Assignment 4

Information Retrieval

CS 834

Fall 2017

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Problem 8.3

1.1 Problem Statement

For one query in the CACM collection (provided at the book website), generate a ranking using Galago, and then calculate average precision, NDCG at 5 and 10, precision at 10, and the reciprocal rank by hand.

1.2 Solution

This problem uses CACM corpus provided in the test collection of the book. The queries ran against the CACM collection have been used from the processed queries section of the CACM test collection.

1.2.1 Query:Interested in articles on robotics motion planning particularly the geometric and combinatorial aspects we are not interested in the dynamics of arm motion

The below XML file shows the query which was used in Galago for batch-search.

The below commands were run on command prompt for generating the ranked list of query with 10 results.

```
F:\Fall2017\InformationRetreival\Assignment3\Galgo\galagosearch-1.04-bin\galagosearch-1.04\
bin>galago.bat batch-search --index=F:\Fall2017\InformationRetreival\Assignment4\index.
idx --count=10 F:\Fall2017\InformationRetreival\Assignment4\onequery.xml > F:\Fall2017\
InformationRetreival\Assignment4\cacm_onequery.list
```

The below list file was generated by batch search from Galago.

```
6 Q0 CACM-0695 1 -164.66490173 galago
6 Q0 CACM-2826 2 -166.82664490 galago
6 Q0 CACM-2828 3 -167.13014221 galago
```

```
6 Q0 CACM-1664 4 -167.29315186 galago
```

The below commands were run on Galago to evaluate the ranked list for query generated from the previous step.

```
F:\Fall2017\InformationRetreival\Assignment3\Galgo\galagosearch-1.04-bin\galagosearch-1.04\bin> galago.bat eval F:\Fall2017\InformationRetreival\Assignment4\cacm_onequery.list F:\Fall2017\InformationRetreival\Assignment4\cacm.rel >F:\Fall2017\InformationRetreiva\Assignment4\onequeryoutput.txt
```

The below text file is the result of eval command by using the relevance judgments file provided in cacm test collections.

num_ret	6	10
num_rel	6	3
num_rel_ret	6	3
map	6	0.4111
ndcg	6	0.5833
ndcg15	6	0.5833
R-prec	6	0.3333

bpref	6 0.0000
recip_rank	6 0.3333
P5	6 0.4000
P10	6 0.3000
P15	6 0.2000
P20	6 0.1500
P30	6 0.1000
P100	6 0.0300
P200	6 0.0150
P500	6 0.0060
P1000	6 0.0030
num_ret	all 10
num_rel	all 3
num_rel_ret	all 3
map	all 0.4111
ndcg	all 0.5833
ndcg15	all 0.5833
R-prec	all 0.3333
bpref	all 0.0000
recip_rank	all 0.3333
P5	all 0.4000

P10	all 0.3000
P15	all 0.2000
P20	all 0.1500
P30	all 0.1000
P100	all 0.0300
P200	all 0.0150
P500	all 0.0060
P1000	all 0.0030

The calculation of precision for the ranking was done with the help of relevance judgments provided with the test collections. It contains list of all the documents relevant to a collection. Below is the list of relevant documents for query 6.

- 6 QO CACM-1543 1
- 6 Q0 CACM-2078 1
- 6 Q0 CACM-2828 1

Precision at 10

$$Precision(1) = 0/1 = 0$$

$$Precision(2) = 0/2 = 0$$

$$Precision(3) = 1/3 = 0.33$$

$$Precision(4) = 1/4 = 0.25$$

$$Precision(5) = 2/5 = 0.4$$

$$Precision(6) = 3/6 = 0.5$$

$$Precision(7) = 3/7 = 0.429$$

$$Precision(8) = 3/8 = 0.375$$

$$Precision(9) = 3/9 = 0.33$$

$$Precision(10) = 3/10 = 0.3$$

where Precision(6) denotes precision at search result 6.

The Precision(10) for this query is 0.3.

Mean Average Precision

$$MAP = 1/3(1/3 + 2/5 + 3/6)$$

$$MAP = 0.41$$

Reciprocal Rank

The first relevant document is at position 3.

$$RR = 1/3$$

$$RR = 0.333$$

Normalized Discounted Cumulative Gain at 5

Relevance Score: 0, 0, 1, 0, 1

$$CG_5 = 0 + 0 + 1 + 0 + 1$$

Table 1.1: Discounted Cumulative Gain

i
$$rel_i$$
 $log_2(i+1)$ $rel_i/log_2(i+1)$

1 0 1 0

2 0 1.585 0

 $3 \quad 1 \quad 2 \quad 0.5$

 $4 \quad 0 \qquad 2.322 \qquad 0$

 $5 \quad 1 \quad 2.585 \quad 0.387$

$$CG_5 = 2$$

$$DCG = \sum_{i=1}^{5} rel_i/log_2(i+1)$$

$$DCG = 0 + 0 + 0.5 + 0 + 0.387 = 0.887$$

Ideal Discounted Cumulative Gain has the document and relevance score as below:

Relevance Score: 1, 1, 0, 0, 0

$$CG_5 = 1 + 1 + 0 + 0 + 0$$

$$CG_5 = 2$$

$$IDCG = \sum_{i=1}^{5} rel_i / log_2(i+1)$$

$$IDCG = 1 + 0.631 + 0 + 0 + 0 = 1.631$$

Normalized Discounted Cumulative Gain (NDCG)= DCG / IDCG

$$NDCG = 0.887/1.631 = 0.544$$

Table 1.2: Ideal Discounted Cumulative Gain

i
$$rel_i log_2(i+1) rel_i/log_2(i+1)$$

1 1 1 1

2 1 1.585 0.631

 $3 \ 0 \ 2 \ 0$

4 0 2.322 0

 $5 \quad 0 \quad 2.585 \quad 0$

Normalized Discounted Cumulative Gain at 10

Relevance Score: 0, 0, 1, 0, 1, 1, 0, 0, 0, 0

$$CG_{10} = 0 + 0 + 1 + 0 + 1 + 1 + 0 + 0 + 0 + 0$$

$$CG_{10} = 3$$

$$DCG = \sum_{i=1}^{10} rel_i / log_2(i+1)$$

$$DCG = 0 + 0 + 0.5 + 0 + 0.387 + 0.356 + 0 + 0 + 0 + 0 = 1.243$$

Ideal Discounted Cumulative Gain has the document and relevance score as below:

Relevance Score: 1, 1, 1, 0, 0, 0, 0, 0, 0, 0

$$CG_{10}=3$$

$$IDCG = \sum_{i=1}^{10} rel_i/log_2(i+1)$$

$$IDCG = 1 + 0.61 + 1 + 0.5 + 0 + 0 + 0 + 0 + 0 + 0 = 2.11$$

Normalized Discounted Cumulative Gain (NDCG)= DCG / IDCG

$$NDCG = 1.243/2.11 = 0.589$$

Table 1.3: Discounted Cumulative Gain

- i $rel_i log_2(i+1) rel_i/log_2(i+1)$
- 1 0 1 0
- $2 \quad 0 \quad 1.585 \quad 0$
- $3 \quad 1 \quad 2 \quad 0.5$
- 4 0 2.322 0
- 5 1 2.585 0.387
- 6 1 2.807 0.356
- 7 0 3 1
- $8 \quad 0 \quad 3.17 \quad 0$
- 9 0 3.22 0
- 10 0 3.459 0

Table 1.4: Ideal Discounted Cumulative Gain

_	_ / /	$rel_i/log_2(i+1)$
 1 1 1 :	IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	1 21. / 1.000 11. + 1.1

- 1 1 1 1
- $2 \quad 1 \quad 1.585 \quad 0.61$
- $3 \quad 1 \quad 2 \quad 0.5$
- 4 0 2.322 0
- 5 0 2.585 0
- $6 \quad 0 \quad 2.807 \quad 0$
- 7 0 3 0
- 8 0 3.17 0
- $9 \quad 0 \quad 3.22 \quad 0$
- 10 0 3.459 0

Problem 8.4

2.1 Problem Statement

For two queries in the CACM collection, generate two uninterpolated recall precision graphs, a table of interpolated precision values at standard recall levels, and the average interpolated recall-precision graph.

2.2 Solution

This problem uses CACM corpus provided in the test collection of the book. The queries ran against the CACM collection have been used from the processed queries section of the CACM test collection.

Table 2.1: Precision and Recall Table for Query 6

Index 1 2 3 7 8 9 6 10 Relevant no yes yes yes no no no no no no Recall 0.670 0 0.330.331 1 1 1 1 0.429Precision 0 0 0.330.250.40.50.3750.330.3

2.2.1 Query: Interested in articles on robotics motion planning particularly the geometric and combinatorial aspects we are not interested in the dynamics of arm motion

The below list file generated from batch search of Galago has ranked result of 10 for the query.

- 6 Q0 CACM-0695 1 -164.66490173 galago
- 6 Q0 CACM-2826 2 -166.82664490 galago
- 6 Q0 CACM-2828 3 -167.13014221 galago
- 6 Q0 CACM-1664 4 -167.29315186 galago
- 6 Q0 CACM-1543 5 -167.67100525 galago
- 6 Q0 CACM-2078 6 -168.12095642 galago
- 6 Q0 CACM-2176 7 -168.47441101 galago
- 6 Q0 CACM-1113 8 -169.13491821 galago
- 6 Q0 CACM-0605 9 -169.25828552 galago
- 6 Q0 CACM-1517 10 -169.68901062 galago

Table 2.2: Precision and Recall Table for Query: Parallel algorithms

1 2 3 7 8 9 Index 4 5 6 10 Relevant no no yes no no yes no yes yes yes Recall 0 0 0.2 0.20.20.40.40.60.8 1 Precision 0 0 0.330.250.20.330.2860.3750.440.5

2.2.2 Query: Parallel algorithms

The below list file generated from batch search of Galago has ranked result of 10 for the query.

- 19 Q0 CACM-2433 1 -10.82248116 galago
- 19 Q0 CACM-1811 2 -10.83561993 galago
- 19 Q0 CACM-2973 3 -10.86207390 galago
- 19 Q0 CACM-2289 4 -10.93358135 galago
- 19 Q0 CACM-0950 5 -10.96933460 galago
- 19 Q0 CACM-2266 6 -11.00977135 galago
- 19 Q0 CACM-2785 7 -11.07456684 galago
- 19 Q0 CACM-3075 8 -11.07782173 galago
- 19 Q0 CACM-3156 9 -11.13305855 galago
- 19 Q0 CACM-2664 10 -11.15161610 galago

Both the queries show their highest precision of 0.5 at recall value of 1. This results in a interpolated recall-precision graph to be a straight horizontal line at precision 0.5. Interpolated recall-precision graph is drawn with curves that are descending in manner,

Recall vs Precision Graph

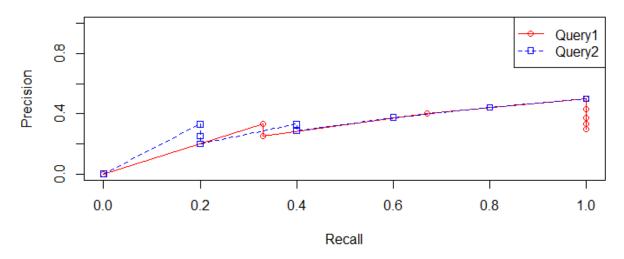


Figure 2.1: Recall vs Precision Graph

starting from highest point towards to lowest. But, for both the queries highest precision was reached at recall 1leading to a straight line.

Table 2.3: Precision values at standard recall levels calculated using interpolation

Recall	0.0	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9	1	
Ranking 1	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	
Ranking 2	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	
Average Ranking	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	

Interpolated Recall vs Precision Graph

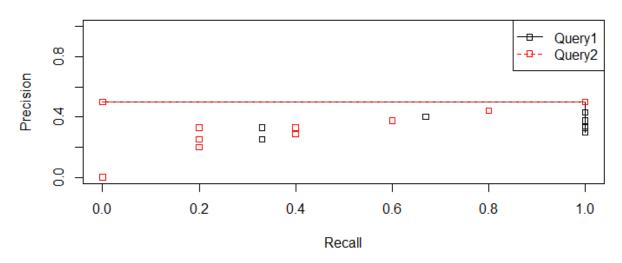


Figure 2.2: Interpolated Recall vs Precision Graph

Average Recall-Precision graph using standard recall levels

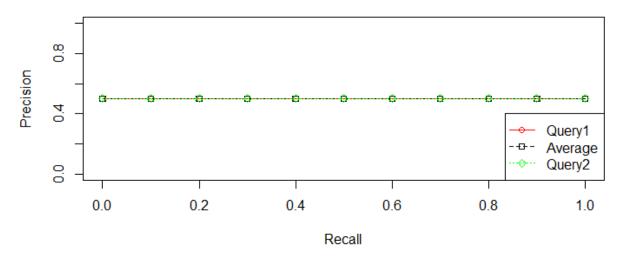


Figure 2.3: Average recall-precision graph using standard recall levels

Problem 8.5

3.1 Problem

Generate the mean average precision, recall-precision graph, average NDCG at 5 and 10, and precision at 10 for the entire CACM query set.

3.2 Solution

The results have been calculated by Galago batch-search for rank result of 10. Galago provides NDCG 5 and 15 but not NDCG10. So, by making a request for 10 results per query the NDCG15 is same as NDCG10 because all the calculations are based on 10 results for each query.

The values for MAP is lesser than Precision at 10 for the CACM query set. Th query sets having most of their relevant results at top indexes (between 1 and 5) have higher or

equal NCDG5 and NCDG10, but for queries with most of their relevant documents lower in the rank (between 6 and 10) have higher NCDG10 than NCDG5.

MAP for CACM Query Set

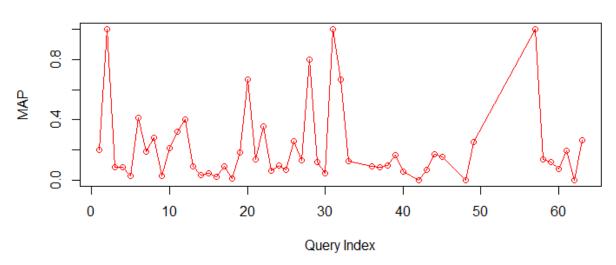


Figure 3.1: MAP values for CACM Query Set

NCDG5 for CACM Query Set

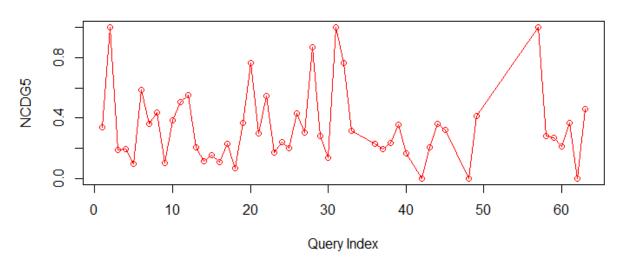


Figure 3.2: NCDG5 values for CACM Query Set

NCDG10 for CACM Query Set

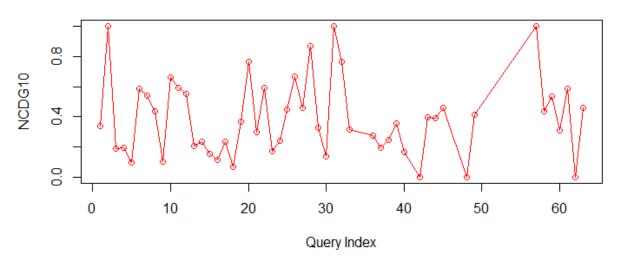


Figure 3.3: NCDG10 values for CACM Query Set

P10 for CACM Query Set

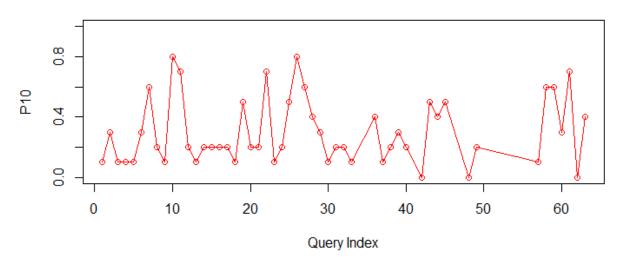


Figure 3.4: Precision at 10 values for CACM Query Set

Recall for CACM Query Set

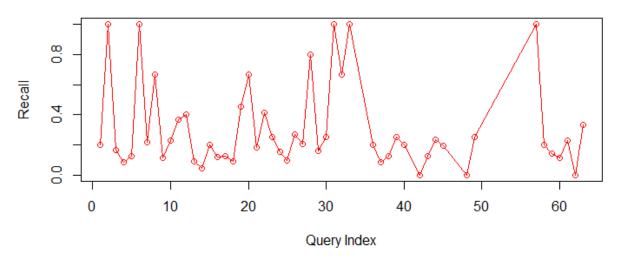


Figure 3.5: Recall values for CACM Query Set

Recall vs Precision for CACM Query Set

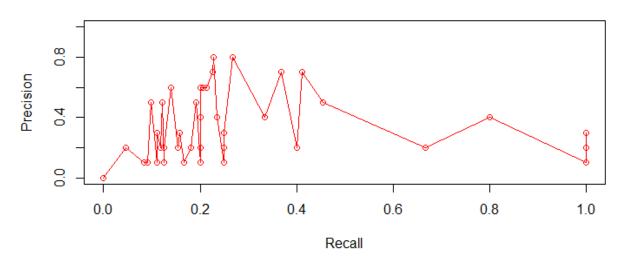


Figure 3.6: Recall vs Precision for CACM Query Set

Problem 8.7

4.1 Problem

Another measure that has been used in a number of evaluations is R-precision. This is defined as the precision at R documents, where R is the number of relevant documents for a query. It is used in situations where there is a large variation in the number of relevant documents per query. Calculate the average R-precision for the CACM query set and compare it to the other measures.

4.2 Solution

R-precision is defined as precision at R documents, where R is the number of relevant documents for a query. In context to the CACM query set, the relevance judgments file suggests that all the queries have varied number of relevant documents. It ranges from 0 to 51. So, for

Table 4.1: CACM Query Set measured against different measures

map	0.2840
ndcg	0.4419
ndcg15	0.4734
R-prec	0.3192
bpref	0.0000
recip_rank	0.7296
P5	0.3922
P10	0.2980
P15	0.2614
P20	0.2373
P30	0.1843

calculating R-precision number of relevant documents per query is set to 51 and evaluated in the Galago. The results for the different measures inclusive of R-precision at relevant document 51 are shown in the table:

Problem 8.9

5.1 Problem

For one query in the CACM collection, generate a ranking and calculate BPREF. Show that the two formulations of BPREF give the same value.

5.2 Solution

The solution is based on relevance results for query, interested in articles on robotics motion planning particularly the geometric and combinatorial aspects we are not interested in the dynamics of arm motion. The Table 5.1 shows the relevance results for the query. The relevance results show the search results had 3 relevant documents and 997 non- relevant documents. The ranked list for the query is generated from Galago without providing any explicit value for the number of ranked results. The output of ranked list for the query

contains 3 relevant and 997 non-relevant documents.

<parameters>

<query>

<number>3</number>

<text> intermediate languages used in construction of multi targeted compilers tcoll

</query>

</parameters>

Below is the list of relevant documents to the query.

- 3 Q0 CACM-1134 1
- 3 Q0 CACM-1613 1
- 3 Q0 CACM-1807 1
- 3 Q0 CACM-1947 1
- 3 Q0 CACM-2290 1
- 3 Q0 CACM-2923 1

5.2.1 Calculating BPREF

$$BPREF = 1/R \sum_{d_r} (1 - (N_{d_r}/R))$$

where R is the number of relevant documents that are considered

N is the number of non-relevant documents that are considered

 d_r is the number of relevant documents

Table 5.1: Relevance table showing R relevant and non-relevant documents

Index Relevance
No
No
No
Yes
No
Yes
Yes
Yes
Yes
Yes

For a query result with 3 relevant documents R is 3 implying that first 3 non-relevant documents are considered. The relevance table for the query is manipulated for BPREF as,

$$BPREF = 1/R \sum_{d_r} (1 - (N_{d_r}/R))$$

$$BPREF = 1/3[(1 - 2/3) + (1 - 3/3) + (1 - 3/3)]$$

$$BPREF = 1/3[(1/3) + 0 + 0] = 1/9 = 0.11$$

5.2.2 Calculating BPREF based on preference

$$BPREF = P/(P+Q)$$

where P is the number of relevant documents

Q is the number of non-relevant documents

For query: Code optimization for space efficiency

P = 3

Q = 3

BPREF = 3/(3+3)

BPREF = 0.5

The value of BPREF is 0.11 and 0.5 by computing respectively with relevant documents formula and the clickthrough preference formula. The values computed are not equal because the BPREF for clickthough formula does not take into account the ranking of relevant results and the value for BPREF with relevant documents formula dips due to the fist relevant result showing after two non-relevant result and the rest two relevant result appear after all the non-relevant documents taken in consideration. Thus, the last two relevant documents do not add any value to the BPREF. Using the clickthrough preference approach for calculating BPREF, with 3 relevant and 3 non-relevant documents the value computes to a 0.5.

5.2.3 Alternate Approach

The bpref approach works best when all the results sets are not judged. So rather than relying on relevance judgments provided with the CACM result, I have decided to manually find relevant and non-relevant documents from the search results generated by the galago search engine for the query to calculate the bpref value using both the formulas.

On mannual validation of the first 10 search results generated for the query (interested in articles on robotics motion planning particularly the geometric and combinatorial aspects we are not interested in the dynamics of arm motion) reveals 4 relevant and 6 non-relevant

Table 5.2: Manual Generated Relevance table showing R relevant and non-relevant documents

Index	Relevance
1	No
2	Yes
3	Yes
4	No
5	Yes
6	Yes
7	No
8	No
9	No
10	No

documents. So, for calculating BPREF, R becomes 4 because it has 4 relevant document.//

$$BPREF = 1/R \sum_{d_r} (1 - (N_{d_r}/R))$$

$$BPREF = 1/4[(1 - 1/4) + (1 - 1/4) + (1 - 2/4) + (1 - 2/4)]$$

$$BPREF = 1/4[(3/4) + (3/4) + (2/4) + (2/4)] = 0.625$$

$$BPREF = P/(P + Q)$$

$$P = 4$$

$$Q = 2$$

$$BPREF = 4/(4+2)$$

BPREF = 0.667

The manual validation of results generates values 0.625 and 0.667 for both the formulas of BPREF which are roughly equal. This result is way much better than previous result generated using relevant judgments.

Problem 9.8

6.1 Problem

Cluster the following set of two-dimensional instances into three clusters using each of the five agglomerative clustering methods: (4, 2), (3, 2), (2, 2), (1, 2), (1, 1), (1, 1), (2, 3), (3, 2), (3, 4), (4, 3) Discuss the differences in the clusters across methods. Which methods produce the same clusters? How do these clusters compare to how you would manually cluster the points?

6.2 Solution

The code for agglomerate clustering methods generates 3 clusters for the data set.

6.2.1 Discussion of the Methods

Single Linkage: It uses the minimum distance between between elements of two clusters to merge them as one cluster. For two clusters A and B, it finds the minimum euclidean distance between the points of each cluster and compares them against the minimum threshold distance of all the other clusters to merge them in one cluster. The disadvantage of this approach is that it does not consider how far spread each cluster is and focusing only on the minimum distance between the clusters to merge them.

Complete Linkage: It uses the maximum distance between elements of two clusters to merge them as one cluster. For two clusters A and B, it finds the maximum euclidean distance between the points of each cluster and compares them against the minimum distance of all the other clusters to merge them in one cluster. This approach creates a more compact and less spread cluster than single linkage clustering technique.

Average Clustering: It uses average distance of all the elements between two clusters to merge them as one cluster. For two clusters A and B, it finds the average distance by calculating the euclidean distance between all the points in each cluster and dividing them by the number of elements in each cluster. The average distance calculated is compared against average distance of other clusters to merge the minimum average distance clusters in to one cluster. The type of cluster formed by average linkage depend heavily on the structure of clusters, since it is based on the average distance between all the elements in the two clusters.

Average Group Clustering: It uses centroid distance between two clusters to merge them as one cluster. For two clusters A and B, it finds the centroid of the two clusters and merges them together by comparing against the centroid distances of other clusters. It forms similar clusters to the average linkage clusters.

Ward's Method: It uses sum of variance between two clusters to merge them as one cluster. It forms minimum spread clusters around the centroid of the cluster.

```
return minDistance
15
16
  def agglomerativeSingleLinkageCluster(clusterValue, clusterPoints):
18
      distanceMatrix = []
19
      finalClusters = []
20
21
      for i in range(0,len(clusterPoints)):
22
           finalClusters.append([i])
23
24
      for c in range (len (cluster Points), cluster Value, -1):
25
           bestClusterA = []
26
           bestClusterB = []
27
           bestCost = math.inf
28
           for i in range(0,len(clusterPoints)):
               temp = []
30
               for j in range(0,len(clusterPoints)):
                   temp.append(0)
               distanceMatrix.append(temp)
           for i in range(0,len(clusterPoints)):
               for j in range((i+1),len(clusterPoints)):
36
                   distance = singleLinkage(clusterPoints[i], clusterPoints[j])
37
                   distanceMatrix[i][j] = (distance)
38
                   if bestCost > distance:
39
                        bestCost = distance
40
```

```
if bestClusterA and bestClusterB:
41
                           bestClusterA.pop()
                           bestClusterB.pop()
                       bestClusterA.append(clusterPoints[i])
44
                       bestClusterB.append(clusterPoints[j])
45
          clusterPoints.remove(bestClusterB[0])
46
          index = clusterPoints.index(bestClusterA[0])
47
          for x in range(0,len(bestClusterB[0])):
48
               clusterPoints [index].append(bestClusterB[0][x])
49
      return (clusterPoints)
50
  def completeLinkage(clusterPointA, clusterPointB):
      maxDistance = 0
53
      for i in range(0, len(clusterPointA)):
54
          for j in range(0,len(clusterPointB)):
               distance = math.sqrt(math.pow((clusterPointA[i][0]-clusterPointB[j
56
     [[0]],2) + math.pow((clusterPointA[i][1]-clusterPointB[j][1]),2))
               if maxDistance < distance:
57
                   maxDistance = distance
      return maxDistance
  def agglomerativeCompleteLinkageCluster(clusterValue, input):
62
      distanceMatrix = []
63
      finalClusters = []
64
65
```

```
clusterPoints = input
66
      for i in range(0,len(clusterPoints)):
           finalClusters.append([i])
69
      for c in range (len (cluster Points), cluster Value, -1):
70
           bestClusterA = []
71
           bestClusterB = []
72
           bestCost = 0
73
           for i in range(0,len(clusterPoints)):
74
               temp = []
75
               for j in range(0,len(clusterPoints)):
                   temp.append(0)
77
               distanceMatrix.append(temp)
79
           for i in range(0,len(clusterPoints)):
80
               for j in range((i+1),len(clusterPoints)):
81
                   distance = completeLinkage(clusterPoints[i], clusterPoints[j])
                   distanceMatrix[i][j] = (distance)
                   if bestCost < distance:</pre>
                        bestCost = distance
                        if bestClusterA and bestClusterB:
                            bestClusterA.pop()
87
                            bestClusterB.pop()
88
                        bestClusterA.append(clusterPoints[i])
89
                        bestClusterB.append(clusterPoints[j])
90
           clusterPoints.remove(bestClusterB[0])
91
```

```
index = clusterPoints.index(bestClusterA[0])
92
            for x in range(0,len(bestClusterB[0])):
                 clusterPoints[index].append(bestClusterB[0][x])
       return (clusterPoints)
96
   \begin{array}{lll} \textbf{def} & average Linkage (\ cluster Point A\ , & cluster Point B\ ): \end{array}
       averageDistance = 0
98
       for i in range(0, len(clusterPointA)):
99
            for j in range(0,len(clusterPointB)):
100
                 averageDistance = averageDistance + math.sqrt(math.pow((
101
       clusterPointA[i][0] - clusterPointB[j][0]),2) + math.pow((clusterPointA[i
       [1] - cluster Point B [j][1], [2]
       averageDistance = averageDistance/(len(clusterPointA) +len(clusterPointB))
102
       return averageDistance
103
104
105
   def agglomerativeAverageLinkageCluster(clusterValue, clusterPoints):
107
       distanceMatrix = []
108
        finalClusters = []
109
110
       for i in range(0,len(clusterPoints)):
111
            finalClusters.append([i])
113
        for c in range (len (cluster Points), cluster Value, -1):
114
            bestClusterA = []
115
```

```
bestClusterB = []
116
           bestCost = math.inf
           for i in range(0,len(clusterPoints)):
               temp = []
119
                for j in range(0,len(clusterPoints)):
120
                    temp.append(0)
                distanceMatrix.append(temp)
           for i in range(0,len(clusterPoints)):
124
                for j in range ((i+1), len (cluster Points)):
                    distance = averageLinkage(clusterPoints[i], clusterPoints[j])
                    distanceMatrix[i][j] = (distance)
127
                    if bestCost > distance:
128
                        bestCost = distance
129
                        if bestClusterA and bestClusterB:
130
                            bestClusterA.pop()
131
                            bestClusterB.pop()
                        bestClusterA.append(clusterPoints[i])
133
                        bestClusterB.append(clusterPoints[j])
           clusterPoints.remove(bestClusterB[0])
135
           index = clusterPoints.index(bestClusterA[0])
136
           for x in range(0,len(bestClusterB[0])):
137
                clusterPoints [index].append(bestClusterB[0][x])
138
       return (clusterPoints)
139
140
def averageGroupLinkage(clusterPointA, clusterPointB):
```

```
centroidX_A = 0
142
        centroidY_A = 0
143
        centroidX_B = 0
144
        centroidY_B = 0
145
        centroidA = []
146
        centroidB = []
147
        for i in range(0, len(clusterPointA)):
148
            centroidX_A = centroidX_A + clusterPointA[i][0]
149
            centroidY_A = centroidY_A + clusterPointA[i][1]
        centroid A. append (centroid X_A/(len(cluster Point A)))
151
        centroidA.append(centroidY_A/(len(clusterPointA)))
152
        for j in range (0, len (clusterPointB)):
153
            centroidX_B = centroidX_B + clusterPointB[j][0]
154
            centroidY_B = centroidY_B + clusterPointB[j][1]
155
        centroidB.append(centroidX_B/(len(clusterPointB)))
156
        centroidB.append(centroidY_B/(len(clusterPointB)))
157
                              \operatorname{math.sqrt}(\operatorname{math.pow}((\operatorname{centroidA}[0] - \operatorname{centroidB}[0]), 2) +
        clusterDistance =
158
       math.pow((centroidA[1]-centroidB[1]),2))
        return clusterDistance
159
160
161
   def agglomerativeAverageGroupLinkageCluster(clusterValue, clusterPoints):
162
163
        distanceMatrix = []
164
        finalClusters = []
165
166
```

```
for i in range(0,len(clusterPoints)):
167
            finalClusters.append([i])
169
       for c in range (len (cluster Points), cluster Value, -1):
170
            bestClusterA = []
171
            bestClusterB = []
172
            bestCost = math.inf
173
            for i in range(0,len(clusterPoints)):
174
                temp = []
                for j in range (0, len (cluster Points)):
176
                    temp.append(0)
177
                distanceMatrix.append(temp)
178
179
            for i in range(0,len(clusterPoints)):
180
                for j in range((i+1),len(clusterPoints)):
181
                    distance = averageGroupLinkage(clusterPoints[i], clusterPoints
182
       [j])
                    distanceMatrix[i][j] = (distance)
                    if bestCost > distance:
                         bestCost = distance
185
                         if bestClusterA and bestClusterB:
186
                             bestClusterA.pop()
187
                             bestClusterB.pop()
188
                         bestClusterA.append(clusterPoints[i])
189
                         bestClusterB.append(clusterPoints[j])
190
            clusterPoints.remove(bestClusterB[0])
191
```

```
index = clusterPoints.index(bestClusterA[0])
192
            for x in range(0,len(bestClusterB[0])):
                clusterPoints [index]. append (bestClusterB[0][x])
194
       return (clusterPoints)
195
196
   def wardMethod(clusterPointA, clusterPointB):
197
       centroidX_A = 0
198
       centroidY_A = 0
199
       centroidX_B = 0
200
       centroidY_B = 0
201
       centroidA = []
202
       centroidB = []
203
       for i in range (0, len (clusterPointA)):
204
            centroidX_A = centroidX_A + clusterPointA[i][0]
205
            centroidY_A = centroidY_A + clusterPointA[i][1]
206
       centroid A. append (centroid X_A / (len (cluster Point A)))
207
       centroid A. append (centroid Y_A / (len (cluster Point A)))
208
       for j in range(0,len(clusterPointB)):
209
            centroidX_B = centroidX_B + clusterPointB[j][0]
210
            centroidY_B = centroidY_B + clusterPointB[j][1]
211
       centroidB.append(centroidX_B/(len(clusterPointB)))
       centroidB.append(centroidY_B/(len(clusterPointB)))
213
                             math.sqrt(math.pow((centroidA[0]-centroidB[0]),2) +
       clusterDistance =
214
      \operatorname{math.pow}((\operatorname{centroidA}[1] - \operatorname{centroidB}[1]), 2))
       variance = len(clusterPointA)* len(clusterPointB)* clusterDistance/ len(
215
       clusterPointA)+ len(clusterPointB)
```

```
return variance
216
   def agglomerativeWardMethod(clusterValue, clusterPoints):
220
       distanceMatrix = []
221
       finalClusters = []
222
223
       for i in range(0,len(clusterPoints)):
224
           finalClusters.append([i])
225
226
       for c in range (len (cluster Points), cluster Value, -1):
227
           bestClusterA = []
228
           bestClusterB = []
229
           bestCost = math.inf
230
           for i in range(0,len(clusterPoints)):
231
                temp = []
                for j in range(0,len(clusterPoints)):
                    temp.append(0)
                distanceMatrix.append(temp)
235
236
           for i in range(0,len(clusterPoints)):
237
                for j in range((i+1),len(clusterPoints)):
238
                    distance = wardMethod(clusterPoints[i], clusterPoints[j])
239
                    distanceMatrix[i][j] = (distance)
240
                    if bestCost > distance:
241
```

```
bestCost = distance
242
                        if bestClusterA and bestClusterB:
                            bestClusterA.pop()
                            bestClusterB.pop()
245
                        bestClusterA.append(clusterPoints[i])
246
                        bestClusterB.append(clusterPoints[j])
247
           clusterPoints.remove(bestClusterB[0])
248
           index = clusterPoints.index(bestClusterA[0])
249
           for x in range(0,len(bestClusterB[0])):
250
               clusterPoints [index]. append (bestClusterB[0][x])
251
       return (clusterPoints)
252
253
  def main():
254
       clusters = 3
255
       256
      1]], [[2, 3]], [[3, 2]], [[3, 4]], [[4,3]]]
       file = open("ClusteringOutput.txt", "w")
257
       clusterPointsSingle = agglomerativeSingleLinkageCluster(clusters,input)
       file.write("Single Linkage" + "\n")
259
       for i in range (0, len (cluster Points Single)):
260
           file.write("Cluster" + str(i+1) + ": " +str(clusterPointsSingle[i]) +
261
       "\n")
        \text{inputComplete } = \! [[[-4 \,, \ -2]] \,, \ [[-3 \,, \ -2]] \,, \ [[-2 \,, \ -2]] \,, \ [[-1 \,, \ -2]] \,, \ [[1 \,, \ -1]] \,, \\
262
       [[1, 1]], [[2, 3]], [[3, 2]], [[3, 4]], [[4,3]]]
       clusterPointsComplete = agglomerativeCompleteLinkageCluster(clusters,
263
      inputComplete)
```

```
file . write ("\nComplete Linkage\n")
264
                           for i in range(0, len(clusterPointsComplete)):
265
                                           file . write ("Cluster" + str(i+1) + ": "+str(clusterPointsComplete[i])
266
                           + "\n")
                           inputAverage = [[[-4, -2]], [[-3, -2]], [[-2, -2]], [[-1, -2]], [[1, -1]],
267
                            [[1, 1]], [[2, 3]], [[3, 2]], [[3, 4]], [[4,3]]]
                           clusterPointsAverage = agglomerativeAverageLinkageCluster(clusters,
268
                        inputAverage)
                            file . write ("\nAverage Linkage\n")
269
                           for i in range (0, len (clusterPointsAverage)):
270
                                            file . write ("Cluster" + str(i+1) + ": " + str(clusterPointsAverage[i])+
271
                            "\n")
                           inputAverageGroup \, = \, \left[ \left[ \left[ \, -4 \, , \, -2 \right] \right], \, \left[ \left[ \, -3 \, , \, -2 \right] \right], \, \left[ \left[ \, -2 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, 1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, -2 \right] \right], \, \left[ \left[ \, -1 \, , \, 
272
                         [-1], [[1, 1]], [[2, 3]], [[3, 2]], [[3, 4]], [[4, 3]]
                           clusterPointsAverageGroup = agglomerativeAverageGroupLinkageCluster(
273
                         clusters ,inputAverageGroup)
                           file.write("\nAverage Group Linkage\n")
274
                           for i in range(0, len(clusterPointsAverageGroup)):
275
                                           file . write ("Cluster" + str(i+1) + ": "+str(clusterPointsAverageGroup
276
                         [i] + "\n")
                          inputWard = [[[-4, -2]], [[-3, -2]], [[-2, -2]], [[-1, -2]], [[1, -1]],
277
                         [[1, 1]], [[2, 3]], [[3, 2]], [[3, 4]], [[4,3]]]
                           clusterPointsWard = agglomerativeWardMethod(clusters,inputWard)
278
                            file . write ("\nWard's Algorithm Linkage\n")
279
                           for i in range(0, len(clusterPointsWard)):
280
```

Single Linkage

Cluster 1: [[-4, -2], [-3, -2], [-2, -2], [-1, -2]]

Cluster 2: [[1, -1], [1, 1]]

Cluster 3: [[2, 3], [3, 2], [3, 4], [4, 3]]

Complete Linkage

Cluster 1: [[-4, -2], [4, 3], [3, 4], [-3, -2], [3, 2], [-2, -2], [2, 3], [-1, -2]]

Cluster 2: [[1, -1]]

Cluster 3: [[1, 1]]

Average Linkage

Cluster 1: [[-4, -2], [-3, -2], [-2, -2], [-1, -2]]

Cluster 2: [[1, -1], [1, 1]]

Cluster 3: [[2, 3], [3, 2], [3, 4], [4, 3]]

Average Group Linkage

Cluster 1: [[-4, -2], [-3, -2], [-2, -2], [-1, -2]]

Cluster 2: [[1, -1], [1, 1]]

Cluster 3: [[2, 3], [3, 2], [3, 4], [4, 3]]

Ward's Algorithm Linkage

Cluster 1: [[-4, -2], [-3, -2], [-2, -2], [-1, -2]]

Cluster 2: [[1, -1], [1, 1]]

Cluster 3: [[2, 3], [3, 2], [3, 4], [4, 3]]

6.2.2 Clustering Results vs Manual Clustering

All the agglomerate clustering techniques except for complete linkage technique produced the same result for the provided data set. The results of the clusters formed by all the clustering methods has been shown above.

Manual Clustering of the data points on euclidean distance results in the same clusters generated by agglomerative clustering.

Cluster 1: (-4,-2), (-3,-2), (-2,-2), (-1,-2)

Cluster 2: (1,-1), (1,1)

Cluster 3: (2,3), (3,2), (3,4), (4,3)

Cluster 1 is easy to create due to it being far away from other data points. Cluster 2 and 3 have a close margin where the distance between point (1,1) and (-1,1) is equal to 2 and the distance between point (1,1) and (2,3) is $\sqrt{5}$. Therefore the points (1,1) and (-1,1) have been clustered together as Cluster 2.

If the data points are clustered on the basis of the quadrants they lie in:

Cluster 1: 1st Quadrant— > (-4,-2), (-3,-2), (-2,-2), (-1,-2)

Cluster 2: 3rd Quadrant ->(1,-1)

Cluster 3: 4th Quadrant -> (1,1), (2,3), (3,2), (3,4), (4,3)

Chapter 7

Problem 9.9

7.1 Problem

Use K-means and spherical K-means to cluster the data points in Exercise 9.8. How do the clustering differ?

7.2 Solution

```
Created on Nov 24, 2017

Quathor: nauman

,,,

import numpy as np

from sklearn.cluster import KMeans
```

```
8 from spherecluster import SphericalKMeans
 def kMeans(num):
11
     file = open("Kmeans.txt", "a+")
12
     13
     1], [2, 3], [3, 2], [3, 4], [4,3]])
     kmeans = KMeans(n_clusters=num, random_state=0).fit(input)
14
     file.write("K means output for cluster size: "+ str(num) + "\n")
     file.write("Clusters index of points" + "\n")
16
     file.write(str(kmeans.labels_) + "\n")
17
     file . write ("Center of Clusters \n")
18
     file.write(str(kmeans.cluster\_centers\_) + "\n")
19
     file.close()
20
21
  def sphericalKMeans(num):
22
     num = 4
23
     file = open("Kmeans.txt", "a+")
24
     [1], [2, 3], [3, 2], [3, 4], [4,3]]
     kmeans = SphericalKMeans(n_clusters=num).fit(input)
     file.write("Spherical K means output for cluster size : "+ str(num) + "\n"
27
     )
     file.write("Clusters index of points" + "\n")
28
     file . write (str(kmeans.labels_+) + "\n")
29
     file . write ("Center of Clusters \n")
30
```

```
file.write(str(kmeans.cluster_centers_) + "\n")
31
      file.close()
35 kMeans (4)
36 sphericalKMeans (4)
  K means output for cluster size : 3
  Clusters index of points
  [2 2 2 2 0 0 1 1 1 1]
  Center of Clusters
  [[ 1.0000000e+00
                      5.55111512e-17]
   [ 3.00000000e+00 3.00000000e+00]
   [ -2.50000000e+00 -2.00000000e+00]]
  Spherical K means output for cluster size : 4
  Clusters index of points
  [3 3 3 1 2 0 0 0 0 0]
  Center of Clusters
  [[ 0.70710678  0.70710678]
   [-0.4472136 -0.89442719]
   [ 0.70710678 -0.70710678]
   [-0.81836024 -0.57470559]]
  K means output for cluster size : 2
```

Clusters index of points

[0 0 0 0 0 1 1 1 1 1]

Center of Clusters

[[-1.8 -1.8]

[2.6 2.6]]

Spherical K means output for cluster size : 4

Clusters index of points

[3 3 3 0 2 1 1 1 1 1]

Center of Clusters

[[-0.4472136 -0.89442719]

[0.70710678 0.70710678]

[0.70710678 -0.70710678]

[-0.81836024 -0.57470559]]

K means output for cluster size : 4

Clusters index of points

[2 2 0 0 3 3 1 1 1 1]

Center of Clusters

[[-1.50000000e+00 -2.00000000e+00]

[3.00000000e+00 3.00000000e+00]

[-3.50000000e+00 -2.00000000e+00]

[1.00000000e+00 5.55111512e-17]]

Table 7.1: Index to Data Point relation

Index 0 1 2 3 4 5 6 7 8 9

Data Point (-4, -2) (-3,-2) (-2,-2) (-1,-2) (1,-1) (1,1) (2,3) (3,2) (3,4) (4,3)

Spherical K means output for cluster size : 4

Clusters index of points

[3 3 3 1 2 0 0 0 0 0]

Center of Clusters

[[0.70710678 0.70710678]

[-0.4472136 -0.89442719]

[0.70710678 -0.70710678]

[-0.81836024 -0.57470559]]

The output of the code shows clusters on the basis of the index of data points and center of each cluster. Table 5.1 shows the index to data point relationship. The results for cluster size 2,3 and 4 are show for both the clustering techniques.

The clusters formed by K-means for cluster size 3 are same as the clusters formed by Single Linkage, Average Linkage and Average Group Linkage from the problem 9.8. The clusters formed by spherical K-means for cluster 3 is different from K-means due to the difference between finding similarity between points for clustering. K-means uses euclidean distance to find similarity while spherical K-mean uses cosine similarity to cluster items together. Spherical K-means clusters items on which fall in the same quadrant approach discussed in the previous problem.

Chapter 8

Problem 9.11

8.1 Problem

The K nearest neighbors of a document could be represented by links to those documents.

Describe two ways this representation could be used in a search application.

8.2 Solution

The primary advantage of K nearest neighbor compared to K means and agglomerative clustering techniques is that a document can be present in multiple clusters unlike the K means and agglomerative clustering technique where each documents is in a single cluster.

The K nearest neighbors of a documents can be used for tell-text search in search application. All the documents in the corpus can be clustered by K nearest neighbor. The

document can be in multiple clusters based on the features extracted from the cluster. For a corpus of documents containing sports news from USA, it can be classified into multiple clusters as football, baseball, basketball, athletics, Olympics, CONCAF Cup etc. A document that is in cluster soccer can simultaneously be clustered with Olympics and CONCAF Cup. Similarly the queries to the documents can also be clustered on the basis of its features. It presents all the documents that are in the cluster soccer for a query of feature soccer. The same set of documents can also be recalled if the query is about CONCAF Cup. It allows for relevant search results.

The K nearest neighbors can also be used for showing clustered search results. It can be helped to refine a very generic query to a more precise query to return relevant result. For example, a system that employs K nearest neighbor clustering on its queries. For a user input query apple, it will show all the possible clusters where the query term apple appears. For apple, it can be a fruit, it can be the company Apple, it can be products of Apple Company etc. A user on the basis of suggestions can select the particular suggestion to find search results that fall under the specified feature cluster.