## DiagnosticsAndCommunityDetection

December 27, 2020

```
[1]: import numpy as np
     from urllib.request import urlopen
     import scipy.optimize
     import random
     from sklearn.decomposition import PCA
     from collections import defaultdict
     from sklearn.linear_model import LogisticRegression
     from sklearn.metrics import accuracy_score
     from sklearn.metrics import mean_squared_error as MSE
     from sklearn.linear_model import RidgeClassifier
     import pandas as pd
[2]: def parseDataFromURL(fname):
       for 1 in urlopen(fname):
         yield eval(1)
     def parseData(fname):
       for 1 in open(fname):
         yield eval(1)
     print("Reading data...")
     # Download from http://jmcauley.ucsd.edu/cse255/data/beer/beer_50000.json"
     data = list(parseData("data/beer_50000.json"))
     print("done")
    Reading data...
    done
[]:
[]:
[3]: #select data
     categoryCounts = defaultdict(int)
     for d in data:
         categoryCounts[d["beer/style"]] += 1
     categories = [c for c in categoryCounts if categoryCounts[c] > 1000]
```

```
catID = dict(zip(list(categories),range(len(categories))))
[4]: catID
[4]: {'American Double / Imperial IPA': 0,
     'Rauchbier': 1,
     'American Pale Ale (APA)': 2,
     'American Porter': 3,
     'Russian Imperial Stout': 4,
     'American IPA': 5,
     'Fruit / Vegetable Beer': 6,
     'American Double / Imperial Stout': 7,
     'Rye Beer': 8,
     'Scotch Ale / Wee Heavy': 9,
     'English Pale Ale': 10,
     'Czech Pilsener': 11,
     'Old Ale': 12}
[]:
[]:
[5]: idx = catID.values()
    ohe = np.array(np.eye(max(idx) +1).astype(int))
[6]:
    ohe_map = dict(zip(list(categories),ohe))
[7]:
    ohe_map
0]),
     'Rauchbier': array([0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]),
     'American Pale Ale (APA)': array([0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0]),
     'American Porter': array([0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0]),
     'Russian Imperial Stout': array([0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0]),
     'American IPA': array([0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0]),
     'Fruit / Vegetable Beer': array([0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0]),
     'American Double / Imperial Stout': array([0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0,
    0]),
     'Rye Beer': array([0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0]),
     'Scotch Ale / Wee Heavy': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]),
     'English Pale Ale': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0]),
     'Czech Pilsener': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0]),
     'Old Ale': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1])}
[]:
```

```
[]:
 []:
 []:
 [8]: for d in data:
          if d['beer/style'] in categories:
              d['beer/style'] = ohe_map[d['beer/style']]
          else:
              d['beer/style'] = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0])
 []:
 []:
 [9]: x = np.array([d['beer/style'] for d in data])
 []:
[10]: ones = np.ones((len(x),1))
      ones
[10]: array([[1.],
             [1.],
             [1.],
             ...,
             [1.],
             [1.],
             [1.]]
[11]: X = np.hstack([ones,x])
      x_1 = [[d['review/appearance'],d['review/palate'], \
             d['review/taste'],d['review/overall'],d['review/aroma']] for d in data]
      x_2 = [len(d['review/text']) for d in data]
      x_2 = x_2 / max(np.array(x_2))
      x_2 = np.reshape(x_2, (-1, 1))
      X = np.hstack([X,x_1])
      X = np.hstack([X,x_2])
      X
[11]: array([[1.
                       , 0.
                                    , 0.
                                                , ..., 1.5
                                                               , 2.
              0.05549672],
             [1.
                        , 0.
                                    , 0.
                                                , ..., 3.
                                                                , 2.5
              0.071595 ],
```

```
, 0.
             [1.
                                    , 0.
                                         , ..., 3. , 2.5
              0.08388053],
                        , 0.
             [1.
                                    , 0.
                                                , ..., 3.5
                                                               , 3.5
              0.13577632],
                                                , ..., 4.
             [1.
                        , 0.
                                    , 0.
                                                                , 4.
              0.20800678],
             [1.
                        , 0.
                                    , 0.
                                                , ..., 4.5
                                                               , 4.
              0.09510697]])
 []:
[12]: y = [d['beer/ABV'] > 7.0 \text{ for d in data}]
 []:
[13]: #shuffle data
      Xy = list(zip(X,y))
      random.shuffle(Xy)
      X = np.array([d[0] for d in Xy])
      y = np.array([d[1] for d in Xy])
 []:
[14]: X = X[:len(x)//2]
      Xtest = X[len(x)//2:]
      ytrain = y[:len(x)//2]
      ytest = y[len(x)//2:]
 []:
```

0.1 1) Train a logistic regressor using this one-hot encoding to predict whether beers have an ABV greater than 7 percent (i.e., d['beer/ABV'] > 7). Train the classifier on the training set and report its performance in terms of the accuracy and Balanced Error Rate (BER) on the test set, using a regularization constant of C = 10. For all experiments use the class weight='balanced' option (2 marks).

```
TN_ = np.logical_and(np.logical_not(pred), np.logical_not(ytest))
FN_ = np.logical_and(np.logical_not(pred), ytest)

TP = sum(TP_)
FP = sum(FP_)
TN = sum(TN_)
FN = sum(FN_)
FPR = FP/(FP+TN)
FNR = FN/(FN+TP)
print("balanced error rate: ", (1/2)*(FPR+FNR))
print('accuracy: ', accuracy_score(ytest, pred))
```

balanced error rate: 0.16068652372986186

accuracy: 0.84876

[]:

0.2 2) Extend your model to include two additional features: (1) a vector of five ratings (review/aroma,review/overall, etc.); and (2) the review length (in characters). The length feature should be scaled to be between 0 and 1 by dividing by the maximum length. Using the same value of C from the previous question, report the BER of the new classifier (1 mark).

balanced error rate: 0.14223937720215946

accuracy: 0.8626

0.3 3) Implement a complete regularization pipeline with the balanced classifier. Split your test data from above in half so that you have 50%/25%/25% train/validation/test fractions. Consider values of C in the range $\{10^-6, 10^-5, 10^-4, 10^-3\}$ . Report (or plot) the train, validation, and test BER for each value of C. Based on these values, which classifier would you select (in terms of generalization performance) and why (1 mark)?

```
[]:
[17]: def mse(model, X, y):
          preds = model.predict(X)
          diffs = [(float(a)-float(b))**2 for (a,b) in zip(preds,y)]
          return sum(diffs) / len(diffs)
[18]: N = len(v)
      Ntrain = N//2
      Nvalid = N//4
      Ntest = N//4
      Xtrain = X[:Ntrain]
      Xvalid = X[Ntrain:Ntrain+Nvalid]
      Xtest = X[Ntrain+Nvalid:]
      ytrain = y[:Ntrain]
      yvalid = y[Ntrain:Ntrain+Nvalid]
      ytest = y[Ntrain+Nvalid:]
[19]:
      C = [10**-6, 10**-5, 10**-4, 10**-3]
      bestModel = None
      bestMSE = None
      for c in C:
          model = LogisticRegression(C = c, class_weight = 'balanced').fit(Xtrain,_
       →ytrain)
          preds = [model.predict(Xtrain), model.predict(Xvalid), model.predict(Xtest)]
          ys = [ytrain, yvalid, ytest]
          mse_tr = mse(model, Xtrain, ytrain)
          mse_v = mse(model, Xvalid, yvalid)
          print("mse train/validation: " + str(mse_tr), str(mse_v))
          print("C value: ", str(c))
          titles = ["train BER: ", "valid BER: ", "test BER: "]
          for m in range(len(preds)):
              TP_ = np.logical_and(preds[m], ys[m])
```

```
FP_ = np.logical_and(preds[m], np.logical_not(ys[m]))
  TN_ = np.logical_and(np.logical_not(preds[m]), np.logical_not(ys[m]))
  FN_ = np.logical_and(np.logical_not(preds[m]), ys[m])

TP = sum(TP_)
  FP = sum(FP_)
  TN = sum(TN_)
  FN = sum(FN_)
  FNR = FP/(FP+TN)
  FNR = FN/(FN+TP)
    print(titles[m], (1/2)*(FPR+FNR))

print("\n")

if not bestModel or mse_v < bestMSE:
    bestModel = model
    bestMSE = (mse_v)</pre>
```

mse train/validation: 0.3216 0.32384

C value: 1e-06

train BER: 0.3167114713429558 valid BER: 0.3203850384473046 test BER: 0.31577497649604813

mse train/validation: 0.3198 0.32184

C value: 1e-05

train BER: 0.314951238004041 valid BER: 0.31842232410872157 test BER: 0.3145511464911047

mse train/validation: 0.29728 0.30072

C value: 0.0001

train BER: 0.29315496535476443 valid BER: 0.29805167770056334 test BER: 0.2905913557276848

mse train/validation: 0.19412 0.19136

C value: 0.001

train BER: 0.19575385402175077 valid BER: 0.19330076277807476 test BER: 0.18964224229757196

0.3.1 I would select the classifier with C value of .001, based on the lowest validation MSE and lowest test BER.

```
[20]: bestModel

[20]: LogisticRegression(C=0.001, class_weight='balanced')

[21]: bestMSE

[21]: 0.19136
```

0.4 4) (CSE158 only) An ablation study measures the marginal benefit of various features by re-training the model with one feature 'ablated' (i.e., deleted) at a time. Considering each of the three features in your classifier above (i.e., beer style, ratings, and length), report the BER with only the other two features and the third deleted (1 mark).

```
[22]: C = [10**-6,10**-5,10**-4,10**-3]
                   Xtrain_ = np.hstack([Xtrain[:,:14],Xtrain[:,-1:]])
                   Xtest_ = np.hstack([Xtest[:,:14],Xtest[:,-1:]])
                   Xvalid_ = np.hstack([Xvalid[:,:14], Xvalid[:,-1:]])
                   toFitX = [Xtrain[:,:-1],
                                                   Xtrain[:,-6:],
                                                   Xtrain 1
                   toPredTr = [Xtrain[:,:-1], Xtrain[:,-6:], Xtrain_ ]
                   toPredV = [Xvalid[:,:-1], Xvalid[:,-6:], Xvalid_]
                   toPredT = [Xtest[:,:-1], Xtest[:,-6:], Xtest_ ]
                   ablations = ["(ohe styles and ratings)", "(ratings and review length)", "(ohe⊔
                     →styles and review length)"]
                   for c in C:
                               print("C value: ", c)
                               for i in range(len(toFitX)):
                                            logreg = LogisticRegression(C=c, class_weight = 'balanced', max_iter = LogisticRegression(C=c, class_weight = LogisticRe
                      →400).fit(toFitX[i],ytrain)
                                            trpred = logreg.predict(toPredTr[i])
                                            vpred = logreg.predict(toPredV[i])
                                            pred = logreg.predict(toPredT[i])
                                            TP = np.logical and(trpred, ytrain)
                                            FP_ = np.logical_and(trpred, np.logical_not(ytrain))
                                            TN_ = np.logical_and(np.logical_not(trpred), np.logical_not(ytrain))
```

```
FN_ = np.logical_and(np.logical_not(trpred), ytrain)
       TP = sum(TP)
       FP = sum(FP_)
       TN = sum(TN)
       FN = sum(FN)
       FPR = FP/(FP+TN)
       FNR = FN/(FN+TP)
       print("training balanced error rate: " + ablations[i], (1/2)*(FPR+FNR))
       TP_ = np.logical_and(vpred, yvalid)
       FP_ = np.logical_and(vpred, np.logical_not(yvalid))
       TN_ = np.logical_and(np.logical_not(vpred), np.logical_not(yvalid))
       FN_ = np.logical_and(np.logical_not(vpred), yvalid)
       TP = sum(TP)
       FP = sum(FP_)
       TN = sum(TN)
       FN = sum(FN_{-})
       FPR = FP/(FP+TN)
       FNR = FN/(FN+TP)
       print("validation balanced error rate: " + ablations[i], (1/
\rightarrow2)*(FPR+FNR))
       TP_ = np.logical_and(pred, ytest)
       FP_ = np.logical_and(pred, np.logical_not(ytest))
       TN_ = np.logical_and(np.logical_not(pred), np.logical_not(ytest))
       FN_ = np.logical_and(np.logical_not(pred), ytest)
       TP = sum(TP)
       FP = sum(FP)
       TN = sum(TN)
       FN = sum(FN)
       FPR = FP/(FP+TN)
       FNR = FN/(FN+TP)
       print("test balanced error rate: " + ablations[i], (1/2)*(FPR+FNR))
       print("\n")
   print("\n")
```

C value: 1e-06 training balanced error rate: (ohe styles and ratings) 0.31654486428633577 validation balanced error rate: (ohe styles and ratings) 0.31945649492166617 test balanced error rate: (ohe styles and ratings) 0.315453673566917

training balanced error rate: (ratings and review length) 0.3402036260149308 validation balanced error rate: (ratings and review length) 0.34117176898721163 test balanced error rate: (ratings and review length) 0.33679075848465934

training balanced error rate: (ohe styles and review length) 0.26728720943897877 validation balanced error rate: (ohe styles and review length) 0.2675420967145031

test balanced error rate: (ohe styles and review length) 0.2668961129439944

C value: 1e-05

training balanced error rate: (ohe styles and ratings) 0.3153848550956065 validation balanced error rate: (ohe styles and ratings) 0.3184022780541963 test balanced error rate: (ohe styles and ratings) 0.3147117979556703

training balanced error rate: (ratings and review length) 0.33924707969313833 validation balanced error rate: (ratings and review length) 0.3409358423454912 test balanced error rate: (ratings and review length) 0.3362474223793813

training balanced error rate: (ohe styles and review length) 0.16274811347004606 validation balanced error rate: (ohe styles and review length) 0.15988861589703524

test balanced error rate: (ohe styles and review length) 0.16148495006798255

C value: 0.0001

training balanced error rate: (ohe styles and ratings) 0.2931954044604063 validation balanced error rate: (ohe styles and ratings) 0.2981570479871705 test balanced error rate: (ohe styles and ratings) 0.29076223747751395

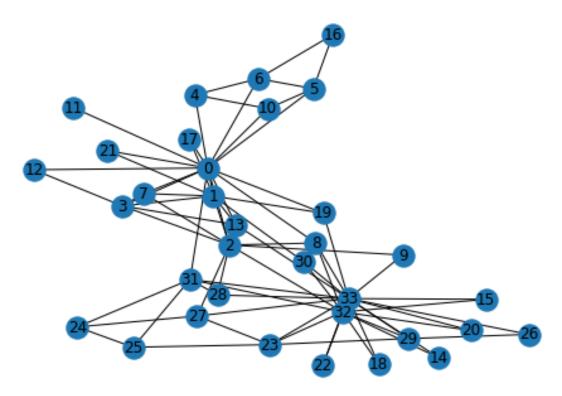
training balanced error rate: (ratings and review length) 0.33476255914717773 validation balanced error rate: (ratings and review length) 0.3388675007196019 test balanced error rate: (ratings and review length) 0.3314922932536461

training balanced error rate: (ohe styles and review length) 0.16274811347004606 validation balanced error rate: (ohe styles and review length) 0.15988861589703524

test balanced error rate: (ohe styles and review length) 0.16148495006798255

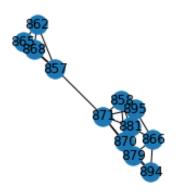
```
C value: 0.001
     training balanced error rate: (ohe styles and ratings) 0.19617014006804112
     validation balanced error rate: (ohe styles and ratings) 0.19406379785353017
     test balanced error rate: (ohe styles and ratings) 0.18983718321073442
     training balanced error rate: (ratings and review length) 0.31866571074436545
     validation balanced error rate: (ratings and review length) 0.32467515111641104
     test balanced error rate: (ratings and review length) 0.3212216009337423
     training balanced error rate: (ohe styles and review length) 0.16274811347004606
     validation balanced error rate: (ohe styles and review length)
     0.15988861589703524
     test balanced error rate: (ohe styles and review length) 0.16148495006798255
 []:
 []:
 []:
 []:
         Tasks (Community Detection)
[23]: import networkx as nx
      import matplotlib.pyplot as plt
      from urllib.request import urlopen
[24]: # Karate club
      G = nx.karate_club_graph()
      nx.draw(G, with_labels = True)
      plt.show()
      plt.clf()
      edges = set()
```

nodes = set()



<Figure size 432x288 with 0 Axes>





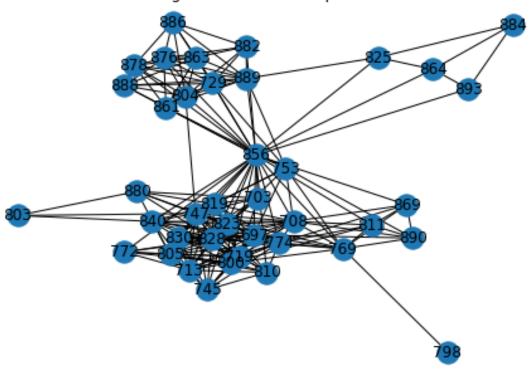


- 1.1 6) How many connected components are in the graph, and how many nodes are in the largest connected component (1 mark)? Next we'll implement a 'greedy' version of normalized cuts, using just the largest connected component found above. First, split it into two equal halves, just by taking the 50% of nodes with the lowest and 50% with the highest IDs.
- 1.1.1 There are 3 connected components and 40 nodes in the largest connected component

```
[ ]:
[25]: num_cc = len(list(nx.connected_components(G)))
    print(num_cc)
    print(len(max(list(nx.connected_components(G)))))

3
    40
[26]: max_cc = max(list(nx.connected_components(G)))
[27]: nx.draw(G.subgraph(max_cc), with_labels = True)
    plt.title("largest connected component")
    plt.show()
```





1.2 7) What is the normalized-cut cost of the 50/50 split you found above (1 mark)? Now we'll implement our greedy algorithm as follows: during each step, we'll move one node from one cluster to the other, choosing whichever move minimizes the resulting normalized cut cost (in case of a tie, pick the node with the lower ID). Repeat this until the cost can't be reduced any further.

```
[29]: #number of edges between low cluster and high cluster
def edges_across_cut(graph, set1, set2):
    return len(list(nx.edge_boundary(graph,set1,set2)))
edges_across_cut(G.subgraph(max_cc), lows, highs)
```

[29]: 92

```
[30]: DegreeView({856: 29, 893: 3, 753: 11, 889: 12, 719: 16, 747: 21, 708: 17, 697:
      17, 800: 16, 830: 14, 745: 11, 805: 16, 774: 17, 804: 11, 729: 11, 890: 6, 769:
      12, 823: 18, 828: 19, 882: 7, 886: 7, 861: 9, 863: 10, 810: 10, 803: 3, 840: 13,
      713: 15, 819: 15, 888: 8, 878: 9, 811: 7, 876: 10, 869: 6, 772: 7, 703: 8, 864:
      4, 880: 7, 884: 3, 798: 1, 825: 4})
[31]: nodes_max_cc = max(list(nx.connected_components(G)))
      lows, highs = sorted(nodes_max_cc)[:len(nodes_max_cc)//
      →2],sorted(nodes_max_cc)[len(nodes_max_cc)//2:]
      lows = set(lows)
     highs = set(highs)
      def calc norm cut cost(lows, highs):
          #calcs number of edges across cut
          num_edges = edges_across_cut(G.subgraph(max_cc), lows, highs)
          #calcs sum of degrees in each cluster
          total_low = sum([G.subgraph(max_cc).degree[v] for v in lows])
          total_high = sum([G.subgraph(max_cc).degree[v] for v in highs])
          cost = (1/2)*((num_edges/total_high) + (num_edges/total_low))
          return cost
      print("normalized 50/50 split cost: ")
      calc_norm_cut_cost(lows, highs)
     normalized 50/50 split cost:
[31]: 0.42240587695133147
[32]: #from piazza:
      #Q8 understanding
      # So if we have nodes [1,2,3,4,5,6] and after question 7 we have our splits:
      \rightarrow [1,2,3] and [4,5,6]
      # Would the algorithm be finding splits:
      #[1,2]/[3,4,5,6], [1,3]/[2,4,5,6], [2,3]/[1,4,5,6], [1,2,3,4]/[1,4,5,6],
      \rightarrow [5,6], [1,2,3,5]/[4,6], [1,2,3,4,6]/[4,5]
      # And then finding the minimum normalized cut cost?
      # And if [1,2]/[3,4,5,6] was the optimal one, would we proceed like:
      # [1], [2,3,4,5,6], [2]/[1,3,4,5,6], [1,2,3]/[4,5,6] [1,2,4]/[3,5,6], u
       \rightarrow [1,2,5]/[3,4,6], [1,2,6]/[3,4,5] ?
```

[30]: G.subgraph(max\_cc).degree()

```
[33]: def greedy(graph, set1, set2):
          min_cost = np.inf
          results1 = {}
            results2 = {}
            cluster1_2 = set1.copy()
            cluster2_2 = set2.copy()
          for node in set1:
              #Create copies to check for every possible combination
              cluster1 = set1.copy()
              cluster2 = set2.copy()
              #move one node from one to the other
              cluster1.remove(node)
              cluster2.add(node)
              #calc normalized cut cost
              cost = calc_norm_cut_cost(cluster1, cluster2)
              #add node that minimizes cost to outcome dictionary
              if cost < min_cost:</pre>
                  min_cost = cost
                  min_node = node
                   results1[min_cost] = min_node
              elif cost == min_cost:
                   if node < min node:</pre>
                       min_cost = cost
                       min node = node
                       results1[min_cost] = min_node
            for node in set2:
      #
                 #Create copies
      #
                 cluster1 = set1.copy()
      #
                 cluster2 = set2.copy()
                 #move one node from one to the other
      #
                 cluster2.remove(node)
                 cluster1.add(node)
      #
                 #calc normalized cut cost
                 cost = calc_norm_cut_cost(cluster1, cluster2)
      #
      #
                 if cost < min cost:</pre>
      #
                     min\_cost = cost
                     min_node = node
      #
                     results2[min cost] = min node
      #
                 elif cost == min_cost:
                     if node < min_node:</pre>
      #
                         min\_cost = cost
      #
                         min_node = node
      #
                         results2[min\_cost] = min\_node
```

```
out = {v:k for k, v in sorted(results1.items(), key=lambda item: item[0])}
          return list(out.items())[0]
[34]: greedy(G.subgraph(max_cc),lows,highs)
[34]: (729, 0.3873319662793347)
[35]: nodes_max_cc = max(list(nx.connected_components(G)))
      lows, highs = sorted(nodes max cc)[:len(nodes max cc)//
       →2],sorted(nodes_max_cc)[len(nodes_max_cc)//2:]
      lows = set(lows)
      highs = set(highs)
      while True:
          prev_cost = calc_norm_cut_cost(lows, highs)
          #move from set1 to set2
          node1, node1_cost = greedy(G.subgraph(max_cc),lows,highs)
          #move from set2 to set1
          node2, node2_cost = greedy(G.subgraph(max_cc),highs,lows)
          #if node cost from cluster 1 to cluster 2 is less than cost from cluster 2
       \rightarrow to cluster 1
          #then move node, vice versa
          if node1_cost < node2_cost and node1_cost < prev_cost:</pre>
              min_node = node1
              min cost = node1 cost
              lows.remove(min_node)
              highs.add(min node)
              print("test1: this is working")
          elif node2_cost < node1_cost and node2_cost < prev_cost:</pre>
              min_node = node2
              min_cost = node2_cost
              highs.remove(min_node)
              lows.add(min_node)
              print("test2: this is working")
          #if tie, pick node with lower id
          elif node1_cost == node2_cost:
              if node1 < node2:</pre>
                  lows.remove(node1)
                  highs.add(node1)
                  min_cost = node1_cost
```

```
print("test3: this is working")
             else:
                 highs.remove(node2)
                 lows.add(node2)
                 min_cost = node2_cost
                 print("test4: this is working")
         else: break
         print("test: reachable statement:")
     print("elements of resulting split: ")
     print("split1: ", lows)
     print("split12: ", highs)
     print("normalized cut cost: ", calc_norm_cut_cost(lows, highs))
    test1: this is working
    test: reachable statement:
    test1: this is working
    test: reachable statement:
    test2: this is working
    test: reachable statement:
    elements of resulting split:
    split1: {769, 772, 774, 798, 800, 803, 805, 810, 811, 819, 823, 697, 828, 830,
    703, 708, 840, 713, 719, 856, 869, 745, 747, 880, 753, 890}
    split12: {804, 825, 729, 861, 863, 864, 876, 878, 882, 884, 886, 888, 889, 893}
    normalized cut cost: 0.09817045961624274
[]:
[]:
[]:
```