

EECS E6893 Big Data Analytics

Homework #1 Report

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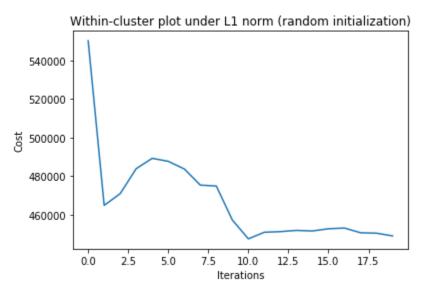
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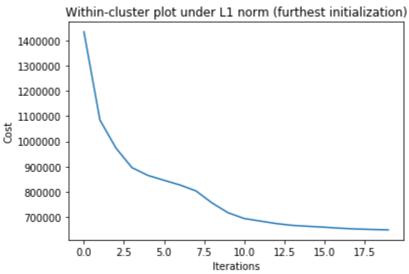
October 2022

Question 1

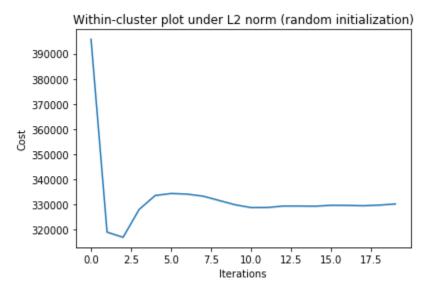
(1) The results are as follows. For random initialization, it can reach a better result within 20 iterations compared to the furthest initialization (compared to the cost function value at iteration 20).

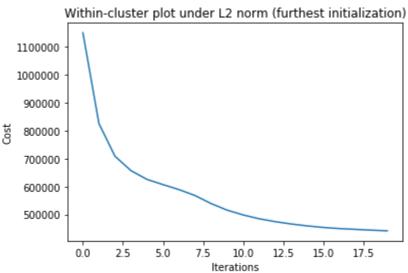
However, the furthest initialization's cost function is always decreasing within 20 iterations while random initialization sometimes failed to minimize the cost function. It might be an indication that the furthest initialization is a bad choice to start, thus leaving lots of room to improve.



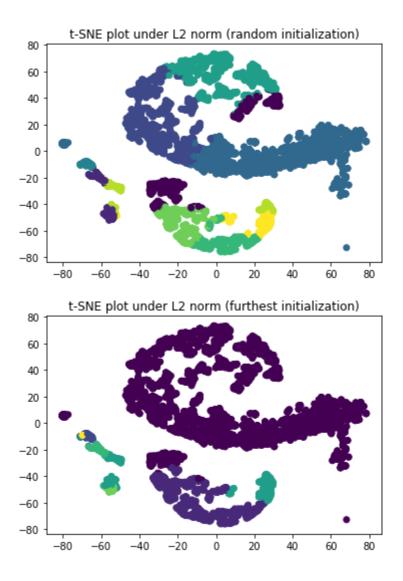


(2) The results are as follows. Still, the situations are very similar to (1). Random initialization can achieve better results but is not able to constantly minimize the cost function.





(3) The results are as follows. It can be seen that the furthest initialization is worse because its plot has fewer colors



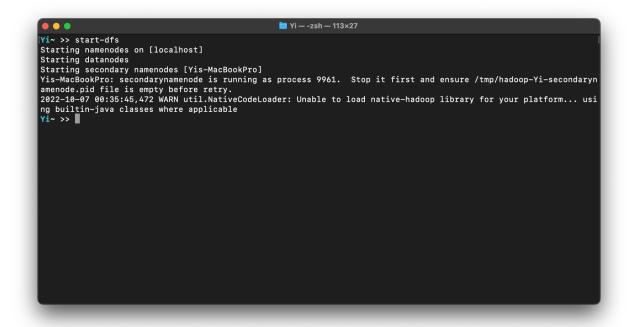
- (4) Yes. Regardless of the similarity measure, random initialization is better in terms of cost. As mentioned before, the *c2.txt* gives us the furthest cluster setting. However, it is a bad starting point because the underlying data clusters are not necessarily far away from each other. Given the limited iterations, a bad starting point will not give us good results, which is the reason for the outperformance of the random initialization.
- (5) Given a fixed number of centroids K and iterations T, suppose we have N data points to cluster, the time complexity is O(KTN)

Question 2

(1) The installation results are as follows.

```
hadoop—rzsh — 113x27

/////sr/local/Cellar/hadoop/3.3.4/libexec/etc/hadoop—rzsh .... ////sr/local/Cellar/hadoop—rzsh .... ////sr/local-race .... ///sr/local-race .... //sr/local-race .... ///sr/local-race .... //sr/local-race .... //sr/lo
```



(2) The HTTP API results are as follows. I think the following metrics are the most important ones.

DataNodes usages

This metrics gives us how heavily we are utilizing the datanodes, with min, max, median and standard deviation, we can have a comprehensive knowledge of the usages.

Missing blocks (obtained through the JSON output using jmx suffix)
A missing block metric by its name, documents the number of meaning blocks in the cluster. It is important for us to evaluate the health of our cluster because we cannot recover the data in that block anymore if it is missed, so it's crucial to keep track of this metric.

Dead Nodes

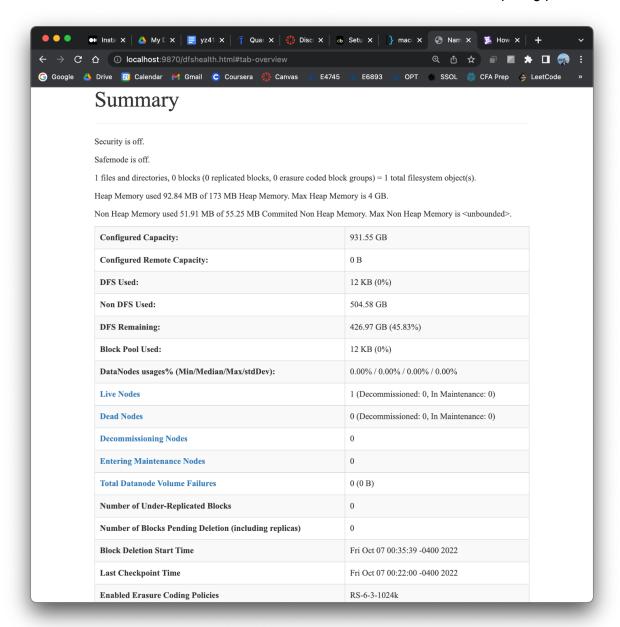
This metric simply count the number of dead nodes, which are nodes that have not communicated with namenode for a while. We need to monitor this because too many dead nodes will result in data loss and will consume lots of cluster recources.

Block Pool Used

This metrics measures the percentage of the the blocks we are using. In order to ensure the continuity of the cluster, we should make sure enough blocks are available at any time.

Total Datanode Volume Failure

It simply tracks the volume of failure when a disk is down. The new version of Hadoop is able to handle the disk failure, i.e. the datanode will continue to work even if disks are failed (up to a preset threshold). Monitoring this metric can help us understand how the failed volume behaves over time, based on this information, we can choose our threshold to maximize our computing power.

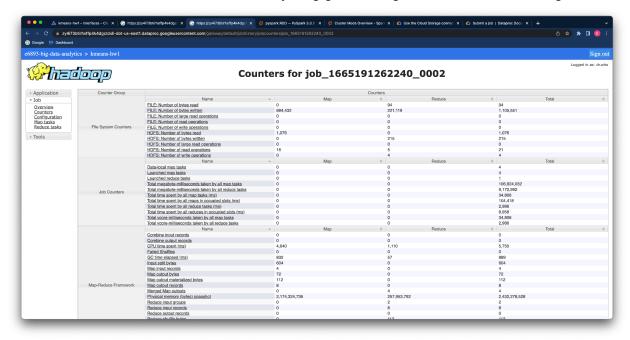


(3) To monitor the tasks, we clicked the counters for a specific Job to see more details. For file system counter, there are numbers regarding read/written for map and reduce operations.

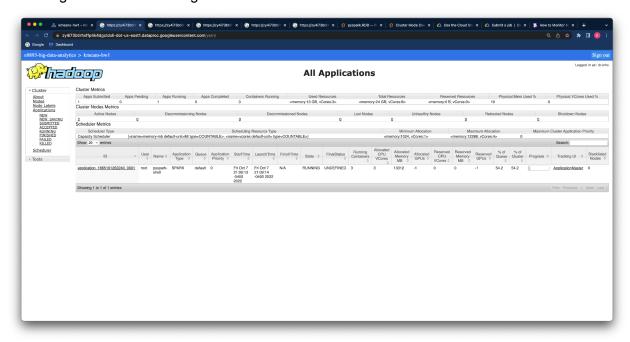
For jobs counter, the API provides total time for mapping and reducing. Through these metrics, we can know which part of the job is harder and takes more time.

For map-reduce framework, we can have information about memory usages, heap usages etc.

For shuffle error counters, we can see if anything goes wrong in the course of shuffling.



(4) The API gives us the following.



There are 3 metric classes shown as cluster metrics, cluster nodes metrics, scheduler metrics. I will discuss the important ones from each category.

Total Resources & Used Resources

It tells the number of cores and the memory we have. Combining these two metrics, we can get a glimpse of the overall performance cluster resources.

Unhealthy Nodes

It tells the number of unhealthy nodes. A node is unhealthy if its disk utilization level is beyond some preset threshold. Because other nodes will allocate their resources to make up for his unhealthy brother, they might become unhealthy later, causing catastrophic chain reaction. Thus, it is important to monitor this metrics.

Minimum & Maximum Allocation

It measures the maximum memory and number of cores used for a job. By monitoring this number, we can learn what jobs consume most of our resources.

Lost Nodes

The usage is as its name suggests. We can look at this metric and add new nodes if number of lost nodes is too large.

Scheduling Resource Type

This metric tells us what kind of resources is being allocated to our jobs. It is again useful for us since we can learn the resources that our jobs consume.

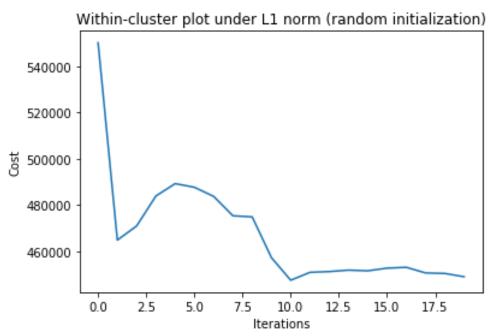
```
In [1]: import operator
         import sys, os
         import numpy as np
         import matplotlib.pyplot as plt
         import seaborn as sns
         from scipy import linalg
         from sklearn.manifold import TSNE
         from pyspark import SparkConf, SparkContext
         /opt/conda/anaconda/lib/python3.7/site-packages/statsmodels/tools/_testing.py:19: FutureWarning: pandas.util.testing is deprecated. U
         se the functions in the public API at pandas.testing instead.
           import pandas.util.testing as tm
 In [2]: # Macros.
         MAX ITER = 20
         DATA_PATH = "gs://6893_course_data/hw1_q1/data.txt"
         C1_PATH = "gs://6893_course_data/hw1_q1/c1.txt"
         C2_PATH = "gs://6893_course_data/hw1_q1/c2.txt"
         L1 = 1
         L2 = 2
In [10]: # Helper functions
         def closest(p, centroids, norm):
             Compute closest centroid for a given point.
             Args:
                 p (numpy.ndarray): input point
                 centroids (list): A list of centroids points
                 norm (int): 1 or 2
             Returns:
                 int: The index of closest centroid.
             closest_c = min([(i, linalg.norm(p - c, norm))
                             for i, c in enumerate(centroids)],
                              key=operator.itemgetter(1))[0]
             return closest_c
         def within_cluster_cost(data, centroids, norm):
             Compute within-cluster cost.
             Args:
                 data (RDD): a RDD of the form (centroid, (point, 1))
                 centroids (list): A list of centroids points
                 norm (int): 1 or 2
             Returns:
                  Float: Within-cluster cost of the current classification.
             cost = sum(data.map(lambda pt: linalg.norm(pt[1][0] - centroids[pt[0]], norm)).collect())
             return cost
         def tSNE_vis(data, norm, init):
             Produce a t-SNE 2D dimensional clustering result.
             Args:
                 data (RDD): a RDD of (centroid, (point, 1))
                 norm (int): 1 or 2
                 init (str): 'random' or 'furthest'
             Returns:
                 None
              11 11 11
             arr = np.array(data.map(lambda pt: list(pt[1][0])).collect())
             labels = data.keys().collect()
             embedded = TSNE(n_components=2, random_state=666).fit_transform(arr)
             x = embedded[:, 0]
             y = embedded[:, 1]
             plt.scatter(x, y, c=labels)
```

plt.title('t-SNE plot under L%d norm (%s initialization)'%(norm, init))

plt.show()

```
Conduct k-means clustering given data and centroid.
            Args:
                data (RDD): RDD of points
                centroids (list): A list of centroids points
                norm (int): 1 or 2
                init (str): 'random' or 'furthest'
            Returns:
                RDD: assignment information of points, a RDD of (centroid, (point, 1))
                list: a list of centroids
            cost_list = np.zeros(MAX_ITER)
            # iterative k-means
            for i in range(MAX ITER):
                # Points assignment
                points = data.map(lambda p: (closest(p, centroids, norm), (p, 1)))
                # Cost Calculation
                cost_list[i] = within_cluster_cost(points, centroids, norm)
                # Updata centroids
                reduced pts = points.reduceByKey(lambda p1, p2: (p1[0] + p2[0], p1[1] + p2[1]))
                centroids = reduced_pts.values().map(lambda c: c[0] / c[1]).collect()
            # cost plot
            plt.plot(cost_list)
            plt.xlabel('Iterations')
            plt.ylabel('Cost')
            plt.title('Within-cluster plot under L%d norm (%s initialization)'%(norm, init))
            plt.show()
            return points, centroids
In [5]: # Spark settings
        /gateway/default/node/conf?host&port = SparkConf()
        sc = SparkContext(/gateway/default/node/conf?host&port=/gateway/default/node/conf?host&port)
        # Load the data, cache this since we're accessing this each iteration
        data = sc.textFile(DATA_PATH).map(
                lambda line: np.array([float(x) for x in line.split(' ')])
                ).cache()
        # Load the initial centroids c1, split into a list of np arrays
        centroids1 = sc.textFile(C1_PATH).map(
                lambda line: np.array([float(x) for x in line.split(' ')])
                ).collect()
        # Load the initial centroids c2, split into a list of np arrays
        centroids2 = sc.textFile(C2_PATH).map(
                lambda line: np.array([float(x) for x in line.split(' ')])
                ).collect()
```

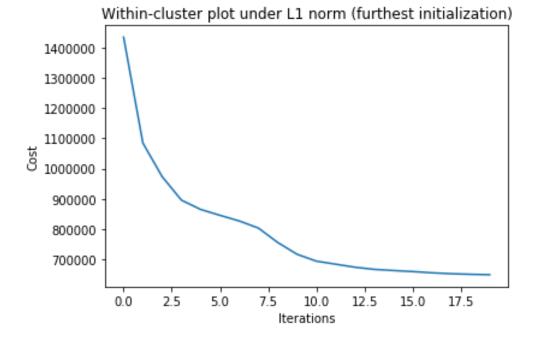




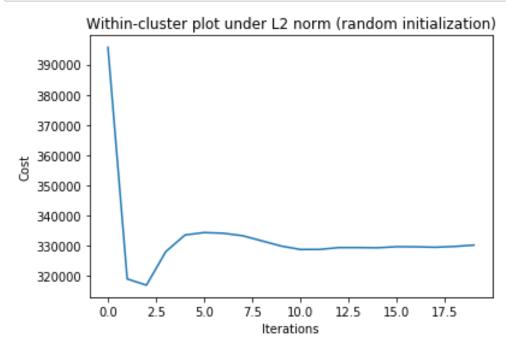
In [4]: # K-means clustering

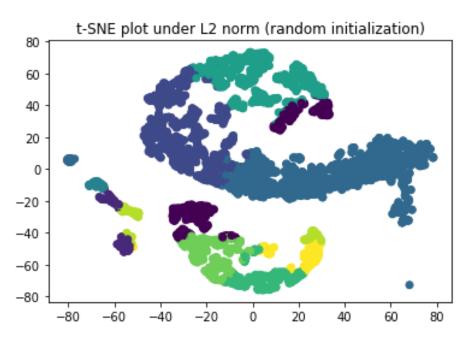
def kmeans(data, centroids, norm, init):

```
In [12]: # run kmeans
clustered_2, updated_centroids_2 = kmeans(data, centroids2, L1, 'furthest')
```



```
In [13]: # run kmeans
    clustered_3, updated_centroids_3 = kmeans(data, centroids1, L2, 'random')
    # t-SNE visualization
    tSNE_vis(clustered_3, L2, 'random')
```





In [14]: # run kmeans
 clustered_4, updated_centroids_4 = kmeans(data, centroids2, L2, 'furthest')
 # t-SNE visualization
 tSNE_vis(clustered_4, L2, 'furthest')

