

EFI Quick Help

How do I get Cytoscape to work (Mac)?

Looking at sequence-similarity networks with Cytoscape may require resetting memory allocation (especially for versions prior to 2.8.2). Here is one way to configure your Mac so that you can open and view sequence similarity networks generated by the **Superfamily/Genome Core**. See also the Cytoscape Wiki.

- Download a property list editor if you don't have one (such as a free trial of PlistEdit Pro from here).
- 2. Open the Applications folder (from Finder, not from the dock) and then open the Cytoscape folder.
- 3. Right click (or hold control key and click mouse) on the Cytoscape Application icon (looks like one at the top of this page) and select "Show Package Options."
- 4. Open Content folder and then the "info.plist" file using the list editor program.
- 5. Examine VMOptions under Java and change the Class (2nd column) to "Array" by clicking on "String."
- Select VMOptions line (will be grey) and click "New Child" at top of window and then "New Sibling." Two new lines will appear below VMOptions.
- 7. Enter –Xmx20M as the Value (3rd column) for the first new line and –Xms2G for the second new line. The result should look like the screen-shot at the right.
- 8. Save file.

