# C.ELEGANS MULTIPLEX CONNECTOME

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## **Reference and Acknowledgments**

This README file accompanies the dataset representing the multiplex neuronal network of the nematode "Caenorhabditis Elegans". If you use this dataset in your work either for analysis or for visualization, you should acknowledge/cite the following papers:

"Wiring optimization can relate neuronal structure and function" Beth L. Chen, David H. Hall, and Dmitri B. Chklovskii PNAS 2006 103 (12) 4723-4728

"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://www.pnas.org/content/103/12/4723.abstract

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

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Visit

PLEXMATH: <a href="http://www.plexmath.eu/">http://www.plexmath.eu/</a>

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

for further details.

# Description of the dataset

Caenorhabditis elegans connectome, where the multiplex consists of layers corresponding to different synaptic junctions: electric ("ElectrJ"), chemical monadic ("MonoSyn"), and polyadic ("PolySyn").

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

- 1. Electric ("ElectrJ")
- 2. Chemical Monadic ("MonoSyn")

### 3. Chemical Polyadic ("PolySyn")

There are 279 nodes, labelled with integer ID between 1 and 279, and 5863 synaptic connections. The multiplex is undirected (with only one direction specified) and unweighted, stored as edges list in the file

celegans\_connectome\_multiplex.edges

with format

layerID nodeID nodeID weight

(Note: weight is 1 for all edges)

The IDs of all layers are stored in

celegans\_connectome\_layers.txt

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

celegans\_connectome\_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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