Lab5: Data classification using K-Nearest Neighbor Classifier

You are given the **Pima Indians Diabetes Database** as a csv file (pima-indians-diabetes.csv). This dataset is originally from the National Institute of Diabetes and Digestive and Kidney Diseases. It consists 768 tuples each having 9 attributes. The last attribute for every tuple signifies the class label (0 for non-diabetes and 1 diabetes). It is a two class problem. Other attributes are input features.

- 1. Write a python program to
 - a. Normalize all the attributes, except class attribute, of pima-indians-diabetes.csv using min-max normalization to transform the data in the range [0-1]. Save the file as pima-indians-diabetes-Normalised.csv
 - b. **Standardize**, all the attributes, except class attribute, of pima-indians-diabetes.csv using **z-normalization**. Save the file as pima-indians-diabetes-Standardised.csv
- 2. Split the data of each class from pima-indians-diabetes.csv into train data and test data. Train data contain 70% of tuples from each of the class and test data contain remaining 30% of tuples from each class. Save the train data as diabetes-train.csv and save the test data as diabetes-test.csv
 - a. Classify every test tuple using **K-nearest neighbor** (**KNN**) method for the different values of K (1, 3, 5, 7, 9, 11, 13, 15, 17, 21). Perform the following analysis:
 - i. Find **confusion matrix** (use 'confusion matrix') for each K.
 - ii. Find the **classification accuracy** (You can use 'accuracy_score') for each K. Note the value of K for which the accuracy is high.
- 3. Split the data of each class from pima-indians-diabetes-Normalised.csv into train data and test data. Train data should contain same 70% of tuples in Question 2 from each of the class and test data contain remaining same 30% of tuples from each class. Save the train data as diabetes-train-normalise.csv and save the test data as diabetes-test-normalise.csv
 - a. Classify every test tuple using **K-nearest neighbor** (**KNN**) method for the different values of K (1, 3, 5, 7, 9, 11, 13, 15, 17, 21). Perform the following analysis:
 - i. Find **confusion matrix** (use 'confusion matrix') for each K.
 - ii. Find the **classification accuracy** (You can use 'accuracy_score') for each K. Note the value of K for which the accuracy is high.
- 4. Split the data of each class from pima-indians-diabetes-Standardised.csv into train data and test data. Train data should contain same 70% of tuples in Question 2 from each of the class and test data contain remaining same 30% of tuples from each class. Save the train data as diabetes-train-standardise.csv and save the test data as diabetes-test-standardise.csv
 - a. Classify every test tuple using **K-nearest neighbor** (**KNN**) method for the different values of K (1, 3, 5, 7, 9, 11, 13, 15, 17, 21). Perform the following analysis:
 - i. Find **confusion matrix** (use 'confusion_matrix') for each K.

- ii. Find the **classification accuracy** (You can use 'accuracy_score') for each K. Note the value of K for which the accuracy is high.
- 5. Plot and the **classification accuracy vs K**. for each cases (original, normalized and standardized) in a same graph and compare & observe how it is behaving.
- 6. Why the value of K is considered as **odd** integer?

Note:

- 1. Note that while splitting the data (original, normalized and standardized) into train and test set, use the same seed value for random split of all 3 cases. This is to ensure that same training and test samples will be there in all 3 cases.
- 2. You can import the KNeighborsClassifier class from the sklearn.neighbors library
- 3. You can use the functions StandardScaler for standardization and MinMaxScaler for min-max normalization in scikit-learn.
- 4. Refer the slide uploaded in moodle about the performance evaluation to know more about confusion matrix and Accuracy.