Computational approaches to the topological and dynamical features of biochemical networks

Adrián López García de Lomana

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Acknowledgements

Abstract

This is the abstract.

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Introduction

systems biology.

Chapter 2

Objectives

The objectives of this thesis are:

- Open a new research line within the group.
- Own a computational framework for the analysis of the dynamics of gene regulatory networks.
- Further knowledge on the quantitative features of biochemical networks.

Chapter 3

Results

3.1 Developmental Biology and Mathematics

Book chapter.

3.2 Topology of Cellular Networks

3.3 Computational Tools: `ByoDyn`

3.4 Applications of `ByoDyn`

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Chapter 4

Discussion

This is the discussion.

Chapter 5

Conclusions

These are the conclusions.