

## Scraping Web of Science

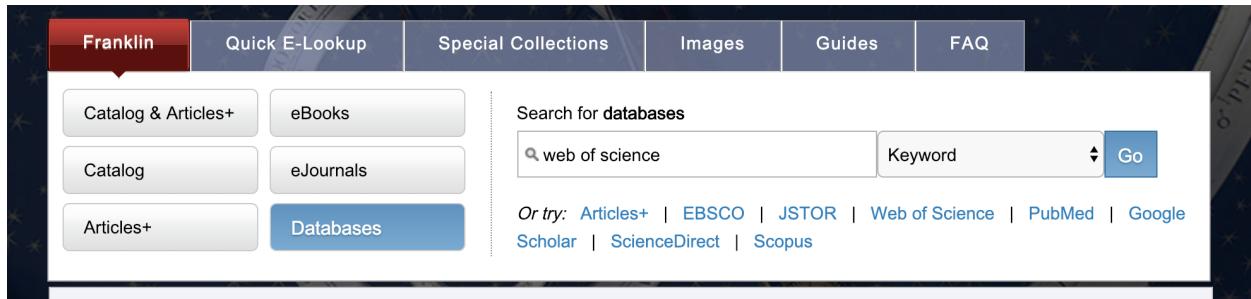
The aim of the protocol is to create a dataset with all the manuscripts published by faculty in your neuroscience department of interest. The dataset of manuscripts will have the name of the authors, the manuscript abstract, and keywords associated with the manuscript. We will then be able to apply natural language processing techniques to ask questions related to the diversity of the science being conducted by individuals and departments on these data.

**Step 1:** Manually create a list of faculty in the neuroscience department in an excel file (if you don't have excel on your computer, a google drive (<https://www.google.com/drive/>) spreadsheet will work. The first column should contain the faculty member's first name and the second column should contain the faculty member's surname.

**Step 2:** Go to web of science and find the web of science institutional code for your neuroscience department.

Go to Penn Libraries: <https://www.library.upenn.edu/>

Search for web of science in “Databases”:



Click on the “ISI web of science” result that comes up:

1. ISI web of science  Bookmark

Publication: Philadelphia, PA : Institute for Scientific Information

Format/Description: Database/Website

Online resource: Connect to resource.  
Connect to citation indexes tutorial.  
Connect to cited reference searching tutorial.

Show Availability Details

And then scroll to the bottom of the page that loads and click on “Connect to Resource”:

Web link: [The Rosengarten Family Fund Home Page](#)



Access Restriction: Restricted for use by site license.

Online: [Connect to resource.](#)

<http://hdl.library.upenn.edu/1017/6959>

[Connect to citation indexes tutorial.](#)

<http://hdl.library.upenn.edu/1017/6959/1>

[Connect to cited reference searching tutorial.](#)

<http://hdl.library.upenn.edu/1017/6959/2>

You should get to the Web of Science splashpage. Click on “Advanced Search” and your page will look like this:

Booleans: AND, OR, NOT, SAME, NEAR

Field Tags:

TS= Topic	SA= Street Address
TI= Title	CI= City
AU= Author [Index]	PS= Province/State
AI= Author Identifiers	CU= Country/Region
GP= Group Author [Index]	ZP= Zip/Postal Code
ED= Editor	FO= Funding Agency
DO= DOI	FG= Grant Number
PY= Year Published	FT= Funding Text
AD= Address	SU= Research Area
OG= Organization-Enhanced [Index]	WC= Web of Science Category
OO= Organization	IS= ISSN/ISBN
SG= Suborganization	UT= Accession Number
	PMID= PubMed ID
	ALL= All Fields

On the right-hand slide, in the “Field Tags” section, click on the [\[index\]](#) after OG=Organization-Enhanced. Search for your department in the search bar that appears (here I’m searching for Johns Hopkins):

## Web of Science

Organizations - Enhanced List

\*\* Use this list to find the preferred name for an organization and the variants we have identified and use the Browse and Find features to locate organizations to add to your query.

Click on a letter or number to browse organizations alphabetically by title

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	0	1	2	3	4	5	6	7	8
---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---

Enter text to find organizations containing or related to the text.

Johns Hopkins

[Back to top](#)

Find your department's tag, click the "Add" button beside it and click "Ok" where it says "Transfer your selected organization(s) below to the Organizations - Enhanced field on the search page" at the bottom of the page.

Add	D Johns Hopkins Oncology Center
Add	D Johns Hopkins University
Add	D Johns Hopkins University Applied Physics Laboratory
Add	D King George's Medical University
Add	D Makerere University
Add	D National Institutes of Health (NIH) - USA
Add	D National University of Singapore
Add	D NIH National Cancer Institute (NCI)

Transfer your selected organization(s) below to the Organizations - Enhanced field on the search page. OK Cancel

Johns Hopkins University

This will bring you back to the advanced search and will give you the code you will need for your manuscript search. Here the code we need for Johns Hopkins is OG=(Johns Hopkins University):

Use field tags, Boolean operators, parentheses, and query sets to create your query. Results will appear in the Search History table at the bottom of the page. ([Learn more about Advanced Search](#))

Example: TS=(nanotub\* AND carbon) NOT AU=Smalley RE  
#1 NOT #2 [more examples](#) | [view the tutorial](#)

OG=(Johns Hopkins University)

**Step 2:** Use excel to create a block of text that you can then copy and paste later when searching to speed up the process.

Paste the code below into the first row of your completed excel sheet. It can be pasted into any column as long as it is not the first two that you have created using the faculty member names. This is creating a search string we will use later consisting of author names (AU=) and department of origin (OG=[change this based on Step 1]). The text in red will fill automatically with the faculty member's surname (=B1) and their first name (=A1).

AU=	=B1	=A1	AND	OG=(Johns Hopkins University)	OR
-----	-----	-----	-----	-------------------------------	----

Highlight all 6 columns, grab the bottom-right corner of the rightmost cell with your mouse, and drag it down. It will autofill all the faculty members you entered as you drag it down:

	A	B	C	D	E	F	G	H	I
1	Marilyn	Albert		AU=	Albert	Marilyn	AND	OG=(Johns Hopkins University)	OR
2	Yeka	Aponte		AU=	Aponte	Yeka	AND	OG=(Johns Hopkins University)	OR
3	Jay	Baraban		AU=	Baraban	Jay	AND	OG=(Johns Hopkins University)	OR
4	Amy	Bastian		AU=	Bastian	Amy	AND	OG=(Johns Hopkins University)	OR
5	Dwight	Bergles		AU=	Bergles	Dwight	AND	OG=(Johns Hopkins University)	OR
6	Seth	Blackshaw		AU=	Blackshaw	Seth	AND	OG=(Johns Hopkins University)	OR
7	Mary	Blue							
8	Antonello	Bonci							
9	Solange	Brown							
10	Pete	Calabresi							
11	Peter	Campochiaro							
12	Michael	Caterina							
13	Pablo	Celnik							
14	Jeremiah	Cohen							
15	Carlo	Colantuoni							
16	Ed	Connor							
17	Susan	Courtney-Faruqee							
18	Kathleen	Cullen							
19	Ted	Dawson							
20	Valina	Dawson							
21	John	Desmond							

Remove the last cell when you get to the very bottom:

112	Mark	Wu		AU=	Wu	Mark	AND	OG=(Johns Hopkins University)	OR
113	King-Wai	Yau		AU=	Yau	King-Wai	AND	OG=(Johns Hopkins University)	OR
114	Eric	Young		AU=	Young	Eric	AND	OG=(Johns Hopkins University)	OR
115	Don	Zack		AU=	Zack	Don	AND	OG=(Johns Hopkins University)	OR
116	David	Zee		AU=	Zee	David	AND	OG=(Johns Hopkins University)	OR
117	Kechen	Zhang		AU=	Zhang	Kechen	AND	OG=(Johns Hopkins University)	OR
118	Feng-Quan	Zhou		AU=	Zhou	Feng-Quan	AND	OG=(Johns Hopkins University)	OR
119									

**Step 3:** Search web of science for all relevant articles. Select all the text in the 6 columns you created, copy it, and paste it into the advanced search in Web of Science, setting the restriction options to search for “English” and “Article” and press Search!

#1 IN01 #2 more examples | view the tutorial

AU= Albert Marilyn AND OG=(Johns Hopkins University) OR
AU= Aponte Yeka AND OG=(Johns Hopkins University) OR
AU= Baraban Jay AND OG=(Johns Hopkins University) OR
AU= Bastian Amy AND OG=(Johns Hopkins University) OR
AU= Bergles Dwight AND OG=(Johns Hopkins University) OR
AU= Blackshaw Seth AND OG=(Johns Hopkins University) OR
AU= Blue Mary AND OG=(Johns Hopkins University) OR
AU= Bonci Antonello AND OG=(Johns Hopkins University) OR
AU= Brown Solange AND OG=(Johns Hopkins University) OR
AU= Calabresi Pete AND OG=(Johns Hopkins University) OR

**Search**

Restrict results by languages and document types:

All languages	All document types
English	<b>Article</b>
Afrikaans	Abstract of Published Item
Arabic	Art Exhibit Review

**Step 4:** Output the resulting data. Click on the search result number (here 3, 674). This will bring you to the articles:

## Search History:

Set	Results	
		<a href="#">Save History / Create Alert</a> <span style="margin-left: 20px;"><a href="#">Open Saved History</a></span>
# 1	<a href="#">3,674</a>	(AU= Albert Marilyn AND OG=(Johns Hopkins University) OR AU= Aponte Yeka AND OG=(Johns Hopkins University) AND OG=(Johns Hopkins University) OR AU= Bastian Amy AND OG=(Johns Hopkins University) OR AU= Berg Hopkins University OR AU= Blackshaw Seth AND OG=(Johns Hopkins University) OR AU= Blue Mary AND OG=(Johns Hopkins University) OR AU= Bonci Antonello AND OG=(Johns Hopkins University) OR AU= Brown Solange AND OG=(Johns Hopkins University) OR AU= Calabresi Pete AND OG=(Johns Hopkins University) OR AU= Campochiaro Peter AND OG=(Johns Hopkins University) OR AU= Celnik Pablo AND OG=(Johns Hopkins University) OR AU= Colantuoni Michael AND OG=(Johns Hopkins University) OR AU= Colantuoni Carlo AND OG=(Johns Hopkins University) OR AU= Connor E

Click on the dropdown option to “Save to Other File Formats”. An option box will show. Web of science restricts exporting to 500 articles at a time so you may have to repeat this process a few times by changing the numbers you place in “Records” - here we are getting articles 1 to 500. Make sure the “Record Content” and “File Format” settings match:

**Send to File**

✖

Number of Records:  All records on page  
 Records  to

Record Content:

File Format:

Click “Send” and the plain text document will download.

## Building Semantic Networks

This section will guide you through the steps to make a network that can be visualized using Gephi and analyzed using MATLAB. If you haven't already, please [download and install Gephi](#).

### **Constructing a network from scientific abstracts**

#### **1. Download word embedding model from the link below (file size: 809 MB)**

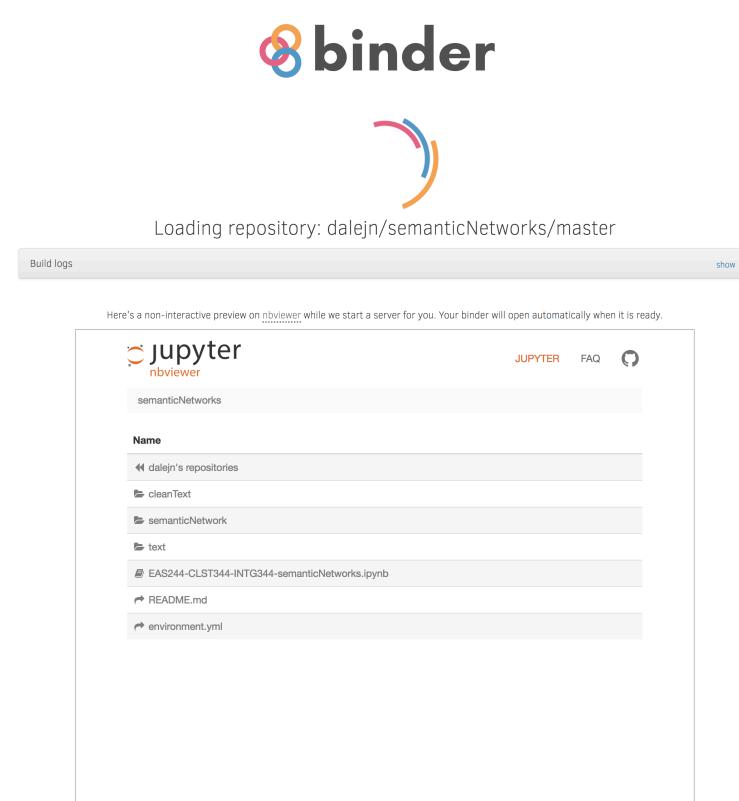
[https://www.dropbox.com/s/je4slqywoh8wnr1/enwiki\\_5\\_ner.txt?dl=1](https://www.dropbox.com/s/je4slqywoh8wnr1/enwiki_5_ner.txt?dl=1)

This is a neural network model of word vector embeddings with 296,630 word vectors representing words' semantic meaning trained on 2,252,637,050 words from Wikipedia. If interested in more details, see last section.

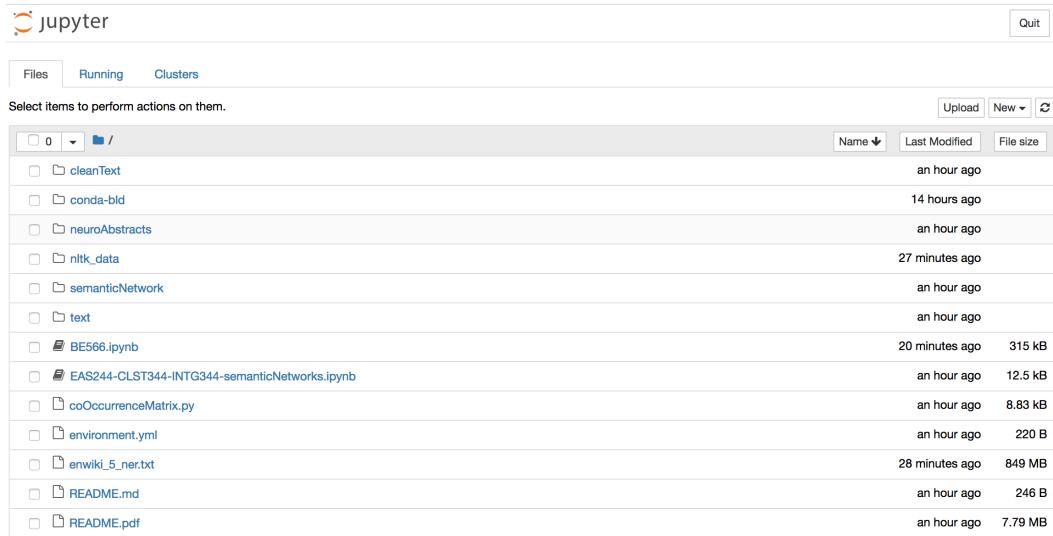
#### **2. Start programming environment**

<https://mybinder.org/v2/gh/dalejn/semanticNetworks/master>

- A. You should see the loading page below. Depending on how many students are trying to access the server, it may take a few minutes or longer to start the programming environment. If it is still loading after 10 minutes, please refresh the page.

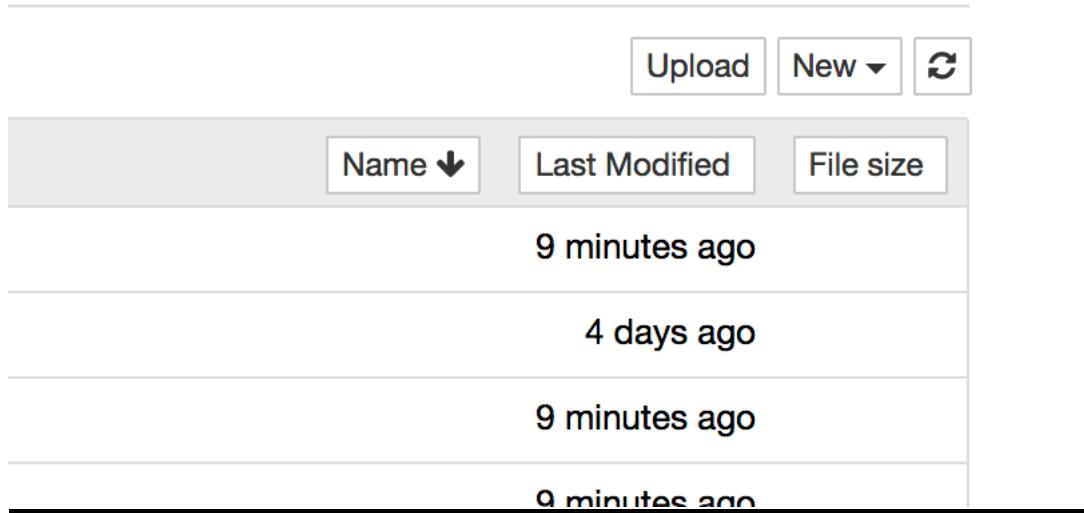


- B.** When the page is loaded, you will see a screen like the one below. Note, that the environment will automatically shut down after 10 minutes of inactivity (if you leave your window open, this will be counted as “activity”).



### 3. Upload the model downloaded from step 1 to the environment

- A.** Click on “Upload,” go to where you downloaded the file from Step 1 and select it.



- B.** Press Ok after selecting the model to upload.

## Large file size warning

The file size is 809 MB. Do you still want to upload it?

- C. Click on the blue Upload button and you should see the percent progress of the upload. Depending on how many students are accessing this environment, this upload can take a few minutes or longer.



4. **Once the upload has completed, click on the text link “BE566.ipynb” as seen below to open a programming notebook.**



This will open a new tab/page where you should see the below:

```
In [1]: from nltk import sent_tokenize, word_tokenize
from nltk.corpus import WordNetLemmatizer as wnl
import nltk, re, string, glob
from itertools import islice, compress
import iter工具
import matplotlib.pyplot as plt
import networkx as nx
nltk.download('punkt')
nltk.download('wordnet')

model = './enwiki_5_ner.txt'
word_vectors = gensim.models.KeyedVectors.load_word2vec_format(model, binary=False)

#####
# Initialising a define helpful Functions#
#####

translator = str.maketrans('', '', string.punctuation.replace('-', '')) #filters punctuation except dash
lemmatiseCondition = 1
lemmatizer = wnl()

# Function for finding index of words of interest, like 'references'

def find(target):
    for i, word in enumerate(sents):
        try:
            i = word.index(target)
        except ValueError:
            continue
        yield i

# Function for handling the input for gensim word2vec

class FileToSent(object):
    def __init__(self, filename):
        self.filename = filename

    def __iter__(self):
        for line in open(self.filename, 'r'):
            ll = line.strip().split(',')
            ll = [c for c in ll if c not in string.punctuation] for s in ll
            ll = [num.strip() for num in ll]
            yield ll

# Function for looking for element x occurs at least n times in list

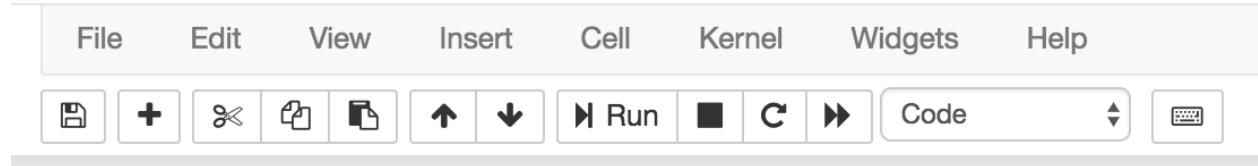
def check_list(list, x, n):
    gen = (True for i in list if i==x)
    return next(islice(gen, n-1, None), False)

In [1]: #####
# Read in .txt file(s) from a specified directory #
#####

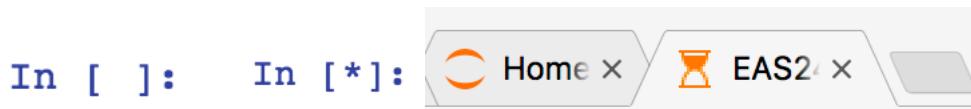
IDs = glob.glob('../Text/*')
IDs_subIDs = []
for ID in IDs:
    IDs_subIDs += glob.glob(ID + '/*.txt')
print(len(IDs)) # Print number of files read
```

## 5. Run the code

- A. 1<sup>st</sup> box: loads the relevant packages, modules, and functions. Click on the first box, which should then become highlighted along the borders. At the top of the screen in the toolbar, click the Run button.



You should see the left-most text change to the right. Your tab in the web browser will also show an hourglass. This means the code in this box is running. This can take a few minutes or more to finish.



- B. 2<sup>nd</sup> box: reads in your text and cleans it. When the code is done running, the asterisk above will be replaced by a number and the hourglass will be replaced by a notebook. Now, click on the second box of code, which should become highlighted along the borders. Run this code by clicking the Run button. This will take a few seconds at most.
  - C. 3<sup>rd</sup> box: constructs the semantic network matrix from the shortest N cosine-distances (the strongest N associations). Select the third box. Depending on the number of nodes in the graph (i.e. number of unique words in the text), you may want to change the total number of edges (i.e. connections between the words) in the constructed semantic network. I've set this number to be 10 times the number of nodes, but this can be changed depending on how dense you want the resulting semantic network to be.

```
# The number of connections we want: either as a factor of the number of words or a set number
num_top_conns = len(my_words) * 19
```

You will see a list of words outputted.

- D. 4<sup>th</sup> box: constructs a co-occurrence network based on 5-gram sliding window  
(represents a count of word pairs that co-occur with each other in 5-word chunks across the entire text)

## **6. Download the relevant outputted files**

- A. You can close the notebook now. Return to the directory screen and click the "semanticNetwork" link.



- B. Check a box (**one at a time**) to download **all** the files in this folder. You must select the files one at a time, or else the Download button doesn't appear.



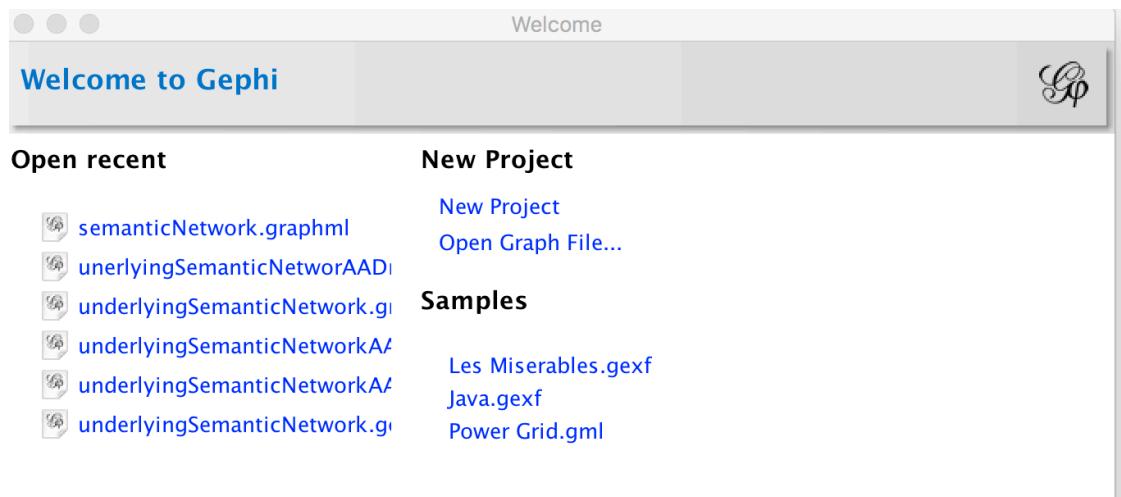
- C. Once you've downloaded **all files in this folder**, press Quit at the top right.



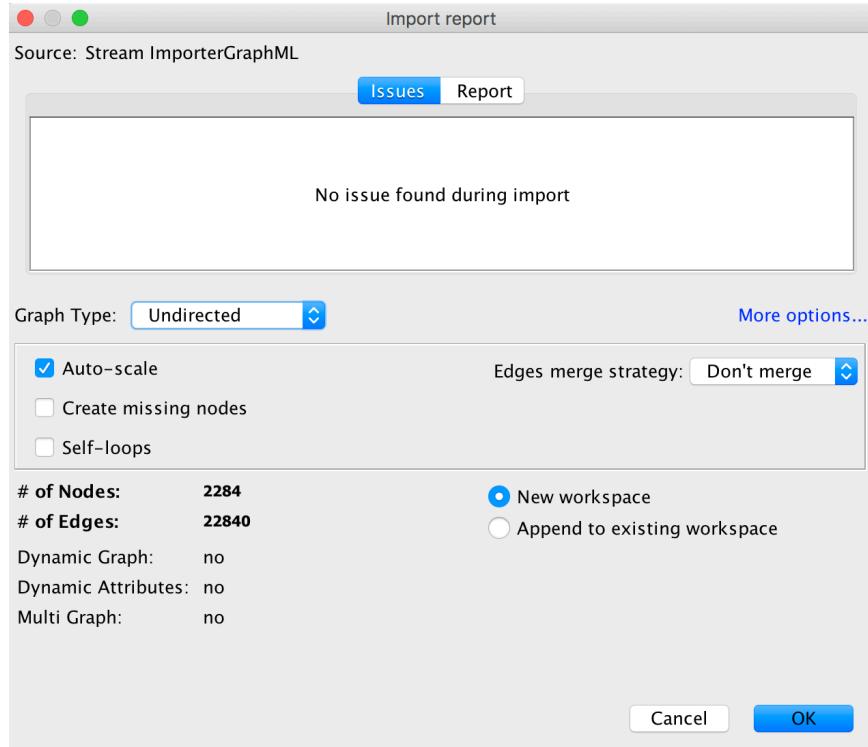
## 7. Visualize the network in Gephi

- A. Open Gephi. If you haven't already downloaded and installed it, [please do so here](#).

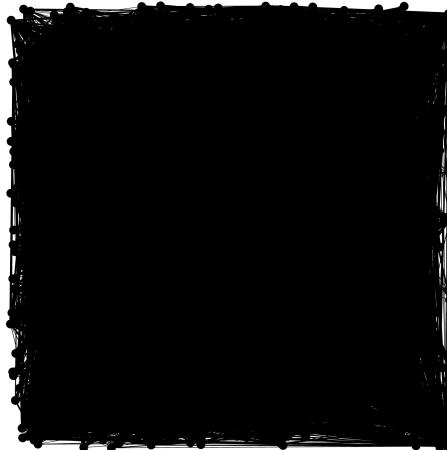
You will see a Welcome screen like below. Click on "Open Graph File..." If you don't see the welcome screen, go to File and Open on your computer's toolbar.



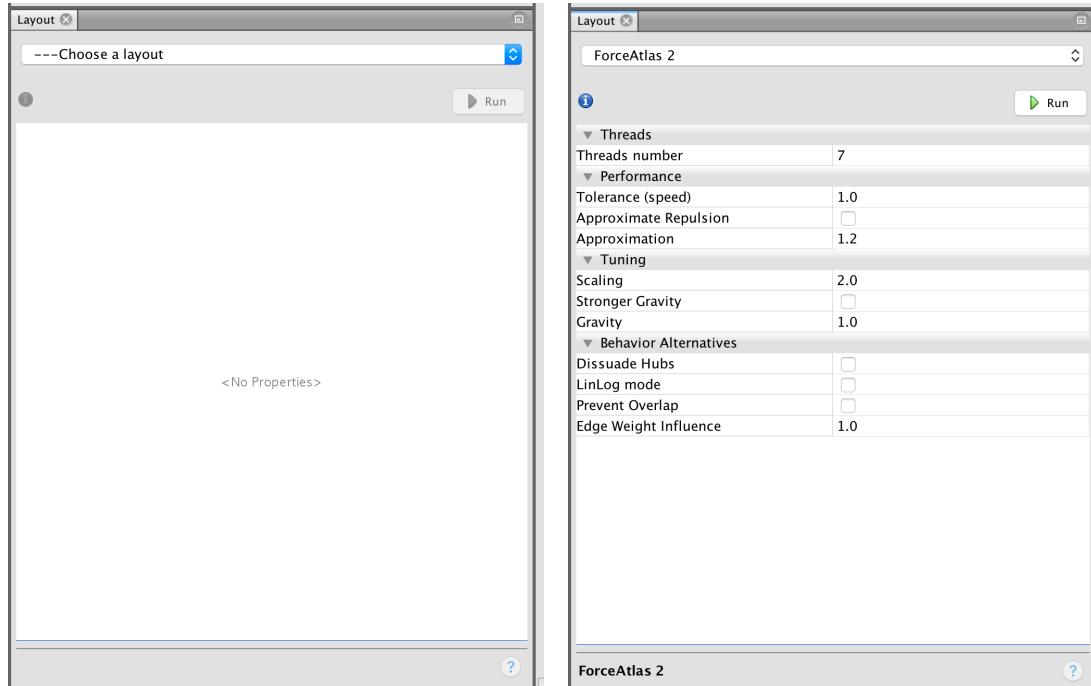
- B. Navigate to where you downloaded semanticNetwork.graphml and open this file.
- C. A new screen should pop up. Default settings should be fine but please check that they match the ones below and then press OK. Note: If you are loading coOccurrenceNetwork.gexf, there will be many "Issues" listed—you may ignore them and press OK.



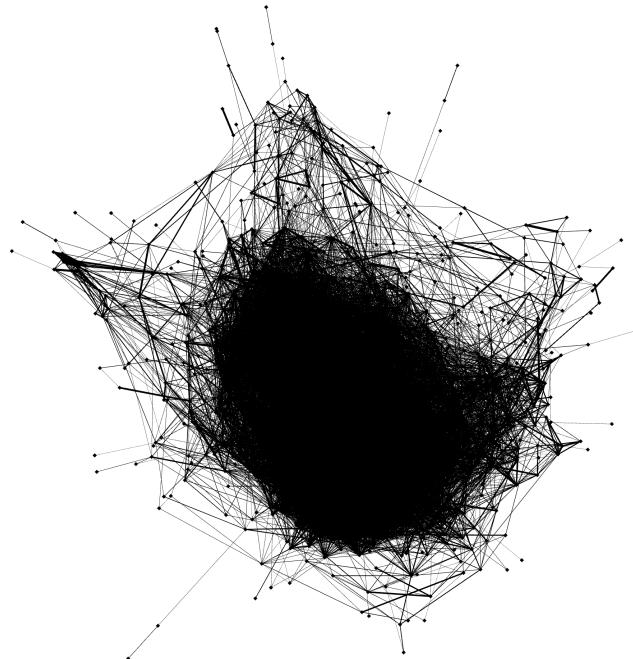
- D. The network should be visible. If not visible, you may need to go to Gephi's menu at the top and select Window > Graph. You can scroll your cursor over nodes to see connections. We'll now clean this up to make it more interpretable.



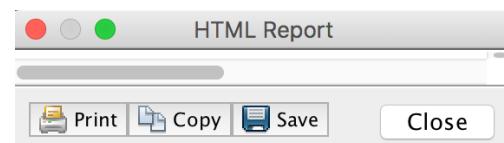
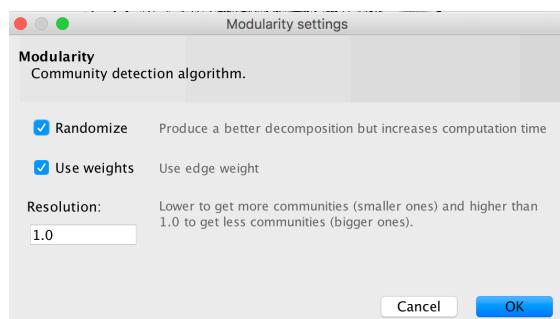
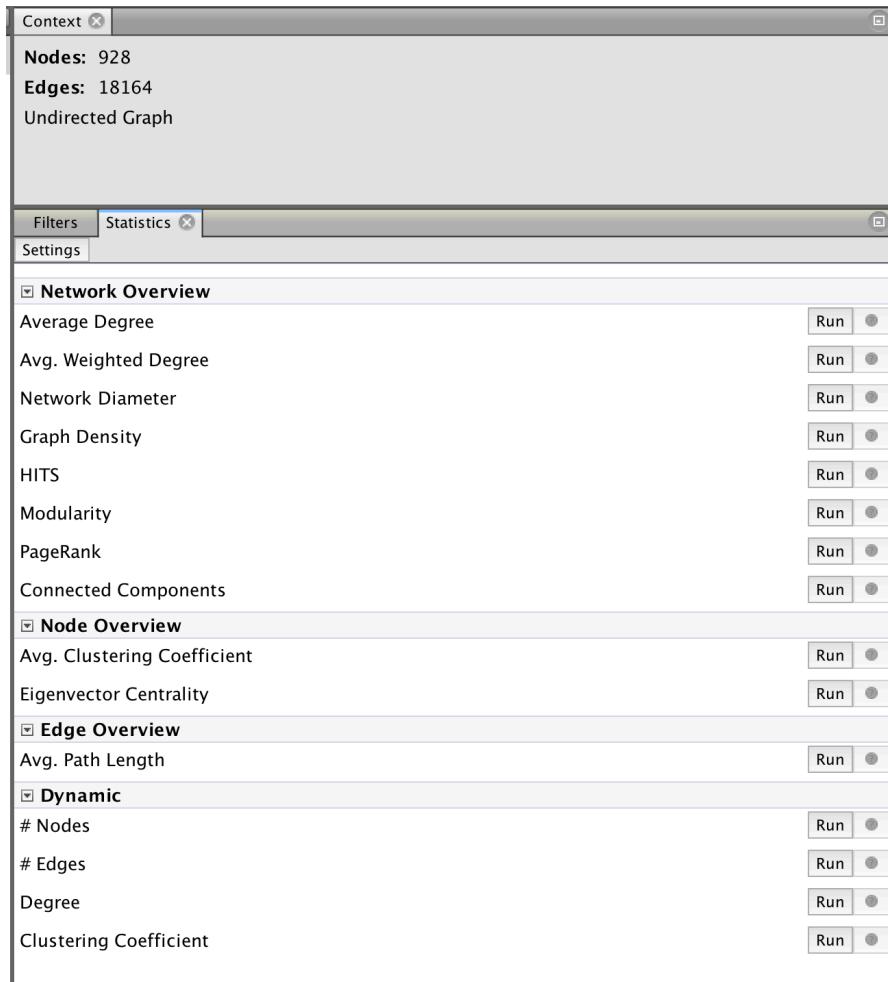
- E. On the bottom left, click on “---Choose a layout” and select “ForceAtlas 2.” Then, press the Run button, which will then be replaced by a Stop button.



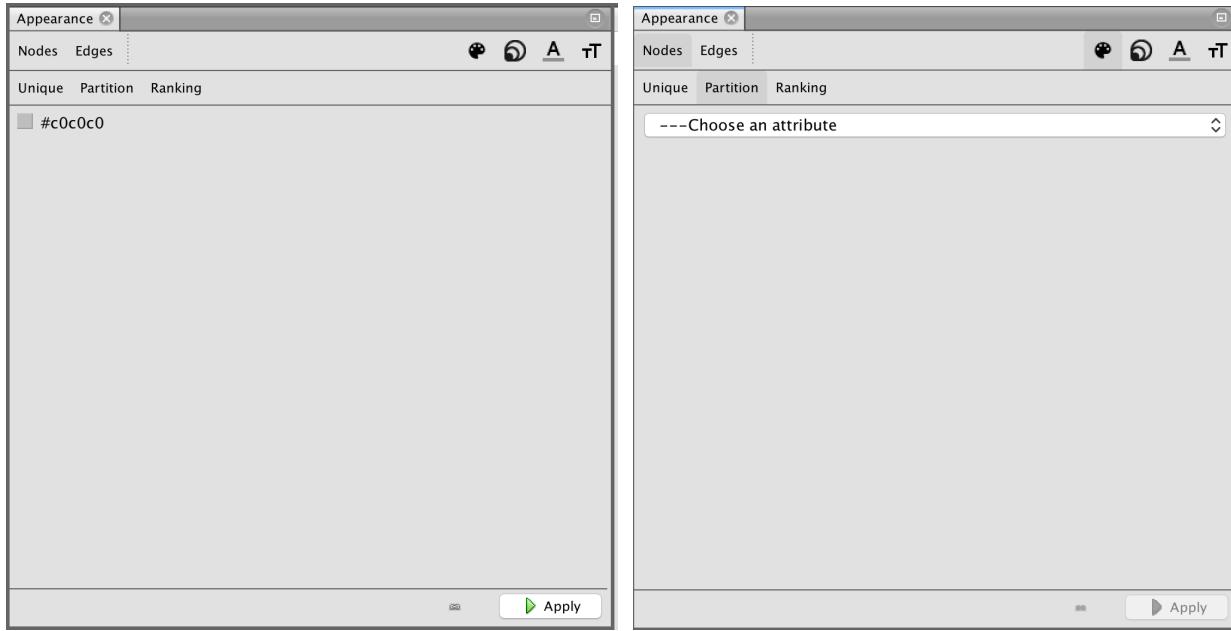
- F. Use your mouse wheel to zoom in and out. Hold right click and drag to move the network around. When the network looks like it has stabilized (not moving much), press the Stop button.



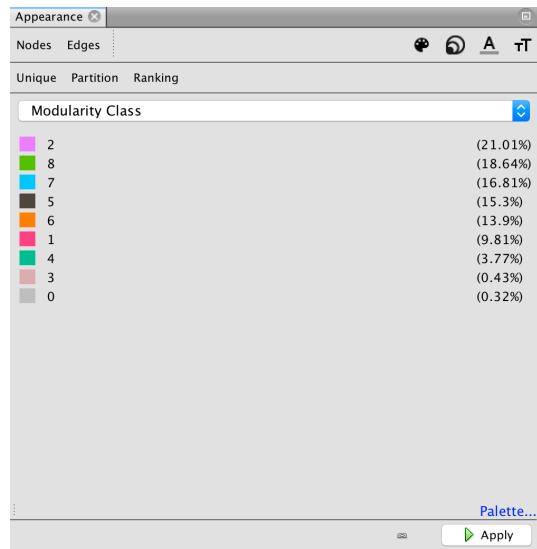
- G. On the right-side toolboxes, click on the “Statistics” tab and then click “Run” for the Modularity measure. A box will pop up; click OK. Another pop-up labeled HTML Report will appear; click Close.



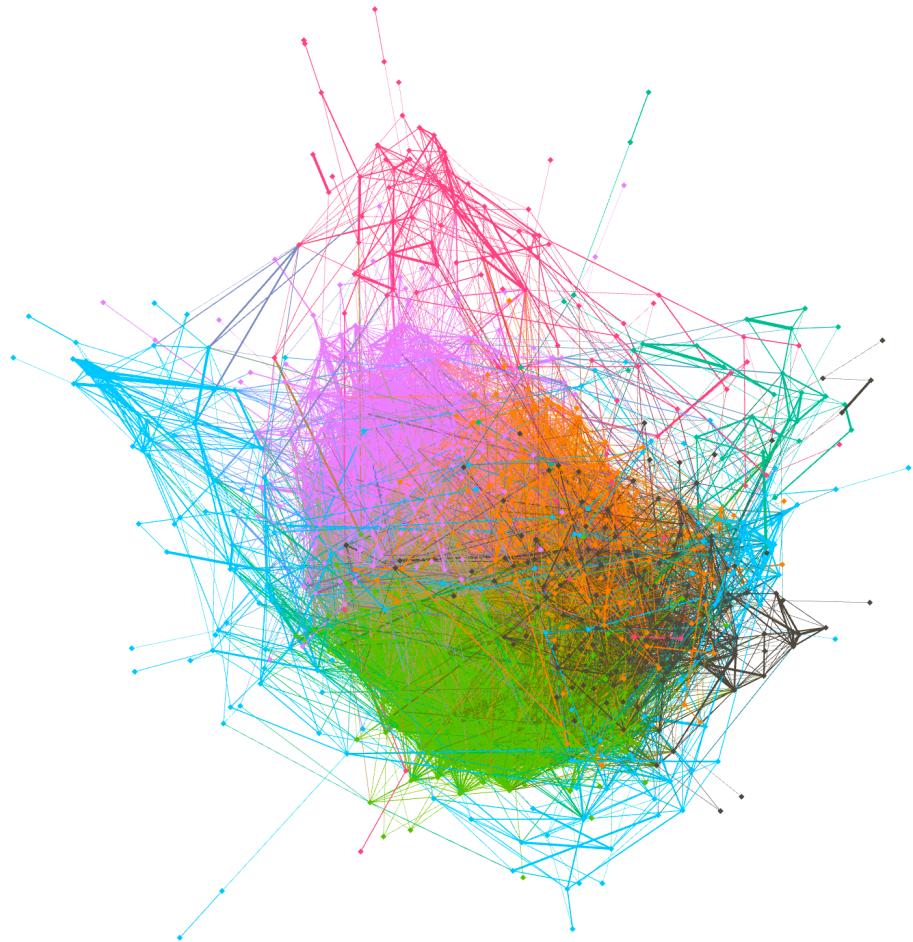
H. At the top-left toolbox labeled Appearance, click on the “Partition” tab.



- I. Click on “—Chose an attribute” and select “Modularity Class.” Click “Apply.”



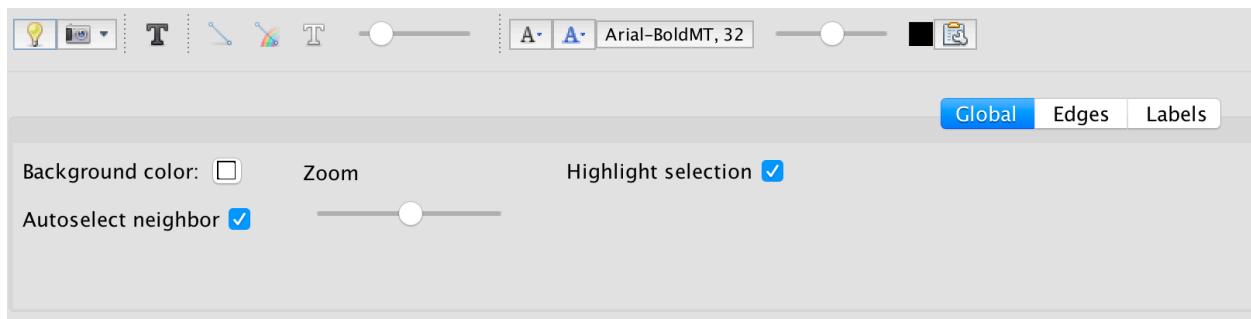
Your graph should now be colored by module.

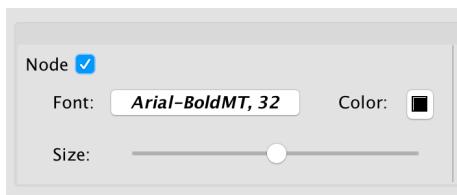
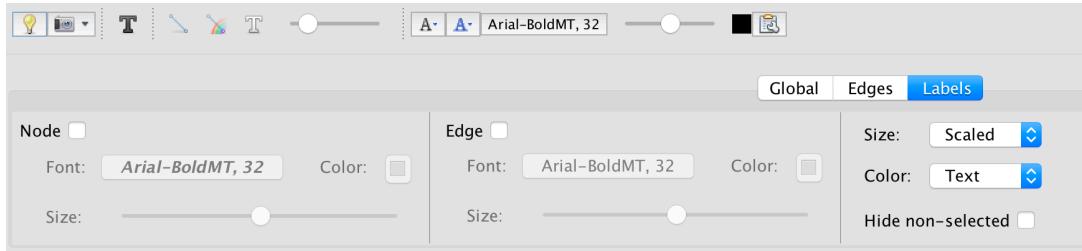


- J. Label each node with the unique word it represents. At the bottom, there is a toolbar; click the small button all the way to the right.

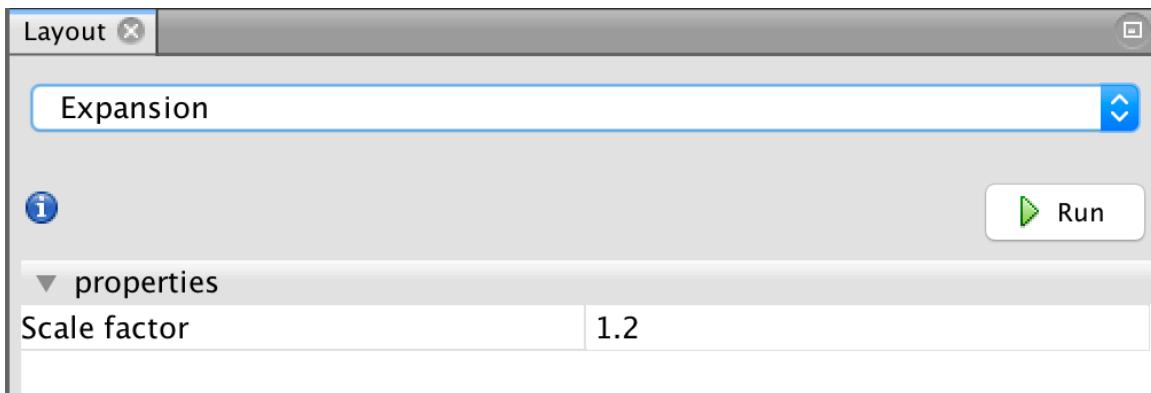


- K. Click on the “Labels” tab and click the checkbox for Node.

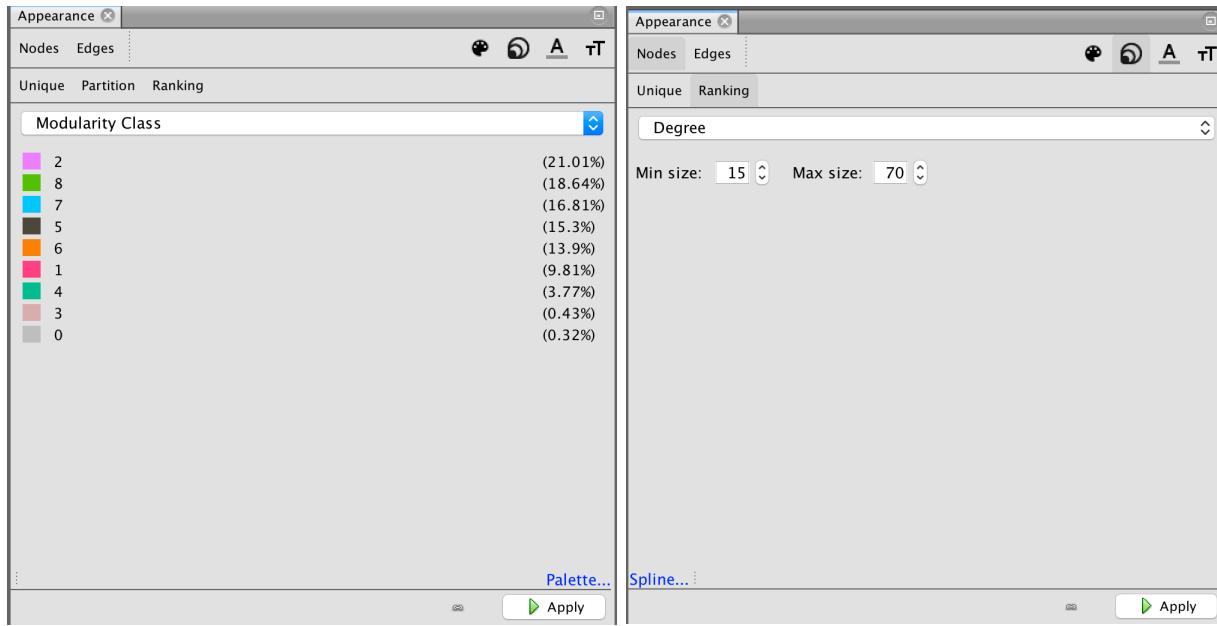




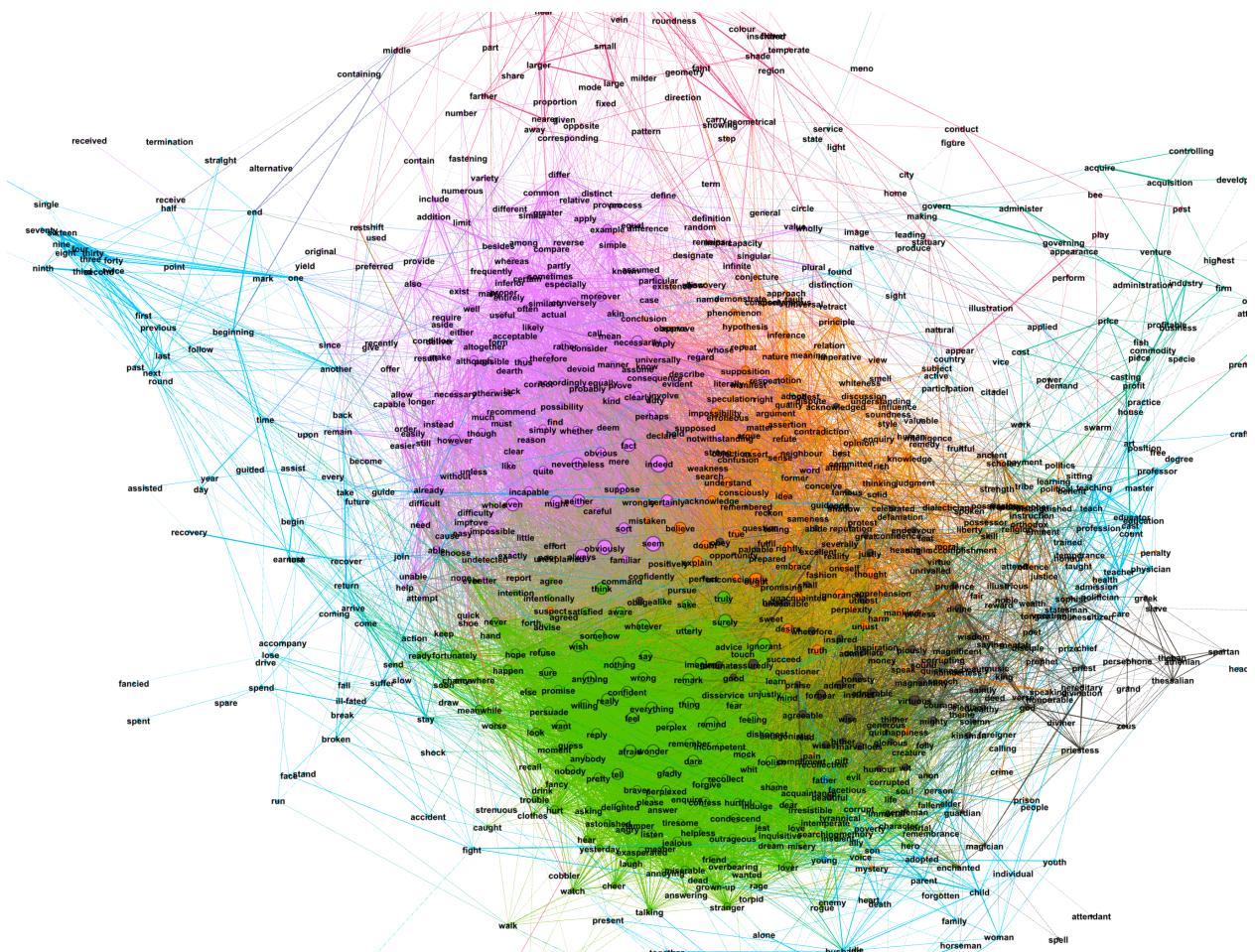
- L. Spread out the graph to read the text more easily by going to the “Layout” pane again (the same place where you applied “ForceAtlas 2”) and select “Expansion.” Click on Run a few times until you’re satisfied with the spread of the text labels. Then, zoom out on the graph using your mouse wheel.



- M. Finally, navigate back to the “Appearance” pane and click the  button. Click on the “Ranking” button. Then, select “Degree” from the dropdown menu and change the “Min size” and “Max size” to the ones below. Click “Apply”.



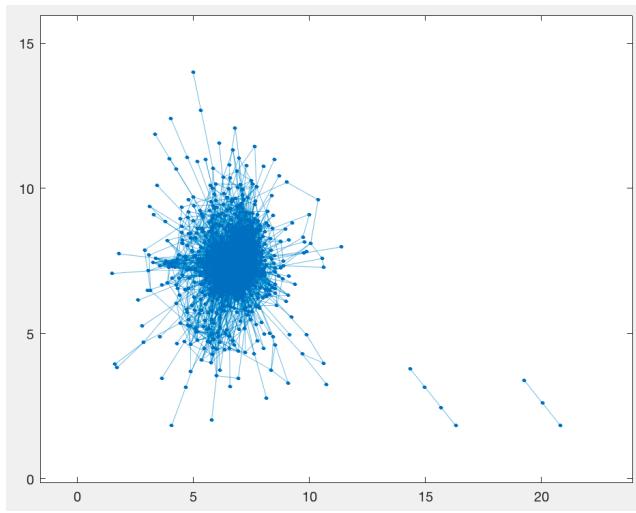
N. Explore your graph! You can scroll over individual nodes to see what words are connected.



O. Repeat step 7 with the `coOccurrenceNetwork.gexf`. How do the networks differ?

**8. Visualize/analyze the network in MATLAB.**

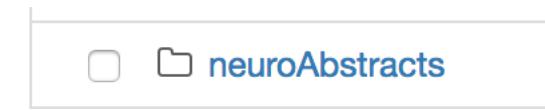
A. Please see `semanticNetworkAnalysisReadme.pdf` for further instructions.



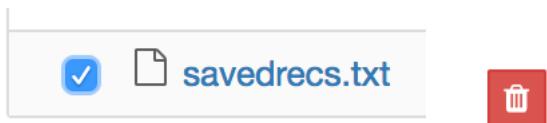
### Constructing a network matrix from your own text

**1. Follow the same steps as above, but on Step 3 also do the following:**

A. Click on the “neuroAbstracts” link to navigate to that directory.



B. Check the box for “savedrecs.txt” and then delete it by clicking the red trashcan button.



C. Click on the “Upload” button and upload your text files (must be in .txt format) and make sure the filename ends with .txt

D. Continue Steps 4 and on as described in the above steps.

## More details

### 1. Word vector embeddings model

Pre-trained model from <http://vectors.nlpl.eu/explore/embeddings/en/models/>  
 Removed all "universal part of speech" tags from the corpus

=====

General model information:

dimensions: 300

window: 5

iterations: 5

vocabulary size: 296630

id: 3

=====

Training corpus:

====

description: English Wikipedia Dump of February 2017

url: <https://dumps.wikimedia.org/>

tool: Wikipedia Extractor

case preserved: True

stop words removal: NLTK

id: 2

tokens: 2252637050

lemmatized: True

tagset: UPOS

tagger: Stanford Core NLP v. 3.6.0

NER: True

public: True

=====

Training algorithm:

name: Continuous Skipgram

url: <https://github.com/RaRe-Technologies/gensim>

tool: Gensim

version: 2.1

command: None

id: 0

**Math underlying word2vec:** <https://arxiv.org/pdf/1411.2738.pdf>,

**Nice visualization/demonstration of some toy training data:** <https://ronxin.github.io/wevi/>

**Google's original word2vec paper:** <https://arxiv.org/pdf/1301.3781.pdf>