ORYCTOLAGUS MULTIPLEX GPI NETWORK

Last update: 1 July 2014

Reference and Acknowledgments

This README file accompanies the dataset representing the multiplex genetic and protein interactions network of the Oryctolagus Cuniculus, also known as European rabbit. If you use this dataset in your work either for analysis or for visualization, you should acknowledge/cite the following papers:

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"Biogrid: a general repository for interaction datasets"

C. Stark, B.-J. Breitkreutz, T. Reguly, L. Boucher, A. Breitkreutz, and M. Tyers.

Nucleic Acids Research 2006 34 (1) D535-D539
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"MuxViz: A Tool for Multilayer Analysis and Visualization of Networks" Manlio De Domenico, Mason A. Porter, and Alex Arenas Journal of Complex Networks 2015 3 (2) 159-176

that can be found at the following URLs:

http://nar.oxfordjournals.org/content/34/suppl 1/D535.abstract

http://comnet.oxfordjournals.org/content/3/2/159

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Visit

PLEXMATH: http://www.plexmath.eu/

ALEPHSYS: http://deim.urv.cat/~alephsys/

for further details.

Description of the dataset

We consider different types of genetic interactions for organisms in the Biological General Repository for Interaction Datasets (BioGRID, thebiogrid.org), a public database that archives and disseminates genetic and protein interaction data from humans and model organisms. BioGRID currently includes more than 720,000 interactions that have been curated from both high-throughput data sets and individual focused studies using over 41,000 publications in the primary literature. We use BioGRID 3.2.108 (updated 1 Jan 2014).

The multiplex network used in the paper makes use of four layers corresponding to:

- 1. Direct interaction
- 2. Association
- 3. Physical association

There are 144 nodes, labelled with integer ID between 1 and 144, and 144 connections. The multiplex is directed and unweighted, stored as edges list in the file

oryctolagus_genetic_multiplex.edges

with format

layerID nodeID nodeID weight

(Note: weight is 1 for all edges)

The IDs of all layers are stored in

oryctolagus_genetic_layers.txt

The IDs of nodes, together with their name can be found in the file

oryctolagus_genetic_nodes.txt

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Contacts

If you find any error in the dataset or you have questions, please contact

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