# CYTOSCAPE — SHORT TUTORIAL

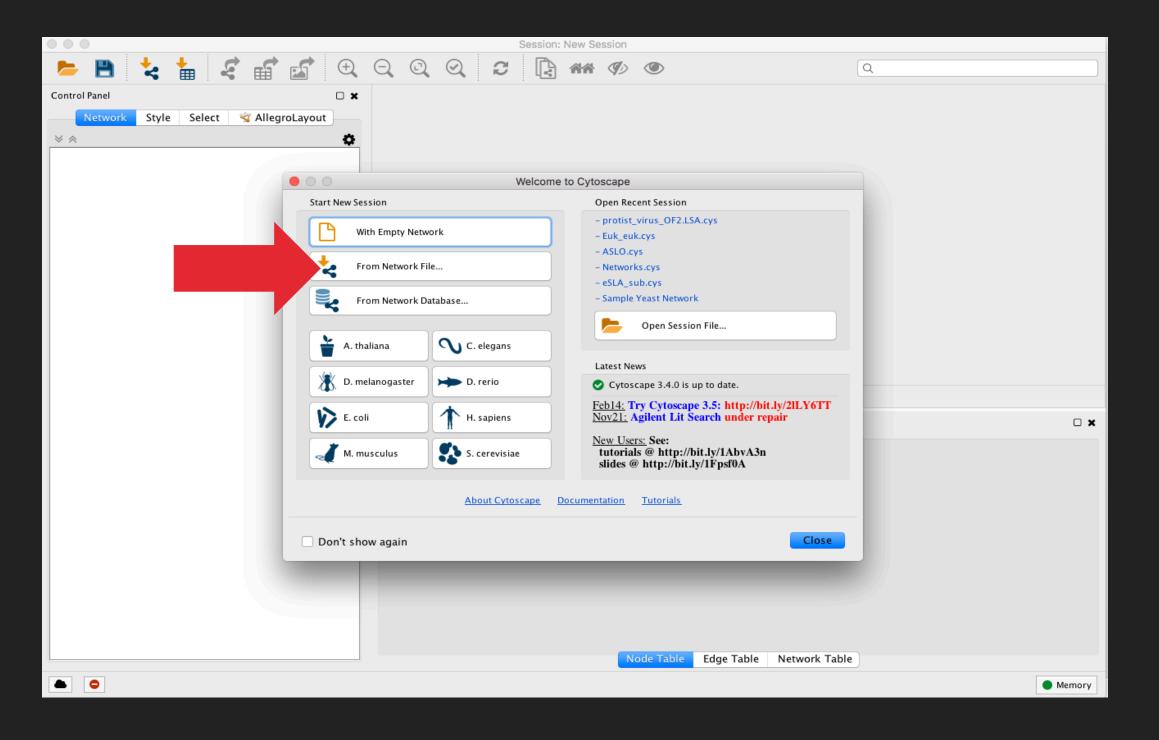
## THE NETWORK FILES

- Intro to Cytoscape (https://cytoscape.org/)
- Files on https://github.com/krabberod/BIO9905MERG1\_V21/tree/main/Networks
- The commands for a typical run
  - eLSA\_starting\_script.sh
- Input data
  - eLSA\_otu\_table.tsv
  - eLSA\_top100\_otus.tsv

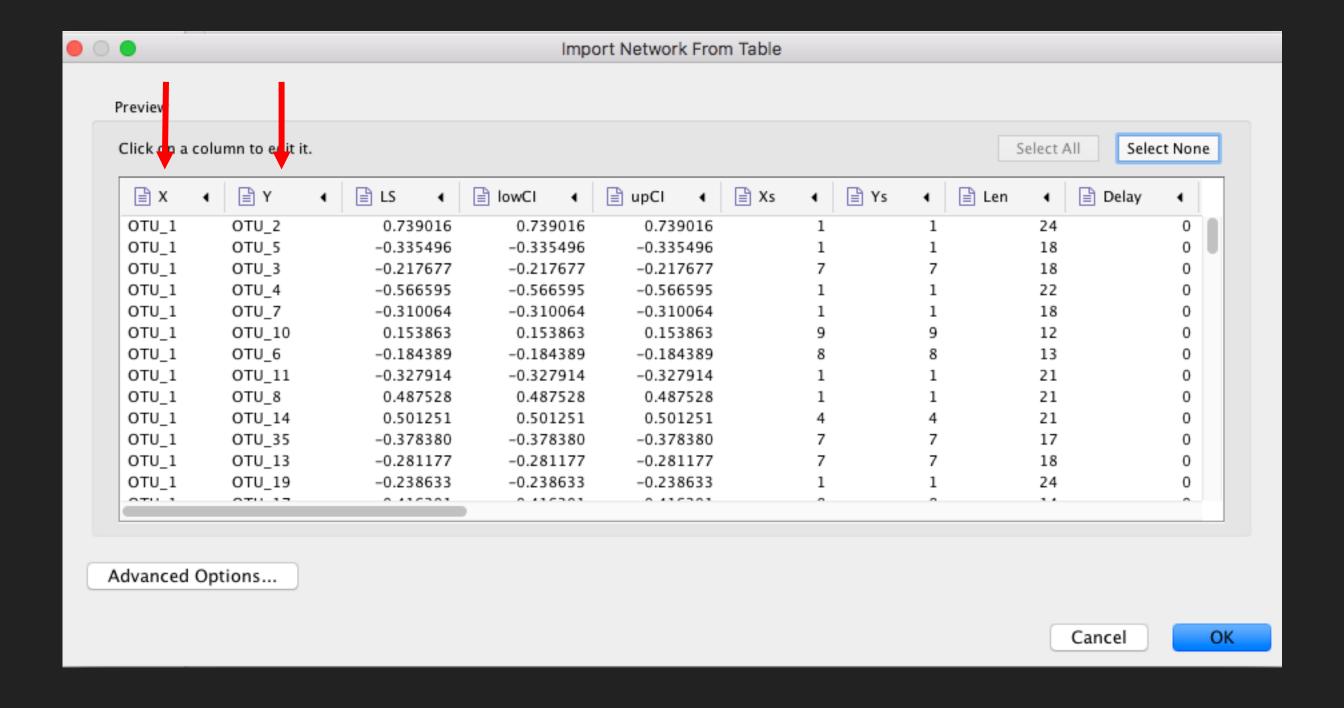
# THE NETWORK FILES

- Output from eLSA
  - eLSA\_network\_top100.perm.d0.tsv eLSA\_network\_top100.perm.d1.tsv eLSA\_network.d0.tsv eLSA\_network.d1.tsv
- eLSA\_node\_annotation\_relabund.tsv
- eLSA\_node\_annotation\_tax.tsv
- eLSA\_for\_cytoscape.cys (can be opened directly in cytoscape)

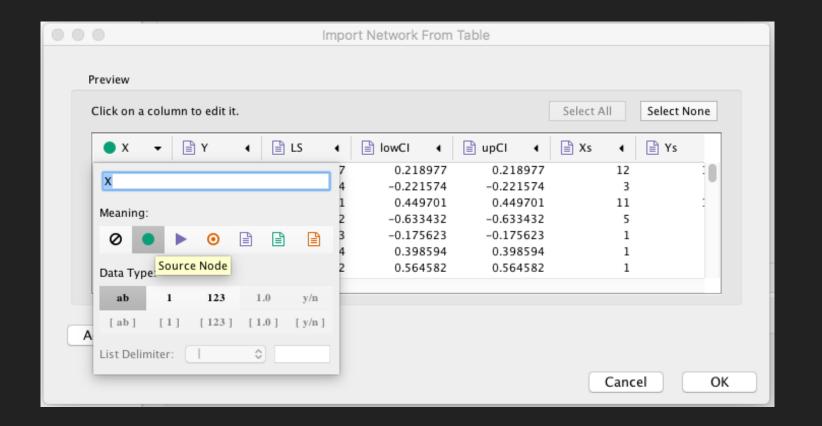
open Cytsocape, import eLSA\_network\_top100.perm.d0.tsv

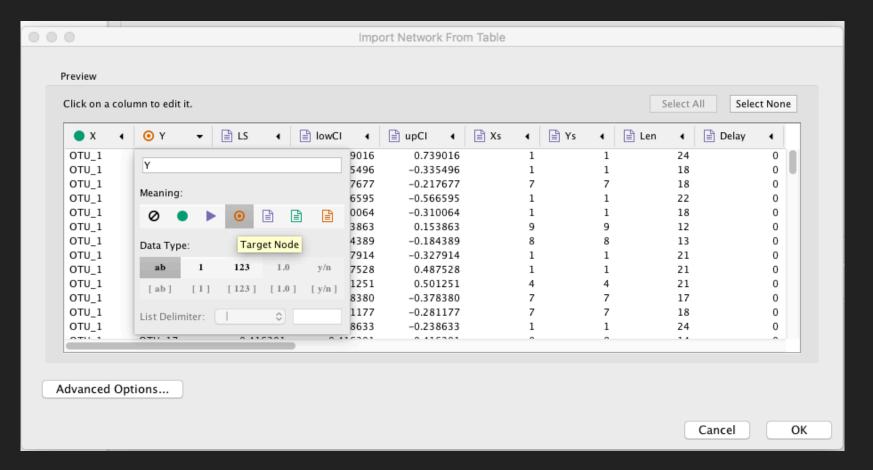


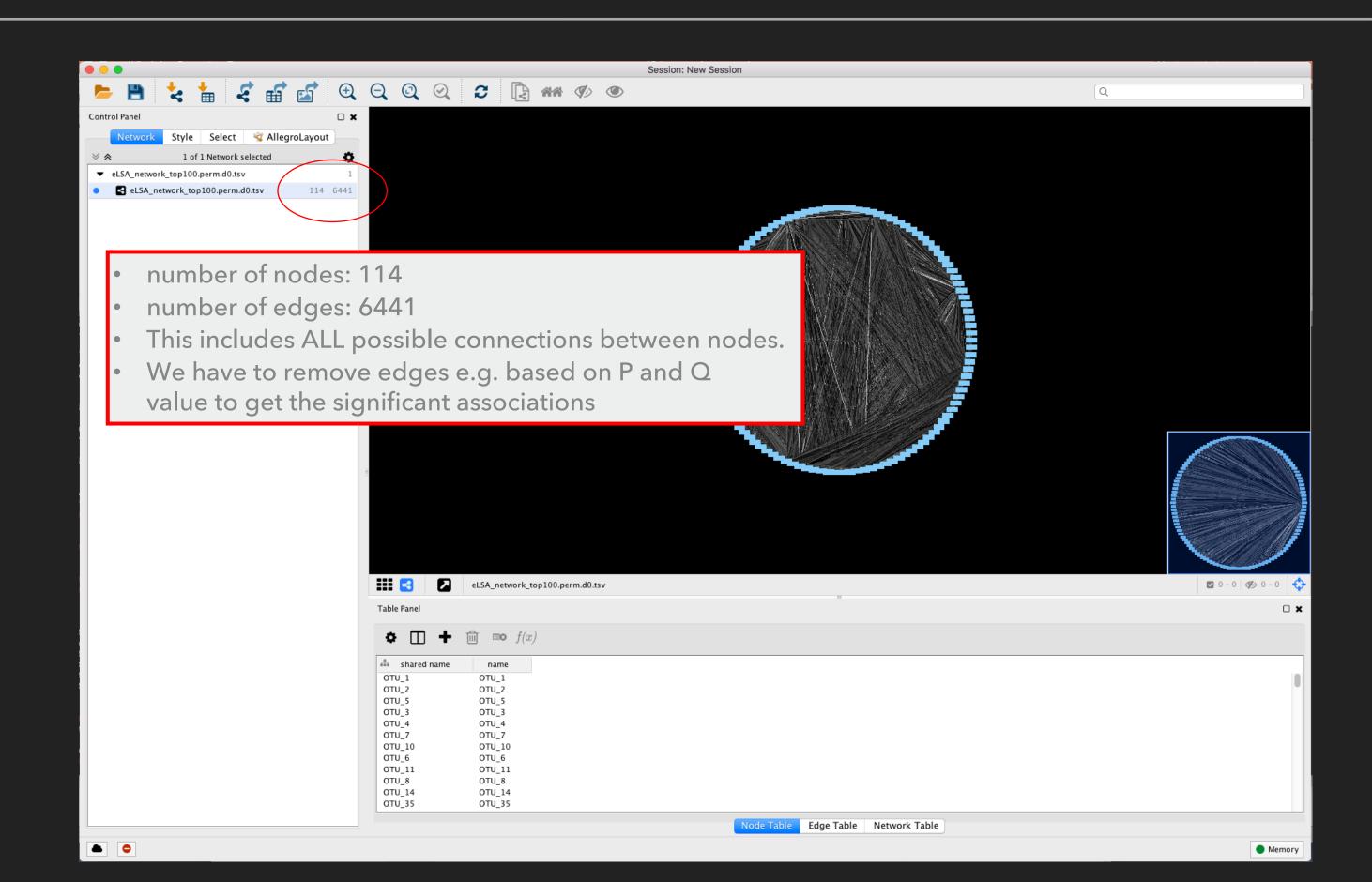
 Define target and source node (this is an undirected networks so the order of the nodes doesn't matter)

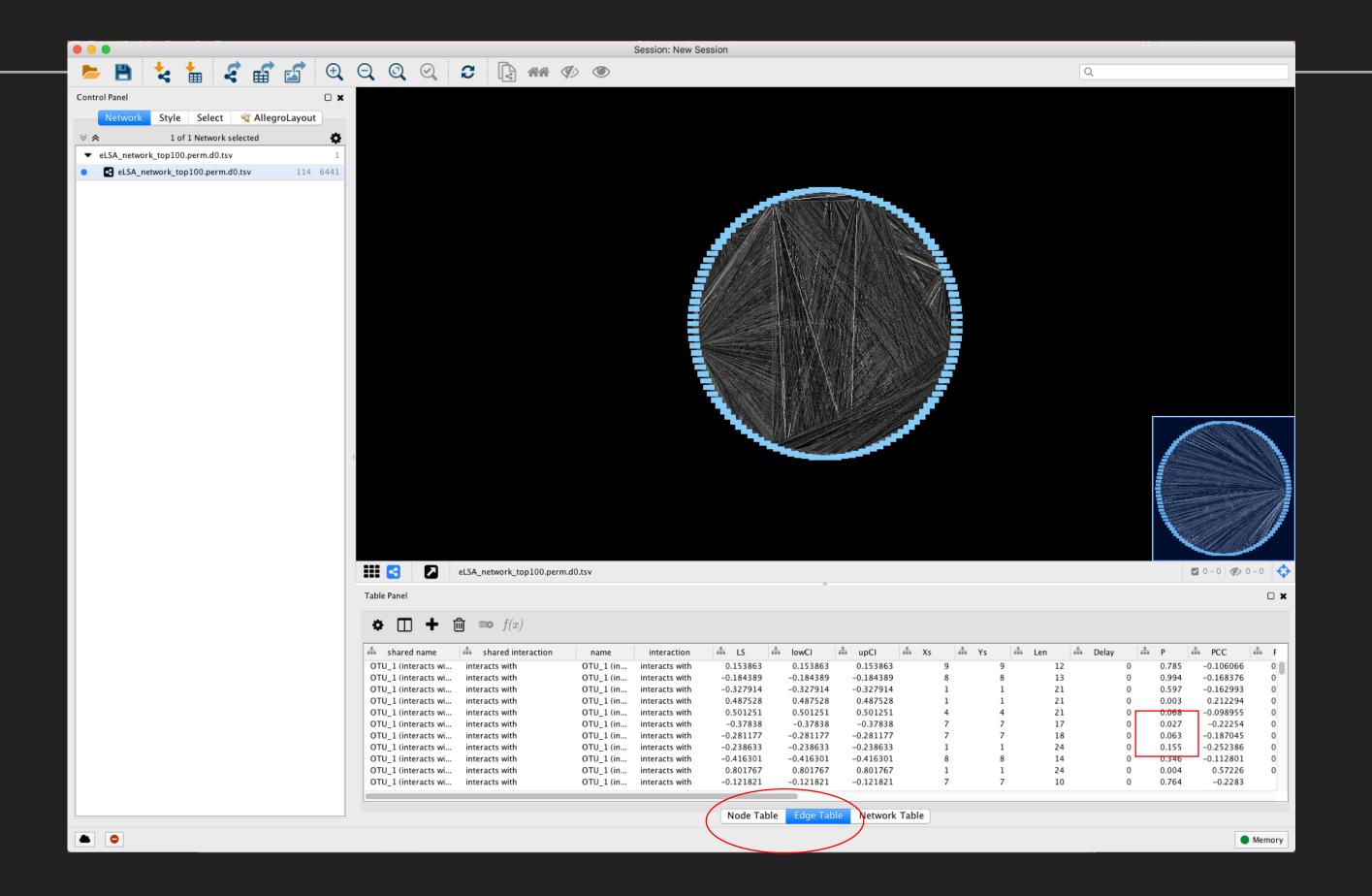


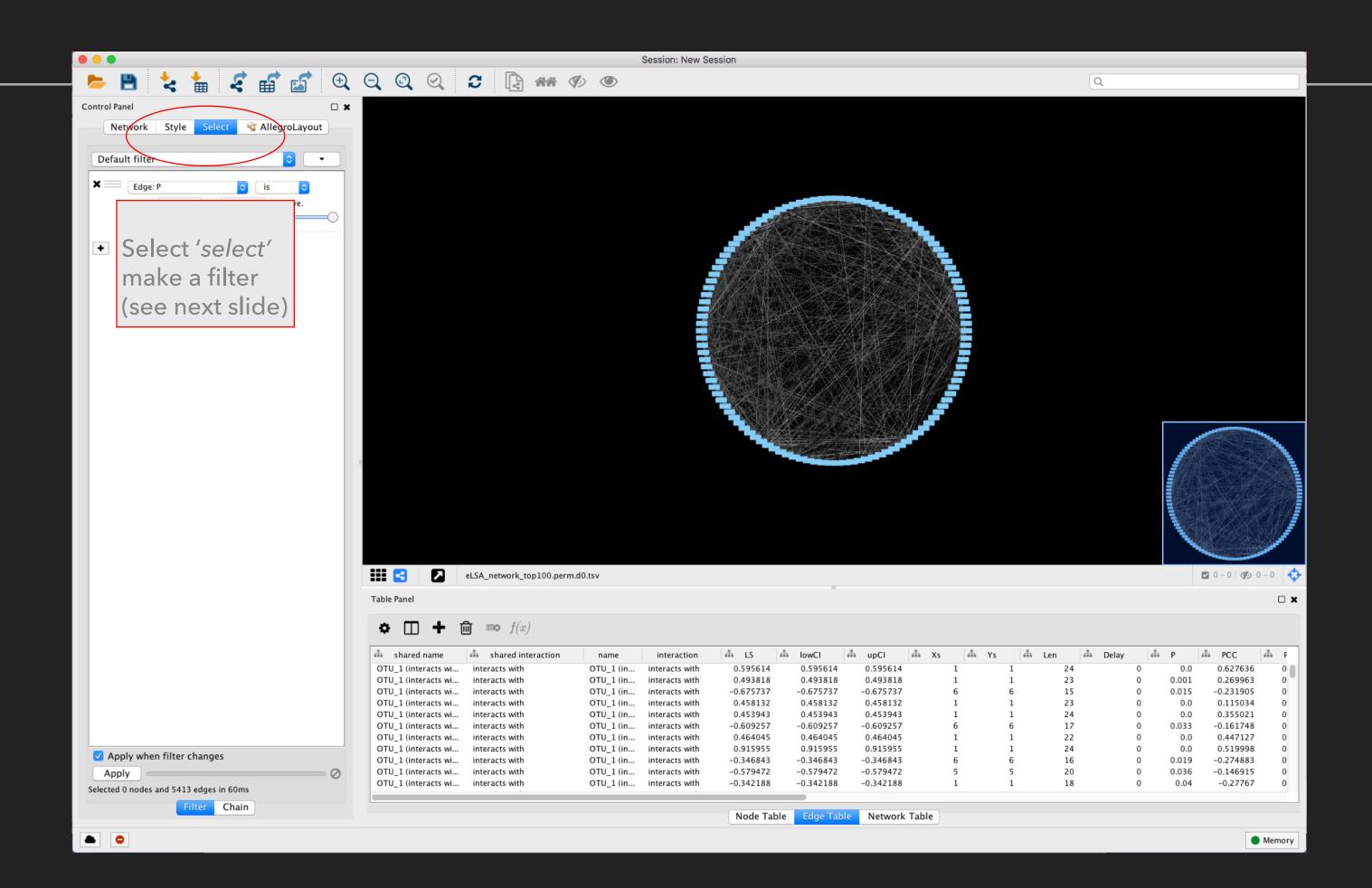
### **USING CYTOSCAPE**



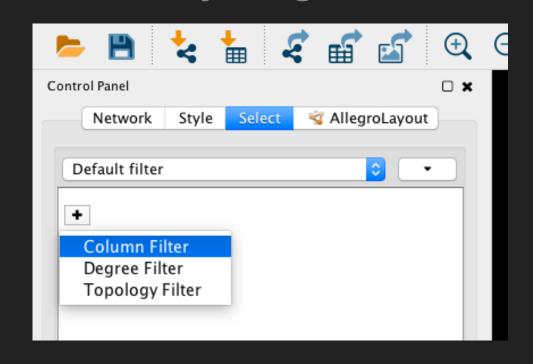


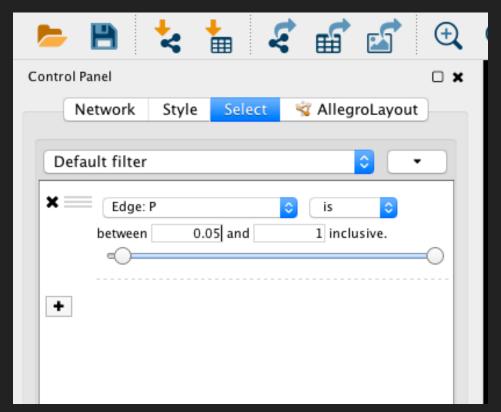


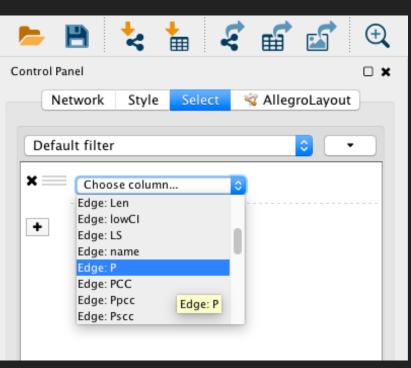


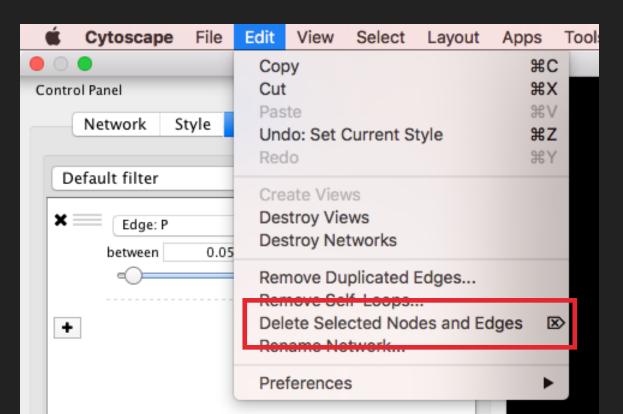


Setting up a filter for P value (repeat for Q value) How many edges are left?

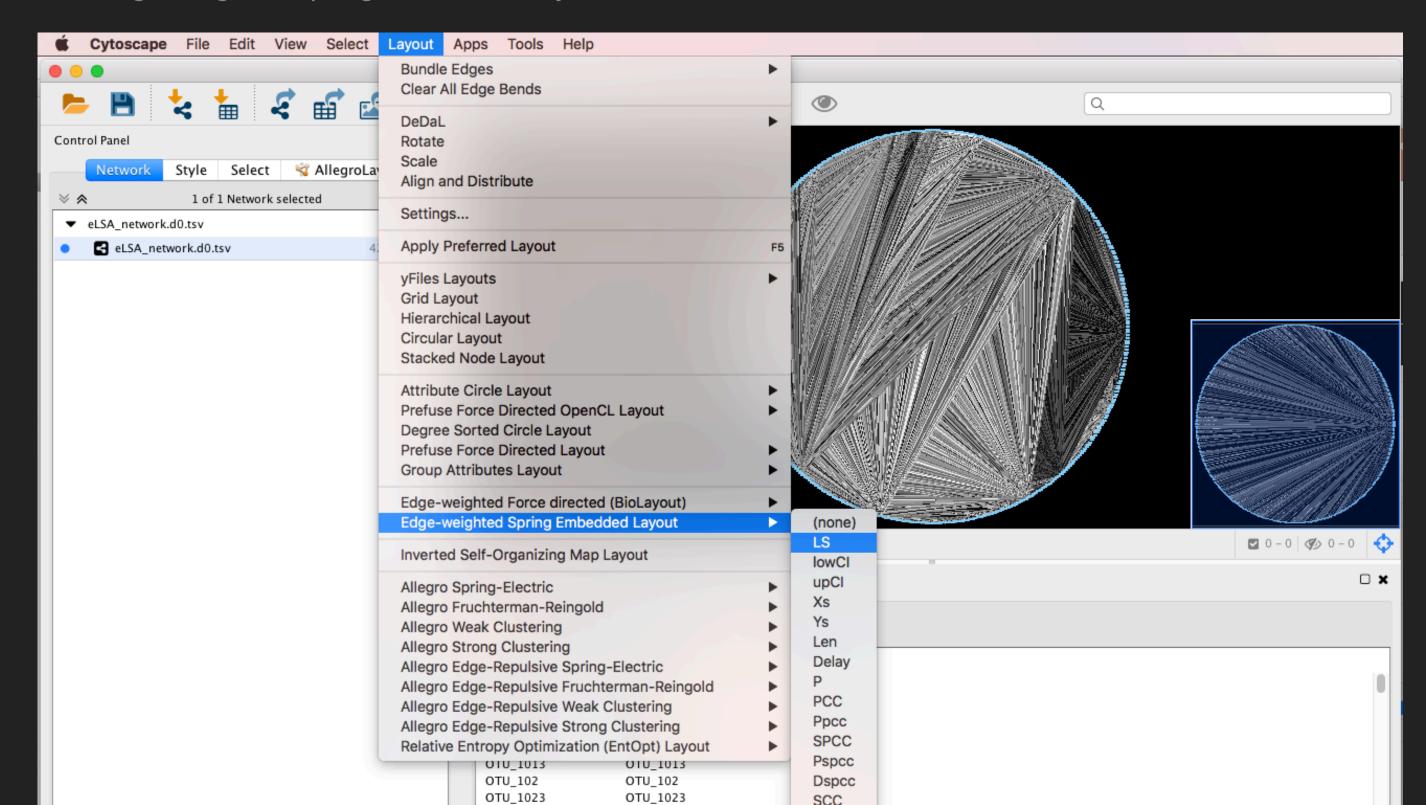




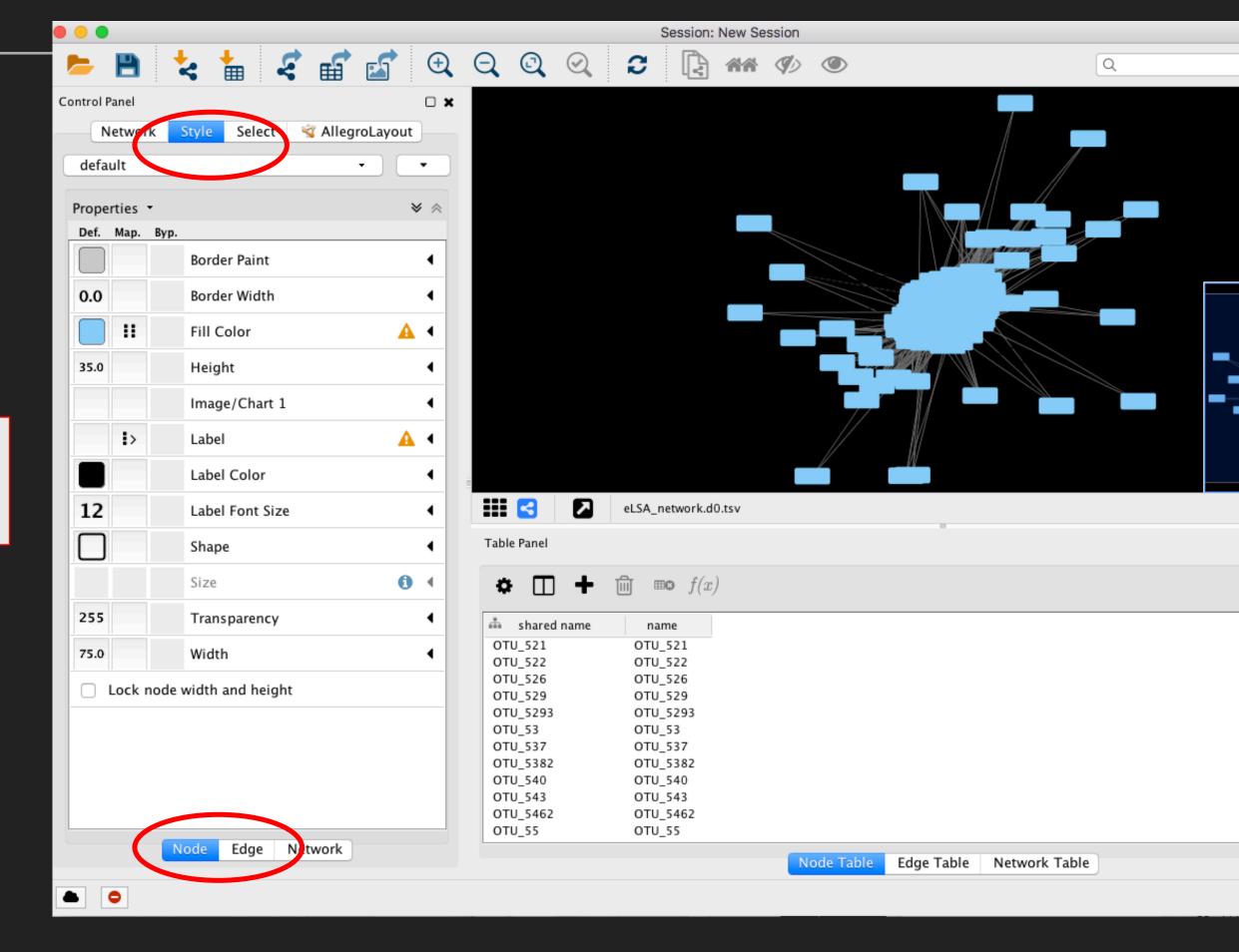




### Choose a layout - Circular is fastest Edge-weighted Spring Embedded Layout (LS) looks better©



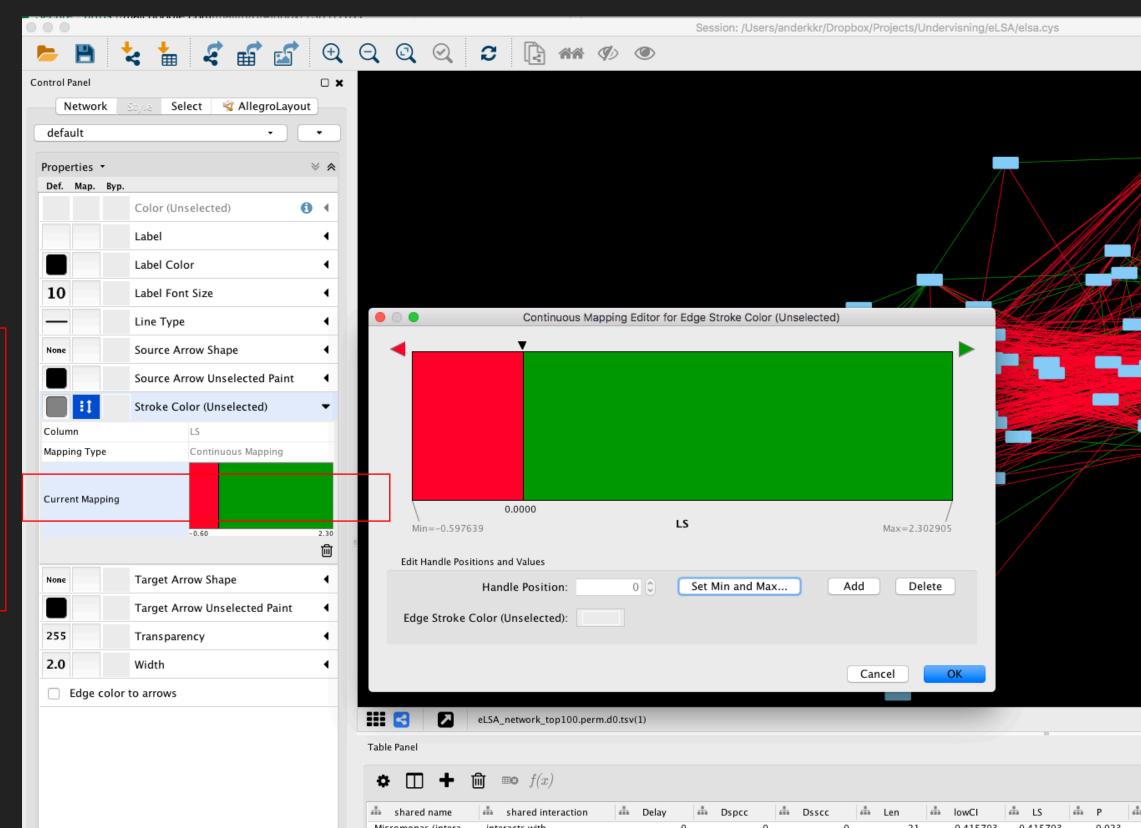
Now choose 'Style' and then 'Edge' to change the color of the edges



Use *Stroke Color* to make the positive edges green and the negative edges red

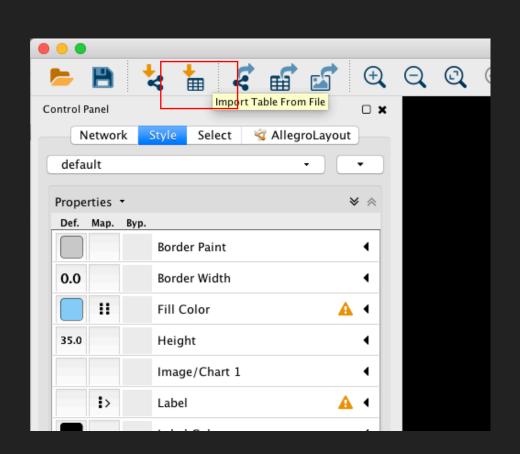
Column: LS

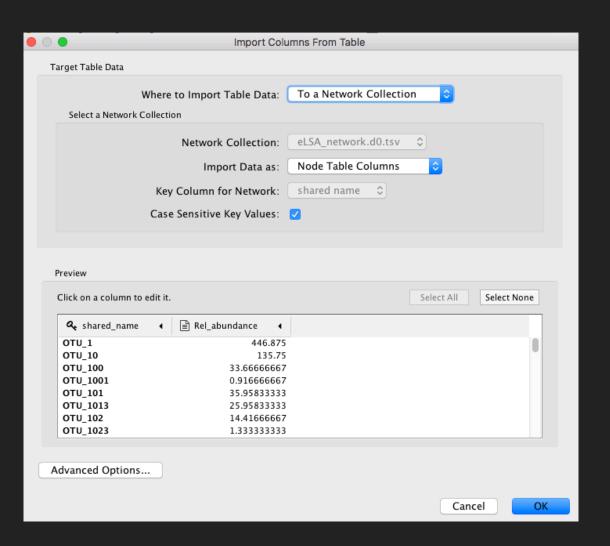
Mapping type: Continuous



# ADD ANNOTATIONS TO THE NODES- IMPORT TABLE

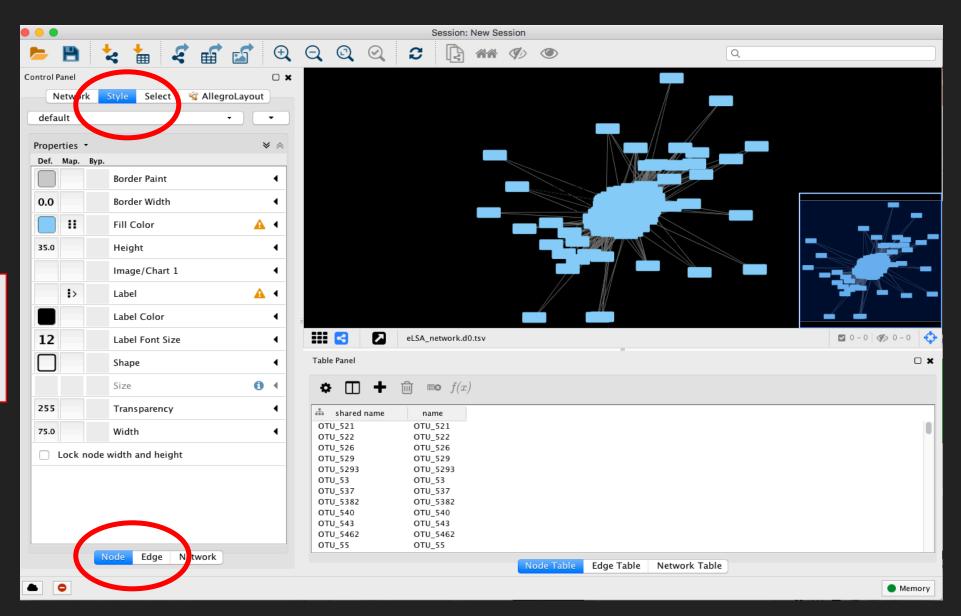
- Two annotation files:
  - 1) Relative abundance (eLSA\_annotation\_relabund.tsv)
  - 2) Taxonomic assignment (blastn vs MAS from UParse pipeline)



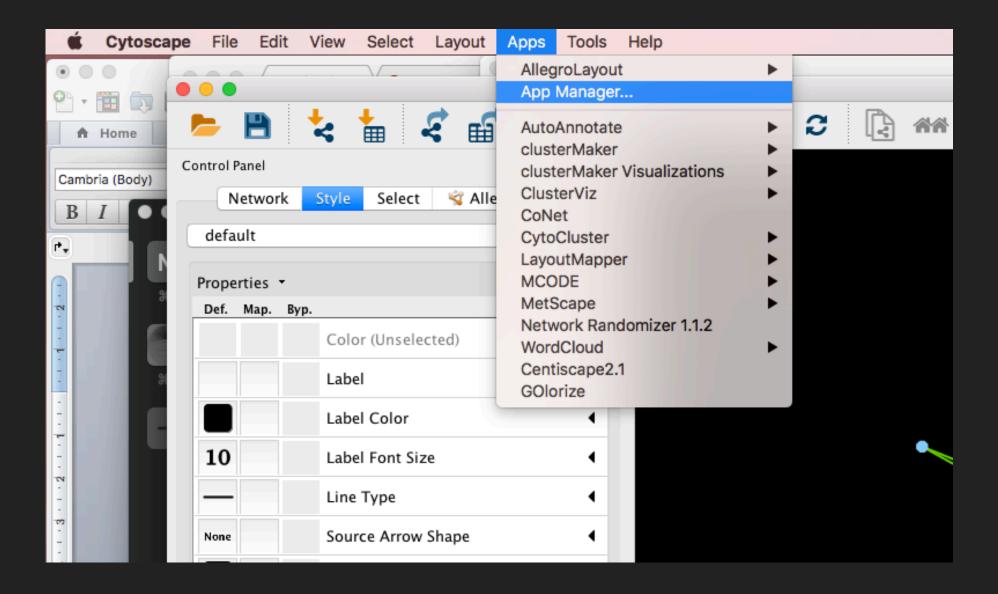


 Change the name, shape, color and/or size of the nodes based on the relative abundance and the taxonomic assignment of the OTUs

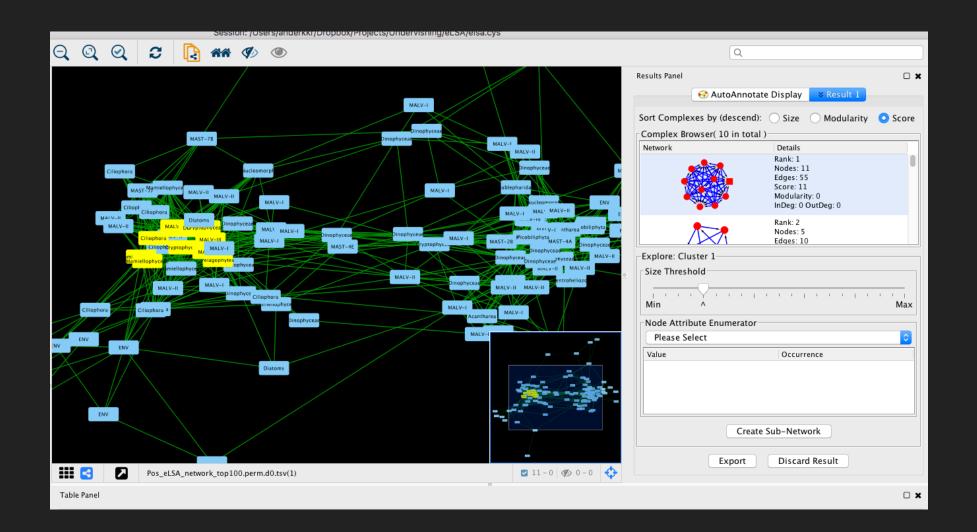
Choose 'select'
and then 'node' to change
the color, shape etc of the nodes



- Searching for Modules:
- Install ClusterViz from the App Manager



- Searching for Modules:
- (Save your project before moving on, just in case)
- Remove the negative edges, then search for Modules using MCODE in the ClusterViz App.



 Searching for Hubs (species that are highly connected, or have a high Degree). MCODE added the Degree for each node to the Node table.

