General Instruction

- Submit uncompressed file(s) in the Dropbox folder via BeachBoard (Not email).
- 1. Using scikit learn, evaluate the classification accuracy of the decision tree, bagging, AdaBoost, and Random forest.
 - (a) Load the Breast cancer data using sklearn.datasets.load_breast_cancer.
 - (b) (2 points) Print out the names of the features (X) and the name of the target (y).
 - (c) (2 points) Allocate the half of the data to *Train* (X_train, y_train) and the remaining half to *Test* (X_test, y_test).
 - (d) The common goal of the classifiers is predicting target using features.
 - (e) The classifiers should be trained using *Train* set and be tested using *Test* set.
 - (f) Use the 'Gini' index as the criterion and fix the maximum depth of trees as 2.
 - (g) (5 points) Write a program that generates a decision tree from X_train, y_train and predict y_pred from X_test. You can compute accuracy of the classifier by comparing y_pred and y_test. Please print out the accuracy.
 - (h) (5 points) Visualize the tree using sklearn.tree.plot_tree. Each node of trees should include feature name.
 - (i) (5 points) Similarly, write a program that generates multiple decision trees using the bagging. This method should record its prediction accuracy at bagging_score by varying the parameter n_estimators. Draw a 2D line plot whose X-axis is n_estimators and Y-axis bagging_score, and the plot should have more than 20 data points of different X-axis values.
 - (j) (5 points) Similarly, write a program that generates multiple decision trees using the AdaBoost. Draw a 2D line plot whose X-axis is n_estimators and Y-axis boost_score, and the plot should have more than 20 data points of different X-axis values.
 - (k) (10 points) Similarly, write a program that generates multiple decision trees using the random forest. Draw a 3D surface plot whose X-axis is n_estimators, Y-axis max_features, and Z-axis forest_score. The plot should have more than 100 data points of different pair of X-axis and Y-axis values.
 - (l) Submit your Assn12.ipynb which includes all the plots.