cogsci 131 - assignment5

March 5, 2020

```
[3]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import random
```

1 CogSci 131 - Assignment 5

Selena Zhang

1.1 Question 1

The first way I could convert these to distances would be to subtract the similarity from 1 and do that for every cell in the row. The second way to convert these into distances would be to take the inverse of the similarity, i.e. 1/similarity, to find the distance. The third way to convert would be to take the logarithm of 1-similarity.

There is an inverse relationship between similarity and the distance: the more similar something is (closer to 1), the less distance there is between then. All of these methods preserve this negative relationship, but taking the inverse and having an arbitrary reference point may not be the most accurate nor optimal way to do so. The inverse may create some hyperbolic and polynomial curves that won't be best for regression fitting, and the logarithm may lead to distorted graphs and scaling issues.

For this assignment, I will use "1 minus similarity" definition, which is the most efficient and consistent way to find the distances because they all retain the same scale. Furthermore, a linear transformation like this will prevent any strange polynomial errors or approximations.

```
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```

1.2 Question 2

```
[14]: #euclidean distance function
      def euclid_distance(x, y):
          \# x = [i, j], y=[u, v]
          return ((x[1]-x[0])**2 + (y[1] - y[0])**2)**0.5
      #computing stress
      #equation: sum of (psi - dist(pi, pj))^2
      def stress(positions):
          summation = \Pi
          #iterating over each row and column
          for i in np.arange(len(positions)):
              #j is 2d
              for j in np.arange(1):
                  #ignoring diagonals
                  if i != j:
                      summation.append((distances[i][j] -__
       →euclid_distance(positions[i], positions[j]))**2)
                  else:
                      pass
          return np.cumsum(summation)[-1]
      #values: 2 dimensions, 21 sports
      testpoints = np.random.uniform(0,1,size=(21,2))
      stress(testpoints)
```

[14]: 2.813157081812429

1.3 Question 3

```
[15]: ##'helper' functions to make the final computation easier
#function to add deltas
def delta_helper(positions, i, j, delt):
#turn matrix into an array
```

```
array = np.array(positions)
  #add the change in delta
array[i, j] = array[i,j] + delt
return array

#gradient for one individual point at position i and j
def gradient(positions, i, j, delta):
  plus_delta = stress(delta_helper(positions, i, j, delta))
  #multiplying delta by -1 will subtract delta off
  minus_delta = stress(delta_helper(positions, i, j, -1*delta))
  gradient_result = (plus_delta - minus_delta)/(2*delta)
  return gradient_result
#test
gradient(testpoints, 5, 0, 0.01)
```

[15]: 0.02488808451643898

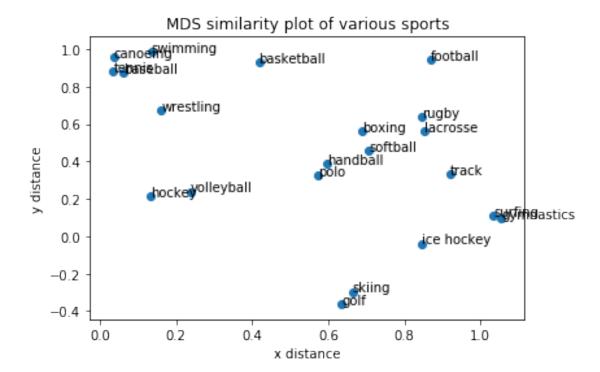
```
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[ 0.17492891, -0.17492891],
[-0.07495241, 0.07495241],
[ 0.60552865, -0.60552865]])
```

1.4 Question 4

```
[18]: #mds function
     def min_mds():
          #generate actual points and distance matrices inside function
         points = np.random.uniform(0,1,size=(21,2))
         distances = [1-x for x in similarities]
         #labels for sports
         sport = ["football", "baseball", "basketball", "tennis", "softball", "

→ "canoeing",

               "handball", "rugby", "hockey", "ice hockey", "swimming", "track", [
      "volleyball", "lacrosse", "skiing", "golf", "polo", "surfing", "
      #following gradient downhill to minimize stress as per lecture
         #scaling it down
         for i in np.arange(100):
             for point in points:
                 points -= gradient_all(points)*0.01
         #appending values
         x_vals=[]
         y_vals=[]
         for position in points:
             x_vals.append(position[0])
             y_vals.append(position[1])
         #plot format
         plt.figure()
         plt.title('MDS similarity plot of various sports')
         plt.xlabel('x distance')
         plt.ylabel('y distance')
         plt.scatter(x_vals, y_vals)
         #adding labels
         for i, sport in enumerate(sport):
             plt.annotate(sport, (x_vals[i], y_vals[i]))
     min_mds()
```



These results somewhat are in alignment with my intution: sports that are more similar to each other tend to be near the same area, while radically different sports are on opposite ends of the graph. While I may not personally agree that some of these sports are more similar than others and where they are placed in relation to each other, it gives a very rough estimate of where things are placed in the MDS space.

1.5 Question 5

```
#following gradient downhill to minimize stress as per lecture
    #scaling it down
   for i in np.arange(100):
        for point in points:
           points -= gradient_all(points)*0.01
##calculating pairwise distance
   x_vals=[]
   y_vals=[]
   for position in points:
       x_vals.append(position[0])
        y_vals.append(position[1])
   pairwise_dist = []
   for i in np.arange(len(x_vals)):
        pairwise_dist.append(dist(x_vals[i], y_vals[i]))
   print(pairwise_dist)
   psych_dist = []
   for i in np.arange(len(points)):
        for j in np.arange(1):
            #ignoring diagonals
            if i != j:
                psych_dist.append((distances[i][j]))
#
      #plot format
   plt.figure()
   plt.title('Pairwise distances vs psychological distances')
   plt.xlabel('psychlogical distances')
   plt.ylabel('pairwise distances')
   plt.scatter(psych_dist, pairwise_dist)
   #adding labels
   for i, sport in enumerate(sport):
        plt.annotate(sport, (psych_dist[i], pairwise_dist[i]))
    #iterating over each row and column
```

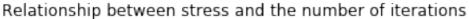
Good plots will show that the difference between the psychological error and the MDS will have a one to one linear relationship.

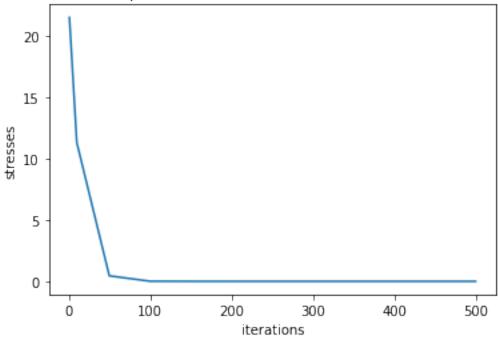
1.6 Question 6

```
[19]: def stress_vs_iterations():
    #blank stress, preset number of iterations
    iterations = [1, 10, 50, 100, 150, 200, 250, 300, 350, 400, 450, 500]
    stresses = []
    for i in np.arange(len(iterations)):
        #setting points and steps again, similar to question 4
        points = np.random.normal(0 ,1 ,size=(21,2))
        for steps in np.arange(iterations[i]):
            points -= gradient_all(points)*0.01
        stresses.append(stress(points))
```

```
print(stresses)
plt.plot(iterations, stresses)
plt.xlabel('iterations')
plt.ylabel('stresses')
plt.title('Relationship between stress and the number of iterations')
plt.figure()
stress_vs_iterations()
```

```
[21.493172058270517, 11.323302725254536, 0.44725609331477933, 0.008356861283938288, 0.00011574382872268488, 2.202869698137527e-06, 3.3712649032372206e-08, 7.982383494005966e-10, 1.4367044351008043e-11, 1.7349680544145147e-13, 2.7298272527611056e-15, 2.6035901405332846e-17]
```



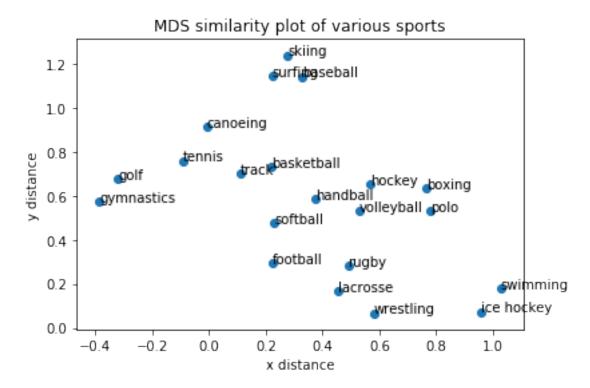


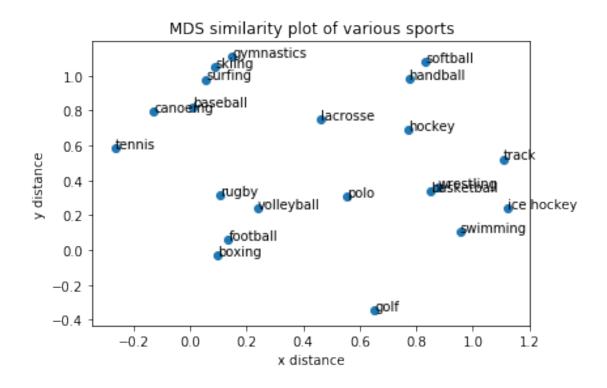
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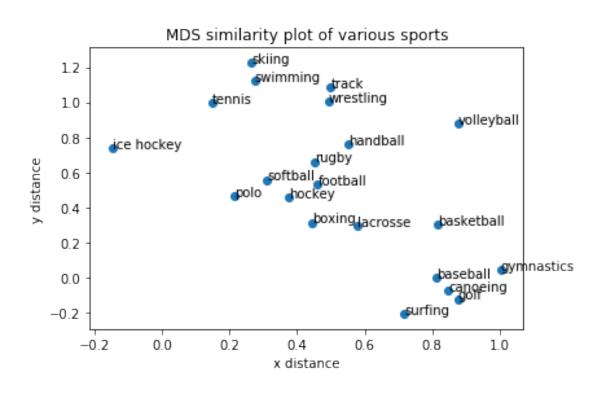
This graph shows what the stress is at various number of iterations. We find the minimum stress on this graph and match it to the number of iterations. We see that between 50 and 100 iterations, the stress drops dramatically from 40+ to 1. This means that the minimum number of iterations we need to perform is around 50 if we would like to minimize stresses while still running an efficient code

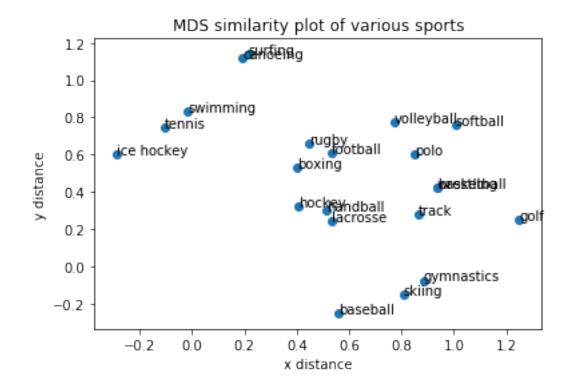
1.7 Question 7

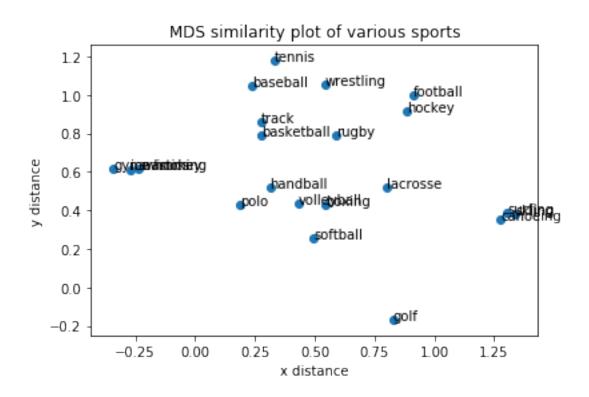
```
[22]: for i in range(9): min_mds()
```

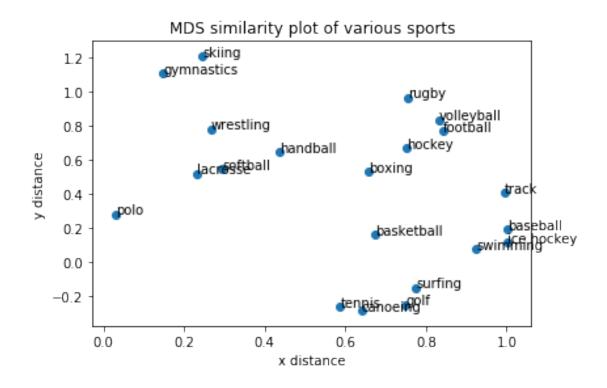


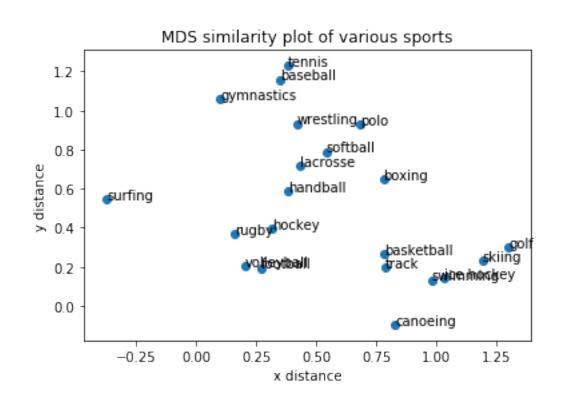


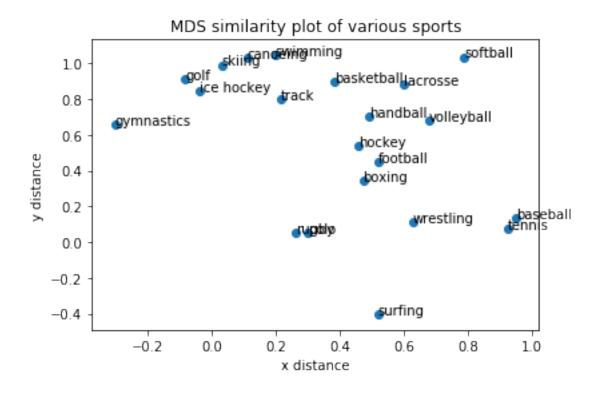


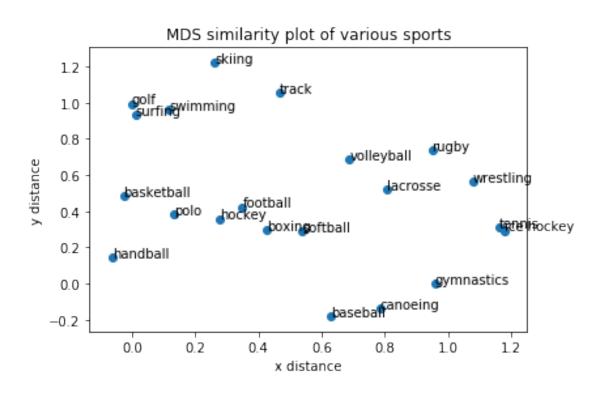




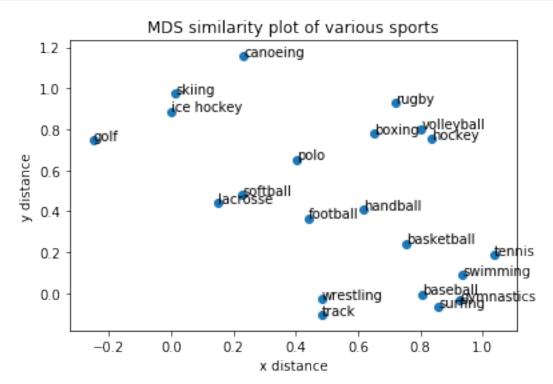








[24]: ##10th graph min_mds()



None of these graphs are the same because the points variable is random and is set within each function, meaning that it will be called when the function is executed, and it will present a random set of starting points each time. As a result, none of these graphs are the same, because the function to minimize stress relies on these random points in the equation.

1.8 Question 8

If I were to choose the best one to use, I would first run different versions of the function and code them with more iterations and a smaller step size to know that I was following the gradient more closely. I could also juse the min() function to find the one that minimizes all stress within the MDS.