

Network analysis and visualization with Cytoscape and STRING

Campinas, April 3, 2019

Fernanda Luz P. Costa



Biocom

Overview

Introduction

- Network analysis in biology
- Types of biological networks
- Protein-protein interaction networks
- Building and analysing PPINs

Practice

- Getting familiar with the STRING database

Biocom

Network analysis in biology

- Biological systems are often represented as networks which are complex sets of binary interactions or relations between different entities;
- **Systems biology** aims to understand biological entities at the systemic level, analysing them not only as individual components, but also as **interacting systems** and their emergent properties;
- **Network biology** which allows the representation and analysis of biological systems using tools derived from graph theory.

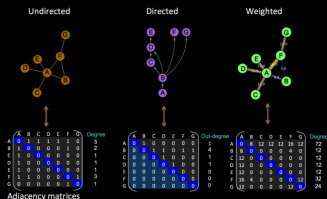


Figure: Example of molecular network types and its respective matrices.

Types of biological networks

- Protein-protein interaction networks
- Metabolic networks
- Genetic interaction networks
- Gene / transcriptional regulatory networks
- Cell signalling networks

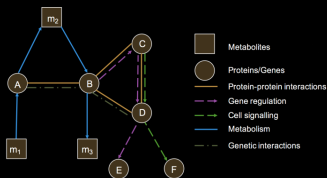


Figure: Example of molecular network[Dr. Natasa Przulj].

Adapted from [Millan, 2011]

Protein-protein interaction networks

Protein-protein interactions (PPIs) are essential to almost every process in a cell, so understanding PPIs is crucial for understanding cell physiology in normal and disease states. It is also essential in drug development, since drugs can affect PPIs.

Protein-protein interaction networks (PPIN) are mathematical representations of the physical contacts between proteins in the cell. These contacts:

- are specific;
- occur between defined binding regions in the proteins;
- and have a particular biological meaning.

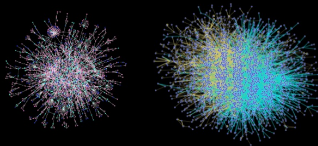


Figure: Yeast and human interactomes[Jeong et al., 2001][Rual et al., 2005].

Adapted from [Millan, 2011]

Building and analysing PPINs

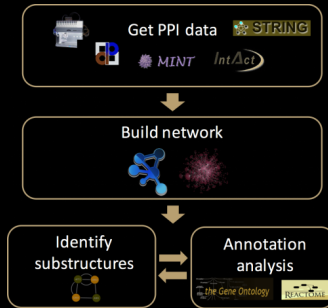


Figure: A potential workflow for building and analysing protein-protein interaction networks.

Adapted from [Millan, 2011]

Getting familiar with the STRING database

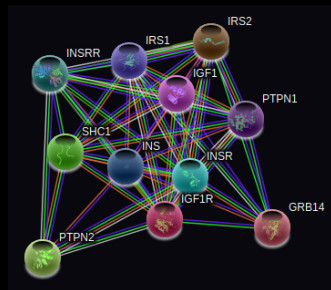
Basic Exercises – STRING database

- ex 1 – 4

Advanced Exercises – STRINGapp for Cytoscape

- ex 5 – 8

STRING [Doncheva et al., 2018]





Biocom

Tecnologia e ciência se encontram aqui

Oferecemos uma vasta gama de serviços especializados em consultoria e análise de dados biológicos. Auxiliamos na extração e interpretação de resultados obtidos a partir de dados biológicos gerados por tecnologias de Nova Geração.

consultoriabiocom.com.br

www.facebook.com/consultoriabiocom

www.linkedin.com/company/consultoriabiocom

References



Pablo P. Millan (2011)

Network analysis of protein interaction data: an introduction.
EMBL-EBI Train Online.



Doncheva NT, Morris JH, Gorodkin J and Jensen LJ (2018)

Cytoscape stringApp: Network analysis and visualization of proteomics data.



Jeong et al. (2001)

Nature, 411 (3).



Rual et al. (2005)

Nature, 437 (4).



Dr. Natasa Przulj

Imperial College, London.