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Final Project
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This project implements the k-means algorithm for the Wisconsin Breast Cancer data set while using Python. The project fills in missing values, adds a predicted class and then applies an error statement to the predicted class.

Phase 1 consisted of downloading the breast cancer data into Python. Certain values were missing in column A7 and needed to be replaced to further analyze. To replace the missing values, the mean imputation method was used. This was done by using the `fillna()` function. Once the missing values were replaced, the mean, median, standard deviation and variance of each of the attributes A2 to A10 were found using built in Python functions. Each column's results were rounded to one decimal place and printed.

In Phase 2, k-means algorithm was implemented. Using the dataset used and adjusted in Phase 1, we used k-means computation on columns A2 to A10. In this phase, two initial centroids were to be chosen at random, which is reflected by the programming. Once the centroids were chosen, they are displayed with their values from column A2 to A10. A new column is created called the Predicted Class. Each of the 699 data points were computed for their Euclidian distance from the initial centroids. Each point would fall into one of the two predicted clusters (or class). If the distance of the data point was closer from μ_2 to μ_4 , it would be assigned to Predicted Class = 2, and vice versa. The phase would assign each data point to a cluster, then update the centroids. This would happen until the centroids did not change from their previous iteration or until the steps were iterated 50 times. The results were then printed.

In Phase 3, the quality of the clustering was analyzed. This was done by calculating the error rate of the clusters. There were two clusters, benign and malign cells. This phase found the error rate for the

benign cells, malign cells, and total error rate. Using the definitions and formulae stated in the instructions, we can calculate the error rates. The results for each phase are shown below.

Phase 1

Attribute A2

Mean: 4.4

Median: 4.0

Variance: 7.9

Standard Deviation: 2.8

Attribute A3

Mean: 3.1

Median: 1.0

Variance: 9.3

Standard Deviation: 3.1

Attribute A4

Mean: 3.2

Median: 1.0

Variance: 8.8

Standard Deviation: 3.0

Attribute A5

Mean: 2.8

Median: 1.0

Variance: 8.2

Standard Deviation: 2.9

Attribute A6

Mean: 3.2

Median: 2.0

Variance: 4.9

Standard Deviation: 2.2

Attribute A7

Mean: 3.5

Median: 1.0
Variance: 13.0 Standard Deviation: 3.6

Attribute A8 -----

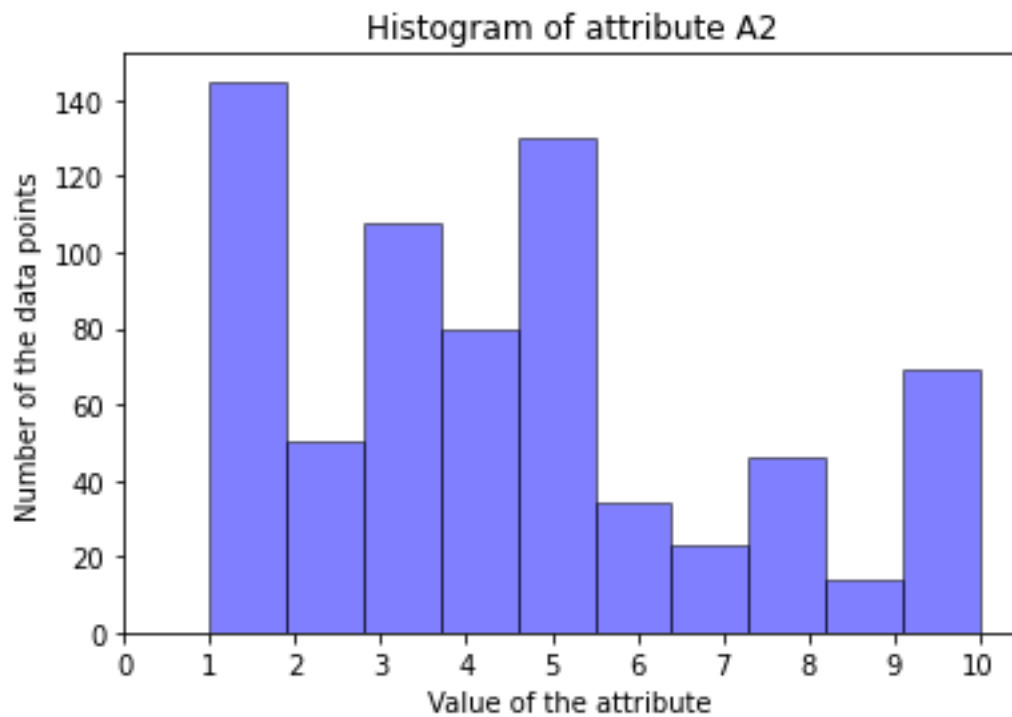
Mean: 3.4
Median: 3.0
Variance: 5.9
Standard Deviation: 2.4

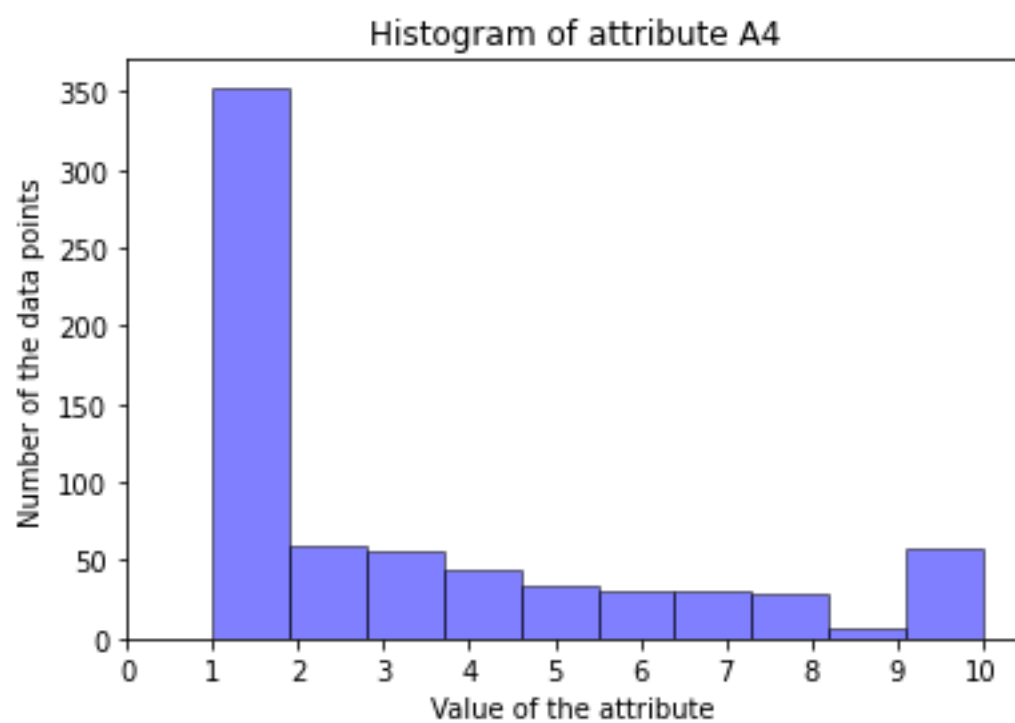
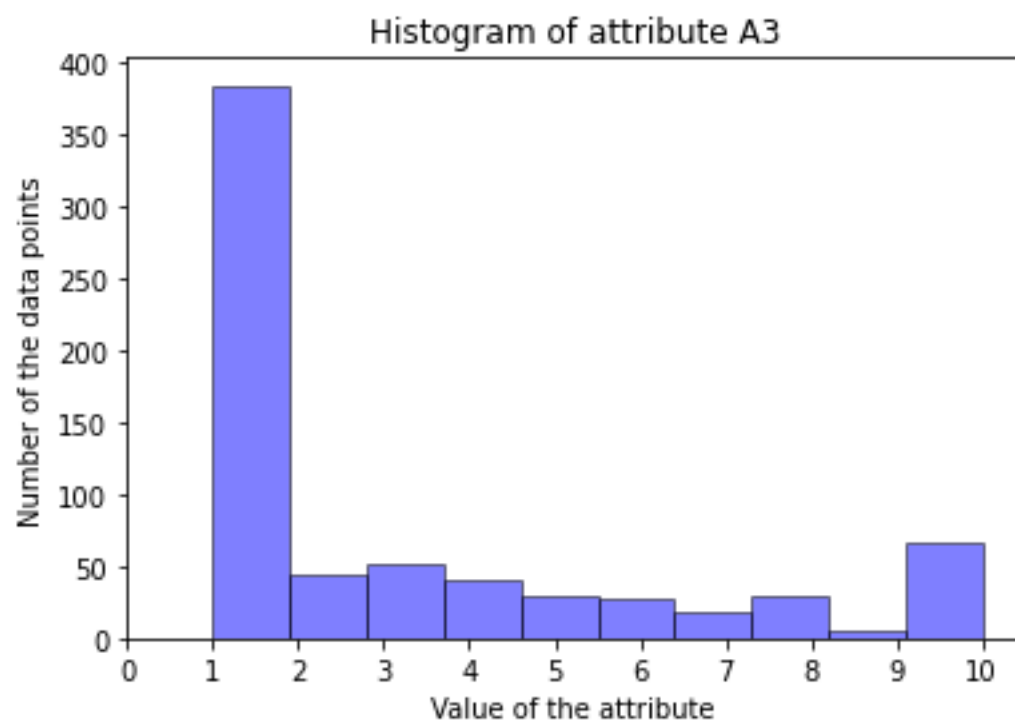
Attribute A9

Mean: 2.9
Median: 1.0
Variance: 9.3
Standard Deviation: 3.1

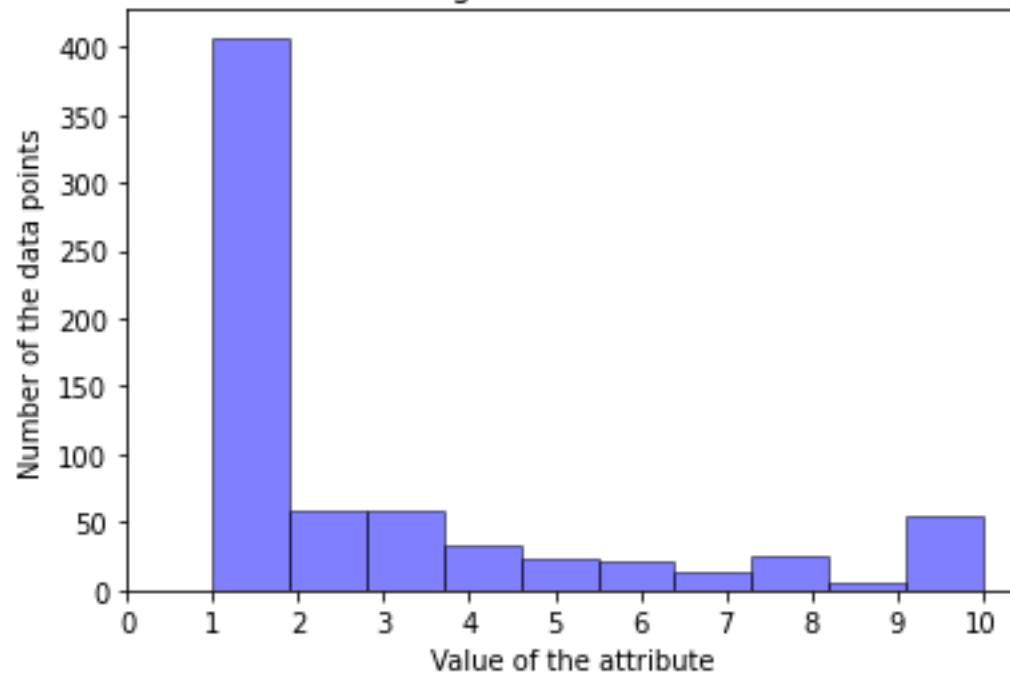
Attribute A10 -----

Mean: 1.6
Median: 1.0
Variance: 2.9
Standard Deviation: 1.7

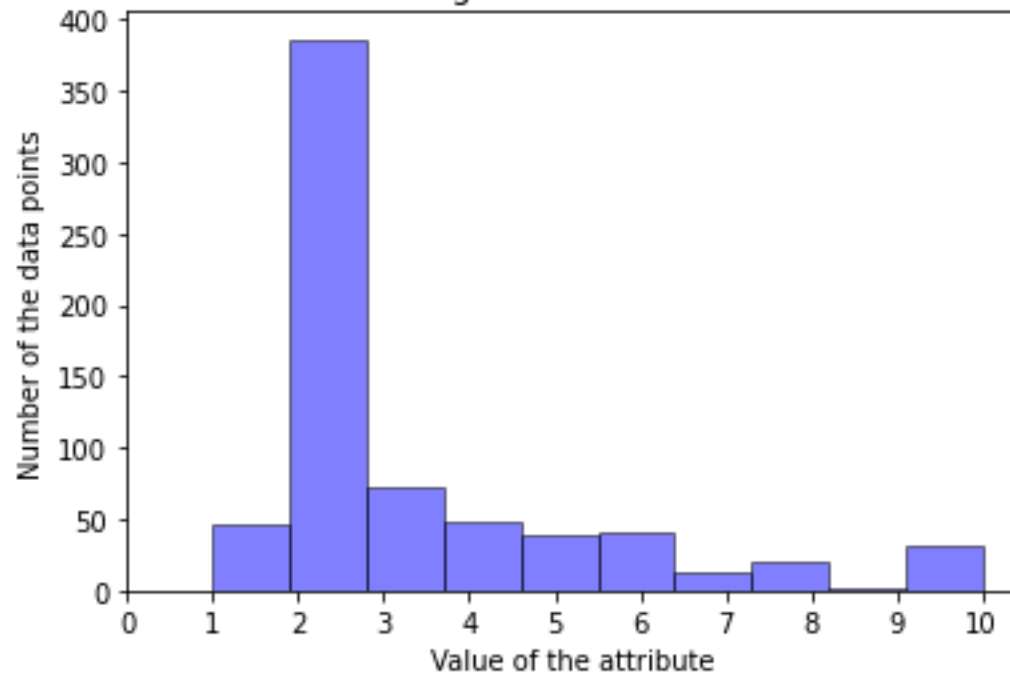




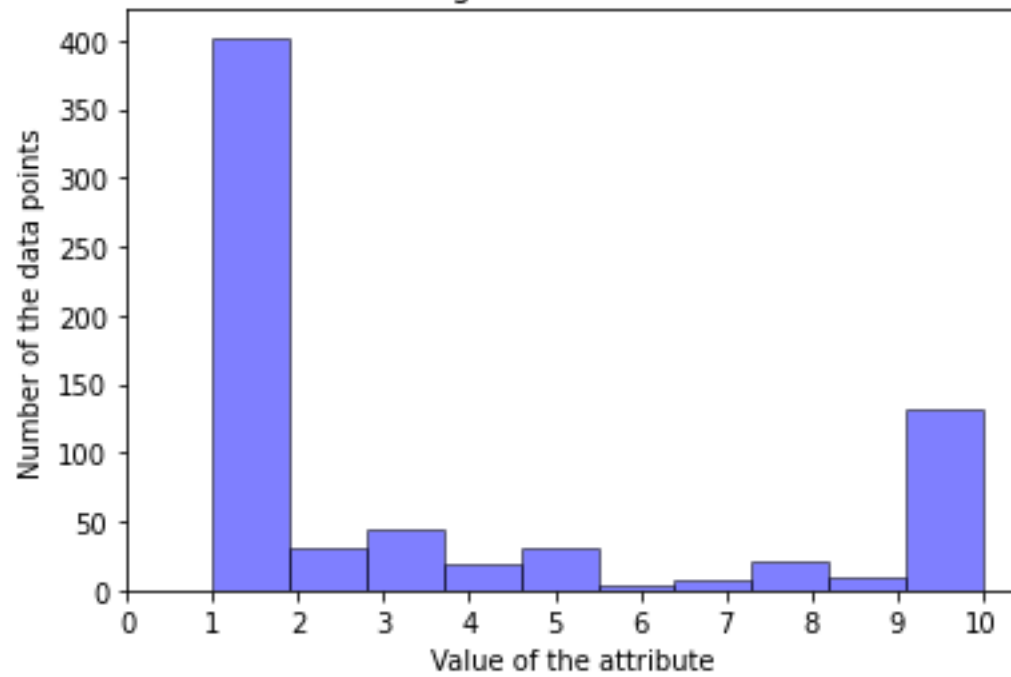
Histogram of attribute A5



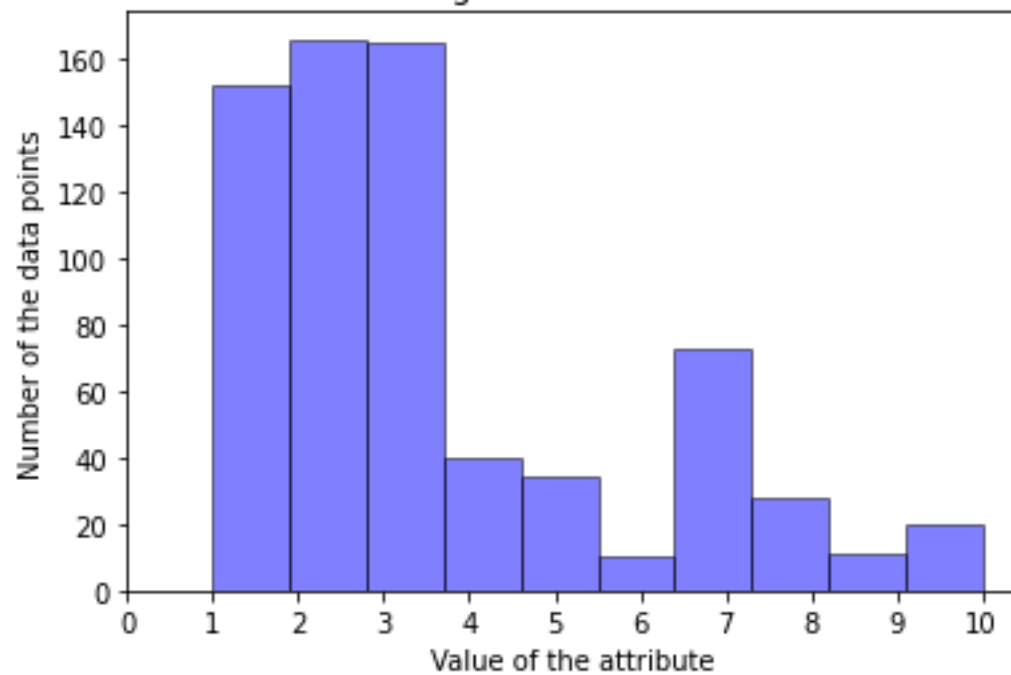
Histogram of attribute A6



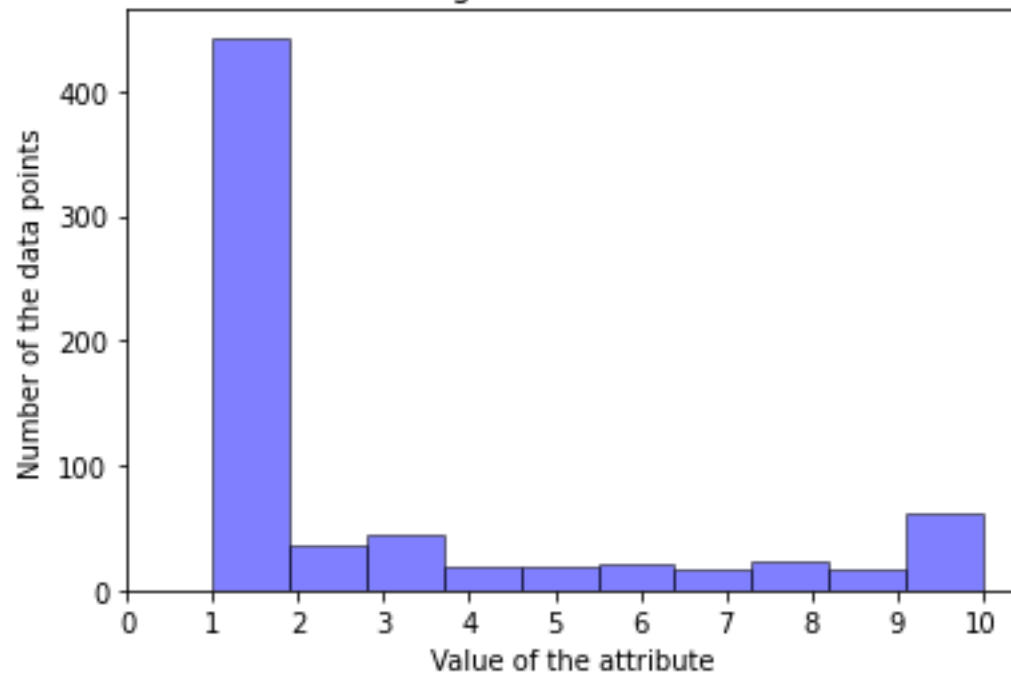
Histogram of attribute A7



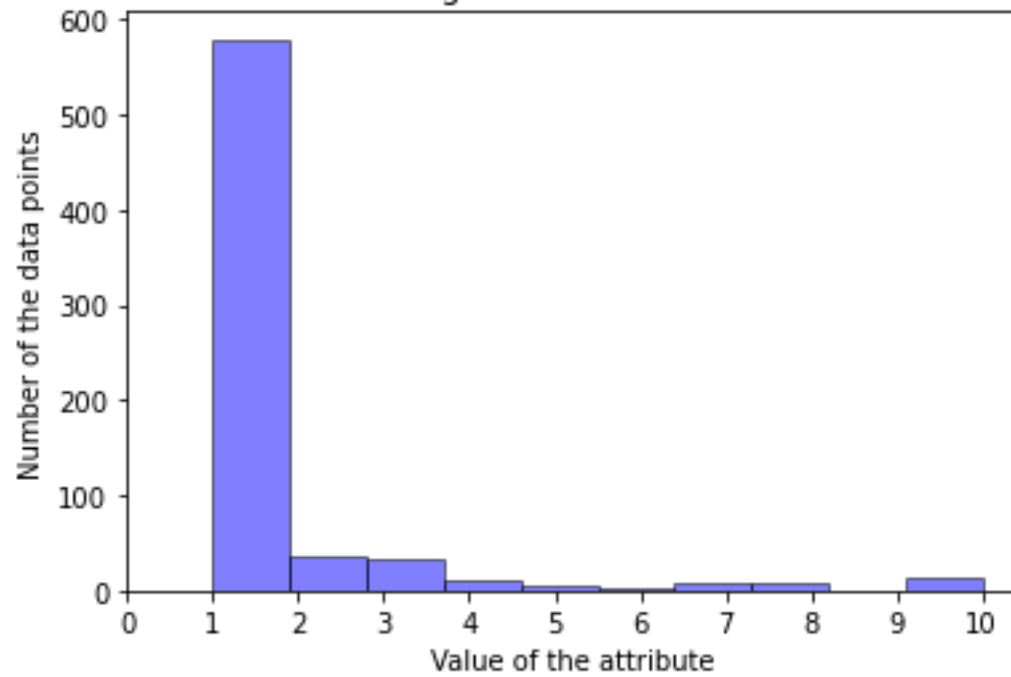
Histogram of attribute A8



Histogram of attribute A9



Histogram of attribute A10



Phase 2

Randomly selected row 310 for centroid mu_2.

Initial centroid mu_2:

A2 2.0
A3 1.0
A4 1.0
A5 1.0
A6 3.0
A7 1.0
A8 2.0
A9 1.0
A10 1.0

Name: 310, dtype: float64

Randomly selected row 592 for centroid mu_4.

Initial centroid mu_4:

A2 10.0
A3 3.0
A4 4.0
A5 5.0
A6 3.0
A7 10.0
A8 4.0
A9 1.0
A10 1.0

Name: 592, dtype: float64

Program ended after 4 iterations.

Final centroid mu_2:

A2 3.0472103004291844
A3 1.3025751072961373
A4 1.446351931330472
A5 1.3433476394849786
A6 2.087982832618026
A7 1.3800011310866602
A8 2.1051502145922747
A9 1.261802575107296
A10 1.109442060085837

Final centroid mu_4:

A2 7.1587982832618025
A3 6.798283261802575
A4 6.7296137339055795
A5 5.733905579399142

A6 5.472103004291846
A7 7.873965526992126
A8 6.103004291845494
A9 6.07725321888412
A10 2.5493562231759657

Final Cluster Assignment:

	Scn	Class	Predicted Class
0	1000025	2	2
1	1002945	2	4
2	1015425	2	2
3	1016277	2	4
4	1017023	2	2
5	1017122	4	4
6	1018099	2	2
7	1018561	2	2
8	1033078	2	2
9	1033078	2	2
10	1035283	2	2
11	1036172	2	2
12	1041801	4	2
13	1043999	2	2
14	1044572	4	4
15	1047630	4	2
16	1048672	2	2
17	1049815	2	2
18	1050670	4	4
19	1050718	2	2

Phase 3

Total errors: 4.3 %

Data points in Predicted Class 2: 466

Data points in Predicted Class 4: 233

Error data points, Predicted Class 2:

	Scn	Class	Predicted Class
12	1041801	4	2
15	1047630	4	2
23	1057013	4	2
25	1065726	4	2
50	1108370	4	2
51	1108449	4	2
57	1113038	4	2
59	1113906	4	2
63	1116132	4	2

65	1116998	4	2
101	1167439	4	2
103	1168359	4	2
105	1169049	4	2
222	1226012	4	2
273	428903	4	2
348	832226	4	2
356	859164	4	2
455	1246562	4	2
489	1084139	4	2

Error data points, Predicted Class 4:

	Scn	Class	Predicted Class
1	1002945	2	4
3	1016277	2	4
40	1096800	2	4
196	1213375	2	4
252	1017023	2	4
259	242970	2	4
296	616240	2	4
315	704168	2	4
319	721482	2	4
352	846832	2	4
434	1293439	2	4

Number of all data points: 699

Number of error points: 30

Error rate for class 2: 4.1 %

Error rate for class 4: 4.7 %

Total error rate: 4.3 %

As shown by the total error rate, the k-means algorithm was a close prediction at a rate of 4.3% total error.