## CSCI 184 Homework 1 Due Date: April 15th, 2022

## 1) One-Hot Encoding

For this part, you will focus on a house price detection dataset that comprises of 1460 rows and 81 columns. You will perform a comparative study between using one-hot encoded categorical columns as opposed to not using them at all.

What do you need to do?

- 1. Load the dataset from 'train.csv' into a pandas DataFrame and print it along with its shape.
- 2. For dealing with missing values, delete a column with any Null value (**Note: not recommended approach, but here focus is one-hot encoding**). Also delete the 'ID' column.
- 3. For simplicity, we will work with all the numerical columns but just a subset of the categorical columns. Among all the remaining categorical columns after step 2, choose the categorical columns that have less than 10 unique values.
- 4. Print the resultant dataframe and its shape after step 3.
- 5. Divide the data into X and Y, where X is the set of features and Y is the target variable. For this dataset the 'SalePrice' is the target variable.
- 6. For step 5, two approaches will be used (as discussed in class):
  - a. Set of ONLY numerical features and dropping all Categorical features
  - b. Set of features comprising both numerical and **one-hot encoded** categorical features.
- 7. Print the original dataframe joined with the one-hot encoded columns.
- 8. Once, the two sets of features are obtained from step 6, train a simple Linear Regression model from sklearn and obtain the Mean Absolute Error for both the cases:
  - a. Mean Absolute Error when Dropping Categoricals:
  - b. Mean Absolute Error with One-Hot Encoding
- 9. Write a report with screenshots of your results and the final results for step 8 a and b.
- 10. Submit your code as an .ipynb file and a document reporting your findings.

## 2) Feature Selection

For this part, you will focus on feature selection for the diabetes dataset.

You can download the data directly using this URL:

https://raw.githubusercontent.com/jbrownlee/Datasets/master/pima-indians-diabetes.dat a.csv

What do you need to do?

- 1. Load the dataset. The link does not have any names for headers. Add them with the following (in order: Pregnancies, Glucose, BloodPressure, SkinThickness, Insulin, BMI, DiabetesPedigreeFunction, Age, Outcome)
- 2. Print the DataFrame and its shape.
- 3. Convert the DataFrame object to NumPy array.
- 4. Segregate the data into X and Y. Here 'Outcome' is your target variable.
- 5. Perform Feature Extraction
  - a. Using Filter-Based Method
    - i. For numerical input and categorical output, refer to slides to figure out which method you want to use.
- 6. Set k value to 3,4 and 5 for step 5
- 7. For each value of k, specify the features that are extracted:

Method	Value of k	Feature Names	
Filter-Based	3	:	
	4	:	
	5	:	

- 8. After you have extracted the features, build a Logistic Regression Model from sklearn using two features:
  - a. Case 1: Two of the features are from the list of extracted features using the Filter-based method for k = 3.4 and 5
    - For example, if Pregnancies, Glucose, BMI and Age are the four features as a result of k = 4, choose any 2 of the features to train your model.

- b. **Case 2:** Two of the features are columns that were not extracted in the feature extraction phase.
  - For example, if Pregnancies, Glucose, BMI and Age are the four features as a result of k = 4, choose any 2 of the features from BloodPressure, SkinThickness, Insulin, DiabetesPedigreeFunction.

For each case, choose two combinations of features and train your model.

- 9. Evaluate your model. You may choose to evaluate the training data itself.
- 10. Report your findings in a table:

Case	Feature Names	Precision	Recall	F1-Score
1				
2				

Write a report with screenshots of your results and the final results for step 10. Submit your code as an .ipynb file and a document reporting your findings.

You may choose to use the same report for two of the parts, but please use two separate .ipynb files for the two parts.