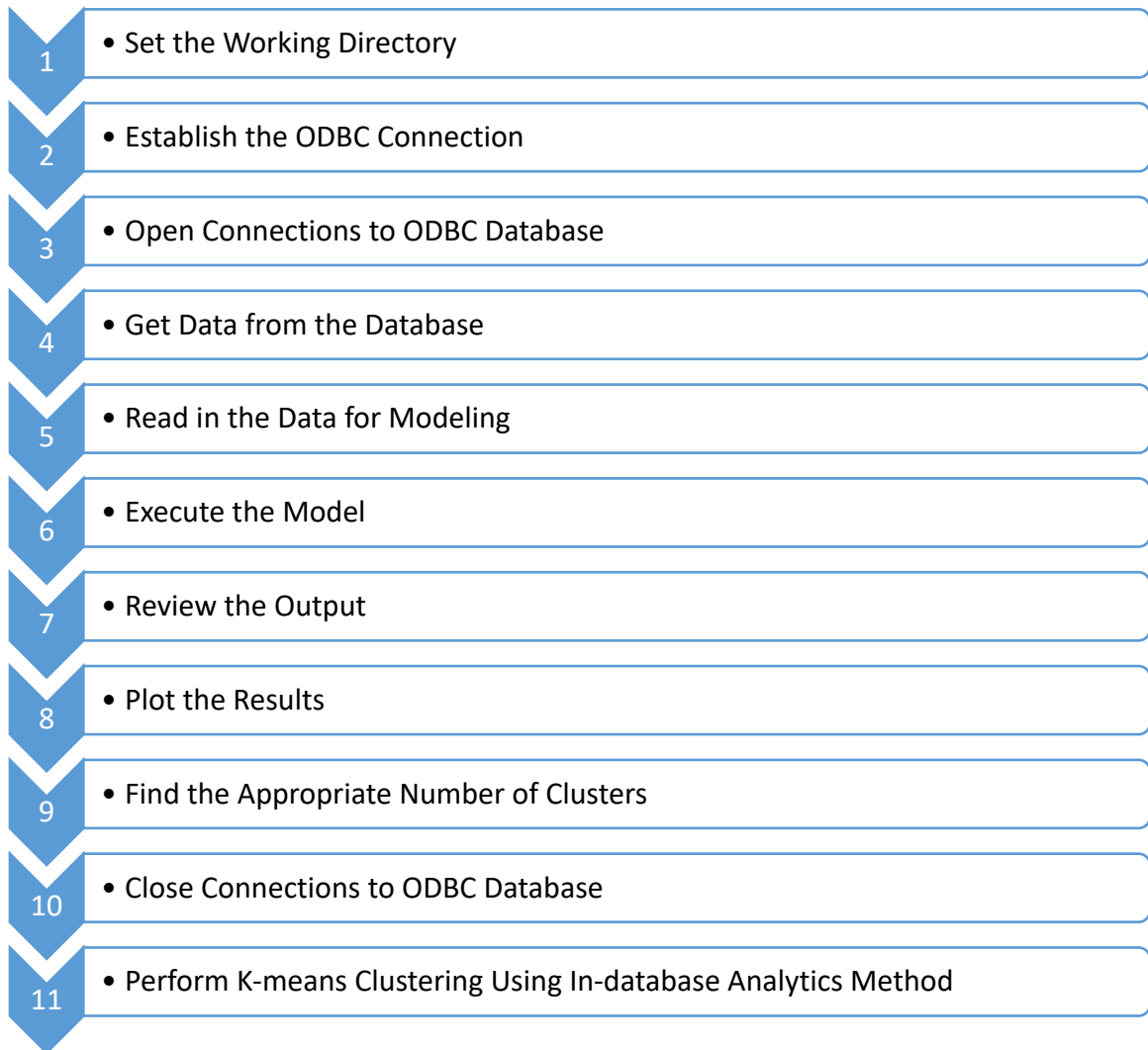


Lab Exercise 4: K-means Clustering

Purpose:	<p>This lab is designed to investigate and practice K-means Clustering. After completing the tasks in this lab you should be able to:</p> <ul style="list-style-type: none">• Use R functions to create K-means Clustering models• Use ODBC connection to the database and execute SQL statements and load datasets from the database in an R environment• Visualize the effectiveness of the K-means Clustering algorithm using graphic capabilities in R• Use MADlib functions for K-means clustering
Tasks:	<p>Tasks you will complete in this lab include:</p> <ul style="list-style-type: none">• Use the R –Studio environment to code K-means Clustering models• Use the ODBC connection in the R environment to create the average household income from the census database as test data for K-means Clustering• Use R graphics functions to visualize the effectiveness of the K-means Clustering algorithm• Use MADlib functions for K-means clustering
References:	<p>References used in this lab are located in your <i>Student Resource Guide Appendix</i>. http://www.statmethods.net/advstats/cluster.html (originally from Everitt & Hothorn).</p>

Workflow overview



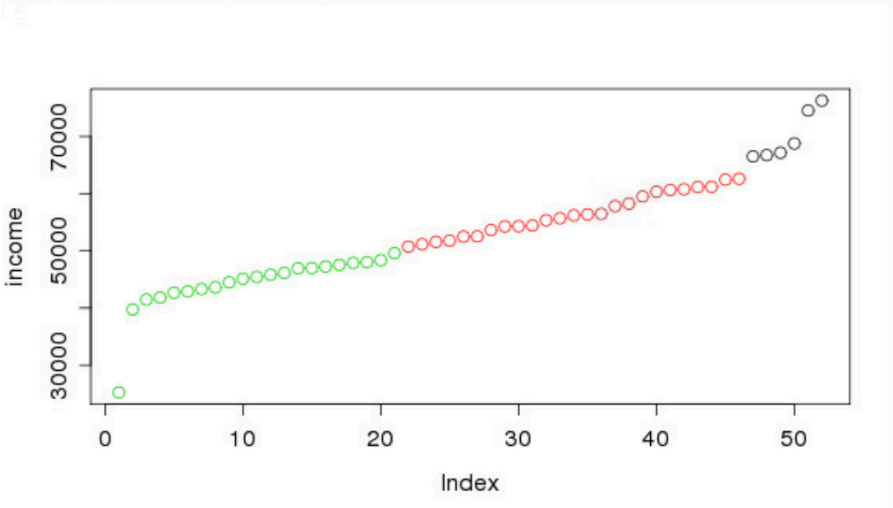
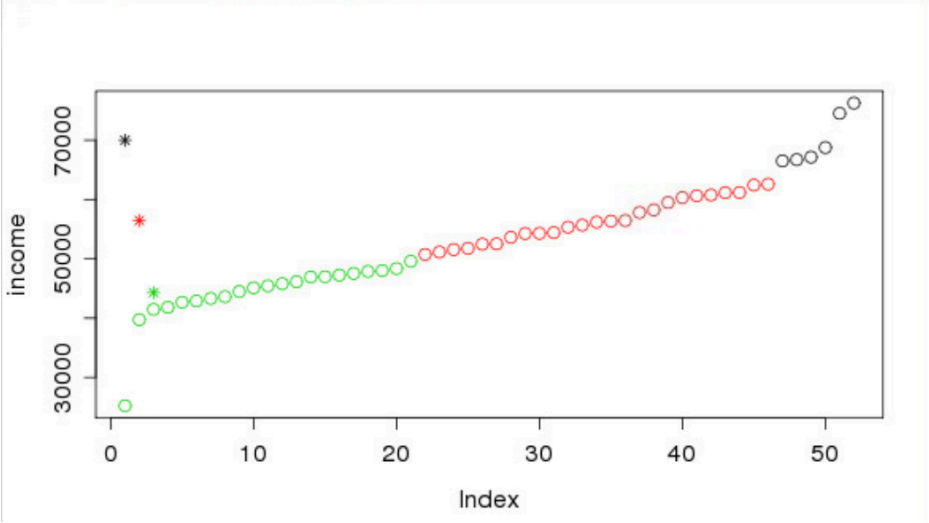
Lab Instructions

Step	Action
1	Log in with GPADMIN credentials onto R-Studio.
2	<p><u>Set the Working Directory:</u></p> <p>1. Set working directory to ~/LAB04/, execute the command:</p> <pre>setwd("~/LAB04")</pre> <ul style="list-style-type: none">• (Or use the “Tools” option in the tool bar in the RStudio environment.)
3	<p><u>Establish the ODBC Connection:</u></p> <p>Load the RODBC package, type in:</p> <pre>library('RODBC')</pre>
4	<p><u>Open Connections to ODBC Database:</u></p> <p>1. Ensure the username(uid) and password (pwd) are provided correctly in the following command:</p> <pre>ch <- odbcConnect("Greenplum",uid="gpadmin", case="postgresql",pwd="changeme")</pre>

Step	Action
5	<p><u>Get Data from the Database:</u></p> <ol style="list-style-type: none"> Before creating the table, "income_state", you must first delete the table, if it already exists. Type in: <pre>sqlDrop(ch,"income_state")</pre> Use the sqlQuery command to create the table, "income_state" : <pre>> sqlQuery(ch, "CREATE TABLE income_state AS SELECT f.name AS state , round(avg(h.hinc),0) AS income FROM housing AS h JOIN fips AS f ON h.state = f.code WHERE h.hinc > 0 GROUP BY f.name DISTRIBUTED BY (income); ")</pre> <p>Note: This code creates the table, "income_state", in database "training2".</p> Inspect this table using the following command: <pre>sqlColumns(ch,"income_state")</pre> Review the output on the console. <p>Note: The SQL Query is available for you to copy and paste into the working directory File name: mod4lab4.sql.</p>

Step	Action
6	<p><u>Read in the Data for Modeling:</u></p> <p>You need the data to be read in as a “matrix”.</p> <ol style="list-style-type: none"> 1. Execute the following statement to read in the database table “income_state”. Use the sqlFetch command. The “rownames” attribute ensures the data is rendered as a matrix and the row names are taken from the column “state”. <pre>income <- as.matrix(sqlFetch(ch,"income_state", rownames="state")) > summary(income)</pre> <ol style="list-style-type: none"> 2. Review the results of “income” on the console window. 3. Ensure that in the “data” window the variable “income” is represented as a 52x1 integer matrix. ***<u>NOTE – Check in workspace windows to see this!</u>

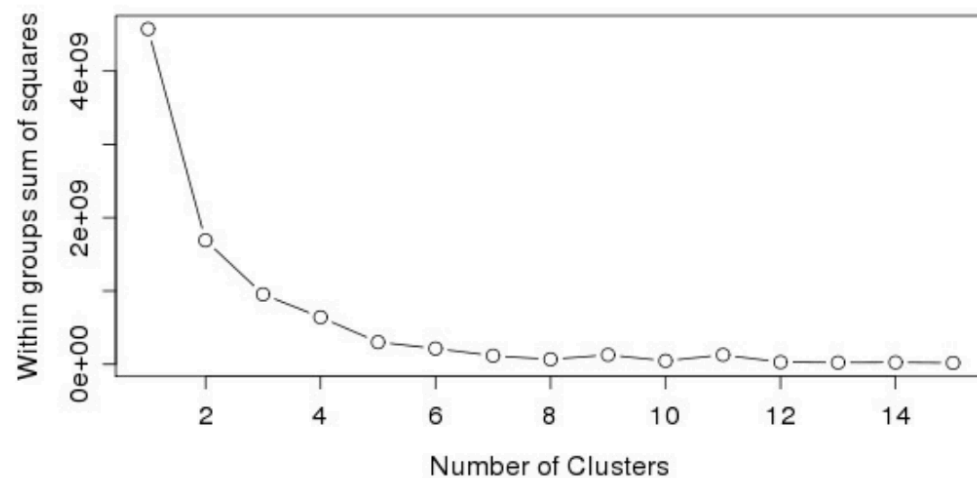
Step	Action
7	<p><u>Execute the Model:</u></p> <ol style="list-style-type: none"> Sort the data "income" before the modeling process. This will make it easier to understand the results and in visualizing. <pre>income <- sort(income)</pre> <p>The K-Means function, provided by the <i>cluster</i> package, is used as follows:</p> <pre>kmeans(x, centers, iter.max = 10, nstart = 1, algorithm = c("Hartigan-Wong", "Lloyd", "Forgy", "MacQueen"))</pre> <p>where the arguments are:</p> <ul style="list-style-type: none"> x: A numeric matrix of data, or an object that can be coerced to such a matrix (such as a numeric vector or a data frame with all numeric columns). centers: Either the number of clusters or a set of initial (distinct) cluster centers. If a number, a random set of (distinct) rows in x is chosen as the initial centers. iter.max: The maximum number of iterations allowed. nstart: If <i>centers</i> is a number, <i>nstart</i> gives the number of random sets that should be chosen. algorithm: The algorithm to be used. It should be one of the following "Hartigan-Wong", "Lloyd", "Forgy" or "MacQueen". If no algorithm is specified, the algorithm of Hartigan and Wong is used by default. Cluster the data into 3 groups (centers = 3) and also specify the number of random sets to start with as, 15. <pre>> # Fit the k-means cluster with 3 initial cluster centers > km <- kmeans (income,3,15)</pre>

Step	Action
9	<p><u>Plot the Results:</u></p> <p>1. Now plot the results using the following commands:</p> <pre data-bbox="407 457 964 527">> # plot clusters > plot(income, col = km\$cluster)</pre>  <p>The scatter plot shows 'income' on the y-axis (ranging from 30,000 to 70,000) and 'Index' on the x-axis (ranging from 0 to 50). The data points are colored based on their cluster assignment. There are three distinct clusters: a green cluster of points at the bottom left (Index 0-20, income 30,000-50,000), a red cluster of points in the middle (Index 20-45, income 50,000-65,000), and a black cluster of points at the top right (Index 45-50, income 65,000-70,000). The points follow a general upward trend.</p> <pre data-bbox="407 1031 1105 1100">> # plot centers > points(km\$centers, col = 1:3, pch = 8)</pre>  <p>This scatter plot is identical to the one above, but it also includes the cluster centers. The centers are marked with asterisks (*) and colored according to their cluster: green for the first cluster, red for the second, and black for the third. The green center is located at approximately Index 3 and income 45,000. The red center is at approximately Index 2 and income 60,000. The black center is at approximately Index 2 and income 75,000.</p> <p>2. Review the output on the graphic window.</p>

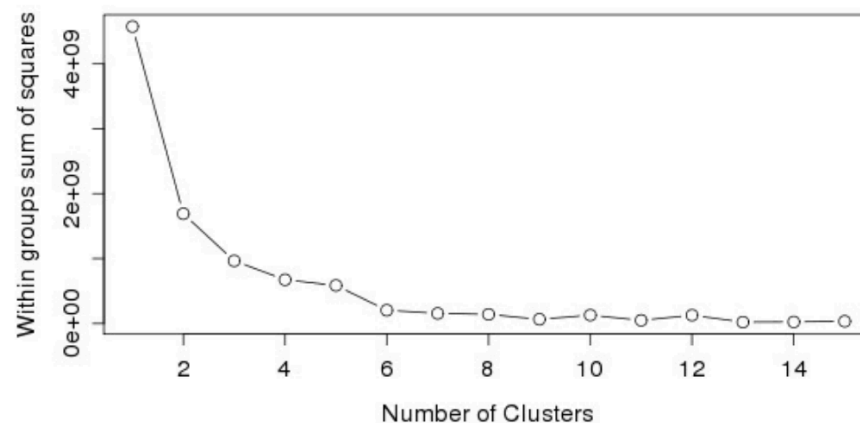
Find the Appropriate Number of Clusters:

1. Plot the within-group-sum of squares and look for an "elbow" of the plot. The elbow (if you can find one) tells you what the appropriate number of clusters probably is. (Adapted from <http://www.statmethods.net/advstats/cluster.html> (originally from Everitt & Hothorn)).

```
wss <- numeric(15)
> for (i in 1:15) wss[i] <- sum(kmeans(income,
  centers=i)$withinss)
> plot(1:15, wss, type="b", xlab="Number of Clusters",
  ylab="Within groups sum of squares")
```



2. Review the output on the graphic window. Is there an elbow to the plot? Yes
3. Repeat the modeling with a few values around the elbow (or 4 and 5 centers if there is no elbow) and review the results.



Step	Action
11	<p><u>Close Connections to ODBC Database:</u></p> <p>Use the following command:</p> <pre>odbcClose(ch)</pre> <p>The R Code for this exercise is available at /home/gpadmin/LAB04/kmeans1.R</p>

ACTION**Perform K-means Clustering Using In-database Analytics Method:**

1. **Open putty and connect to your BE IP**
2. **Navigate to cd LAB04**
3. **Run the sql command:** `psql -d training2 -f kmeansmadlib.sql`

Below is what the command is doing

```
DROP TABLE IF EXISTS myschema.data;
CREATE TABLE myschema.data (
  pid INT
  , position FLOAT8[])
DISTRIBUTED BY (pid);
```

```
INSERT INTO myschema.data (pid,position[1])
SELECT
  h.state
, round(avg(h.hinc),0)
FROM
  housing AS h
WHERE
  h.hinc > 0
GROUP BY
  h.state
;
```

```
SET SEARCH_PATH to madlib,public,myschema;
```

```
SELECT madlib.kmeans_random('myschema.data', 'position',
null, 'km_p', 'km_c','l2norm', 15,0.001,True, True, 6);
```

```
SELECT * FROM madlib.km_c;
SELECT * FROM madlib.km_p;
```

The first part of the code is similar to the one created in step 5 of this lab. The K-means function is called by:

using *random* centroid seeding method for a provided k :

```
SELECT * FROM kmeans\_random(
  'src_relation', 'src_col_data', 'src_col_id',
  'out_points', 'out_centroids',
  'dist_metric',
  max_iter, conv_threshold,
  evaluate, verbose,
```

```
k  
);
```

The centroid locations are stored in `kmeans_out_centroids_(run_id)`:

In the example above

`input_table` is `myschema.data`,

number of clusters is 6

4. The output will be of `km_c` and `km_p` from schema
`myschema.data`

*****provide screenshot**

This is similar to what you had done in step 7.

End of Lab Exercise