-Marquardt
- Using geometry
  - Nelder-Mead (simplex)
  - Hooke-Jeeves

- Based on genetics
  - Genetic algorithm
  - Evolutionary programing
  - Evolution strategy
- Other stochastic
  - random search
  - particle swarm
  - simulated annealing

# Specification of a parameter estimation problem

- What kind of information is needed for the computer to do a parameter estimation?
  - The model
  - the experimental data
  - the mapping between experimental data and model simulation results
  - the ranges of possible values for the unknown parameters
  - the optimization algorithm



possibly several experiments per file

	$\Theta \bigcirc \Theta$	\varTheta 🔿 🙆 Experimental Data		
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	fitting2.txt		periment	
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## Important:

- The result of a parameter fitting always needs to be inspected afterwards!
- Having a good result for a fit does **not** mean that the parameter value is the "true" one. This depends on the assumptions about the errors and the correctness of the model.
- For the stochastic algorithms the result is not reproducible!

# Using several experiments for parameter estimation

- The more data available, the better.
  - So if you have data from several experiments it should be used for parameter estimation simultaneously
- COPASI can deal with an arbitrary number of experiments, also of different kinds (combined time course/steady state, different variables, different time points, etc.)

### several experiments...

- Adding data from several experiments is straight forward in COPASI. Several data files can be specified and each can contain several experiments
- Important information: What is the same for all experiments and what is different between experiments? For the things that are different, are they known or unknown?

### several experiments...

- Simplest case: Repeated experiments.
  - nothing special needs to be done in COPASI
- Several experiments under different conditions. The conditions are known.
  - Example: Different stimulations in several experiments.
  - In COPASI: The stimulation needs to be a parameter in the model. In the experimental data specification this value is selected as an *independent* parameter. *Independent* data is known data that is provided in the data file. Dependent data is data that is used for fitting.

### several experiments...

- Experiments where some conditions are different, but not known
  - Example: *In vivo* experiments, even if the experiment is repeated with the same preparation, the initial conditions (inside the organism) are typically different.
  - In COPASI: The user can specify that some parameters are fitted for all experiments, and some are fitted for a specific subset of experiments.