Identification of cell signaling pathways based on biochemical reaction kinetics repositories

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Introduction

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Understanding the functioning of cell signaling is important in many biological areas.

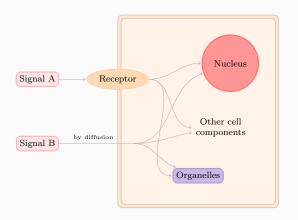


Figura 1: A general cell signaling mechanism.

Cell Signaling Pathways

A cell signaling network can be characterized by a sequence of chemical reactions that allows the presence of a signal to modify the state or behaviour of a cell.

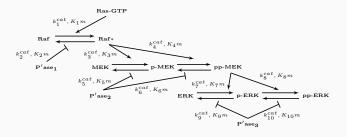


Figura 2: An example of a signaling pathway.

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Using biochemical and enzymatic kinetics, we can write equations that represent the rate of change of concentration for a chemical species.

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Repeating this procedure for all reactions of a pathway allows us to derive a system of ordinary differential equations that can model the signaling pathway.

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As the input, a description of a biological experiment and a set of experimental measurements are given. A possible output to the problem is composed by:

- a model composed by a set of chemical reactions that are relevant for the biological experiment;
- information about the reaction rate constants of the model.

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Hence, it is desirable to construct a method that can systematically modify these models and choose the one that better represents the experiment.

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On her work, the problem of identification of cell signaling pathways is treated as a feature selection problem.

Feature Selection Problem

The feature selection problem is a combinatorial optimization problem:

Given a set of features S and a cost function c, find subset $X \in \mathcal{P}(S)$, with minimum cost c(X).

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Figura 3: An example of feature selection search space with 5 features.

Feature Selection for Identification of Signaling Pathways

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Wu's Search Algorithm for Feature Selection

The search algorithm used by Wu is the Sequential Forward Selection (SFS).

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where R(M) is a regularization function.

Fundamental Concepts

Model Selection

Experiments on Model Selection

Next Steps