

**National University of Singapore
School of Computing**

**SWS3018 Predictive Analytics
Lab 7**

Learning Objectives

- Perform k-means clustering on R
- Perform hierarchical clustering on R

1. In this exercise, we will work with the Breast Cancer Wisconsin dataset. It can be download from:

Data:

<https://archive.ics.uci.edu/ml/machine-learning-databases/breast-cancer-wisconsin/breast-cancer-wisconsin.data>

Description of the data:

<https://archive.ics.uci.edu/ml/machine-learning-databases/breast-cancer-wisconsin/breast-cancer-wisconsin.names>

- a) Generate a K-means model for doing clustering
- b) Calculate the accuracy of the model generated in (a)
- c) Do you get different accuracy for different k-means model (different seed values)?
- d) Generate the hierarchical clustering models the different dissimilarity measures (complete, single, average, centroid). Show the dendrogram for each case
- e) Generate the clusters for 4 different hierarchical clustering models in (d) and measure the accuracy of the models for each case. Which linkage approach is better?
- f) Determine the classification accuracy for the decision tree, bagging, and forest approach and compare the accuracy with the above clustering approaches. How about Logistic Regression and SVM?