EE412 Foundation of Big Data Analytics, Fall 2022 HW2

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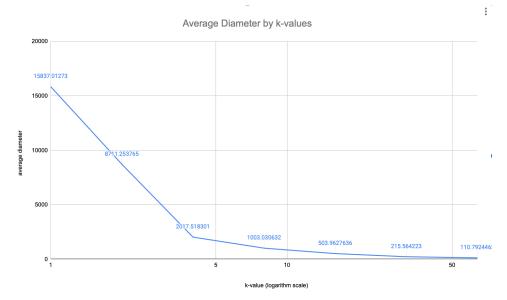
Discussion Group (People with whom you discussed ideas used in your answers):

On-line or hardcopy documents used as part of your answers:

Answer to Problem 1

(a) [5 pts] Solve the following problem, which is based on the exercises in the Mining of Massive Datasets 2nd edition (MMDS) textbook.

- (b) [20 pts] Implement the k-Means algorithm using Spark
 - The number-of-clusters vs. average diameter plot.



• The k value with an explanation why it is good for this data.

Please choose the best k value based on your thought. You will get full points if the reasoning is logical.

Changing the x-axis to logarithmic scale results in a point where the graph is clearly bent. The point is where $\mathbf{k} = \mathbf{4}$. And when k is larger than 4, as k is doubled, the average diameter is halved, which means total diameter is not changing that much.

Answer to Problem 2

Exercise 11.1.7

```
import numpy as np
THRESHOLD = 1e-5
A = np.array([
   [1, 1, 1],
   [1, 2, 3],
   [1, 3, 6],
1)
eig_vec1 = np.array([1, 1, 1]).reshape((3, 1))
diff = 1
while diff > THRESHOLD:
   new_eig_vec = A @ eig_vec1
  new_eig_vec = new_eig_vec / np.linalg.norm(new_eig_vec)
   diff = np.linalg.norm(eig_vec1 - new_eig_vec)
   eig_vec1 = new_eig_vec
print("(a) first eigen vector")
print(eig_vec1, end="\n\n")
eig_val1 = (eig_vec1.T @ A @ eig_vec1)[0,0]
print("(b) first eigen value")
print(eig_val1, end='\n\n')
A2 = A - eig_val1 * (eig_vec1 @ eig_vec1.T)
print("(c) new matrix")
print(A2, end='\n\n')
def get_eig_pair(M):
   eig_vec = np.array([1, 1, 1]).reshape((3, 1))
   diff = 1
   while diff > THRESHOLD:
       new_eig_vec = M @ eig_vec
       new_eig_vec = new_eig_vec / np.linalg.norm(new_eig_vec)
       diff = np.linalg.norm(eig_vec - new_eig_vec)
       eig_vec = new_eig_vec
   eig_val = eig_vec.T @ M @ eig_vec
   new_M = M - eig_val * (eig_vec @ eig_vec.T)
   return eig_vec, eig_val[∅,0], new_M
```

```
eig_vec2, eig_val2, A3 = get_eig_pair(A2)

print("(d) second eigen pair")
print(eig_vec2)
print(eig_val2, end='\n\n')

eig_vec3, eig_val3, A4 = get_eig_pair(A3)

print("(e) third eigen pair")
print(eig_vec3)
print(eig_val3, end='\n\n')

V = np.concatenate([eig_vec1, eig_vec2, eig_vec3], axis=1)
U = np.diag([eig_val1, eig_val2, eig_val3])

print("Verification: reconstructed matrix")
print(V @ U @ V.T)
```

(a) first eigen vector

[[0.19382289]

[0.4722474]

[0.85989248]]

(b) first eigen value

7.872983346206856

(c) New matrix

[[0.70423318 0.27936729 -0.31216529]

[0.27936729 0.24418608 -0.19707675]

[-0.31216529 -0.19707675 0.1785974]]

(d) second eigen pair

[[0.81649634]

[0.40824695]

[-0.40825011]]

1.0000000000042972

(e) third eigen pair

[[0.54384155]

[-0.78122828]

[0.30646167]]

0.1270166537934505

Exercise 11.3.1

```
import numpy as np
RANK = 2
M = np.array([
  [1,2,3],
  [3,4,5],
  [5,4,3],
  [0,2,4],
  [1,3,5],
1)
S1 = M @ M.T
S2 = M.T @ M
print("(a) M@M.T and M.T@M")
print("- M@M.T", S1, sep='\n')
print("- M.T@M", S2, sep='\n', end='\n\n')
w1, v1 = np.linalg.eig(S1)
w2, v2 = np.linalg.eig(S2)
print("(b) eigenpairs for matrices of part (a)")
print("- w1", w1, "- v1", v1, "- w2", w2, "- v2", v2, sep='\n', end='\n\n')
U = v1[:, np.argsort(w1)][:, -RANK:]
V = v2[:, np.argsort(w2)][:, -RANK:]
S = np.sqrt(np.diag(sorted(w1)[-RANK:]))
print("(c) SVD for the matrix M")
print("- U", U, "- V", V, "- S", S, sep="\n", end='\n\n')
sv2 = S[0,0]
S[0,0] = 0
approximate = U @ S @ V.T
print("(d) approximation to the matrix M")
print(approximate, end='\n\n')
sv1 = S[-1, -1]
retained energy = sv1**2 / (sv1**2 + sv2**2)
```

```
print("(e) retained energy")
print(retained_energy)
  (a) M@M.T and M.T@M
      - M@M.T
      [[14 26 22 16 22]
      [26 50 46 28 40]
      [22 46 50 20 32]
      [16 28 20 20 26]
      [22 40 32 26 35]]
      - M.T@M
      [[36 37 38]
      [37 49 61]
      [38 61 84]]
  (b) eigenpairs for matrices of part (a)
      -w1
      [ 1.53566996e+02 -2.16919039e-15 1.54330035e+01 -2.69964138e-15
      -1.63115212e-16]
      - v1
      [[ 0.29769568  0.94131607 -0.15906393 -0.57735012 -0.21094872]
      [ 0.57050856 -0.17481584 0.0332003 -0.22666834 0.06716429]
      [ 0.52074297 -0.04034212  0.73585663  0.10591706 -0.13512315]
      [ 0.32257847 -0.18826321 -0.5103921 -0.27280206 -0.68074095]
      [ 0.45898491 -0.21515796 -0.41425998  0.72776982  0.68507159]]
      [1.53566996e+02 1.54330035e+01 2.99519331e-15]
      - v2
      [[-0.40928285 -0.81597848 0.40824829]
      [-0.56345932 -0.12588456 -0.81649658]
      [-0.7176358 0.56420935 0.40824829]]
  (c) SVD for the matrix M
      - U
      [[-0.15906393 0.29769568]
      [ 0.0332003  0.57050856]
      [ 0.73585663  0.52074297]
      [-0.5103921 0.32257847]
      [-0.41425998 0.45898491]]
```

[[-0.81597848 -0.40928285]

```
[-0.12588456 -0.56345932]
[ 0.56420935 -0.7176358 ]]
- S
[[ 3.92848616 0. ]
[ 0. 12.39221516]]
```

(d) approximation to the matrix M

```
[[-1.509889 -2.0786628 -2.64743661]

[-2.89357443 -3.98358126 -5.0735881 ]

[-2.64116728 -3.63609257 -4.63101787]

[-1.63609257 -2.25240715 -2.86872172]

[-2.32793529 -3.20486638 -4.08179747]]
```

(e) retained energy

0.9086804524257934

• The reason approximation of M is negative

Since we collect vectors U and V through eigen pairs from M_T M and MM_T, it is impossible to distinguish the sign of matrix M. Which means, we will get the same SVD decomposition result from matrix M and -M.

Answer to Problem 3

(a) [15 pts] Solve the following problems, which are based on the exercises in the MMDS textbook.

Exercise 9.3.1

(a) Jaccard distance

$$J(A, B) = \frac{1}{2} = 0.5$$

$$J(A, C) = \frac{1}{2} = 0.5$$

$$J(B, C) = \frac{1}{2} = 0.5$$

(b) Cosine distance

$$C(A, B) = \frac{5 \times 3 + 5 \times 3 + 1 \times 1 + 3 \times 1}{\sqrt{4^2 + 5^2 + 5^2 + 1^2 + 3^2 + 2^2} \times \sqrt{3^2 + 4^2 + 3^2 + 1^2 + 2^2 + 1^2}} = 0.60$$

$$C(A, C) = \frac{4 \times 2 + 5 \times 3 + 3 \times 5 + 2 \times 3}{\sqrt{A^2 + 5^2 + 5^2 + 5^2 + 5^2 + 2$$

$$C(B, C) = \frac{4 \times 1 + 3 \times 3 + 2 \times 4 + 1 \times 5}{\sqrt{3^2 + 4^2 + 3^2 + 1^2 + 2^2 + 1^2} \times \sqrt{2^2 + 1^2 + 3^2 + 4^2 + 5^2 + 3^2}} = 0.514$$

(c) Rounded Jaccard distance

$$J(A, B) = \% = 0.6$$

$$J(A, C) = 4/6 = 0.66$$

$$J(B, C) = \% = 0.833$$

(d) Rounded Cosine distance

$$C(A, B) = \frac{2}{\sqrt{4} \times \sqrt{3}} = 0.577$$

$$C(A, C) = \frac{2}{\sqrt{4} \times \sqrt{4}} = 0.5$$

$$C(B, C) = \frac{1}{\sqrt{3} \times \sqrt{4}} = 0.289$$

(e) Normalized Matrix

(f) Normalized Cosine distance

$$C(A, B) = \frac{1.67 \times 0.67 + 1.67 \times 0.67 + 2.33 \times 1.33 + 0.33 \times 1.33}{\sqrt{0.67^2 + 1.67^2 + 2.33^2 + 0.33^2 + 1.33^2} \times \sqrt{0.67^2 + 1.67^2 + 0.67^2 + 1.33^2 + 0.33^2 + 1.33^2}} = 0.584$$

$$C(A, C) = \frac{\frac{-0.67 \times 1 - 0.33 \times 2}{\sqrt{0.67^2 + 1.67^2 + 1.67^2 + 2.33^2 + 0.33^2 + 1.33^2} \times \sqrt{1^2 + 2^2 + 1^2 + 2^2}} = -0.115$$

$$C(B, C) = \frac{\frac{-1.67 \times 2 - 0.33 \times 1 - 1.33 \times 2}{\sqrt{0.67^2 + 1.67^2 + 0.67^2 + 1.33^2 + 0.33^2 + 1.33^2} \times \sqrt{1^2 + 2^2 + 1^2 + 2^2}} = -0.740$$

Exercise 9.3.2

(a) Clustering

i. (f, h) : J(f,h) = 0ii. $(b,d) : J(b,d) = \frac{1}{3}$ iii. $(b,d,g) : J(d,g) = \frac{1}{3}$ iv. $(b,d,g,a) : J(a,g) = \frac{1}{2}$

Final clusters: (f,h), (b,d,g,a), (c), (e)

(b) User-Cluster Matrix

	(f,h)	(b,d,g,a)	(c) (e)
A	1	4.25	
В	1	1.75	4
C	3.5	2.5	1

(c) Cosine distance

Cosine distance
$$C(A, B) = \frac{1 \times 1 + 4.25 \times 1.75}{\sqrt{1^2 + 4.25^2} \times \sqrt{1^2 + 1.75^2 + 4^2}} = 0.431$$

$$C(A, C) = \frac{1 \times 3.5 + 4.25 \times 2.5}{\sqrt{1^2 + 4.25^2} \times \sqrt{3.5^2 + 2.5^2 + 1^2}} = 0.733$$

$$C(B, C) = \frac{1 \times 3.5 + 1.75 \times 4.25 + 4 \times 1}{\sqrt{1^2 + 1.75^2 + 4^2} \times \sqrt{3.5^2 + 2.5^2 + 1^2}} = 0.6$$

(b) [20 pts] Implement collaborative filtering

(c) [20 pts] Movie Recommendation Challenge (incomplete)

Intialization

Train-Test Split

To check the performance of the algorithm and prevent over-fitting, I splitted nonzero elements into a train and test(validation) set.

```
nonzeros = utility_matrix.nonzero()
random_idx = np.random.permutation(num_data)
splitting_point = int(num_data * 0.8)
train_set = (
    nonzeros[0][random_idx[:splitting_point]],
    nonzeros[1][random_idx[:splitting_point]]
)
validation_set = (
    nonzeros[0][random_idx[splitting_point:]],
    nonzeros[1][random_idx[splitting_point:]]
)
```

Normalization

Normalize utility matrix with both user-mean and movie-mean matrix.

```
user_mean_matrix = np.apply_along_axis(
  lambda row: row[row > 0].mean() if sum(row) > 0 else 0.,
  axis=1,
  arr=utility_matrix,
)
movie_mean_matrix = np.apply_along_axis(
  lambda col: col[col > 0].mean() if sum(col) > 0 else 0.,
  axis=0,
  arr=utility_matrix,
)
mean_matrix = (user_mean_matrix[:, None] + movie_mean_matrix[None, :]) / 2
normalized_matrix = (utility_matrix - mean_matrix)
M = normalized_matrix * create_mask(train_set, utility_matrix.shape)
validation_matrix = normalized_matrix * create_mask(validation_set,
  utility_matrix.shape)
```

Perturbation

Each element of V and U is random normal distribution with mean 0 and standard deviation 1.

```
u, v = num_users+1, num_movies+1
U = np.random.normal(0, 1, (u, m))
V = np.random.normal(0, 1, (m, v))
```

Optimization

Permutation

In order to apply optimization randomly, I permute the indices of U and V

```
U_idx = np.random.permutation(np.array(U.nonzero()).T).T
V_idx = np.random.permutation(np.array(V.nonzero()).T).T
```

Optimization

Since the number of movies is much bigger than the number of users, optimization is conducted proportional to its size. (e.g. adjust U optimization 1 times, and then adjust V optimization 244 times, and repeat it)

```
for i in tqdm(range(u * m)):
    # Adjust U
    r, s = U_idx[:, i]
    x1 = (V[s, :] ** 2).sum()
    x2 = (V[s, :] * M[s, :]).sum()
    x3 = (U[r, :] * (V @ V[s, :])).sum()
    U[r, s] += (x2 - x3) / x1
# Adjust V

for j in range(uv_rate):
    r, s = V_idx[:, i * uv_rate + j]
    y1 = (U[:, r] ** 2).sum()
    y2 = (U[:, r] * M[:, s]).sum()
    y3 = (V[:, s] * (U.T @ U[:, r])).sum()
    V[r, s] += (y2 - y3) / y1
```