CSCI4022 F21 HW7

November 11, 2021

1 CSCI4022 Homework 7; Undirected Graphs

1.1 Due Monday, November 8 at 11:59 pm to Canvas and Gradescope

Submit this file as a .ipynb with all cells compiled and run to the associated dropbox.

Your solutions to computational questions should include any specified Python code and results as well as written commentary on your conclusions. Remember that you are encouraged to discuss the problems with your classmates, but you must write all code and solutions on your own.

NOTES:

- Any relevant data sets should be available on Canvas. To make life easier on the graders if they need to run your code, do not change the relative path names here. Instead, move the files around on your computer.
- If you're not familiar with typesetting math directly into Markdown then by all means, do your work on paper first and then typeset it later. Here is a reference guide linked on Canvas on writing math in Markdown. All of your written commentary, justifications and mathematical work should be in Markdown. I also recommend the wikibook for LaTex.
- Because you can technically evaluate notebook cells is a non-linear order, it's a good idea to
 do Kernel → Restart & Run All as a check before submitting your solutions. That way
 if we need to run your code you will know that it will work as expected.
- It is **bad form** to make your reader interpret numerical output from your code. If a question asks you to compute some value from the data you should show your code output **AND** write a summary of the results in Markdown directly below your code.
- 45 points of this assignment are in problems. The remaining 5 are for neatness, style, and overall exposition of both code and text.
- This probably goes without saying, but... For any question that asks you to calculate something, you must show all work and justify your answers to receive credit. Sparse or nonexistent work will receive sparse or nonexistent credit.
- There is not a prescribed API for these problems. You may answer coding questions with whatever syntax or object typing you deem fit. Your evaluation will primarily live in the clarity of how well you present your final results, so don't skip over any interpretations! Your code should still be commented and readable to ensure you followed the given course algorithm.

Shortcuts: Problem 1 | Problem 2 | Problem 3 | Problem 4 | Extra Credit

```
[1]: import matplotlib.pyplot as plt
     import numpy as np
     import pandas as pd
     import networkx as nx
     # import scipy.stats as stats
     # import statsmodels.api as sm
```

```
[2]: PLOT_SIZE_X = 10
     PLOT_SIZE_Y = 6
     FONT_SIZE
                 = 16
```

Back to top # Problem 1 (Theory: Network MLEs; 5 pts)

Suppose graphs are generated by picking a probability p and choosing each edge independently with probability p, as if our entire network were coming from a Community Affiliation Graph Model (AGM) with only one universal community.

- What value of \hat{p} gives the maximum likelihood of seeing a specific graph with n total nodes and e total edges?
- What is the probability that your observed graph is the one generated if in fact $p = \hat{p}$?

Hint/Intuition: if you have 4 nodes and observe 3 total edges, what should P(edge) be? What would be the resulting P(graph)?

Solution:

Back to top # Problem 2 (Initializing BigCLAM: PageRank; 10 pts)

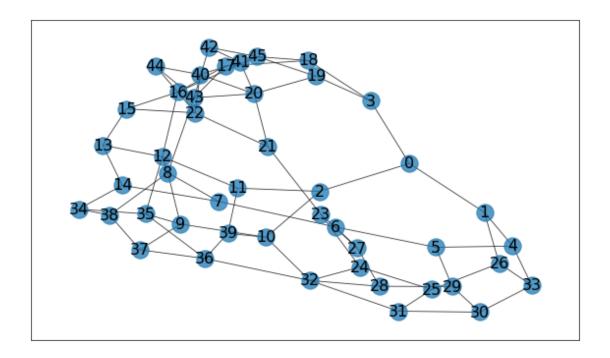
Below is some code to load in and plot a modified variation of the "Tutte" graph, an otherwise 3-way symmetric graph. Note that you are given both

- edges, a list of the edges of the graph
- neighbors, a dictionary holding the lists of the neighboring vertices for any provided integer vertex-key

which can be used for desired algorithms.

```
[3]: | #NB: this is a well known graph known as the "Tutte" graph
     edges=[(0, 1), (0, 2), (0, 3), (1, 4), (1, 26), (2, 10), (2, 11), (3, 18), (3, __
      →19), \
             (4, 5), (4, 33), (5, 6), (5, 29), (6, 7), (6, 27), (7, 8), (7, 14), (8, 1)
      \rightarrow 9), (8, 38), \
             (9, 10), (9, 37), (10, 39), (11, 12), (11, 39), (12, 13), (12, 35), (13, 12)
      \rightarrow14), (13, 15),\
             (14, 34), (15, 16), (15, 22), (16, 17), (16, 44), (17, 18), (17, 43), \dots
      \hookrightarrow (18, 45), (19, 20), \
             (19, 45), (20, 21), (20, 41), (21, 22), (21, 23), (22, 40), (23, 24),
      \hookrightarrow (23, 27), (24, 25), \
```

```
(24, 32), (25, 26), (25, 31), (26, 33), (27, 28), (28, 29), (28, 32),
 \rightarrow (29, 30), (30, 31), \
       (30, 33), (31, 32), (34, 35), (34, 38), (35, 36), (36, 37), (36, 39), 
\hookrightarrow (37, 38), (40, 41), \
       (40, 44), (41, 42), (42, 43), (42, 45), (43, 44)]
#add a few edges for fun and value and to break symmetry
newedges=[(32,36), (32,10), (20,40), (20,43), (8,12), (8,43), (12,16), (35,9), 
\rightarrow (16,45), (16,22)]
for newedge in newedges:
    edges.append(newedge)
nodes=list(range(46))
neighbors={key: [] for key in range(len(nodes))}
for edge in edges:
    neighbors[edge[0]].append(edge[1])
    neighbors[edge[1]].append(edge[0])
G=nx.Graph()
G.add_nodes_from(nodes)
for i in range(len(edges)):
    G.add_edge(edges[i][0],edges[i][1])
np.random.seed(4022)
pos = nx.spring_layout(G) #algorithm that tries to "cluster" node plotu
→ locations; it saves the xy coords of each vertex
labels=dict(zip(range(46),range(46))); #label 0-45
fig, ax = plt.subplots(figsize=(PLOT_SIZE_X, PLOT_SIZE_Y))
nx.draw_networkx_nodes(G, pos, node_size=300, alpha=.75); #node colors easy tou
⇒see labels over
nx.draw_networkx_edges(G, pos, width=0.8, alpha=0.8); #draw edges, using_
\rightarrow xy-node layout provided
nx.draw_networkx_labels(G, pos, labels, font_size=FONT_SIZE); #label vertices_
 →by index/number
```



Our goal is to come up with a sensibile initialization before we run BigCLAM. The original paper for BigCLAM suggests using "locally minimal neighborhoods." Rather than figure out a new algorithm, let's try initializing by:

- 1) Run PageRank on the Graph, treating each edge (a,b) as both the directed edges $a \to b$ and $b \to a$.
- 2) Extract the node with the highest PageRank as an initial "center" to get community affiliation of "1" for community A.
- 3) For community B's intial center, get the node with the highest PageRank that's not connected to the node in 2).
- 4) For community C's intial center, get the node with the highest PageRank that's not connected to either of the nodes in 2) and 3).

Plot your resulting initialization, as in the example code below.

```
[4]: #Here's some colors on this
#random example initialization
centers=set([19,28,34])
colorf=lambda x: 'green' if (x in centers) else 'yellow'
colors=[colorf(node) for node in nodes]

fig, ax = plt.subplots(figsize=(PLOT_SIZE_X,PLOT_SIZE_Y))

#draw the nodes, using xy-node layout 'pos' provided
```

```
nx.draw_networkx_nodes(G, pos, node_color=colors, node_size=300, alpha=.75);

#node colors easy to see labels over

nx.draw_networkx_edges(G, pos, width=0.8, alpha=0.8); #draw edges, using

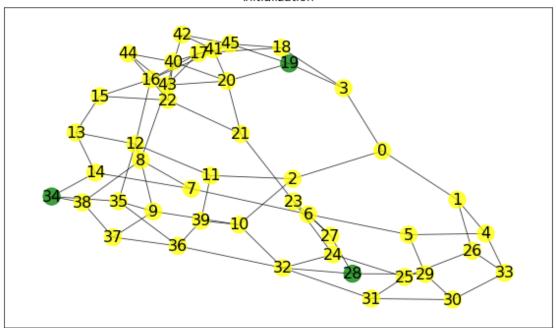
xy-node layout provided

nx.draw_networkx_labels(G, pos, labels, font_size=FONT_SIZE); #label vertices

by index/number

ax.set_title('Initialization');
```

Initialization



[]: #Your pagerank and plot here

```
def get_neighbors(nodes, edges):
   neighbors= {node: [] for node in nodes}
   for edge in edges:
      neighbors[edge[0]].append(edge[1])
      neighbors[edge[1]].append(edge[0])
   return neighbors
def dist L2(x,y):
   return np.sqrt(np.sum((x-y)**2))
def rank_links(links, ranks):
   l = list()
   for i, link in enumerate(links):
      l.append((link, ranks[i]))
   return sorted(l, key=lambda x: x[1], reverse=True)
def sparse_pagerank(Compact_Matrix):
   BETA
          = 0.8
   all_links = get_all_links(Compact_Matrix)
           = len(all links)
   r_old = np.repeat(1/n, n)
   tolerance = 10e-6
   converged = False
   iterations = 0
   while not converged:
      r_{new} = np.repeat((1-BETA)/n, n)
      # distribute importance payments between the nodes
      for link, outlinks in Compact_Matrix.items():
          d i = len(outlinks)
          for outlink in outlinks:
                     = all links.index(outlink)
                     = all links.index(link)
             r_new[j] += ((BETA * r_old[i]) / d_i)
      r new /= np.sum(r new)
      if dist_L2(r_new, r_old) <= tolerance:</pre>
          converged = True
      r_old = r_new.copy()
      iterations += 1
   r = rank_links(all_links, r_new)
   return r, iterations
```

```
def get_all_links(Compact_Matrix):
   s = set(Compact_Matrix.keys())
   for value in Compact_Matrix.values():
      for item in value:
          s.add(item)
   return list(s)
def PageRank(M, indices_to_names):
       = M.shape[0]
   BETA = 0.8
   Nmat = np.ones((n,n))/n
   A1 = BETA*M + (1-BETA)*Nmat
   r_old = np.repeat(1/n, n)
   r_new = np.ones(n)
   tolerance = 10e-6
         = 0
   iters
   converged = False
   while not converged:
      r_new = np.matmul(A1, r_old)
      if dist_L2(r_new, r_old) <= tolerance:</pre>
          converged = True
      r_old = r_new.copy()
      iters += 1
   r = rank_links(indices_to_names, r_new)
   return r, iters
def initialize_communities(r, neighbors, n):
   1 = []
   if n == 0:
      return 1
   else:
      1.append(r[0][0])
       for _ in range(n-1):
          all_neighbors = [item for i in 1 for item in neighbors[i]]
          max val = 0
          for node in r:
              if node[0] not in all_neighbors and node[0] not in 1 and_
\rightarrownode[1] > max_val:
                 max_val = node[1]
                 1.append(node[0])
      return 1
```

```
= len(nodes)
    = UndirectedColumnStochastic(edges, n)
    total = mat.size
    zeros = total - np.count_nonzero(mat)
    print(f'The matrix is {n} X {n}\nThere are {zeros} zeros out of {total} total_u
     →entries: {np.round((zeros/total) * 100, 2)}%')
    ######################################
    num to display = 20
    ####################################
    sparse r, sparse iters = sparse pagerank(neighbors)
    print(f'Top {num_to_display} node ranks:\n{sparse_r[:num_to_display]}')
    print(f'Number of iterations: {sparse iters}')
    ###################################
    r, iters = PageRank(mat, nodes)
    print(f'Top {num_to_display} node ranks:\n{r[:num_to_display]}')
    print(f'Number of iterations: {iters}')
    ####################################
    The matrix is 46 X 46
    There are 1958 zeros out of 2116 total entries: 92.53%
    Top 20 node ranks:
    0.02862912315461888), (8, 0.02845200889064079), (12, 0.02825149775459592), (43,
    0.028126760798845896), (10, 0.02430181518015813), (36, 0.024076127556250423),
    (45, 0.023971411058442832), (35, 0.02358603663265301), (9, 0.02353851311589241),
    (22, 0.02342870936847766), (40, 0.023337409433088766), (4,
    0.021348369572199023), (33, 0.021329164328575814), (26, 0.021247073667926088),
    (1, 0.021220371606918132), (5, 0.021200492745102693), (30, 0.02107862126585504),
    (29, 0.021044721497828923)]
    Number of iterations: 17
    Top 20 node ranks:
    [(16, 0.032977677108899525), (32, 0.03025347335618299), (20,
    0.028629123154618866), (8, 0.02845200889064078), (12, 0.028251497754595914),
    (43, 0.028126760798845886), (10, 0.024301815180158125), (36,
    0.024076127556250413), (45, 0.023971411058442825), (35, 0.023586036632653004),
    (9, 0.0235385131158924), (22, 0.02342870936847765), (40, 0.023337409433088756),
    (4, 0.021348369572199), (33, 0.021329164328575786), (26, 0.021247073667926063),
    (1, 0.021220371606918108), (5, 0.021200492745102672), (30, 0.02107862126585502),
    (29, 0.021044721497828905)]
    Number of iterations: 17
```

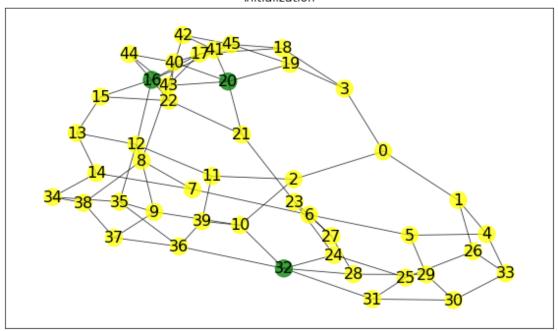
```
[7]: communities = initialize_communities(r, neighbors, 3)
    A = communities[0]
    print(f"Neighbors of {A}: {neighbors[A]}")
    B = communities[1]
    print(f"Neighbors of {B}: {neighbors[B]}")
    C = communities[2]
    print(f"Neighbors of {C}: {neighbors[C]}")

    print(f"Centers are: {A}, {B}, {C}")

Neighbors of 16: [15, 17, 44, 12, 45, 22]
    Neighbors of 32: [24, 28, 31, 36, 10]
    Neighbors of 20: [19, 21, 41, 40, 43]
    Centers are: 16, 32, 20
```

I don't love these centers... But I don't doubt my pagerank solution, so I'm not sure!

Initialization



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Run BigCLAM on the Tutte graph from Problem 2.

- \square Use 3 communities AND a background community such that the probability of any 2 members sharing an edge from the background community is 1%.
- □ Initialize your communities with your solution to Problem 2. If you are unable/did not yet to solve problem 2, you may use the (incorrect) values in the provided code of [19,28,34].
- ☑ You may iterate over the nodes in the order of their index instead of shuffling them.
- \boxtimes Run until convergence by the L_2 norm (element-wise over F) to within 10^{-6} .
- \square At the end, create 3 side-by-side plots of the original graph colored by the 3 community affiliation columns.

```
[]: #Do the thing
```

```
F = np.zeros((num_nodes, num_communities + 1))
         for j, i in enumerate(centers):
             F[i, j] = 1
         F[:,-1] = 0.01
         return F
     F = initialize_F(centers, neighbors, len(nodes))
     print(F)
     c = 3
     comm sums=F.sum(0) #sum over all communities, cached for updating in later
     \rightarrow version
     nu=.1 #step size
     for lezgo in range(100): #number of iterations
         for i in nodes: #loop over rows of F
             F_grad=np.zeros(c) #initialize gradient
             comm_sums=F.sum(0) #should be cached, will deal with later since it's_
      →annoying to update correctly with the O-check
             for j in neighbors[i]: #loop over nbrs of that row's node
                     IP=np.exp(-np.inner(F[i], F[j])) #so we don't have to double_
      \rightarrow compute
                     print("IP: ", IP)
                     F_grad+=F[j][:c]*(IP/(1-IP)+1) ##"first sum" below.
                     print("F_grad: ", F_grad)
             step=nu*(F_grad-comm_sums[:c]-F[i][:c]) #
             print(step)
             F[i][:c]+=step #update row; doing as two steps so can use "step" tou
      →update comm_sums later
             F[i,F[i,:]<0]=0 #0-check
             #print('i',i,'step:', F_grad, 'baseline:',comm_sums) # sanity checks
     print(F)
[]: '''
     My own attempt
     111
     class bigCLAM:
         def __init__(self, edges, k, centers, background_prob=None, step_size=0.1):
             # edge list
             self.edges
                            = edges
             # node list
             self.nodes
                           = list(set([node for edge in edges for node in edge]))
             # dictionary that contains the neighbors of node[i]
             self.neighbors = self.get_neighbors(self.nodes, self.edges)
             # number of nodes/people/items
                            = len(self.nodes)
             self.n
             # number of communities
```

```
self.k
               = k
    if not background_prob:
        self.epsilon\_comm = ((2 * len(edges)) / (self.n * (self.n - 1)))
        self.epsilon_comm = background_prob
    print(((2 * len(edges)) / (self.n * (self.n - 1))) ** 2)
    #initializing F (membership preferance matrix)
    self.F = self.initialize_F(centers)
    # defining our term for convergence check
    self.tolerance = 10e-6
    # step-size nu or sometimes eta
    self.eta = step_size
    # background_prob is called epsilon
# initialize F with the given centers if
def initialize_F(self, centers):
    F = np.zeros((self.n, self.k + 1))
    for j, i in enumerate(centers):
        F[i, j] = 1
    F[:,-1] = self.epsilon_comm
    print(F)
    return F
def get_neighbors(self, nodes, edges):
    neighbors= {node: [] for node in nodes}
    for edge in edges:
        neighbors[edge[0]].append(edge[1])
        neighbors[edge[1]].append(edge[0])
    return neighbors
def get_A(self, edges, nodes):
    A = np.zeros((self.n, self.n))
    for edge in edges:
        i = nodes.index(edge[0])
        j = nodes.index(edge[1])
        A[i,j] = 1
        A[j,i] = 1
    return A
def gradient(self, u):
    sum Fv all
                         = self.F.sum(0)
      print(sum_Fv_all)
                         = self.F[u]
    Fu
      print(Fu)
                         = np.zeros(self.k)
    sum_neighbors
    SIGMA_neighbors
                         = np.zeros(self.k)
    for neighbor in self.neighbors[u]:
```

```
print(self.F[neighbor])
                             = self.F[neighbor]
           print("Fu: ", Fu, "Fv: ", Fv)
           dot_product
                           = -1. * Fu.dot(Fv)
           print("dot_product: ", dot_product)
           print(Fv[:self.k])
           sum neighbors += Fv[:self.k]
           print("np.exp(dot_product): ", np.exp(dot_product))
           print("(1 - np.exp(dot_product)): ", (1 - np.exp(dot_product)))
           print("np.exp(dot_product)/(1 - np.exp(dot_product)): ", np.
→exp(dot_product)/(1 - np.exp(dot_product)))
           print("Fv * (np.exp(dot_product)/(1 - np.exp(dot_product))): ", Fv⊔
→* (np.exp(dot_product)/(1 - np.exp(dot_product))))
           SIGMA_neighbors += Fv[:self.k] * (np.exp(dot_product) / (1 - np.
→exp(dot_product)))
       print(sum_Fv_all, Fv, sum_neighbors)
       SIGMA_non_neighbors = sum_Fv_all[:self.k] - Fu[:self.k] - sum_neighbors
       print("SIGMA_neighbors: ", SIGMA_neighbors)
       print("SIGMA_non_neighbors: ", SIGMA_non_neighbors)
       print("SIGMA_neighbors - SIGMA_non_neighbors: ", SIGMA_neighbors -_
→SIGMA_non_neighbors)
       return SIGMA_neighbors - SIGMA_non_neighbors
         sum Fv neighbors
                             = np.sum([self.F[i] for i in self.neighbors])
#
         sum_non_neighbors = sum_Fv_all - Fu - sum_Fv_neighbors
         F[i, F[i, :] < 0] = 0
#
         print(sum__non_neighbors)
   def LL(self):
       pass
   def run(self):
       converged = False
       while not converged:
           F_old = self.F.copy()
           for node in self.nodes:
                grad = self.gradient(node)
               print("grad: ", grad)
               print("self.F[node][:self.k]: ", self.F[node][:self.k])
                self.F[node][:self.k] += self.eta * grad[:self.k]
               print(self.F[node])
                self.F[i, self.F[i,:] <= 0] = 0
            if np.linalg.norm(F_old - self.F, ord=2) < self.tolerance:</pre>
               converged = True
       print("F: ", self.F)
```

```
[]: k_communities
     epsilon_community = 0.1
     print(np.sqrt(epsilon_community))
     bc = bigCLAM(edges, k_communities, centers, epsilon_community)
     bc.run()
[]: '''
     An attempt at taking someone else's bigclam to at least see if I could get it_{\sqcup}
     \hookrightarrow to work
     111
     def sigm(x):
         return np.divide(np.exp(-1.*x),1.-np.exp(-1.*x))
     def log likelihood(F, A):
         """implements equation 2 of
         https://cs.stanford.edu/people/jure/pubs/biqclam-wsdm13.pdf"""
         A_soft = F.dot(F.T)
         # Next two lines are multiplied with the adjacency matrix, A
         # A is a {0,1} matrix, so we zero out all elements not contributing to the
      → sum
         FIRST_PART = A*np.log(1.-np.exp(-1.*A_soft))
         sum_edges = np.sum(FIRST_PART)
         SECOND PART = (1-A)*A soft
         sum_nedges = np.sum(SECOND_PART)
         log_likeli = sum_edges - sum_nedges
         return log_likeli
     def gradient(F, A, i):
         """Implements equation 3 of
         https://cs.stanford.edu/people/jure/pubs/bigclam-wsdm13.pdf
           * i indicates the row under consideration
         The many forloops in this function can be optimized, but for
         educational purposes we write them out clearly
         n n n
         N, C = F.shape
         neighbours = np.where(A[i])
         print(neighbours)
         nneighbours = np.where(1-A[i])
         print(nneighbours)
         sum_neigh = np.zeros((C,))
```

```
for nb in neighbours[0]:
       print("Fu: ", F[i], "Fv: ", F[nb])
        dotproduct = F[nb].dot(F[i])
        print("dotproduct: ", dotproduct)
        print("sigm(dotproduct): ", sigm(dotproduct))
        sum_neigh += F[nb]*sigm(dotproduct)
   sum_nneigh = np.zeros((C,))
    #Speed up this computation using eq.4
   for nnb in nneighbours[0]:
       print("nnb: ",nnb)
       sum_nneigh += F[nnb]
   print(sum_neigh)
   print(sum_nneigh)
   print(sum_neigh - sum_nneigh)
   grad = sum_neigh - sum_nneigh
   return grad
def initialize_F(centers, n, k, neighbors):
   F = np.zeros((n, k))
   for j, i in enumerate(centers):
       F[i, j] = 1
   return F
def get_neighbors(nodes, edges):
   neighbors= {node: [] for node in nodes}
   for edge in edges:
       neighbors[edge[0]].append(edge[1])
       neighbors[edge[1]].append(edge[0])
   return neighbors
def train(A, C, edges, nodes, centers, iterations = 100):
    # initialize an F
   N = A.shape[0]
    F = np.random.rand(N,C)
   neighbors = get_neighbors(nodes, edges)
   F = initialize_F(centers, len(nodes), C, neighbors)
   print(F)
   for n in range(iterations):
        for person in range(N):
            grad = gradient(F, A, person)
            F[person] += 0.005*grad
            F[person] = np.maximum(0.001, F[person]) # F should be nonnegative
        11 = log_likelihood(F, A)
```

```
print('At step %5i/%5i ll is %5.3f'%(n, iterations, ll))
return F

F = train(mat, 3, edges, nodes, centers)
F_argmax = np.argmax(F,1)

for i,row in enumerate(F):
    print(row)
```

```
[]: #just some testing

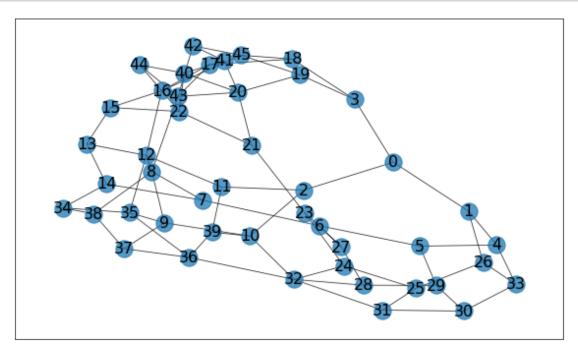
M = [[1,1,1],[0,0,0],[1,0,1],[0,1,0]]
A = np.matrix(M)
for i in range(A.shape[0]):
    print(f"{i}, 1: ", np.where(A[i]))
    print(f"{i}, 0: ", np.where(1-A[i]))
```

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Consider doing cuts or finding non-overlapping communities on the same graph as we used in problems 2-3.

```
[9]: | #NB: this is a well known graph known as the "Tutte" graph
     edges=[(0, 1), (0, 2), (0, 3), (1, 4), (1, 26), (2, 10), (2, 11), (3, 18), (3, 18)]
      →19), \
             (4, 5), (4, 33), (5, 6), (5, 29), (6, 7), (6, 27), (7, 8), (7, 14), (8, 1)
      →9), (8, 38), \
             (9, 10), (9, 37), (10, 39), (11, 12), (11, 39), (12, 13), (12, 35), (13, u)
      \rightarrow14), (13, 15),\
             (14, 34), (15, 16), (15, 22), (16, 17), (16, 44), (17, 18), (17, 43),
      \rightarrow (18, 45), (19, 20), \
             (19, 45), (20, 21), (20, 41), (21, 22), (21, 23), (22, 40), (23, 24),
      \rightarrow (23, 27), (24, 25), \
             (24, 32), (25, 26), (25, 31), (26, 33), (27, 28), (28, 29), (28, 32),
      \rightarrow (29, 30), (30, 31), \
             (30, 33), (31, 32), (34, 35), (34, 38), (35, 36), (36, 37), (36, 39), 
      \hookrightarrow (37, 38), (40, 41), \
             (40, 44), (41, 42), (42, 43), (42, 45), (43, 44)
     #add a few edges for fun and value and to break symmetry
     newedges=[(32,36), (32,10), (20,40), (20,43), (8,12), (8,43), (12,16), (35,9), 
     \rightarrow (16,45), (16,22)]
     for newedge in newedges:
         edges.append(newedge)
     nodes=list(range(46))
     neighbors={key: [] for key in range(len(nodes))}
```

```
for edge in edges:
    neighbors[edge[0]].append(edge[1])
    neighbors[edge[1]].append(edge[0])
G=nx.Graph()
G.add_nodes_from(nodes)
for i in range(len(edges)):
    G.add_edge(edges[i][0],edges[i][1])
np.random.seed(4022)
pos = nx.spring_layout(G) #algorithm that tries to "cluster" node plot_
→ locations; it saves the xy coords of each vertex
labels=dict(zip(range(46), range(46))); #label 0-45
fig, ax = plt.subplots(figsize=(PLOT_SIZE_X,PLOT_SIZE_Y))
nx.draw_networkx_nodes(G, pos, node_size=300, alpha=.75); #node colors easy to_
⇔see labels over
nx.draw_networkx_edges(G, pos, width=0.8, alpha=0.8); #draw edges, using_
\rightarrow xy-node layout provided
nx.draw_networkx_labels(G, pos, labels, font_size=FONT_SIZE); #label vertices_
 \hookrightarrow by index/number
```



1.1.1 Part 1:

Find the Fiedler vector for the graph. Plot the sorted Fiedler vector. Use the edges array defined above.

```
[10]: class Partition:
          # init function: edges is either a standard square edge-matrix, or a list_{\sqcup}
       \hookrightarrow of edges
          def __init__(self, edges):
              self.edges = edges
              if isinstance(edges, list):
                  self.nodes
                               = list(set([edge[0] for edge in edges] + [edge[1]]
       →for edge in edges]))
                  self.n
                                  = len(self.nodes)
                   self.compute_A = True
              elif isinstance(edges, np.matrix):
                   self.nodes
                                = list(range(edges.shape[0]))
                                  = len(self.nodes)
                  self.n
                  self.compute_A = False
              else:
                  self.nodes
                                = None
                                  = 0
                  self.n
                  self.compute A = False
                  print("Wrong Edges format!")
          def is_power2(self, n):
              return True if (n \& (n-1) == 0) and n != 0 else False
          # Retruns the standard square matrix of edges for a graph, as a list of \Box
       →edges as a list of numerical nodes
          # NOTE: maybe one day it would be helpful to rewrite this class to use_
       → dicts so nodes can be anything
          def get_A(self, edges, nodes):
              n = len(nodes)
              A = np.zeros((n, n))
              for edge in edges:
                  i = nodes.index(edge[0])
                  j = nodes.index(edge[1])
                  A[i,j] = 1
                  A[j,i] = 1
              return A
          # Computes the edge degree for each node returned as a square matrix_
       \hookrightarrow diagonally
          def get_D(self, A):
              n = A.shape[0]
              D = np.diag([np.sum(A[:,i]) for i in range(n)])
```

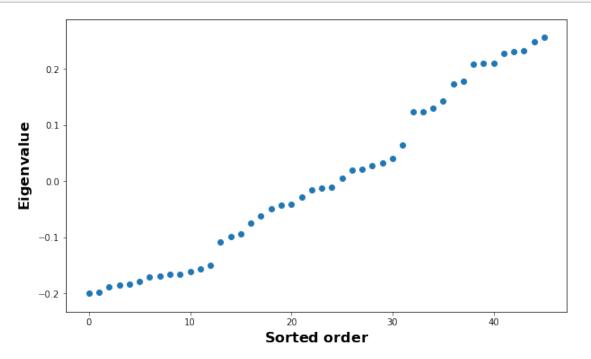
```
return D
   def get_L(self, edges):
       # fairly straightforward... Computing the Laplacian (I believe...),
→stripped straight from lecture/notebook
       A = self.get A(edges)
       D = self.get_D(A)
       L = D - A
       return L
   def get_nodes_from_edges(self, edges):
       s = set()
       for edge in edges:
           s.add(edge[0])
           s.add(edge[1])
       return list(s)
   def get_edges(self, g1, g2, edges):
       e1, e2 = [], []
       partitioned_edges = []
       for edge in edges:
           if edge[0] in g1 and edge[1] in g1:
               e1.append(edge)
           elif edge[0] in g2 and edge[1] in g2:
               e2.append(edge)
       return e1, e2
   def get_all_edges(self, edges, graphs):
       return [edge for edge in edges for graph in graphs if edge[0] in graph_
→and edge[1] in graph]
   def get_fiedler(self, edges, nodes):
       # basically, if edges is given as a matrix, we already have A... So I_{\sqcup}
\rightarrow figured I'd make this a
       # class variable
       if self.compute_A:
           A = self.get_A(edges, nodes)
       else:
           A = edges
       D = self.get_D(A)
       L = D - A
       # get the eigen-stuff
       eigen_vals, eigen_vecs = np.linalg.eig(L)
       # sort the eigenvalues so we can find the correct eigenvector
       sorted_eigen_vals = np.sort(eigen_vals)
       # I should've used the syntax from the notebook, because this gave me_
\hookrightarrowsome issues... However, I figured how
```

```
# to make this work!
       index, = np.where(eigen_vals == sorted_eigen_vals[1])
       # NOTE: :, index[1] makes it work... We're looking for the second
→smallest eigen-pair (Not sure if this is
       # correct mathematical-jargon). Thus, due to Numpy docs, we find the
\rightarrow corresponding eigenalue v[i] for
       # eigenvector w[:,i]:
       # https://numpy.orq/doc/stable/reference/generated/numpy.linalq.eig.html
       fiedler_vec = eigen_vecs[:,index[0]]
       # sort the eigenvector found in previous line
       sorted_fiedler_vec = np.sort(fiedler_vec)
       # sort the indices so we can get the correct order of nodes
       sorted_indices = np.argsort(fiedler_vec)
       true_indices = [nodes[i] for i in sorted_indices]
       return sorted_fiedler_vec, true_indices
   def plot_fiedler(self):
       # plots a sorted fiedler vector
       sorted_fiedler_vec, sorted_indices = self.get_fiedler(self.edges, self.
→nodes)
       fig, ax = plt.subplots(1,1,figsize=(PLOT_SIZE_X,PLOT_SIZE_Y))
       plt.scatter(self.nodes, sorted_fiedler_vec)
       plt.xlabel(r'$\bf{Sorted\ order}$', fontsize=FONT SIZE)
       plt.ylabel(r'$\bf{Eigenvalue}$', fontsize=FONT_SIZE)
   def Bi_partition_helper(self, edges, graph):
       eig_vec, indices = self.get_fiedler(edges, graph)
       g1, g2 = [], []
       for i in range(len(indices)):
           index = indices[i]
           if eig_vec[i] <= 0:</pre>
               g1.append(index)
           else:
               g2.append(index)
       return g1, g2
   def Bi_partition(self, k):
       graph list = []
       edge_list = []
       for i in range(k // 2):
           if i == 0:
               g1, g2 = self.Bi_partition_helper(self.edges, self.nodes)
               graph_list.append(g1)
               graph_list.append(g2)
               e1, e2 = self.get_edges(g1, g2, self.edges)
               edge_list.append(e1)
               edge_list.append(e2)
```

```
else:
               for ii in range(len(graph_list)):
                    g1, g2 = self.Bi_partition_helper(edge_list[ii],__
graph_list.append(g1)
                    graph list.append(g2)
                    e1, e2 = self.get_edges(g1, g2, edge_list[ii])
                    edge list.append(e1)
                    edge_list.append(e2)
       return graph_list[-k:], edge_list[-k:]
   def K_partition(self, k):
       fiedler_vec, indices = self.get_fiedler(self.edges, self.nodes)
       # simply getting the eigenvalue difference for each value in the
\rightarrow eigenvector
       eig_diffs = [np.abs(fiedler_vec[i] - fiedler_vec[i + 1]) for i in_
→range(len(indices) - 1)]
       # initializing the indices of where to cut the graph
       # the crazy line: sorting the indices of the k-1 largest eigenvalue _{\sqcup}
\hookrightarrow differences
       # while I could've done some iterative method, this could be a_{\sqcup}
→relatively incredibly helpful numpy hack
       # thus I googled a good way to do this.
       # https://stackoverflow.com/questions/6910641/
\rightarrow how-do-i-get-indices-of-n-maximum-values-in-a-numpy-array
       k_indices = [0] + list(np.sort(np.array(eig_diffs).argsort()[-(k-1):][::
\hookrightarrow-1])) + [len(indices)]
       graphs
                 = [list(indices[k_indices[i]:k_indices[i + 1]]) for i in_
→range(len(k_indices) - 1)]
                 = self.get_all_edges(self.edges, graphs)
       return graphs, edges
   def Partition(self, k):
       if self.is power2(k):
           graphs, edges = self.Bi_partition(k)
       else:
           graphs, edges = self.K_partition(k)
       return graphs, edges
   def get_pos(self):
       G=nx.Graph()
       G.add_nodes_from(self.nodes)
       for i in range(len(self.edges)):
           G.add_edge(self.edges[i][0], self.edges[i][1])
       np.random.seed(4022)
```

```
pos = nx.spring_layout(G) #algorithm that tries to "cluster" node plot⊔
→ locations; it saves the xy coords of each vertex
       labels=dict(zip(self.nodes, self.nodes))
       for i in range(len(self.edges)):
           G.remove_edge(self.edges[i][0], self.edges[i][1])
       return pos
   def get_colors(self, graphs):
       color_dict = {0: 'yellow', 1: 'red', 2: 'green', 3: 'orange', 4:
return [color_dict[i] for node in self.nodes for i in_
→range(len(graphs)) if node in graphs[i]]
   def get_uncut_edges(self, edges):
       return [edge for 1 in edges for edge in 1]
   def get_cuts(self, uncut_edges):
       cut_edges = list(set(self.edges).difference(set(uncut_edges)))
       nodes = set()
       for cut_edge in cut_edges:
           nodes.add(cut_edge[0])
           nodes.add(cut_edge[1])
       return list(nodes), cut_edges
   def Graph_Partitions(self, k):
       graphs, edges = self.Partition(k)
       pos = self.get_pos()
       colors = self.get_colors(graphs)
       if k % 2 == 0:
           uncut_edges = self.get_uncut_edges(edges)
       else:
           uncut_edges = edges
       cut_nodes, cut_edges = self.get_cuts(uncut_edges)
       G=nx.Graph()
       G.add_nodes_from(self.nodes)
       for i in range(len(uncut_edges)):
           G.add_edge(uncut_edges[i][0], uncut_edges[i][1])
       fig, ax = plt.subplots(figsize=(PLOT_SIZE_X,PLOT_SIZE_Y))
       nx.draw_networkx_nodes(G, pos, node_color=colors, node_size=300, alpha=.
\rightarrow75); #node colors easy
       nx.draw_networkx_edges(G, pos, width=1, alpha=0.8); #draw edges, using_
\rightarrow xy-node layout provided
       nx.draw_networkx_labels(G, pos, labels, font_size=FONT_SIZE); #label_
→vertices by index/number
       G2=nx.Graph()
       G2.add_nodes_from(cut_nodes)
```

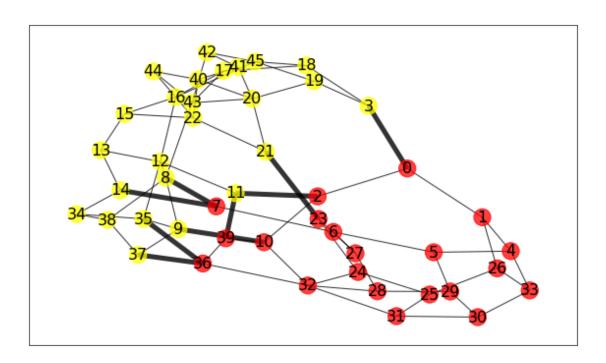
```
[11]: Q4_P1 = Partition(edges)
Q4_P1.plot_fiedler()
```



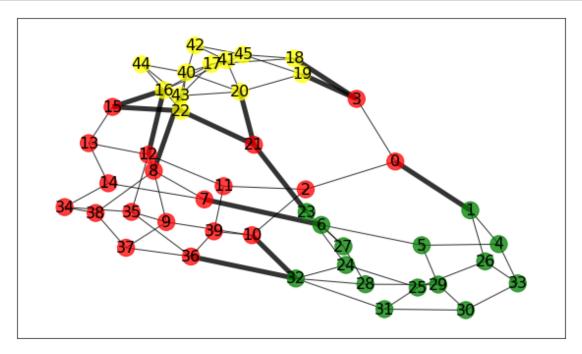
1.1.2 Part 2: Partitions

Part 2a: Partition the graph into 2 graphs using this vector. Include a picture of the associated cut of the graph.

```
[12]: Q4_P2a = Partition(edges)
Q4_P2a.Graph_Partitions(2)
```



Part 2b: Partition the graph into 3 subgraphs using the Fiedler vector. One natural way to do so is to partitition at the two largest jumps in the vector and not use y = 0. Include both associated *cuts* in another picture of the graph.



1.1.3 Part 2c:

Partition the graph into 4 subgraphs using bipartitioning, where you cut once and then cut the resulting subgraphs according to *their* Fiedler vectors.

```
[14]: Q4_P2c = Partition(edges)
Q4_P2c.Graph_Partitions(4)

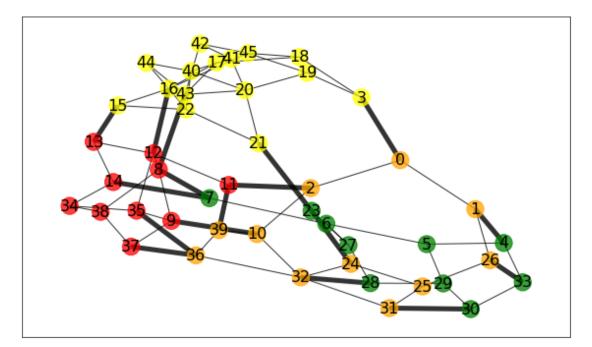
# partitioned graphs

# [3, 19, 18, 41, 45, 20, 42, 21, 40, 17, 44, 22, 43, 16, 15],

# [13, 12, 8, 11, 14, 35, 9, 38, 34, 37],

# [7, 6, 27, 5, 23, 29, 28, 4, 30, 33],

# [24, 25, 26, 31, 1, 32, 0, 36, 10, 2, 39]
```



1.1.4 Part 2d:

Of the 3 partitions above, which do you think were most appropriate. Why?

While my graph partition with 4 graphs makes some sense, I'm not 100% sure it's correct. Assuming it is correct though, I think partitioning the graph with either two or three subgraphs makes the most sense in this example. Two or three subgraphs still produce relatively 'clustered' graphs, and four subgraphs has a tough time clustering two of the subgraphs. This made me think there might

be some sort of bug in my code when I bipartition twice, however, when looking closely at the cuts in the original graph, my four subgraph's cuts seem reasonable.

Back to top # Extra Credit (Rupping RigCI AM on Rig Data: up to ±25 pts). This is

Back to top # Extra Credit (Running BigCLAM on Big Data: up to +25 pts). This problem may be turned in either with HW7 or HW8.

Consider the data set Marvel_Network. This set consists of two columns, hero1 and hero2. Every row is filled with co-occurrence of two such marvel characters in a comic.

1.1.5 Part A) (10 pts) Cleaning and setup

(i) Use item baskets to count how many times each character appears. You may also want to count how many times each *edge* between pairs of characters appears (use your code from A Priori!) Print the most popular 5 characters' names.

```
[15]: #Names and counts
df = pd.read_csv('Marvel_Network.csv', encoding='unicode_escape')
df.head()
```

```
[15]: hero1 hero2

0 LITTLE, ABNER PRINCESS ZANDA

1 LITTLE, ABNER BLACK PANTHER/T'CHAL
2 BLACK PANTHER/T'CHAL PRINCESS ZANDA
3 LITTLE, ABNER PRINCESS ZANDA
4 LITTLE, ABNER BLACK PANTHER/T'CHAL
```

```
[16]: class APriori:
          def __init__(self, cleaned_df, basket_column, item_column, k_tuples,_
       →support_threshold):
                                 = cleaned df
              self.df
              self.k
                                 = k_tuples
              self.n
              # trying to dip my toes in error/exception handling... kinda messy at\Box
       \rightarrow the moment
              try:
                  length
                                 = len(cleaned df[basket column].unique())
              except KeyError as a:
                  raise NameError("Basket Column is invalid!") from a
              else:
                  self.n_baskets = length
                           = int(support_threshold * length)
                  self.s
                                 = basket column
                  self.b_col
              try:
                  _ = cleaned_df[item_column]
              except KeyError as b:
```

```
raise NameError("Item Column is invalid!") from b
       else:
           self.i_col
                          = item_column
   def first_pass(self):
       counts = {}
       # counting step that I am resuing from before
       for basket in (self.df.groupby(self.b_col)[self.i_col].apply(list).
→values):
           for item in basket:
               if item not in counts.keys():
                   counts[item] = 0
               counts[item] += 1
       return prune(counts)
   def prune(counts):
       for key in counts.copy():
           if counts[key] < self.s:</pre>
               try:
                   del counts[key]
               except KeyError:
                   print("Couldn't remove key: {}".format(key))
       self.n = len(counts)
       return counts
   def performAPriori(self, use_hashing=False):
       def second_pass(single_counts):
                         = single_counts.keys()
           double_counts = {}
           # loop through each basket
           for i, basket in enumerate((self.df.groupby(self.b_col)[self.i_col].
→apply(list).values)):
               # loop through each item only if its a frequent item
               length = len(basket)
               for i1 in (range(length)):
                   # loop through each item only if its a frequent item and
                   # it's not equal to item in the loop above
                   for i2 in (range(i1+1, length)):
                        # weird stuff to make sure duplicate keys are not added_
\rightarrow to dict
                       ks = double_counts.keys()
                       t = (basket[i1], basket[i2])
                       if t not in ks:
                            double_counts[t] = 0
                       double_counts[t] += 1
```

```
return self.prune(double_counts), confidence(single_counts,_
→double_counts)
       def confidence(old counts, new counts):
           confidences = []
           for s old, c old in old counts.items():
               set_old = set([s_old]) if type(s_old) != tuple else set(s_old)
               for s_new, c_new in new_counts.items():
                   set_new = set(s_new)
                   if set(set_old).issubset(set_new):
                                = set_new.difference(set_old)
                       difference = list(d_set)[0] if len(d_set) == 1 else_
→tuple(d_set)
                                  = tuple([(c_new / c_old), "{} -> {}".
                       tup
→format(s_old, difference), c_new])
                       confidences.append(tup)
                                  = old_counts[difference]
                                  = tuple([(c_new / c_diff), "{} -> {}".
→format(difference, s_old), c_new])
                       confidences.append(tup)
           return confidences
       if use hashing:
           pass
       frequent_items
                                      = self.first_pass()
       frequent_pairs, confidences
                                      = second_pass(frequent_items)
       sorted_freq_pairs = [(key, val) for key, val in sorted(frequent_pairs.
→items(), key=lambda x: x[1], reverse=True)]
       sorted_confidences = [(i[0], i[1]) for i in sorted(confidences,__
→key=lambda x: x[2], reverse=True)]
       return sorted_freq_pairs, sorted_confidences
```

[]:

(ii) Remove any characters having degree = 1. What do these nodes represent?

(iii) For speed of computation, you should also remove any *edges* with count 1. What do these *edges* represent?

[]:

(iv) Create an adjacency matrix (or probably a compact/sparse representation of such a matrix!) for the data after accounting for parts (ii), (iii). How many characters are there? How many edges?

[]:

1.1.6 Part B) (10 pts) Detect communities in the graph generated above using Big-CLAM

For this problem, take the number of communities to find as 4 (plus a background community!). Use your choice of initialization. If you have **prior knowledge** of any comic book characters, consider setting 4 characters from "different" Marvel universes as your seeds. If not, either use PageRank or properties of your adjacency matrix to decide where to initialize affiliations.

Are the final communities ones you've seen in popular culture? Verify by checking the community affiliation scores of two characters that you believe should share a dominant community, like Wolverine and Professor X.

[]:

1.1.7 Part C) (5 pts) Detect communities in the graph generated above using weighted BigCLAM

It turns out that some edges occurred multiple times in the data set. We can adjust our model to count the edges proportionately to the number of times they occurred: we just have to weight the partial derivatives in our gradient calculation by multiplying each term v in ∇F_u by the number of times u and v were seen together.

Run the model accordingly, and report the community affiliations of the same pair of characters you looked at in Part B). They should be even closer now, right?

[]: